The Ocular Surface 15 (2017) 438–510

Contents lists available at ScienceDirect

The Ocular Surface

journal homepage: www.theocularsurface.com

TFOS DEWS II pathophysiology report

Anthony J. Bron, FRCOph, FMedSci [a, b, 1, *], Cintia S. de Paiva, MD, PhD [c, 1], Sunil K. Chauhan, DVM, PhD [d, 1], Stefano Bonini, MD [e], Eric E. Gabison, MD [f], Sandeep Jain, MD [g], Erich Knop, MD, PhD [h], Maria Markoulli, PhD, MOptom [i], Yoko Ogawa, MD [j], Victor Perez, MD [k], Yuichi Uchino, MD, PhD [j], Norihiko Yokoi, MD, PhD [l], Driss Zoukhri, PhD [m], David A. Sullivan, PhD [d]

[a] Nuffield Department of Clinical Neurosciences, University of Oxford, Oxford, UK
[b] Vision and Eye Research Unit, Anglia Ruskin University, Cambridge, UK
[c] Department of Ophthalmology, Baylor College of Medicine, Houston, TX, USA
[d] Schepens Eye Research Institute & Massachusetts Eye and Ear, Harvard Medical School, Boston, MA, USA
[e] Department of Ophthalmology, University Campus Biomedico, Rome, Italy
[f] Department of Ophthalmology, Fondation Ophtalmologique Rothschild & Hôpital Bichat Claude Bernard, Paris, France
[g] Department of Ophthalmology and Visual Sciences, University of Illinois at Chicago, Chicago, IL, USA
[h] Departments of Cell and Neurobiology and Ocular Surface Center Berlin, Charité – Universitätsmedizin Berlin, Berlin, Germany
[i] School of Optometry and Vision Science, University of New South Wales, Sydney, Australia
[j] Department of Ophthalmology, Keio University School of Medicine, Tokyo, Japan
[k] Department of Ophthalmology, Bascom Palmer Eye Institute, University of Miami, Miami, FL, USA
[l] Department of Ophthalmology, Kyoto Prefectural University of Medicine, Kyoto, Japan
[m] Tufts University School of Dental Medicine, Boston, MA, USA

ARTICLE INFO

Article history:
Received 26 May 2017
Accepted 26 May 2017

Keywords:
TFOS
DEWS II
Dry eye workshop
Dry eye disease
Pathophysiology
Glycocalyx
Hyperosmolarity
Inflammation
Vicious circle

ABSTRACT

The TFOS DEWS II Pathophysiology Subcommittee reviewed the mechanisms involved in the initiation and perpetuation of dry eye disease. Its central mechanism is evaporative water loss leading to hyperosmolar tissue damage. Research in human disease and in animal models has shown that this, either directly or by inducing inflammation, causes a loss of both epithelial and goblet cells. The consequent decrease in surface wettability leads to early tear film breakup and amplifies hyperosmolarity via a Vicious Circle. Pain in dry eye is caused by tear hyperosmolarity, loss of lubrication, inflammatory mediators and neurosensory factors, while visual symptoms arise from tear and ocular surface irregularity. Increased friction targets damage to the lids and ocular surface, resulting in characteristic punctate epithelial keratitis, superior limbic keratoconjunctivitis, filamentary keratitis, lid parallel conjunctival folds, and lid wiper epitheliopathy. Hybrid dry eye disease, with features of both aqueous deficiency and increased evaporation, is common and efforts should be made to determine the relative contribution of each form to the total picture. To this end, practical methods are needed to measure tear evaporation in the clinic, and similarly, methods are needed to measure osmolarity at the tissue level across the ocular surface, to better determine the severity of dry eye. Areas for future research include the role of genetic mechanisms in non-Sjögren syndrome dry eye, the targeting of the terminal duct in meibomian gland disease and the influence of gaze dynamics and the closed eye state on tear stability and ocular surface inflammation.

1. Goals

To:

- Summarize current understanding of tear physiology as it relates to dry eye disease (DED).
- Provide an etiological classification of DED.
- Identify the core mechanisms of DED, especially ocular surface hyperosmolarity, tear instability and the inflammatory response.
- Consider the Vicious Circle of DED and chronic DED as a self-perpetuating disease.
- Discuss asymptomatic and symptomatic DED and the basis of DED symptoms.

* Corresponding author. Nuffield Department of Clinical Neurosciences, University of Oxford, Oxford, UK.
E-mail address: anthony.bron@eye.ox.ac.uk (A.J. Bron).
[1] Subcommittee Co-Chair

http://dx.doi.org/10.1016/j.jtos.2017.05.011
1542-0124/

- Review the role of environment in precipitating DED in at-risk subjects and influencing DED severity.

2. Definition of dry eye disease

TFOS DEWS II has redefined dry eye as: "Dry eye is a multifactorial disease of the ocular surface characterized by a loss of homeostasis of the tear film, and accompanied by ocular symptoms, in which tear film instability and hyperosmolarity, ocular surface inflammation and damage, and neurosensory abnormalities play etiological roles" (see TFOS DEWS II Definition & Classification Subcommittee report [1221]).

3. Introduction

The purpose of this report is to review our understanding of the pathophysiology of DED, highlighting those advances that have occurred since the TFOS DEWS report [1]. Our general thesis is that DED is initiated by desiccating stress and perpetuated by a Vicious Circle of ocular surface inflammation.

The *raison d'etre* of the eye is sight and the precorneal tear film and the cornea provide the first refractive element of the eye that focuses an image of the visual world upon the retina. To maintain optical quality, the tear film must be constantly replenished by blinking and tear secretion. Without this, the tear film would destabilise and the surface of the eye would be exposed to damaging desiccation. Various mechanisms are in place to achieve homeostasis.

4. Anatomy and physiology of the ocular surface and lacrimal system

4.1. Ocular surface

The ocular surface is covered by a continuous sheet of epithelium, lining the cornea, the anterior globe and tarsi and extending to the mucocutaneous junctions (MCJs) of the lid margins. Hydration of the ocular surface is maintained by the tears, which bathe it continuously and provide an unbroken film over its exposed surface. The tears are secreted chiefly by the lacrimal glands, with additional contributions from the conjunctiva, including the goblet cells and Meibomian glands.

The open eye is constantly subjected to desiccating stress through evaporation of the tears, but is protected from damage by homeostatic mechanisms that regulate tear secretion and distribution in response to signals from the ocular surface. In DED, a failure of these mechanisms leads to a quantitative or qualitative deficiency of tears that typically induces tear film instability, wetting defects and hyperosmolar stress, increased friction and chronic mechanical irritation at the ocular surface. This initiates a chain of inflammatory events and surface damage that characterise the disease.

4.2. Main and accessory lacrimal glands

The main lacrimal gland is a tubule-acinar, serous gland composed primarily of acinar, ductal, and myoepithelial cells, with the acinar cells comprising 80% of the total. It develops by a process of branching, involving reciprocal interactions between the epithelium and surrounding mesenchyme [2,3] to produce a three-dimensional tubular network [4]. In humans, the main gland consists of a larger orbital lobe, and a smaller palpebral lobe that abuts the conjunctival sac. The ducts from the orbital lobe pass through, and join with, those of the palpebral gland, to open into the superior fornix [5], via 6 to 12 orifices [6]. In addition, there are about 40 accessory glands of Krause located in the upper fornix and 6 to 8 in the lower fornix. The accessory lacrimal glands of Wolfring, located in the upper (2–5 glands) and lower (1–3 glands) lids, are slightly larger than those of Krause. The accessory lacrimal glands are tubular glands which do not contain acini in humans [7], but do in rabbits [8]. The accessory glands constitute about 10% of the total lacrimal tissue mass [9] and are innervated similarly to the main gland [10]. They are therefore assumed to respond in a similar way to reflex stimulation.

4.2.1. Resident immune cells of the lacrimal gland

The lacrimal gland is richly supplied by immune cells that occupy the interstitial space. They include: plasma cells, B and T cells, dendritic cells, macrophages, bone marrow-derived monocytes, and mast cells [11] (Table 1).

Plasma cells predominate (53.9% of the total), mainly immunoglobulin (Ig) A+ and with a few IgG+, IgM + or IgD+. The IgA + cells synthesize and secrete IgA, which is transported into acinar and ductal cells, combined with J-piece and secretory component (SC) and secreted as dimeric, secretory IgA (sIgA) [12,13]. A similar event may occur in the conjunctiva and in other Eye-Associated Lymphoid Tissues (EALT) [14].

T cells are the next most common cell, (40.3% of total), dispersed with plasma cells in the interstitium, in follicles and aggregates and occasionally between acinar cells. T cell aggregates are typically related to intra-lobular ducts. Overall, T suppressor/cytotoxic cells (T8) are more numerous than T helper cells (T4), distributed almost equally between acini, ducts and interstitium. The T4/T8 ratio is 0.26 in the interstitium. However, T4 cells predominate in follicles and lymphocytic aggregates. Dendritic cells, macrophages, bone marrow-derived monocytes and mast cells are also present.

B-cells are found exclusively in the centre of primary follicles and aggregates and in solitary, secondary follicles, surrounded by T helper cells and a lesser number of suppressor/cytotoxic cells. They are not found in the interstitium. They make up 5.7% of the mononuclear population. B-cells and the dendritic cells of follicles and aggregates express human leukocyte antigen D-related (HLA-DR) as do duct lining cells and the vascular endothelium. Macrophages and dendritic cells are uncommon.

4.2.2. Regulation of lacrimal secretion

The acinar cells are arranged in lobules around a central lumen, with tight junctions surrounding each cell on the apical (luminal) side [12,15]. This configuration permits the unidirectional, basal-to-apical, secretion of water, electrolytes, proteins and mucins [12,15]. The basal portion of the cell contains a large nucleus, rough endoplasmic reticulum, mitochondria, and Golgi apparatus while the apical portion is filled with secretory granules [12,15]. The acinar cells synthesize, store, and secrete proteins and mucins in response to neural and hormonal stimuli [13,15]. They also secrete electrolytes and water. Many of the proteins secreted have either growth factor or bactericidal properties, which are crucial to the health of the ocular surface. Several mucins, both secreted as well as membrane-bound have been detected in the lacrimal gland including MUC1, MUC4, MUC5B, MUC5Ac, MUC6, MUC7 and MUC16 [16–18]. Some of them perform local roles but otherwise their functions are not known.

Like the acinar cells, the duct cells are polarized by apically located tight junctions [12]. Importantly, the ductal cells modify the primary fluid secreted by the acinar cells by absorbing or secreting water and electrolytes [19,20]. The duct cells secrete a KCl-rich solution so that the finally secreted lacrimal gland fluid is rich in K^+ ions. It has been estimated that as much as 30% of the volume of

Table 1
Resident immune cells of the normal human lacrimal gland.

Tissue Layer	Plasma Cells	T Cells	T cell Phenotype	B-Cells	Macs	DCs	pDCs
Acinar	53.9%	40.3%	Generally, suppressor/cytotoxic	5.7%	0.01%	5.6%	+
Ductal			T cells dominate				+
Interstitium	++++						+
Follicles & Aggregates		Espec. peri-ductal	Generally, helper cells dominate	++			
Notes	Mainly IgA + Some IgG, M, D		Activated T cells 0.01%				

Macs = Macrophages; DCs = Dendritic Cells; pDCs = Plasmacytoid cells; Data from Ref. [11].

the final lacrimal gland fluid is secreted by the duct cells [19,20].

The *myoepithelial cells* lie scattered between the acinar and ductal cells and the basal lamina and are interconnected by gap junctions and desmosomes [21] They synthesize basal lamina and their multiple processes form a functional network around the acinar and ductal cells, separating them from the basal lamina and the mesenchymal, stromal cells [22]. Myoepithelial cells contain contractile muscle proteins (α smooth muscle actin, myosin, tropomyosin) [21], and are assumed to assist in expelling fluid from the acini and the ducts.

The lacrimal gland is innervated by the parasympathetic and sympathetic nervous system [23,24]. Nerve terminals are located in close proximity to acinar, ductal, and myoepithelial cells as well as blood vessels, and hence can control a wide variety of lacrimal gland functions [23,24]. Stimulation of lacrimal gland secretion occurs in part through a neural reflex arc originating from the ocular surface [13,15,23,25] with a further trigeminal input arising from the nasal mucosa [26]. Neurotransmitters and neuropeptides released by innervating nerves include acetylcholine, vasoactive intestinal peptide (VIP), norepinephrine, neuropeptide Y (NPY), substance P (SP), and calcitonin gene related peptide (CGRP). Each of these neuromediators interacts with specific receptors present on the surface of lacrimal gland cells to elicit a specific response [13,15,25]. Acetylcholine and norepinephrine are the most potent stimuli of lacrimal gland protein, mucin, water, and electrolyte secretion [13,15].

4.2.3. Lacrimal gland stem cells

The lacrimal glands, like the salivary and mammary glands, retain their ability to regenerate through their whole life span. For epithelial cells of the salivary glands the reported cell turnover is 40–65 days for serous acini and 95 days for duct cells [27]. Since the lacrimal glands share many characteristics in common with the salivary glands, it is possible that lacrimal epithelial cells have a similar cell turnover rate.

Stem cells are present in the lacrimal glands of mice [28], rats [29] and humans [28] and their involvement in repair has been studied in mice [30]. In a lacrimal gland injury model, stem cells participated in lacrimal gland regeneration [31] and those isolated from murine glands by Ackermann et al. had the ability to differentiate into all three germ layers [28].

4.2.4. Mechanisms of gland damage and repair

When the lacrimal gland is damaged acutely, (eg. following radiation exposure) or chronically (eg. in Sjögren syndrome and other autoimmune diseases) [32] the lacrimal gland is infiltrated by lymphocytes and other immune cells, with a predilection for the peri-ductal areas. This leads to a loss of acinar, ductal and myoepithelial cells, probably by both apoptosis and autophagy.

Remodeling following injury often recapitulates events that govern embryonic tissue development and it is therefore not surprising that programmed cell death and a number of growth factors and cytokines known to regulate tissue development play a role during lacrimal regeneration [30,32]. A key mechanism in the murine gland is epithelial-mesenchymal transition (EMT), which, during embryogenesis, helps epithelial cells to acquire migratory and/or invasive properties [33]. During EMT, epithelial cells lose cell-cell and cell-matrix attachments, polarity and epithelial-specific markers, undergo cytoskeletal remodeling, and gain a mesenchymal phenotype [33]. Induction of EMT generates cells with mesenchymal stem-like properties, which can play a significant role in tissue repair [34,35].

4.3. The meibomian glands

The meibomian glands are modified sebaceous, holocrine glands whose acini discharge their entire contents in the process of secretion. Their secretory product (meibomian lipid or meibum) is delivered into a shallow reservoir on the skin of the lid margin, just anterior to the mucocutaneous junction, and is spread onto the preocular tear film with each blink. The embryology, anatomy, histology and physiology of the glands were reviewed fully in the report of the TFOS Meibomian Gland Dysfunction Workshop (2011) [36] and elsewhere [37] and only selected aspects are discussed here.

The development of the meibomian glands has features in common with that of the pilosebaceous unit [38]. The luminal cells of the meibomian ducts, corresponding to the keratinized lining of the lash shaft, express keratohyalin granules and have been regarded as a modified, keratinized epithelium [39]. The glands of Zeiss, which satisfy the sebaceous needs of the cilia, are analogous to the meibomian glands. It appears that the capacity of the meibomian duct cells to keratinize is amplified in certain conditions, such as meibomian gland dysfunction (MGD) where keratinization of the terminal duct is a key feature, in metaplastic trichiasis where dystopic cilia may arise from meibomian orifices, in distichiasis where a row of aberrant lashes replaces that of the meibomian glands, and in follicular ichthyosis where both the meibomian glands and pilosebaceous units of the skin are affected together.

The human meibomian gland is richly innervated with sensory, sympathetic, and parasympathetic nerves [40,41]. These nerve fibers express substance P (SP), vasoactive intestinal peptide (VIP), dopamine β-hydroxylase, acetylcholinesterase, nitric oxide synthase, tyrosine hydroxylase, somatostatin, neuropeptide Y (NPY), and calcitonin gene-related peptide (CGRP) [40,41]. Human meibomian gland epithelial cells also express functional muscarinic and VIP receptors, and respond to an acetylcholine analog, carbamyl choline, and/or VIP with alterations in cyclic adenosine monophosphate (cAMP) and intracellular [Ca^{2+}] levels and cellular proliferation [42]. During differentiation these cells also have increased expression of genes coding for proteins with neuron remodeling and axon guidance activities (e.g. netrin 4 and collagen, type V, alpha2) [43]. In addition to humans, the mouse meibomian gland contains mRNAs of receptors for cholinergic, adrenergic, NPY, serotonin, CGRP, dopamine, γ-aminobutyric acid, glutamate, neurotensin, and somatostatin [36,44].

Multiple factors are known to regulate the meibomian gland. The meibomian gland *in vivo* [36], (see TFOS DEWS II Sex, Gender and Hormones Subcommittee report [1222]), and human

meibomian gland epithelial cells *in vitro* [42,43,45–61], respond to numerous agents with alterations in proliferation, differentiation, cAMP accumulation, signaling pathways, gene expression and/or lipogenesis. These compounds include androgens, estrogens, progesterone, glucocorticoids, insulin, pituitary hormones, mineralocorticoids, growth factors, bacterial toxins, antibiotics, cationic amphiphilic drugs, omega fatty acids, retinoic acid, high glucose, cyclosporine A, an IL-1 receptor antagonist, rebamipide, bimatoprost, pilocarpine and timolol [42,43,45–54,56–60,62,63].

Chemical analysis of expressed meibomian lipid shows it to consist of about 95% nonpolar lipids (mainly wax and cholesterol esters, with a small amount of triglycerides) and 5% polar lipids, (the amphipathic lipid, O-acyl-ω-hydroxy-fatty acid (OAHFA) [64] and phospholipids (PL)) [65]. The concentration of OAHFA exceeds that of PL in meibum but the ratio is reversed in the tear film [66]. The lipid composition of meibum and tears is discussed fully in the TFOS DEWS II Tear Film Subcommittee report [1223].

The key building block for cholesterol and fatty acid synthesis is cytosolic acetyl-CoA, a product of carbohydrate, fatty acid or amino acid metabolism [67]. Cholesterol biosynthesis involves the successive conversion of acetyl-CoA to acetoacetyl-CoA, 3-hydroxy-3-methylglutaryl-CoA (HMG-CoA) and mevalonate, catalyzed respectively by acetoacetyl-CoA-synthase, HMG-CoA synthase 1 and HMG-CoA reductase. Cholesterol itself is utilized in the synthesis of sex steroid hormones and the enzymes regulating this process are present in the human meibomian gland [68].

Fatty acid biosynthesis involves the initial conversion of cytosolic acetyl-CoA into malonyl-CoA, catalyzed by the rate-limiting enzyme, acetyl-CoA carboxylase. Malonyl-CoA is then converted into palmitoyl-CoA in the presence of the enzyme fatty acid synthase and ultimately, palmitoyl-CoA is elongated into longer chain, saturated fatty acids by the addition of 2-carbon units. Production of unsaturated fatty acids requires the action of fatty acid desaturases. The fatty acids are utilized to create neutral and polar lipids. Messenger RNAs for each of the above-mentioned enzymes and others involved in cholesterol and fatty acid synthesis, have been demonstrated in the murine meibomian gland in addition to mRNAs for the sterol regulatory element binding proteins (SREBPs) 1 and 2, which play a critical role in regulating their activity at a transcriptional level [69] SREBP 1 has also been identified in human meibomian gland epithelial cells [52].

The SREBPs, together with the membrane binding transcription factor proteases, (MBTPs), site-1 and 2, (otherwise known as site 1 and site 2 proteases - S1P and S2P) are key regulators of cholesterol and fatty acid synthesis and homeostasis [70].

SREBP-1 and SREBP-2 are membrane-bound transcription factors located in the endoplasmic reticulum (ER). When the cellular demand for lipid rises, SREBPs, complexed with the escort protein, Scap, are transported within coated vesicles to the Golgi apparatus, where they undergo activation within the Golgi membrane. This occurs in two stages. In the first step, the site-1 serine protease, S1P, cleaves the SREBP protein within the Golgi membrane. In the second step, the amino terminal fragment, containing the transcription factor, is rapidly released by the site 2 protease and migrates into the cell nucleus where it activates the transcription of genes needed for cholesterol uptake and synthesis as well as those involved with fatty acid metabolism [67,71,72].

There are additional membrane-bound transcription factors within the ER that act as so-called ER 'stress sensors'. A deficiency in either function, sterol biosynthesis or the ER stress response, may be the basis of the X-linked syndrome of ichthyosis follicularis, atrichia, and photophobia (IFAP syndrome), in which there is a failure of pilosebaceous development in the skin and lids resulting from mutations in the MBTPS2 gene [73].

Of relevance to the role of hormones on meibomian gland function and dysfunction, mRNAs for each of the above-mentioned genes have been shown to be upregulated by testosterone in the castrated mouse model, including adenosine triphosphate (ATP)-citrate lyase and acetyl-CoA synthase, enzymes which are critical for the initiation of lipogenesis [44,69]. Schirra et al. have suggested that the enhanced expression of genes for SREBPs 1 and 2 in response to androgen exposure may explain the hormonal induction of meibomian lipids [68]. SREBP 1 is known to be controlled by androgens at other sites [72].

4.4. The conjunctiva

The conjunctiva is a mucous membrane with a *lamina propria* (stroma) of loose connective tissue, covered by an epithelium that is kept permanently moist. The conjunctiva acts as a barrier against the outer environment and secretes a variety of products into the tear film. It also takes up antigens selectively for immune protection. Several regions of the conjunctiva can be identified [74,75]. The 'marginal' zone extends from the subtarsal fold to the MCJ on the lid margin [76] and includes the mucosa of the lid wiper zone [36]. Proximal to this, the tarsal conjunctiva is tightly attached to the tarsal plate and then continues as a loose orbital zone towards the fornix.

4.4.1. Conjunctival epithelium

The conjunctival epithelial cells, tightly connected by adherens junctions which provide strength against shear stress and the most superficial (i.e. layer 1) cells, are sealed by tight junctions which act as a barrier against the outside world. This barrier is less tight than that of the corneal epithelium [77]. Alterations of conjunctival and corneal integrity are associated with ocular surface disease (OSD) [78]. Between the conjunctival epithelial cells are considerable intercellular spaces [79] that are assumed to be associated with a role in water transport across the epithelium. The conjunctival epithelium consists of two cell types - epithelial cells and goblet cells, both deriving from the same conjunctival stem cell [80].

The conjunctival epithelial cells produce, apart from water, electrolytes and mucins [81], functional proteins such as lubricin [82]. The layer 1 cells produce integral membrane mucins that constitute the superficial glycocalyx of the cell, necessary for wetting by the aqueous tears [83]. Conjunctival epithelial cells contain transmembrane water channels (aquaporins) concerned with water movement between the conjunctiva and the aqueous phase of the tear film [84]. A further epithelial function may be an SC-mediated transcytosis of IgA, from plasma cells in the lamina propria, but this has yet to be shown [85].

4.4.2. Conjunctival epithelial stem cells

Stem cells may be defined as progenitor cells with a high capacity for cell division and the ability to generate a terminally differentiated progeny [86,87]. The stem cells of the corneal epithelium are located at the limbus and the subject has been reviewed extensively [88–91]. The location of the conjunctival stem cells in the human is more controversial. Wei et al., using tritiated thymidine in the rabbit concluded that the fornix was a major site of conjunctival stem cells [92,93]. Pellegrini et al., however, using clonal analysis of cells from various sites, reported that conjunctival stem cells are uniformly distributed in the human bulbar conjunctiva [80]. Pe'er and colleagues, in the mouse, using a tritiated thymidine label, identified conjunctival progenitor cells at both the limbus and the mucocutaneous junction (MCJ), with the MCJ giving rise to cells that streamed toward the fornix [94]. Wirtschafter et al., reported a similar finding in the rabbit, with a focus of label-retaining cells at the MCJ of the lid margin, whose transient amplifying progeny migrated over time towards the fornix [95]. They inferred that conjunctival stem cells were located

mainly at the MCJ. Most recently, in human cadaveric tissue, Stewart et al. have reported the expression of stem cell markers scattered throughout the conjunctiva, with the highest levels in the medial canthal and inferior fornical areas [96].

4.4.3. Conjunctival goblet cells

Human conjunctival goblet cells are present as single cells, scattered throughout the conjunctival epithelium save for a small temporal, perilimbal patch. Their numbers increase from the superior temporal region to the inferior nasal region of the conjunctival sac [97]. They package and secrete the gel-forming mucin, MUC5AC [83], which, when fully glycosylated has a mass of up to 40 MDa [98,99].

Gel mucins have an enormous water-binding capacity and thereby transform the aqueous tears into a mucoaqueous gel that makes up the main volume of the preocular tear film and maintains moisture at the ocular surface [100]. Mucins also have a lubricative function at the lid-globe interface that is important for movements of the eyeball relative to the lids. This lubricative function is needed in particular at the elevated epithelial lip of the lid-wiper, where the posterior border of each lid comes into close contact with the globe. Here, goblet cells are supplied within mucus crypts [101], similar to those in the tarsal conjunctiva [102]. The mucin of the mucoaqueous layer has other protective properties, binding microorganisms and inhibiting their attachment to the epithelium and also binding sIgA, several antimicrobial proteins and peptides [103]. In this way it serves as an integral component of the ocular surface surveillance system [104]. The role of T helper cell (Th1 and Th2) cytokines in goblet cell homeostasis [105] is discussed in a later section.

Release of the secretory mucin, MUC5AC, can be induced by either parasympathetic or sympathetic nerve stimulation [106–108]. In the rat, the parasympathetic neurotransmitters acetylcholine and vasoactive intestinal peptide (VIP) stimulate conjunctival goblet cell secretion *in vivo*, in both cells and organ culture [108–110]. In addition, nucleotides that activate the $P2Y_2$ receptor, such as ATP and uridine triphosphate (UTP), and also $P2Y_2$ receptor agonists, can stimulate goblet cell mucin secretion in rat and human conjunctiva [111,112]. Additionally, the epidermal growth factors (EGF) and brain-derived growth factor (BDNF) stimulate a slow release of MUC5AC from cultured rat conjunctival goblet cells [110,113].

4.4.4. Resident immune cells of the conjunctiva

A study of the resident leukocytes of the human conjunctiva by Hingorani et al. showed a greater number in the bulbar than tarsal conjunctiva [114], although other distributions have been reported [115].

T cells ($CD3^+$) were the dominant cell population, 75% of which were memory, or primed T cells (CD45Ro+) rising to 75–100% in the epithelium. $CD8^+$ T cells were more common than $CD4^+$ T cells in the epithelium whereas their numbers were roughly equal in the stroma. Macrophages ($CD68^+$) were the second most frequent conjunctival leukocyte present in both epithelium and stroma, accounting, with Langerhans cells, for those cells expressing HLA-DR. The exact numbers of leukocytes in general and lymphocytes in particular vary in different studies [114,116,117], but authors agree that T-lymphocytes dominate over B-lymphocytes, and that, of the plasma cells, those that produce IgA by far outnumber those that produce IgM. Neutrophils and occasionally, B cells, were present in the epithelium of both bulbar and tarsal conjunctiva while plasma cells, natural killer cells and mast cells, present in small numbers, were confined to the stroma. As Hingorani et al. conclude, T cells, macrophages and occasional B cells and neutrophils in the epithelium and T cells, B cells, macrophages, plasma cells, NK cells, mast cells and neutrophils in the substantia propria may be considered normal. A fuller review of the cellular, ocular surface immune defence system can be found elsewhere [117–119].

Hingorani et al.found only a single example of a lymphoid aggregate compatible with the presence of CALT (conjunctiva-associated lymphoid tissue), part of the MALT system of mucosa-associated lymphoid tissue, but did not examine fornical tissue, where CALT aggregates are most likely to be found [120]. Wotherspoon et al. [121] who examined the entire human superior and inferior fornical conjunctiva in autopsy material, found organised lymphoid tissue in only 31% of cases. The resident immune cells of the conjunctiva are summarised in Table 2 and those of the cornea in Table 3.

4.5. The glycocalyx of the ocular surface epithelia

The apical membranes of the layer 1 cells of the ocular surface epithelia present microvilli and microplicae which project into the tears and increase the interactive surface area at the tear/cell interface. Contiguous layer 1 cells are connected by tight junctions which restrict the entry of water-soluble solutes into the epithelium, and a further barrier is provided by the dense, apical glycocalyx [122], rich in transmembrane mucins [83]. Heavy glycosylation of the mucin exodomains converts the plasma membranes from a hydrophobic to a hydrophilic surface, which confers wettability to the epithelium [123–125]. The glycocalyx also acts as a lubricant that reduces friction at the ocular surface [126,127] and as a anti-adhesive that combats microbial colonization [128,129].

4.5.1. The transmembrane mucins

The transmembrane mucins of the human corneal and conjunctival epithelial glycocalyx [130], are MUC1 [131], MUC4 [132] and MUC16 [133], with galectin-3 playing additional roles [134]. The membrane-associated mucins possess short cytoplasmic tails, a single transmembrane domain and highly O-glycosylated extracellular, ectodomains with a variable number of tandem repeats (VNTR) [135,136] which extend at least 200–500 nm above the plasma membrane, far beyond other cell surface glycoproteins [137,138], thereby projecting into the tear film.

MUC1 is the smallest of the three glycocalyx mucins, with a molecular weight of approximately 120–300 kDa, roughly doubling in size after full glycosylation [139]. MUC1 exhibits anti-adhesive, cell-cell and cell-extracellular matrix properties [140,141]. The cytoplasmic tail of MUC1 (MUC1-CT) engages in signaling activities involving phosphorylated serine and tyrosine residues that act as binding sites for molecules such as NF-kB [142]. These can regulate the transcription of proinflammatory cytokines, abrogate the interaction of β-catenin with E-cadherin and upregulate the expression of the epithelial mesenchymal transducers (EMTs) [143].

MUC4 has a molecular mass of 900 kDa, several times greater than that of MUC1 [139]. It is expressed predominantly by conjunctival epithelium and to a lesser extent by the epithelium of the limbus and peripheral cornea. Very little is expressed in the central cornea [132,144].

MUC16 is the largest mucin yet identified in the human body, with a molecular mass of 2.5 MDa and a potential glycosylated mass of approximately 20 MDa [136,145,146]. The ectodomain of MUC16 is heavily O-glycosylated and longer than that of the other transmembrane mucins. Its cytoplasmic tail binds to the ezrin/radixin/moesin (ERM) family of proteins, which anchor the mucin to the actin cytoskeleton of the microvilli [128]. Knockdown of MUC16 expression in human corneal epithelial limbal cells, resulted in increased rose bengal dye penetration, increased *Staphylococcus aureus* binding to the epithelium [128] and disruption of tight junctions [147]. This, and other evidence [148] confirms MUC16 as a

Table 2
Resident immune cells of normal human conjunctiva.

Layer	T cells	Macs	LCs	PMNs	B-cells	Plasma cells	NK cells	Mast cells
Epithelium	++++ T cells $CD8^+$ > $CD4^+CD8+/CD4+ = 3.3$	+++	+	+	±			
Stroma	++++ T cells $CD8^+ \cong CD4^+CD4+/CD8+ = 1.3$	+++			±	+	+	+

Macs = macrophages; LCs = Langerhans cells; PMNS = neutrophils; NK = natural killer; Data from Refs. [114,121,1104,1105].

Table 3
Resident immune cells of normal human cornea.

Corneal Layer	Cell type (Phenotype)[a]	Peripheral Cornea[b]	Central Cornea
Epithelium	**Langerhans cells** ($CD45^+$ $CD11c^+$ $CD11b^{lo}$ MHC II^+ $Langerin^+$)	++++	++
Stroma	**Bone marrow-derived DCs**[a] ($CD45^+CD11c^+CD11b^+CD8\alpha^-MHCII^{+/-}CD80/86^{+/-}$)	++++	++
	Non-LC DCs[a] ($CD11c^+$ $Langerin^+$ $CD11b^+$ $CD103^{lo}$)	+++	++
	Macrophages ($CD45^+$ $CD11b^+$ $CD11c^-$)	+++	++
	Monocytic precursor cells $CD14^+$ $MHCII^-$ $B7^-$ $CD40^-$ $GR1^-$ $CD3^-$)	+++	+++
	Tissue PMNs ($CD45^+$ $Ly6G^+$)	+	–

[a] DC, dendritic cells; LC, Langerhans cells; PMNs, polymorphonuclear cells.
[b] Including limbal region; Data from Refs. [474,1106–1111].

key component of the human epithelial glycocalyx barrier, also contributing to the tight-junctional, paracellular barrier at the ocular surface. By contrast, knockdown of MUC1 does not lead to decreased barrier function and moreover, significantly increases the barrier to dye penetrance and bacterial invasion [149].

The soluble lectin galectin-3, the most highly expressed carbohydrate-binding protein of the human conjunctival epithelium [150], is a component of the epithelial glycocalyx. Its carbohydrate recognition domains (CRD) [151] bind to the β-galactoside-containing glycans [152,153], MUC1 and MUC16, to form a polymeric, galectin-glycoprotein lattice which serves various biological functions, such as the regulation of receptor turnover and modulation of cell-cell, cell-matrix and cell-pathogen interactions [153]. Additionally it contributes to the barrier function of the glycocalyx. Downregulation of O-glycan synthesis by human corneal epithelial cells in culture, reduces the barrier to rose bengal penetration, by decreasing galectin-3 binding within the glycocalyx [134] and a similar loss of barrier function occurs, in the absence of galectin-3 expression [134]. These findings indicate that mucin O-glycan interaction with galectin-3 creates a protective lattice barrier within the apical glycocalyx [134,154]. Galectin-3 concentration in the tears may be of future interest as a marker for DED severity since its affinity for glycans of the glycocalyx may be reduced by alterations in glycosylation, or it may be released from the ocular surface inflammatory cells [155].

4.5.2. Other mucin species

The gel-forming mucin MUC5AC is the chief secretory mucin of the human ocular surface [132,156,157] although the soluble mucin MUC7 is detected in the lacrimal gland and in the conjunctival epithelium [17,158]. In human tears, MUC1, MUC4, MUC16 and MUC5AC are present, and $MUC2^{159}$ is also detected at very low levels in the conjunctiva [157]. MUC20, which is the most highly expressed mucin in human conjunctiva [150], is localized along the cell membranes of the intermediate cell layers of the corneal and conjunctival epithelia [160]. mRNA transcripts of the transmembrane mucins MUC13, MUC15 and MUC17 have been identified in human conjunctiva [150,158,160]. Their functions here, as well as that of MUC20, have not yet been elucidated.

4.6. The tear compartments

When the eyes are open, the tears are distributed in 3 compartments. The fornical compartment, occupies the fornix and retrotarsal space and the tear menisci and tear film form the preocular tears. The fornical compartment is assumed to be narrowest at the lid wiper region of the lid margin, which is directly apposed to the globe. The preocular tear film overlies the exposed conjunctiva and cornea [161]. The precorneal tear film follows the contours of the cornea, is roughly 3 μm in thickness and is highly stable [162]. The prebulbar film follows the varying contours of the bulbar conjunctiva but its thickness is unknown.

4.6.1. The tear menisci

Tear menisci are strips of aqueous tear fluid, lying in the angle between the globe and apposed lid margins, which are formed by surface tension forces as the lids separate, in a few hundred milliseconds, in the upstroke of the blink. A negative hydrostatic pressure within the nascent menisci draws water from the forming tear film, causing the two compartments to separate over a region of meniscus-induced thinning. [163,164] This is observable as a black line of reduced fluorescence in the fluorescein-stained tear film, where the aqueous layer is of minimum thickness while the lipid layer remains intact (Fig. 1) [165]. There is no evidence for a gel-like layer in the menisci. Instillation of an aqueous drop expands the

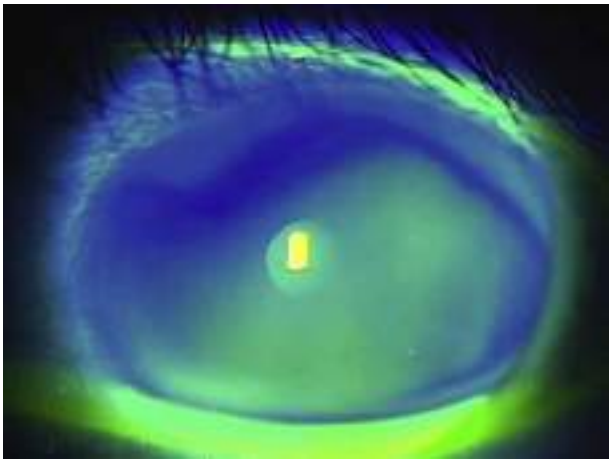

Fig. 1. A fluorescein–stained lower tear meniscus, following drop instillation, The meniscus is broad and full and is segregated from the stained precorneal tear film by a black line of meniscus-induced thinning.

volume of the meniscus and tear film and transiently obliterates the black line [166].

The negative hydrostatic pressure within the menisci is responsible for their concave external surface and opposes the outflow of aqueous into the puncta, so that drainage is limited to roughly the first 2s of the blink interval [167,168]. This effect is enhanced as meniscus volume falls and may play a conserving role in aqueous-deficient dry eye (ADDE).

4.6.1.1. Tear volume and secretion. The volume of the menisci is directly related to the total volume of the tear fluid [169] and to the lacrimal secretory rate [170]. Since the height and radius of curvature of the tear menisci are reduced in ADDE their measurement is of diagnostic value in DED diagnosis [168,171,172]. The volume of the tears has been estimated to be about 7 μl [173] and secretory rate 1.03 ± 0.39 μl/min, with a tear turnover (TTR) of 16.19 ± 5.10%/min [174]. The LG is responsible the bulk of the tear volume and flow [170] with a smaller portion secreted by the conjunctiva [81]. Lacrimal fluid is distributed to and mixed with the preocular film during the blink and is then lost by drainage from the tear menisci via the nasolacrimal system. It is further lost by evaporation from the exposed preocular tears [175–177].

4.6.2. The precorneal tear film

The precorneal tear film has a superficial lipid layer and a mucoaqueous layer which occupies the bulk of the tear thickness and interacts directly with the glycocalyx of the epithelium (see TFOS DEWS II Tear Film Subcommittee report [1223]). There is also a slender, superficial aqueous layer. The tear film is highly stable and its layers cohere during movements of the eye [178].

4.6.3. Tear film lipid layer

The tear film lipid layer (TFLL) derives from the meibum reservoir at the lid margins and is spread onto the tear film with each blink, driven by surface tension forces. It has a mean thickness of 42 nm (15–157 nm) [179]. It plays a significant role in stabilizing the tear film and has been considered until recently to provide a barrier to tear evaporation [36,180,181]. However, some previous and more recent studies have suggested that it reduces evaporation from the mucoaqueous subphase by no more than 10% [182]. This question is critical to the designation of some forms of DED as evaporative dry eye (EDE) – ie. dependent on an excessive evaporative loss from the ocular surface and is discussed further in the report of the TFOS DEWS II Tear Film Subcommittee [1223].

The meibomian glands secrete a lipid mixture (meibum) which is liquid at body temperature, with a melting range between 19.5 and 32.9 °C according to Tiffany [181], or 10–40 °C according to Butovich et al. [183] The clear oil can be expressed from the meibomian orifices by pressing over the glands through the closed lids. Expressibility is greatest nasally and least temporally [184]. Delivery of oil to the lid margin occurs in part through secretion and in part by the expression of small aliquots with each blink. The lid reservoir contains at least 30 times the amount of lipid present on the surface of the tear film (approximately 300 μg vs 10 μg, respectively [185,186].

It is likely that excretion of meibum occurs by flow of lipid from the reservoirs over the lid margin skin and lashes. This would serve to resist tear film contamination by sebaceous skin lipids (sebum).

In keeping with the earlier proposal of Holly [187] and further studies by McCulley [188], the TFLL is considered to organize itself into a layer, a few molecules thick, rich in polar lipids and some long-chain fatty acids and a superficial layer of non-polar lipids. Some proteins and glycoproteins, such as lipocalins, lysozyme, and mucin, are thought to be intercalated with the lipid layer and enhance its stability [189–191].

4.6.3.1. Spreading of the lipid layer. The tear film lipid layer is formed in the upstroke of each blink, when lipid from the lower meibomian reservoir spreads upward over the aqueous subphase of the preocular tear film [192,193]. It has been suggested that thinning of the lipid layer superiorly creates a local rise in surface tension, which is the driving force for spreading [193]. It has been proposed that spreading initially involves an interaction between the polar meibomian lipids and the aqueous phase of the tear film [190] and that the polar lipid layer then acts as a carrier for the nonpolar lipid fraction. In the normal eye, spreading of the tear film lipid layer can be observed clinically by interference video microscopy, when it is seen as an upwardly moving front of horizontally-disposed, colored fringes. The lipid film spreads rapidly at first (about 10 mm/s), lagging markedly behind the upper lid, whose excursion is completed over a few hundred milliseconds [194]. Spreading slows and stabilizes after 1 s or more, with the interference pattern showing remarkable stability over the remainder of the blink interval [194,195].

Spreading of the TFLL is slower in patients with a tear film lipid deficiency (Goto 2003) and also in aqueous tear deficiency [194], which is attributed to the thinness of the aqueous phase. Also, in the former condition the spreading TFLL pattern has been reported to take on a more vertical arrangement [195].

The pattern of colored fringes shown by interferometry is due to topographic variations in thickness of the lipid layer across the film and reflects its intermolecular organization. This can be shown to be remarkably stable during blinking and eye movements. Over a series of blinks the pattern can retain its gross features from blink to blink, only degrading by degrees in a stepwise fashion until it changes abruptly and the process begins again [197]. In this situation it appears that the TFLL is stripped from the mucoaqueous layer and compressed during the downstroke of the blink and restored in the upstroke, with only a moderate disturbance of its intermolecular organization between consecutive blinks. The period over which this can be observed may be greatly shortened in patients with a tear lipid layer deficiency suggesting that intermolecular stability is lost [198]. This is the basis of a clinical test [199].

Similarly, the interference pattern shows great stability during a series of horizontal saccades, again showing only a moderate stepwise degradation over a series of saccades. In this case the TFLL and mucoaqueous subphase behave as a fluid shell that moves with the cornea during each saccade [197]. The influence of a DED state on this behavior would merit study.

4.6.4. The aqueous layer and mucoaqueous subphase

Deep to the TFLL is a mucin-rich layer that is conveniently referred to as the mucoaqueous subphase [200]. The presence of a superficial aqueous layer at its surface, as proposed by Wolff has been debated [201], but it is reasonable to assume that in the process of tear film formation, as aqueous tears are drawn off into the menisci, some residual fluid is retained at the surface of the mucoaqueous layer. (See TFOS DEWS II Tear Film Subcommittee report [1223]) This fluid layer can be expanded transiently by the instillation of a saline drop [166].

Observation of the fluorescein-stained tear film indicates that the mucoaqueous layer of the precorneal film is freshly deposited with each blink and has the physical properties of a gel, due to the presence of the goblet cell mucin [197]. Its mucin component is presumed to be a product chiefly of the tarsal goblet cells, whereas that of the *prebulbar film* is likely to be an admixture of mucins from both tarsal and bulbar glands. An additional coating is received by the peripheral cornea as its passes behind the lids during eye movements in any direction of gaze [197].

The mucoaqueous subphase performs a lubricating function

between the lids and globe [99] and probably maintains wettability of the ocular surface where the glycocalyx is defective, for instance after an abrasion [201]. It also traps shed epithelial cells, inflammatory cells, debris and microorganisms, which are collected into a mucous thread in the lower conjunctival sac and ultimately lost via the punctum [202,203].

The mucoaqueous layer contains salts and numerous proteins derived from the LG, conjunctiva and meibomian gland. Proteins include growth factors such as epidermal growth factor (EGF) and hepatocyte growth factor (HGF), that are essentials for the maintenance of the epithelium [204,205]. There are also defense proteins, such as lysozyme, lactoferrin, surfactant protein-D, and trefoil peptide, concerned with innate immunity, and sIgA [205,206]. Those proteins of lacrimal origin, such as lysozyme and lactoferrin, are decreased in ADDE, making the eye more vulnerable to infection. It has been predicted that the level of these proteins will be normal in EDE where lacrimal function is normal and it would be of value to test this prediction [207].

Plasma proteins, such as albumin, may leak into the tears in DED as a result of inflammation, due to an increase in conjunctival vascular capillary permeability [205,208,209] and, probably, also conjunctival epithelial permeability. The LG cannot be excluded as an additional source.

4.7. Closed eye tears

Overnight eye closure causes a number of physiological changes at the ocular surface. The pO_2 falls and there is a shift towards anaerobic tissue metabolism [210,211]. The tear pH and tear osmolarity fall [212,213], the anterior cornea becomes relatively hypoxic, epithelial permeability increases and corneal edema occurs [214,215]. There is no change in tear glucose levels [216].

Jordan and Baum proposed that, in the waking state, tear secretion is driven in part, by sensory stimuli from the ocular surface, with the expectation that it will be at it lowest when ambient stress is at a minimum [217]. This was born out by the studies of Sack and colleagues, who demonstrated that lacrimal secretion was negligible after an extended period of sleep or eye closure, a change accompanied by a sharp rise in the level of tear sIgA, from around 2% in reflex tear samples, compared to 58% in closed eye tears [218]. Conversely, the levels of lysozyme, lactoferrin and lipocalin, proteins of lacrimal origin which account for about 85–88% of the total protein in basal and reflex tear samples, decrease in closed eye tears to less than 30% of the total [218]. The rise in sIgA concentration may reflect that, unlike the lacrimal-specific proteins, lysozyme, lactoferrin, lipocalin and peroxidase [219], the secretion of sIgA, of plasma cell origin, is not directly coupled to lacrimal secretion. Therefore the rise could be explained by the continued delivery of IgA at the same rate into a lacrimal fluid of greatly reduced volume. This fall in volume secretion also explains, in part, the rise in tear concentration of certain plasma proteins such as vitronectin, fibronectin, α1-antiprotease, α2-antiplasmin, α1-antichymotrypsin and IgG [220,221], which enter the tears by diffusion across conjunctival capillary and epithelial barriers. Such proteins are present in closed eye tears at 2–4% of the serum level, well above the level found in reflex tears. Sack also refers to an increase in vascular permeability in the closed eye state [209].

A striking feature of closed eye tears is a massive accumulation of activated PMNs within the tear fluid several hours after eye closure [218]. Their appearance is preceded by 1–2 h [222], by very high levels of two potent leukotaxic mediators, IL-8 and LTB4. Up to 70% of this leukotaxic activity is removed by immunoprecipitation with antibodies to IL-8, indicating that it is not substantially of PMN origin. Degranulation of the PMNs releases several potent proteases, such as protease-3, elastase, capthepsin G, MMP-9 and urokinase, which, due to the simultaneous presence of a wide range of antiproteases, do not lead to autolytic digestion. Also, despite the presence of the potent angiogenic agent, 12 (R)- hydroxyeicosatrienoic acid [223], and of IL-8, which might stimulate corneal neovascularization, a build-up of α 2-macroglobulin (α2-M) and the conversion of plasminogen to angiostatin, appears to prevent this outcome.

Closed eye tears are also extremely rich in reactive complement products, normally absent [224] from open eye tears. Closed eye tears contain all of the complement components required for the classical and alternate pathways of complement activation, in a concentration of about 2–4% of that of serum. Factors B and C3, however, reach levels approaching one-third of that in serum, suggesting a local source [224]. A significant proportion of the C3 in closed eye tears is converted to C3c and Sack et al. inferred that, since closed eye tears contain two inhibitors of complement conversion (lactoferrin and sIgA), C3 conversion probably occurs through the alternative pathway or by plasmin cleavage [218]. It was also proposed that modulators of complement activation divert the complement system away from membrane-attack complex formation towards opsonization.

In summary, powerful defense and scavenging mechanisms come into play during prolonged eye closure, which serve to remove microbial threats to the ocular surface. These events are highly regulated so that no harm comes to the ocular surface itself. However, this is a potentially risky strategy that could be destabilized in the DED state and the Subcommittee recommends the investigation of closed eye tears and of conjunctival impression cytology specimens following prolonged eye closure, in DED patients. There will be added interest to study patients with Sjögren syndrome, since a genetically-determined, dysfunctional response to inflammatory triggers might create a dysfunctional closed eye tear response.

4.8. Extracellular DNA and NETs in dry eye

A new mechanism leading to tissue damage in DED has been identified since the TFOS DEWS report [1], involving the release of DNA into the tears from desquamating ocular surface epithelial cells and invading neutrophils. This extracellular DNA (eDNA) can, on its own, or combined with molecular components of neutrophil origin, cause direct damage to the ocular surface.

4.8.1. Extracellular DNA of epithelial origin

A source of eDNA is from desquamated ocular surface epithelial cells of conjunctival [114,225], and presumably corneal, origin. Extracellular DNA, facilitated by cathelicidin binding [226], is able to enter cells and stimulate an inflammatory signaling pathway [227], by binding to TLR9 within the cell and initiating a signaling cascade through MyD88. This has two consequences: i. the initiation of a type 1 IFN response [228], and ii the generation of a powerful neutrophil recruitment signal [229–231]. In support of this concept, topical application of a synthetic bacterial DNA mimic to injured corneal epithelium results in the recruitment of neutrophils to the corneas of wild-type, but not TLR9 $^{-/-}$ mice [232]. Increased expression of mRNA for TLR9, MyD88, and interferon (IFN)-type I pathway genes has been found in the exfoliated conjunctival cells from patients with severe ADDE [114,225] and patients with Sjögren syndrome [233], suggesting that epithelial cells contribute to the inflammatory response directly, and also engage in PMN recruitment. Type 1 IFNs (IFN-α/β) augment dendritic cell maturation and activate the adaptive immune system. Increased expression of mRNA for IL-6 and TNF-α was also demonstrated in exfoliated conjunctival cells. The expression of these inflammatory cytokines increases in the corneal and

conjunctival epithelium in some forms of experimental DED [234,235].

4.8.2. Extracellular DNA of neutrophil origin

Neutrophils are key players in the host innate immune response and constitute a first line of defense. While they are present only in small numbers in the normal conjunctiva [114] they are recruited to the ocular surface in profusion in inflammation and are abundant on the ocular surface and in the tears of patients with severe ADDE [114,225].

One strategy adopted by neutrophils, in their defense against microorganisms, is to release cellular contents into the extracellular space to form Neutrophil Extracellular Traps or NETs [236]. These comprise extracellular webs or scaffolds containing decondensed chromatin, histones, neutrophil elastase and antimicrobial peptides such as cathelicidin, each of which individually may be toxic for epithelial cells [237]. Extracellular histones are major mediators of cell death in sepsis [238], cathelicidin fragments are considered to cause erythema, inflammation and telangiectasia in patients with rosacea [239], and neutrophil elastase induces epithelial cell apoptosis [240]. NETs, with all their molecular components, have been demonstrated in mucoid films at the ocular surface in DED [114,225] (Fig. 2). It has been suggested that their association with mucin relates to the action of neutrophil elastase in cleaving the extracellular domains of membrance-associated mucins [241]. In other studies, it has been shown that mucins may induce neutrophil activation [242].

In the healthy eye, NETs may play a physiological role in the defense against pathogens by means of an antimicrobial action and by confining pathogens to a local site of infection [243]. Additionally, immobilization of neutrophil granules within NETs may prevent the diffusion of potentially noxious proteins and proteases to the ocular surface. However, in patients with severe ADDE, eDNA and NETs are present at the ocular surface in excessive amounts [225] and evidence suggests that they participate in the pathogenesis of the disease [225]. There are two explanations for their high levels:

i. Exposure of neutrophils to hyperosmolar stress is a stimulus for NET formation, and the quantitative release of NETs increases exponentially with increase in hyperosmolarity. This is relevant to the situation in severe DED, where, for reasons discussed elsewhere, high levels of osmolarity may be achieved in the tears [244]. Hyperosmolar stress also has an inhibitory effect on some critical neutrophil functions such as migration and degranulation. Thus, in a hyperosmolar milieu, the classical neutrophil-related innate defense mechanisms may be compromised.
ii. In physiological conditions, the level of eDNA and NETs in the tears is regulated by tear nucleases of lacrimal origin, DNase I, and lipocalin, (an endonuclease with a lower level of activity). Nucleases hydrolyze eDNA and enable its clearance from the ocular surface. The concentration of DNase I in tear fluid is similar to that in serum and saliva. Importantly, tear fluid nuclease activity was shown to be low or absent in ADDE patients [114,225], providing an additional basis for the rise in eDNA and NETs in the tears in the DED state, including Sjögren syndrome DED (SSDE), non-Sjögren syndrome DED (NSDE) and graft-versus-host disease (GVHD) [245].

Thus it appears that in DED, NET production is stimulated by tear hyperosmolarity and clearance of both eDNA and NETs is impaired by the tear nuclease deficiency. Both may participate in further neutrophil recruitment [225,244]. The use of topical DNase I therapy for DED is being explored [245].

The Subcommittee recommends that this be considered as an important area for future research, exploring its involvement in lesser degrees of DED and also any interaction with the PMN response in closed eye tears.

4.9. Homeostasis of the tears at the ocular surface

4.9.1. The lacrimal functional unit (LFU)

The production of aqueous tears is regulated to maintain tear osmolarity within narrow limits at all times [246]. Tear homeostasis is achieved reflexly by the lacrimal functional unit (LFU), which consists of the ocular surface, its secretory appendages and the connecting innervation, (Fig. 3) [247]. The trigeminal innervation of the ocular surface epithelia, including the cornea, conjunctiva and lid margins, provides the afferent limb of the feedback loop. The parasympathetic, secretomotor innervation of the ocular appendages, including the lacrimal gland (main, palpebral and accessory), meibomian glands and the conjunctival goblet cells, provides the efferent limb of this loop. The

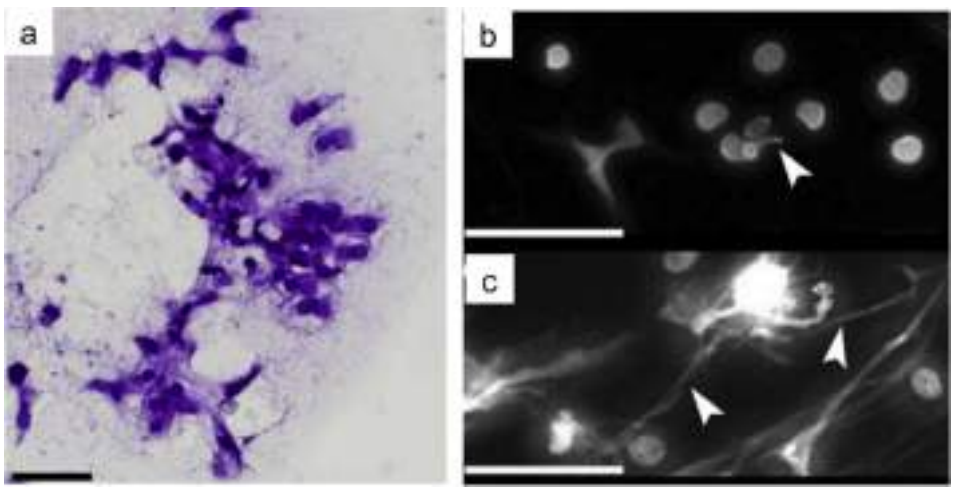

Fig. 2. a. H&E staining of exfoliated surface cells. b. Wide-field fluorescent microscope image after DAPI staining of conjunctival impression material reveals short, sparse eDNA strands (arrowhead) in normal subjects and c. numerous long eDNA strands in a DED patients (arrowheads). (from Sonawane, S., et al. (2012). "Ocular surface extracellular DNA and nuclease activity imbalance: a new paradigm for inflammation in DED." Invest Ophthalmol Vis Sci **53**(13): 8253–8263. – with permission) [225].

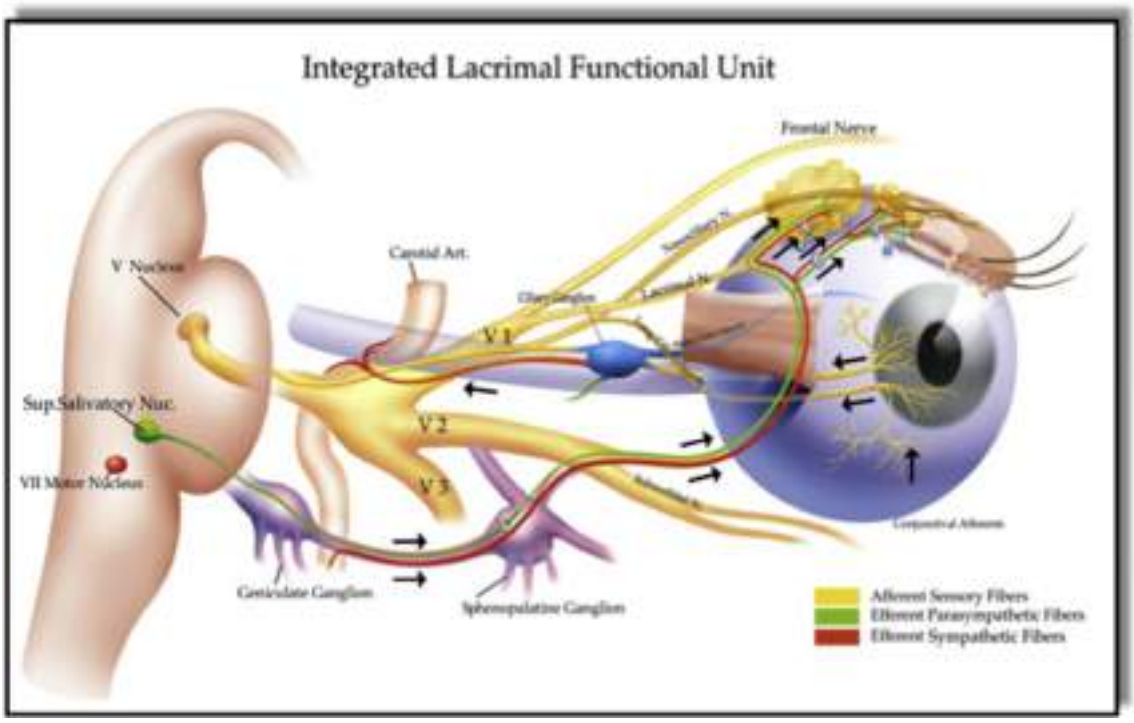

Fig. 3. Representation of the Lacrimal Functional Unit. In the waking state aqueous tear flow is modulated by reflex impulses from the ocular surface and nasal passages which travel in the trigeminal nerve to synapse in the superior salivatory nucleus. (from Dry Eye and Ocular Surface Disorders, Pflugfelder, Beuerman, Stern, 2004 – with permission) [1102].

nasolacrimal passage is also considered to contribute to this reflex system [248]. Another reflex arc that serves to protect the ocular surface is that subserving the blink.

4.9.2. The secretory reflex arc

The afferent limb of the reflex arc arises in the trigeminal nerve, whose central endings synapse with neurons in the superior salivatory nucleus in the brain stem, probably located caudal to the nucleus of the seventh cranial nerve [77]. In the rabbit, the sensory innervation of the central cornea is about 10–20 times that of tooth pulp, while that of the conjunctiva generally is of a lower degree [249]. However, the sensitivity of the posterior lid margin is similar to that of the central cornea [250], which is of relevance to the symptoms of blepharitis.

The efferent limb of the reflex arc is a parasympathetic pathway whose secretomotor, preganglionic fibres arise in the superior salivatory nucleus. These fibres exit the pons by the nervus intermedius of the seventh cranial nerve and reach the pterygopalatine ganglion via the nerve of the pterygoid canal. Here, they relay and the postganglionic fibres reach the lacrimal gland via the lacrimal nerve. An alternative postganglionic pathway has been described, reaching the gland via the retro-orbital nerve plexus [251].

The nature of the relay between the afferent and efferent fibres, in the superior salivatory nucleus, the involvement of interneurons and the interaction with other inputs and supranuclear pathways, is not known, nor is the level of central cross-connectivity between ipsilateral inputs and contralateral outputs fully established. Current studies have not excluded their existence [252]. This contrasts with observations concerning the drive to lacrimal secretion from the nasolacrimal mucosa where cross-connectivity has been demonstrated [26] – ipsilateral anesthesia of the nasal mucosa reduces lacrimal secretion on both sides.

4.9.3. Afferent inputs from the ocular surface

4.9.3.1. Lacrimal secretion and the blink. The trigeminal afferents from the cornea serve a range of sensory modalities which include pain, mechanoreception and temperature and full details are given in the TFOS DEWS II Pain and Sensation report [1224]. Here, it may be noted that sensory inputs from the ocular surface regulate tear production and the blink response and are the basis of sensations of discomfort in DED.

4.9.3.2. Sensory drive to lacrimal secretion. Evidence suggests that in everyday conditions, lacrimal secretion is driven by sensory impulses from corneal, cold-modality, thermoreceptors. Additionally, it appears that, in DED, surface desiccation, stimulating this subset of receptors, in response to hyperosmolarity and surface cooling, determines the compensatory increase in lacrimal secretion, increase in blink rate and sensation of awareness of the eye, rising to the level of discomfort. This compensatory response to drying, occurring in MGD-related DED, where the lacrimal gland is healthy, may explain why some patients with DED experience epiphora and appear to have a “wet dry eye” [253].

Bilateral topical anesthesia causes a reduction in reflex tear secretion of up to two-thirds [217], providing a value which is sometimes referred to as ‘basal tear secretion’. This is a reasonable term as long as it is recognised that it refers to a measurement made in particular environmental conditions and does not exclude inputs to lacrimal secretion from non-ocular sources. Jordan and Baum [217] proposed that the lacrimal secretory rate was adjusted in response to environmental conditions and, as mentioned, tear production is at its lowest after an extended period of eye closure, as in overnight sleep [218]. Also, as Cross and Krupin observed in subjects with normal eyes, basal lacrimal secretion measured after topical anesthesia, (average Schirmer: 12.8 mm) falls markedly after 1 h of general anesthesia (to 1.2 mm), suggesting suppression of

an input from higher central nervous centres [254]. Heigle et al. concluded, 'Perhaps lacrimal gland stimulation results from the sum of sensory inputs from the ipisilateral adnexal skin, cornea, nasal mucosa, contralateral eye and even central stimulation' [255].

There are other sensory inputs to the outflow pathway, from the nasal mucosa, retina and skin, which arise from pain and other noxious stimuli, such as intense cold or bright light, whose quantitative nature is unknown. When unilateral stimulation of the nasal passage on one side results in an increase in Schirmer wetting of both anesthetized eyes, (the nasolacrimal reflex) this cannot be taken as evidence for a reflex response or of cross-connections between trigeminal inputs from the nasal cavity to the superior salivatory nucleus. They could reflect a response of higher centres to the painful stimulus. Tearing in response to painful injury or retinal stimulation with bright lights could have a similar basis. Emotional tearing is under the control of higher centres [256] and there is a hypothalamic influence on autonomic centres in the brain stem [257]. In steady state conditions, most of the tear volume is derived from the lacrimal gland and its osmolarity therefore reflects that of the lacrimal secretion, modified by exposure to the environment when the eyes are open. A longer blink interval is predicted to result in a greater rise in the osmolarity of the preocular tear film and meniscus than a shorter blink interval.

4.9.3.3. Sensory drive to blinking. Spontaneous blinks are thought to arise through the activity of a brainstem 'blink generator', modified by reflex inputs from the ocular surface and inputs from higher centres. Details of the blink generator are not fully known but it may lie in the pontomedullary reticular formation and the medullary reticular nucleus which serve both the facial nucleus and third nerve nuclei. Blink rate falls after bilateral, topical, ocular anesthesia [258], and also following LASIK surgery [259].

4.9.3.4. The blink cycle and tear dynamics. The tear film is regularly refreshed by spontaneous blinking, [258,260] whose rate is adapted to environmental conditions and varies with personal behavior. Blinking plays a key role in tear dynamics by spreading, mixing, and distributing the tears and clearing cellular and other debris. The blink cycle consists of the blink itself (around 200–300 ms) and the blink interval, during which evaporative water loss occurs [261]. The blink rate is expressed in blinks per minute.

4.9.3.5. The blink rate. Wide variations in blink rate have been reported in normal adults, probably reflecting individual variation and the influence of environmental and experimental conditions. It is strongly influenced by mental state, attention, physical activity, eye exposure, and environment. Factors in the environment that are important are relative humidity, temperature and airflow over the eye. Blink rate is increased by low humidity, cold and high wind speeds.

In standard room conditions (eg. 22 °C with humidity of 40.0%), blink rate in normal adults ranges between 15 and 20 per min [261–263]. Blink rate increases in DED, where it is thought to play a compensatory role in refreshing the tear film more frequently [264,265]. Blink rate falls during a number of common visual tasks requiring mental concentration, and it is considered that the increased evaporative loss may act as a trigger for DED [261].

4.10. Optical performance of the tear film

Wavefront aberrometry studies show that, in healthy eyes, optical quality of the tear film decreases steadily during the blink interval. The period over which this occurs is shorter in DED, with the aberration minimum just preceding the breakup of the tear film [266].

4.11. Tear osmolarity

4.11.1. Introduction

Tear film osmolarity is a central factor in the pathogenesis of both ADDE and EDE. Tear hyperosmolority resulting from decreased lacrimal flow or tear film breakup contribute to ocular surface damage both directly, and indirectly, through a cascade of inflammatory events. This hyperosmolar inflammatory environment favors corneal and conjunctival epithelial, and goblet cell apoptosis which further contributes to tear film instability. Inflammation induced by tear film instability and hyperosmolarity also contributes to neurogenic chronic inflammation and increased disease severity [267,268].

In subjects with normal eyes, in standard conditions, tear osmolarity, measured in lower meniscus samples, lies within narrow limits and is remarkably stable in healthy eyes [269]. Evaporation during the blink interval causes a measureable thinning of the tear film, and a consequent rise in tear film osmolarity is predicted [177]. Tomlinson reported a value of 302 ± 9.7 mOsm/L based on data from several studies [270] and importantly, variation between right and left eyes is small (6.9 ± 5.9 mOsm/L) [271]. The narrow range of values in individuals reflects the influence of homeostatic mechanisms, with the blink interval, as the chief modifier of evaporation, likely determining the set point of tear osmolarity between the two eyes [79].

Mathematical modeling suggests that there is a small osmolar differential between the tears and menisci so that in the steady state, the osmolarity of the tear film is higher than that of the menisci [176]. This may relate to the ratio of the tear film thickness to its surface area, compared with that of the menisci, and to tear mixing and flow in the menisci in the early phase of the interblink interval [272]. Modeling considerations also suggest that in DED, this differential is greater. Thus a tear sample taken from the meniscus may under-estimate that of the tears over the surface of the eye and hence of the underlying tissues [176].

While the highest values for tear meniscus osmolarity measured in DED are below 500 mOsm/l, it is likely that levels achieved at the ocular surface are much higher than this, particularly at the site of tear film breakup. Begley and colleagues studied the relationship between tear film breakup and DED and have suggested that local fluctuations in the tear film thickness will induce hyperosmolarity "hot-spots" with significantly higher concentrations than the average tear value [273–275]. Liu et al. [276], compared the character and intensity of symptoms associated with tear film breakup with those induced by instilled hyperosmolar solutions. These studies indicated a threshold of 450 mOsm/l for the induction of symptoms, with a value of 800–900 mOsm/l required to mimic symptoms induced by tear film breakup, that is, far higher than that detected in the meniscus in DED patients. Recent mathematical modeling also predicts major spikes in osmolarity within regions of tear breakup [277–279].

Tear film thickness was studied by using the self-quenching of fluorescein (FL), the reduction of fluorescent efficiency with increasing concentration shown at high concentrations [280]. Close match between FL imaging and a mathematical model incorporating evaporation and osmosis predicted the osmolarity of the uniformly thinning tear film to be as high as 3000 mOsM. The peak values of osmolarity varied depending on the evaporation rate applied in the model. The mathematical model simulated osmolarity within and around areas of tear breakup yielding a peak osmolarity value of approximately 1900 mOsM, close to modeling results of Peng et al. [279] (Fig. 4). These local spikes of hyperosmolarity within areas of tear breakup are considered to be a major source of repeated stress to the ocular surface.

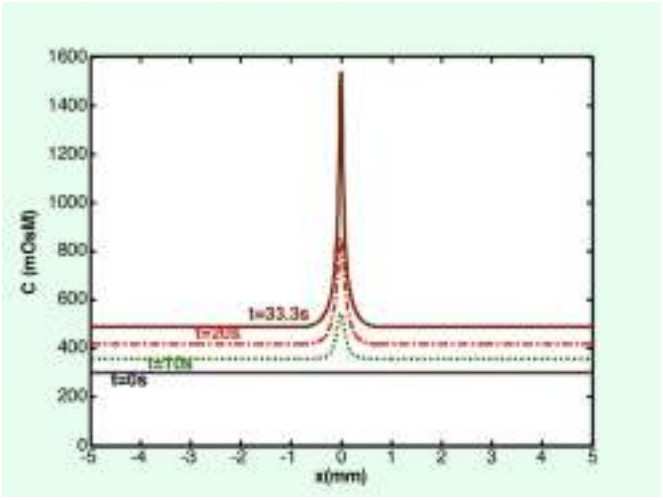

Fig. 4. Predicted evolution of a spike of hyperosmolarity during an extended blink interval based on modeling considerations. Surface osmolarity increases from 300 mOsM to 545 and 850 after 10 and 20 s, respectively, and skyrockets to 1534 mOsM after a 33-s interblink corresponding to tear-film breakup. (from Peng, C. C., et al. (2014). "Evaporation-driven instability of the precorneal tear film." Advances in colloid and interface science **206**: 250–264. – with permission) [279].

4.11.2. Tear osmolarity in dry eye

Tear osmolarity threshold values that discriminate a healthy eye from an eye with DED varies in the literature from 308 mOsm/L to 316 mOsm/L [269,270]. One reported reason for variability in tear osmolarity threshold values is tear film instability, a characteristic of the disease. Normal, mild/moderate, and severe dry eyes have average tear osmolarity values of approximately 302 ± 8 mOsm/L, 315 ± 10 mOsm/L and 336 ± 022 mOsm/L, respectively [281]. Currently, 308 mOsm/L is proposed as a sensitive threshold for discriminating between normal eyes and those presenting with early stages of DED. Conversely, the 316 mOsm/L threshold would better discriminate between mild and moderate/severe DED. In addition to the absolute level of tear osmolarity, variability over time and, particularly, variability between the two eyes, can be a diagnostic indicator and appear to increase with the severity of DED [269,282].

4.11.3. Factors influencing tear osmolarity

Tear osmolarity is influenced by the following intrinsic and extrinsic factors: i. Body hydration, ii. Tear film lipid layer (TFLL) characteristics, iii Palpebral aperture width, iv. Blink interval, v. Tear film stability and, vi. Environmental conditions.

4.11.4. Body hydration

The tears on waking are slightly hypotonic, their tonicity rising over the course of the day, due to evaporation from the tear film. There is a positive relationship between whole body hydration, measured as plasma osmolarity and tear osmolarity and both are raised in patients with DED. Also, tear osmolarity follows plasma osmolarity in subjects with imposed systemic dehydration [283–285]. Consequently measurement of tear osmolarity has been proposed as a possible surrogate for plasma osmolarity, of potential use in the rapid detection of dehydration in the elderly or in sports medicine [283].

4.11.5. The tear film lipid layer

The rate of water loss from the eye is influenced by the quality and thickness of the TFLL. Expression of meibum in normal eyes leads to thickening of the tear film lipid layer [286] and to a reduction of evaporation in both healthy individuals and patients with DED [287]. When the quality or integrity of the TFLL is deficient, as judged by interferometry, evaporative loss may be increased and tear osmolarity increased [175]. A similar outcome may be predicted when TFLL spreading is retarded by a marked aqueous tear deficiency [194].

4.11.6. Palpebral aperture width

As would be anticipated, evaporative loss from the eye is influenced by tear film area. Tsubota and Nakamori examined the effect of gaze position on evaporation rate (at 40% humidity and a blink rate of 30 per min) and showed that evaporative loss is 3.4 and 2.5 times greater when looking up and straight ahead than when looking down, not only per eye, but also per unit area of the ocular surface [288], perhaps suggesting that as the area to be covered increases, the TFLL is thinned.

4.11.7. Blink interval

The tear film is refreshed by blinking [258] and the blink rate is adapted to environmental and social circumstance and to personal behavior. The blink interval, and hence the blink rate, is a determinant of tear osmolarity, with the expectation that prolongation of the interval (a slower blink rate) will raise it. Blink behavior may be constrained while performing selected visual tasks in ways that influence tear stability and evaporative loss. A fall in blink rate has been documented during everyday visual tasks, such as working at a video display terminal, reading in downgaze [289], operating monitor-based and handheld video games, and performing surgery [290,291]. In these situations, both gaze position and difficulty of the visual task are determinants of blink rate.

The effect of a fall in blink rate on evaporative stress while performing downgaze tasks is difficult to predict on basic principles. Both blink rate and palpebral aperture area are decreased, the former tending to increase and the latter to decrease tear evaporation. Also, when viewing computers with the eyes in the primary position, the head may tilt backwards, thus narrowing the palpebral aperture.

4.11.8. Tear film breakup

The importance of a stable tear film for retinal imaging is well known [292], and many approaches are used to study its influence on visual function. Tear film breakup within the blink interval is a cause of visual degradation and its character and time course have been studied in detail in contact lens wearers [293]. The effect of precorneal tear film breakup on vision is due to variations in film thickness, rupture of the film and, in DED, exposed epithelial irregularities at the site of breakup and the presence of light-scattering, epithelial opacities.

Although visual acuity is the standard clinical measure of visual function, it does not provide a full account of visual performance and broader measures of visual function are used, such as contrast sensitivity [292], glare disability [294], and scatter index [295], all of which have been shown to be disturbed in DED [296]. A functional measure of visual acuity has also been developed [297,298].

Tear film breakup time is the most frequently used measure of tear film stability and becomes of pathological importance when it falls below the interblink interval. In most healthy individuals, the tear film is extremely stable and values reported for TBUT are well beyond the normal blink interval [299]. However, tear breakup in the blink interval does occur in some healthy individuals.

The relationship between the blink interval and breakup time can be captured as the Ocular Protection Index (OPI), the breakup time divided by the blink interval [300]. An OPI of $\geq$1, indicates that the breakup time exceeds the blink interval and therefore that the eye is protected from desiccation throughout the blink cycle. An OPI

of <1 indicates that breakup is occurring within the interblink interval and that the eye is exposed to damaging desiccation. In early DED, OPI is initially >1 and nears 1 as disease severity increases, independent of the cause of the DED. Later, as the disease progresses and the OPI falls below 1, hyperosmolarity is amplified locally in the epithelium subjacent to the breakup by the local increase in evaporation. For a given blink interval, the lower the OPI, the greater the imposition of evaporative hyperosmolarity at the ocular surface. In the regions outside the area of breakup, osmolarity is also increased, by diffusion and tear mixing, but to a more modest level.

It is evident that the measurement of tear osmolarity in tear meniscus samples underestimates the level of hyperosmolar stress delivered to the ocular surface in an individual dry eye, and the Subcommittee identified a need to develop techniques for the measurement of osmolarity across the ocular surface, at the tissue level. Some success has been reported in the mouse, measuring surface cation levels by fluorescence ratio imaging [301] and attempts have been made in the clinical situation, measuring tear [302] and tissue conductivity [303], but currently no clinical instrument is available.

Local tear instability, initiated by a loss of ocular surface wettability, such as occurs in xerophthalmia and chronic topical preservative use, can be an independent starting point for tear hyperosmolarity and DED, acting through the mechanism described above. The resulting DED was earlier referred to as an "extrinsic" form of EDE, but a better term is ocular surface-related EDE.

4.11.9. Effect of ambient environment

Certain environmental conditions increase evaporative loss and are risk factors for DED. Evaporation is increased in conditions of low humidity and increased airflow over the surface of the eye [261,304,305]. Such conditions may be combined and may also occur in natural, outdoor conditions. The effect of the environment on evaporation is the basis for providing goggles or water-conserving spectacles for the prevention or treatment of DED states. Exposure to low humidity environment for as little as 90 min has been shown to increase blink rate, ocular discomfort and the presence of cytokines and matrix metalloproteinases (MMPs) in tears [264,306].

4.12. Corneal epithelial barrier disruption

4.12.1. Matrix metalloproteinases and EMMPRIN

Disruption of the epithelial barrier at the ocular surface is a characteristic feature of DED. Exposure of the corneal epithelium to increased osmolarity promotes inflammation, abnormal differentiation, programmed cell death (e.g. apoptosis) and accelerated desquamation [307], with early activation of mitogen activated protein kinase (MAPK) and nuclear factor $_{K}$B (NFκB) stress signaling pathways [308,309]. These pathways initiate a cascade of events, including transcriptional activation of genes encoding inflammatory matrix metalloproteinases (MMPs) (particularly MMP-9) and pro-apoptotic factors [310–312].

MMPs are proteolytic enzymes, involved in wound healing and inflammation, which play a key role in DED pathogenesis by disrupting intercellular epithelial tight junctions, leading to a breakdown of the epithelial barrier. Expression and production of MMPs −1, −3, −9, and −13 by human corneal epithelial cells correlates positively with increasing osmolarity [310–312], acting, at least in part, through the c-Jun N-terminal kinase (JNK) pathway [308]. This activity is inhibited by doxycycline [313]. Among these proteases, MMP-9 is considered to be of central importance in the response to hyperosmolar stress [311,314]. Occludin, a component of the tight junction is a known substrate of this protease and, in a murine model of DED, increased tear levels of MMP-9 were associated with a loss of epithelial barrier function and surface epithelial regularity [314,315]. Increased MMP-9 levels were also observed in the tear fluid of patients with DED, with the concentration of MMP-9 in tears correlating with DED severity. Therefore its quantification has been proposed as a biological marker of disease activity [316–318]. It is relevant that MMP-9 knockout mice exposed to desiccating stress are more resistant to alterations of the corneal epithelial barrier than wild-type animals [315].

The membrane-spanning molecule EMMPRIN (Extracellular MMP inducer; also termed $CD^{+}147$) is an inducer of MMP expression that participates in the pathogenesis of DED through MMP-mediated cleavage of occludin [314]. The molecule is also involved in the pathogenesis of corneal ulceration, stromal melting and stromal remodeling [314,319]. EMMPRIN expression is increased at the ocular surface in DED patients and correlates with MMP-9 levels in tears and in corneal epithelial cell cultures [314]. Increased osmolarity or addition of recombinant EMMPRIN in corneal epithelial cell conditioned medium was responsible for increasing production of both EMMPRIN and MMP-9 and resulted in the disruption of epithelial junctions through the cleavage of occludin. Conversely, selective inhibition of EMMPRIN by siRNA in this system, leads to the inhibition of both MMP-9 induction and epithelial barrier disruption. Moreover, an inverse relationship between the distribution of occludin and EMMPRIN as a function of differentiation and stratification of epithelial cells, both in culture, and in the stratified corneal epithelium *in vivo*, suggests a physiological function for this molecule in epithelial cell barrier homeostasis. Interestingly, unpreserved artificial tears have been shown to decrease EMMPRIN cell surface expression in patients with DED, and addition of cyclosporine to corneal epithelial cell conditioned medium, inhibited cell surface expression of EMMPRIN selectively, in an *in vitro* model of ocular surface preservative toxicity [314,320].

Galectin-3 is required to maintain the barrier function of epithelial glycocalyx [321]. Its levels are increased in the tears of DED patients, associated with increased levels of MMP-9 [155]. In an *in vitro* corneal epithelial cell system, the cell–cell detachment and redistribution of occludin induced by exogenous galectin-3 is considered to involve the induction of MMP-9, a process dependent on the clustering and interaction of galectin-3 with EMMPRIN on the cell surface [321].

4.13. Frictional events at the surface of the eye

Friction between the lids and globe as they move in relation to each other during blinking and eye movements is considered to be a cause of symptoms in DED and its sources have been reviewed by Pult [322]. Frictional symptoms will occur only at times of such relative movement between the lids and the globe.

When two apposed surfaces are in relative motion, the degree of friction between them depends on the nature of the surfaces, the speed of movement, the load applied and the presence of lubrication. Where the surfaces are separated by a fluid layer, lubrication is referred to as hydrodynamic, whereas where they are in direct contact, this is termed boundary lubrication [101,323–325] An intermediate, mixed state also exists [326]. Suitable lubrication can diminish the degree of damage or wear engendered by frictional forces.

4.13.1. Boundary lubrication

Boundary lubrication usually applies when the relative motion between apposed surfaces is slow, which, for the ocular surface, is during the interblink interval when the eyes are stationary, or directed at slow-moving objects. It probably also occurs at the

beginning, end and return points of the blink cycle [325]. At this time the epithelial glycocalyces of the apposed tarsus and globe are in contact, with a variable amount of the mucoaqueous phase intervening. In these circumstances, the cross-linked mucin exodomains of the healthy glycocalyx act as hydrophilic polymer brushes, which greatly lower the coefficient of friction between the apposed surfaces of lid and globe and minimise frictional damage [327,328].

4.13.1.1. Lubricin

Lubricin, or proteoglycan 4, is an amphiphilic glycoprotein expressed by synovial and cartilage cells of the joints, as well as in the major viscera and in muscle, tendon and bone and the eye and brain [82]. In the joints, cooperating with hyaluronic acid, it acts as an efficient boundary lubricant, reducing friction between the apposed joint surfaces [82,329–331]. In other tissues it may serve other physiological functions involving cell proliferation and attachment and matrix binding.

In the eye, lubricin is expressed by trabecular meshwork cells and by the corneal and conjunctival epithelium. Lubricin messenger RNA is also present in lacrimal and meibomian glands [82,330]. Laboratory studies suggest that lubricin may function as a boundary lubricant between the apposed surface of the cornea and lid wiper region [82]. Absence of lubricin in PRG4 knockout (KO) mice is associated with a significant increase in corneal fluorescein staining. Recombinant lubricin has been synthesized successfully [332] and has recently been tested in a clinical trial for the treatment of DED [333].

4.13.2. Hydrodynamic lubrication

Hydrodynamic lubrication applies in conditions of high relative velocity where a fluid layer separates the apposed surfaces. For the tarsus and globe, this occurs during the blink and during saccades. During the downphase of the blink, the upper lid moves chiefly in a vertical direction but also slightly nasally, across the exposed globe with an average velocity of 17–28 cm s^{-1} and a maximum velocity of around 40 cm s^{-1} [322,334]. The lower lid moves nasally, about 4.5 ± 0.9 mm and slightly upwards. The width of the palpebral aperture is also reduced [322]. During saccades, movement is between the tarsus and the unexposed surfaces of the globe. In vertical gaze there is more limited relative movement between the upper lid and globe.

The relationship between friction and velocity, load and viscosity, is described by the Stribeck curve [200,335,336]. This empirical relationship was originally characterized for steel journal bearings using oil lubrication, with the friction coefficient on the y axis and the Hersey number (viscosity*sliding velocity/normal pressure) on the x axis. Many biological surfaces are soft, complex, heterogeneous hydrated materials (such as the cornea and lid tissue) and therefore may not follow classic Stribeck behavior [337], yet the curve can still provide a framework for discussion and interpretation.

According to the Stribeck curve, where tear volume is sufficient, then friction during blinking depends on the rate of relative movement of the apposed surfaces and the tear viscosity. Given that $S = (v \cdot \eta)/t$ where S = shear friction, v is the velocity of the upper lid during the blink, η is the viscosity of the tears and t is the thickness of the tear layer between the apposed surfaces.

The above equation suggests that for a given tear viscosity, the thicker the tear film, the lower friction will be. For a Newtonian fluid whose viscosity is independent of shear rate, hydrodynamic friction increases with increasing viscosity and this may be relevant to events in DED (see below), but normal tears behave as a non-Newtonian fluid [338–340] whose viscosity falls with increasing shear rate (i.e. it shear-thins), so that this consideration does not apply. Therefore, according to this relationship, while acknowledging that the Stribeck curve classically applies to non-porous stiff materials, assumptions can be made. With friction during blinking/saccades depending on the rate of relative movement and tear viscosity, when tear volume is sufficient, the coefficient of friction between lid and globe may be assumed to be low. It is suggested that there is a rapid transition from brush-to-brush or boundary lubrication, to hydrodynamic lubrication with increasing velocity during the blink [322]. The lid margin profile may also be important in the transition from boundary to hydrodynamic lubrication [322].

Friction is greatly increased in DED states due to a failure of lubrication [341], with the loss of mucin gel and glycocalyx or, in ADDE, of fluid volume. This may cause damage to specific sites, as observed in lid wiper epitheliopathy (LWE) [342], parallel conjunctival folds, (LIPCOF) [343], and superior limbic keratoconjunctivitis (SLK).

4.13.3. Frictional forces at the lid-wiper region of the lid

The lid-wiper region of the lid was originally described as that portion of the upper eyelid that comes into intimate contact with the globe and wipes over it during blinking [342,344]. This role had earlier been conceived by Parsons [345] and by Ehlers [323], based on the recognition of a 'stratified squamous' epithelium at this site. It is now recognised to be a feature of both the upper and lower lid [346].

The upper lid wiper consists of an elevated strip of marginal conjunctival epithelium, 100 μm thick, varying in width from 0.3 to 1.5 mm and extending the full length of the lid margin at the level of Riolan's muscle. According to Knop it is composed of a stratified cuboidal epithelium [40], which is closely applied to the globe during the blink [326,347] and is probably the closest region of contact between the upper, and presumably, the lower, lid and globe. In the upper lid, the tarsal mucosa lying proximal to this zone, and separated from the globe by a mucoaqueous layer of unknown thickness, (within 'Kessing's space') [36,97] is thought to be less closely applied. The presence of both goblet cells and goblet cell crypts in the lid wiper epithelium [40,101] is presumed to provide a local, mucinous lubrication system at this point of primary contact, of importance during blinking and to a lesser extent during eye movements where the forces applied are lower.

4.13.4. The consequences of shearing forces at the ocular surface

At a conservative estimate, considering a blink rate of 12 times a minute over a 16 h day, an individual would blink 11,000 times in the course of a day and, assuming a palpebral aperture of 10 mm in height, the lid wiper would have travelled a distance of at least 100 m over the surface of the cornea [101]. Notwithstanding the presence of a lubricating system of high quality, this is a source of shear stress at the ocular surface. It is presumed to play a role in epithelial desquamation, in the punctate epithelial staining found at the ocular surface in the normal eye and in the enhanced punctate epithelial staining of DED. Additionally, as noted, it contributes to other clinical features of DED such LWE LIPCOF and SLK, any of which may occur, in lesser degree, in the absence of DED.

4.14. Epithelial desquamaton

The following scenario, referring to the cornea, may be proposed for the shedding of epithelial cells. A similar process is assumed for the conjunctiva [348]. Epithelial cells arise by the division of stem cells at the corneal limbus and are increased in number by division of transient amplifying cells, in the periphery [349]. Newly formed cells undergo terminal differentiation as they migrate centripetally and to the surface and after a period of residence, undergo a process that leads to shedding, which may be preceded by apoptosis [350]. Epithelial desquamation involves uncoupling of layer 1 cells from

neighbouring cells, with a loss of junctions, including tight junctions and adherens junctions, and dissolution of the apical glycocalyx. At some point, having lost adherence to surrounding cells, the cell destined for shedding is easily displaced by frictional forces. Its place is taken by a younger cell, already equipped with a maturing glycocalyx, which is rapidly integrated with neighbouring cells by means of tight junction formation, thereby restoring the functional integrity of the surface. It is this process that most likely explains the infrequent occurrence of punctate epithelial staining in the normal eye.

4.14.1. Physiological punctate epithelial staining

The subject of punctate epithelial staining has been recently reviewed [348]. A low degree of punctate epithelial staining is a regular finding on the normal cornea and conjunctiva after the instillation of dyes such as fluorescein, lissamine green and rose bengal and may be regarded as a physiological phenomenon. Based on reports in the literature it occurs with a frequency of 4–78% [351], varying with methods of assessment, particularly with volume and concentration of dye instilled and the period of observation. The number of staining points increases with time in the post-instillation period. Punctate epithelial staining on the cornea and conjunctiva, with a characteristic horizontal, interpalpebral pattern, is a diagnostic feature of DED.

In normal subjects, a proportion of corneas show a low level of punctate staining immediately after fluorescein instillation [351–356]. A 'clinically significant' grade of fluorescein staining has been reported to occur in about 12% of non-contact lens wearers [357–359] but the figure rises if 'clinical significance' is ignored – eg. from 37% to 58%, in the study of Korb [353]. Similarly, in a study of normal subjects (median 22 years; range 18–50 years) after the instillation of fluorescein from a an impregnated strip, 79% of subjects showed some degree of corneal staining. Less information is available about the conjunctiva.

Norn reported the frequency of punctate epithelial staining in the normal cornea, read at 1–2 min following the instillation of a 10 μl of 0.125% fluorescein. Punctate staining was present in 4% of subjects under 40 years of age, increasing to 20% above the age of 50 years, after which the frequency became stable. The mean frequency for the group overall (n = 411) was 17% [360]. Similarly, the number of dots per cornea rose with age, although in most subjects the number of dots per cornea was small, with only 1% of subjects showing over 100 dots per cornea, compared to 35% with over 1000 staining dots in patients with DED (Table 4).

In general, the prevalence of staining and dot frequency, increases with dye concentration [360], with time after drop instillation (Korb and Korb 1970) and with age of the subject [360,361]. Caffery and Josephson showed that the regional pattern of corneal staining was individual to the subject, similar in fellow eyes, and, importantly, varied from day to day [356], which was confirmed by Schwallie et al. [362] They concluded that variation might relate to the natural turnover of the epithelium.

Table 4
Micropunctate staining dots per cornea after instillation of fluorescein.

Dots per cornea	Percent with 0.125% fluorescein	Percent with 1.0% fluorescein
Zero	83	27
1–4	9	16
5–9	4	2
10–25	3	4
25–99	1/2	0
100–999	1	16
≥1000	0	35

Percentage of normal corneas showing a given number of micropunctate staining dots per cornea, after instillation of 10 μl of either 0.125% or 1% fluorescein (in combination with 1% rose bengal) (n = 411, including fellow eyes). Staining was read more than 1–2 min following dye instillation (From references 348, 360).

The physiological occurrence of corneal and conjunctival staining suggests that in clinical trials, for instance of DED therapies, zero corneal staining is not a reasonable criterion to define recovery to full corneal health. Also, the time-dependent staining of the normal corneal epithelium and dependence on instilled concentrations, emphasizes the need for standardization of staining routines to assess ocular surface damage.

4.14.2. Mechanism of punctate epithelial staining in normal and in dry eyes

The mechanism that determines punctate epithelial staining has been in debate for over half a century and has been addressed by several recent reviews [79]. There appears to be no direct evidence that punctate dots of stain represent pools of dye lying within spaces left by shed cells, hence the term punctate epithelial erosion is not appropriate [363–365]. Rather, it seems that each dot of stain represents the uptake of dye into a surface epithelial cell.

4.14.3. Staining in the normal eye

Epithelial cells are shed daily from the ocular surface and about 75% of collected cells are corneal [366]. Cells are shed in a diurnal pattern, with more cells shed in the morning and the latter part of the day [367]. About 23% are ghost cells, lacking nuclei, considered to be at a late stage of cell differentiation [368,369]. This is consistent with an earlier study showing the presence of both viable (calcein-positive only) and nonviable (ethidium-positive only) epithelial cells, as well as an intermediate cell type that stained with both calcein and ethidium [367].

The vast majority of layer 1 epithelial cells do not take up dye, whereas shed epithelial cells, trapped in the fornical mucus thread, stain with rose bengal [203], as do immature human limbal corneal epithelial cells grown in culture [370,371]. Similarly, cultured rabbit corneal epithelial cells stain avidly with fluorescein [372,373]. Argueso et al. resolved this issue by demonstrating that exclusion of rose bengal from entry into surface epithelial cells in the intact eye, depends on the presence of a mature glycocalyx expressing MUC1 and MUC 16 mucins, crosslinked by galectin-3 [128,134,148]. The mature glycocalyx forms a barrier to transmembrane dye entry into the layer 1 epithelial cells, while entry into the paracellular space is restricted by intercellular, tight junctions. Bandamwar et al. [350,374] have presented evidence that staining cells are those undergoing apoptosis while preparing for shedding. Such cells possess a defective glycocalyx layer, which is permeable to dyes in clinical use. Once shed, all epithelial cells are incompletely clad with glycocalyx and hence stain readily.

It is hypothesized that the permeability of a cell that is preparing for shedding, increases over time, due to chemical and structural changes in its glycocalyx, so that those cells that are just about to be shed take up stain almost immediately whereas those that are at an earlier stage of preparation, take up stain more slowly. This is presumed to be the basis of the effect of dye concentration or of period of observation, on the frequency of physiological staining dots.

5. The pathology of dry eye disease

These introductory remarks are intended to arm the reader to understand the events responsible for the many forms of DED.

5.1. Introduction

The TFOS DEWS [1] report confirmed tear hyperosmolarity, along with tear instability, as the core drivers of DED. This allowed two major subtypes to be defined, EDE, where tear hyperosmolarity

is the result of an excessive evaporation from the tear film in the presence of normal lacrimal function and ADDE, where hyperosmolarity results from a reduced lacrimal secretion in the presence of a normal rate of tear evaporation (Table 5). The tear film lipid deficiency that accompanies MGD is cited as a typical cause of EDE and the reduced tear secretion due to lacrimal gland damage in age-related DED provides a typical example of ADDE. It was recognised that these subtypes of DED may coexist and this is the case in Sjögren syndrome where lacrimal deficiency frequently coexists with MGD [375,376,1201]. Also, in any form of cicatricial conjunctivitis, DED may be secondary to a lacrimal tear deficiency, a tear lipid deficiency and loss of ocular surface wettability.

Other forms of hybrid DED may also be conceived, in which organic disease of one type may be combined with a functional form of DED of another type [207]. For example in severe EDE, loss of corneal sensitivity could remove the compensatory drive to lacrimal secretion and lead to a secondary, functional aqueous deficiency. Or in ADDE, a severe reduction in tear film thickness could impair TFLL spreading and give rise to a secondary, functional EDE. Additionally and importantly, it may be observed that in any form of DED, once tear breakup occurs within the blink interval, an additional evaporative component is added to the dry eye, regardless of the initiating cause. A consequence of this is that a dry eye that is initiated by a lacrimal tear deficiency becomes an ADDE + EDE as it evolves. It follows that, where comparisons of tear evaporation rate are made between forms of DED classically defined as ADDE and EDE, the OPI should be taken into account. This also has implications for therapy and for subgroup selection and analysis in clinical trials. This Subcommittee recommends that the terms EDE and ADDE be retained to describe the *initiating* basis of a dry eye but that it should be recognised that with progression any form of DED may take on additional evaporative features.

It should be kept in mind that, in a sense, all forms of DED are evaporative, since without evaporation, tear hyperosmolarity cannot occur. Consequently, environment and personal behavior are contributors to ocular surface hyperosmolarity, including external factors such as ambient humidity, temperature and wind speed and personal factors such as blink rate and lid aperture size, gaze position and the influence of systemic medication on tear secretion. The Subcommittee discussed the term, 'hyperevaporative dry eye' as a better way of indicating the role of increased evaporation in DED.

A major contribution of the TFOS DEWS report [1] was the proposition that every kind of dry eye, however initiated, enters a final common pathway in which tear hyperosmolarity and a chain of inflammatory events create a Vicious Circle that perpetuates the DED state [377]. According to this approach, *any etiology of DED will have one or more entry points* into the Vicious Circle. The concept of the Vicious Circle is illustrated in (Fig. 5) and is elaborated in the text that follows.

5.2. The Vicious Circle of dry eye

In the simplest model of DED, with tear hyperosmolarity as its the starting point, the pathological process is propagated by a chain of events that lead to ocular surface damage (Fig. 5). Initially this gives rise to symptoms and compensatory responses, but it also generates inflammatory responses that ultimately lead to chronic ocular surface damage and self-perpetuated disease [377].

This may be summarised as follows:

As noted earlier, tear hyperosmolarity stimulates a cascade of events in the epithelial cells of the ocular surface, involving MAP kinases and NFκB signaling pathways [311] and the generation of inflammatory cytokines (IL-1 [IL-1α; IL-1β]); tumor necrosis factor-α [TNF-α]) and proteases, such as MMP9 [378]. These activate and

Table 5
Causes of dry eye disease.

- **AQUEOUS-DEFICIENT DRY EYE (ADDE)**
  - Sjögren Syndrome Dry Eye (SSDE)
    - associated systemic diseases
      - Rheumatoid arthritis
      - Polyarteritis nodosa
      - Systemic lupus erythematosis
      - Wegener granulomatosis
      - Systemic sclerosis
      - Primary biliary cirrhosis
      - Mixed connective tissue disease
  - Non- Sjögren Syndrome Dry Eye (NSDE)
    - Intrinsic Lacrimal Gland Deficiency
      - Lacrimal gland ablation
        - Congenital alacrima
        - Triple A syndrome
      - Age-related ADDE dry eye
      - Inflammatory and Other Lacrimal Gland infiltration
        - Sarcoidosis
        - Lymphoma
        - Viral Infection
        - Radiation Injury
      - Lacrimal Gland Obstruction
        - Cicatricial Conjunctivitis
          - GVHD
          - Stevens-Johnson Syndrome/TEN
          - Mucous Membrane Pemphigoid
          - Cicatricial pemphigoid
          - Pemphigus
          - Trachoma
          - Chemical injury
      - Hyposecretory States – Failure of the Lacrimal Functional Unit
        - Reflex Afferent Block
          - Topical anesthesia
          - Trigeminal nerve injury
            - Refractive surgery
            - Neurotrophic keratitis
        - Secretomotor Block
          - Parasympathetic damage
          - Pharmacological inhibition
        - Combined Afferent and Efferent Block
          - Familial dysautonomia
      - Other Disorders
        - Meige Syndrome
        - Diabetes Mellitus
        - Pseudoexfoliation
- **EVAPORATIVE DRY EYE**
  - Meibomian Gland Diseases
    - Lid-Related
      - Meibomian Gland Dysfunction (MGD)
        - Primary
          - Meibomian seborrhea
          - Obstructive MGD
            - Cicatricial/non-cicatricial
        - Secondary to Local Disease
          - Anterior blepharitis
          - Ocular surface inflammation
          - Contact lens wear
        - Secondary to Systemic Dermatoses
          - Rosacea
          - Seborrheic dermatitis
          - Atopic dermatitis
          - Icthyosis
          - Psoriasis
        - Secondary to Chemical Exposure
          - 13-cis retinoic acid
          - Polychlorinated biphenols
          - Antiandrogens
      - Genetically Determined Meibomian Gland Diseases
        - Meibomian Agenesis and Dystichiasis
        - Anhydrotic Ectodermal Dysplasia
        - Ectrodactyly Syndrome
        - Epidermolysis Bullosa
        - Ichthyosis Follicularis
        - Turner Syndrome;
      - Disorders of Lid Aperture, Congruity, Dynamics
        - Blink-Related

(*continued on next page*)

Table 5 (*continued*)

Parkinson's Disease
Ocular Surface-Related Evaporative Dry Eye
Allergic Eye Disease
Vitamin A Deficiency
Short Breakup Time Dry Eye
Iatrogenic Disease

recruit inflammatory cells to the ocular surface, which become an additional source of inflammatory mediators [379]. Such mediators, acting with tear hyperosmolarity itself, lead to a reduced expression of glycocalyx mucins, to apoptotic death of surface epithelial cells [380] and to a loss of goblet cells. Hyperosmolarity also induces corneal epithelial cell death through non-apoptotic processes [62]. Goblet cell loss is a feature of every form of DED [381,382], reflected by reduced tear levels of MUC5AC [383,384]. Altered expression of glycocalyx mucins is a likely basis for ocular surface staining in DED and by compromising ocular surface wetting, leads to early tear film breakup. This amplifies or initiates ocular surface hyperosmolarity, which completes the Vicious Circle

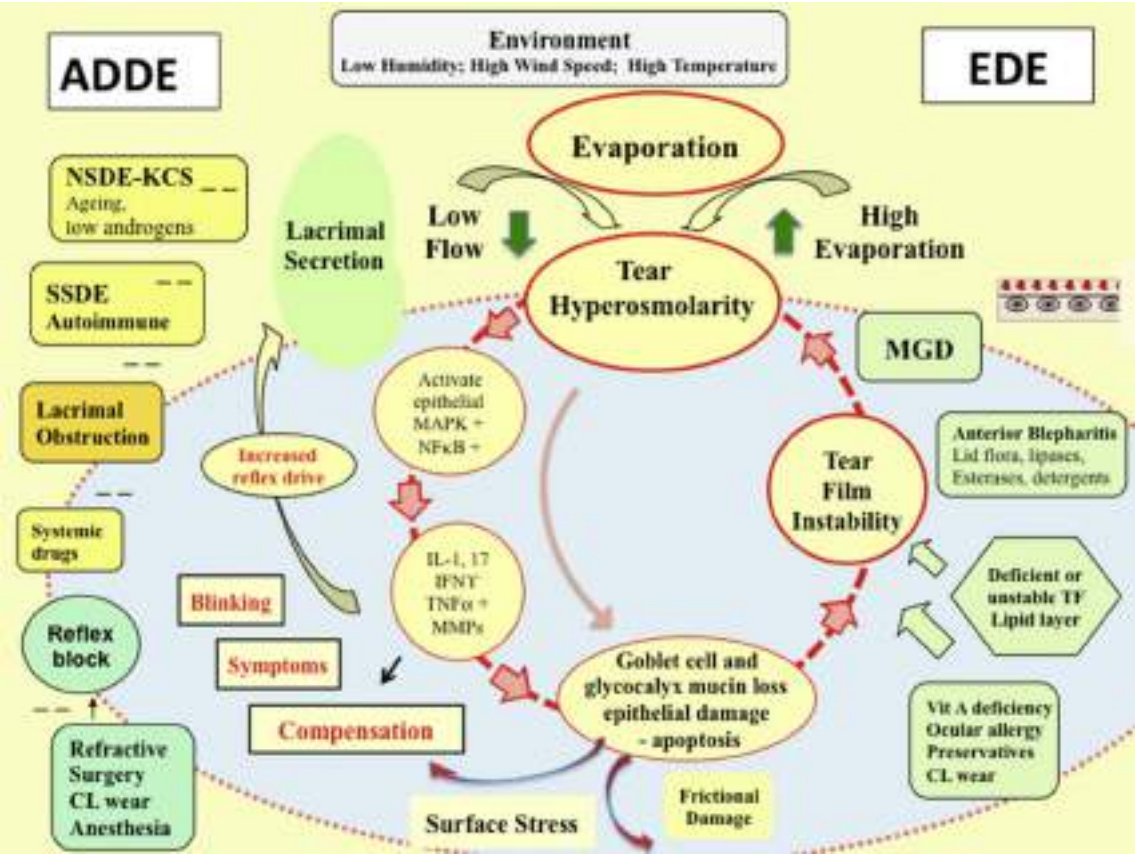

Fig. 5. The Vicious Circle of Dry Eye Disease. The core mechanism of DED is tear hyperosmolarity, which is the hallmark of the disease. It damages the ocular surface both directly and by initiating inflammation. The cycle of events is shown at the centre of the figure. Two forms of DED are recognised, ADDE and EDE. In ADDE, tear hyperosmolarity results when lacrimal secretion is reduced, in conditions of normal evaporation from the eye. In EDE, tear hyperosmolarity osmolarity is caused by excessive evaporation from the exposed tear film in the presence of a normally functioning lacrimal gland. Since tear osmolarity can only rise as a result of tear evaporation in both ADDE and EDE, tear hyperosmolarity is due to evaporation from the ocular surface and in that sense, all forms of DED are evaporative. EDE is a hyper-evaporative state. In DED, tear hyperosmolarity is considered to set up a cascade of signaling events within surface epithelial cells, that leads to the release of inflammatory mediators and proteases. Such mediators, together with the tear hyperosmolarity itself, are conceived to cause goblet cell and epithelial cell loss and damage to the epithelial glycocalyx. Damage is reinforced by inflammatory mediators from activated T-cells, recruited to the ocular surface. The net result is the characteristic punctate epitheliopathy of DED and a tear film instability which leads at some point to early tear film break-up. This break-up exacerbates and amplifies tear hyperosmolarity and completes the Vicious Circle of events that lead to ocular surface damage. Ultimately this is thought to lead to self-perpetuation of the disease. Tear film instability can be initiated without the prior occurrence of tear hyperosmolarity, by conditions that affect the ocular surface, including xerophthalmia, ocular allergy, topical preservative use and contact lens wear. In this case, early tear film breakup (an Ocular Protection Index <1) is the primary basis for tear film hyperosmolarity at first experienced locally at the site of breakup and increasing in severity, at some point being detectable in tear meniscus samples. This represents an ocular surface–related form of EDE. In MGD-related EDE tear hyperosmolarity results from a tear film lipid layer deficiency. In ADDE the onset of early break-up during the evolution of the disease, may add an evaporative element to the dry eye. There are various causes of ADDE. It may result from blocking the sensory drive to the lacrimal gland that is essential to maintain osmolar homeostasis. Bilateral topical anesthesia can cause both a reduction in tear secretion and blink rate. Dry eye due to such a reflex block can be caused by chronic abuse of topical anesthetics, trigeminal nerve damage and refractive surgery including LASIK surgery. The delivery of aqueous tears to the tear sac can also be reduced by obstruction of the lacrimal ducts, which can occur in any form of cicatricial conjunctival disease, such as trachoma, erythema multiforme, graft-versus-host-disease and chemical burns. A number of drugs in systemic use, such as antihistamines, beta-blockers, antispasmodics, diuretics and some psychotropic drugs, cause a reduction in lacrimal secretion and are risk factors for DED. Tear secretion rate falls in later life. The anti-glaucoma drugs pilocarpine and timolol also have direct effects on human meibomian gland epithelial cells that may influence their morphology, survival and/or proliferative capacity, and possibly promote MGD [61,1103]. In the western world the most common cause of ADDE is inflammatory infiltration of the lacrimal gland, encountered most severely in autoimmune disorders such as Sjögren syndrome dry eye (SSDE) and, with lesser severity, in non-Sjogren dry eye (NSDE). Inflammation causes both acinar and ductal epithelial cell dysfunction and/or destruction and a potentially reversible neurosecretory block. A receptor block may also be caused by circulating antibodies to the muscarinic, M3 receptor. Inflammation is favoured by low tissue androgen levels. Epithelial injury and a defective glycocalyx, loss of tear volume and of goblet cell mucin, lead to increased frictional damage and friction-related symptoms. The tear hyperosmolarity and epithelial injury caused by DED, stimulates corneal nerve endings, leading to symptoms of discomfort, increased blink rate and, potentially, a compensatory reflex increase in lacrimal tear secretion. This compensatory secretion is more likely in EDE, since lacrimal function is potentially normal. Adapted from Bron, Definition of dry eye disease' in Chan 2015 - Springer [79].

and establishes the mechanism that perpetuates the disease.

It has been emphasized by Baudouin et al. that the Vicious Circle offers *entry points* for any cause of DED [385]; tear hyperosmolarity need not be the starting point. Thus the chain of events leading to tear film instability may be initiated by several different disorders, including, but not limited to, ocular surface inflammation due to allergic eye disease, topical preservative toxicity and loss of conjunctival goblet cells or altered mucin expression, due to xerophthalmia.

5.3. Compensatory events in dry eye

It is inherent to our current understanding of DED that exposure of the ocular surface to desiccating stress sets up a compensatory, secretory tear response, via the lacrimal functional unit that tends to offset a rise in tear osmolarity and slow disease progression. As summarised in the TFOS DEWS II Pain and Sensation report [1224], both tear hyperosmolarity and surface cooling may trigger this response. The cold modality fibres of the cornea are stimulated by hyperosmolarity and could both increase the secretory drive to the lacrimal gland and lead to an increased blink rate. Evaporative cooling in EDE or in relation to early tear film breakup [279,386], could add to this sensory drive. The finding of a reduced threshold to sensory stimulation, in some DED patients [387], could amplify these responses. Other authors have reported decreased corneal sensitivity in DED [388,389], which may imply that, as DED progresses in severity, corneal sensation becomes impaired. In keeping with this, a number of studies have reported a reduction in subepithelial nerve density in DED [390]. Such a sequence could impact unfavourably on compensatory responses and could contribute to discrepancies between symptom intensity and objective signs of DED. However, this possibility, which would be important to our understanding of DED progression, has not been addressed in long-term studies.

5.4. Symptoms

Any symptomatic disease goes through a subclinical phase in which the features of the disease are not obvious and the patient is symptom free. DED is no exception. (see the TFOS DEWS II Definition and Classification and Diagnostic Methodology report [1225]) But the burden of DED for the patient relates to symptoms and there is now a greater understanding of their causes. DED affects both vision and comfort of the eye. Potential sources of symptoms in DED are listed in Table 6.

There is evidence to support a direct role for hyperosmolarity as one basis for ocular discomfort in DED. As noted, instillation of hyperosmolar drops causes pain of an intensity related to the level of hyperosmolarity, but at levels far higher than that detected in tear meniscus samples in DED patients [276]. Modeling considerations have suggested that the levels of hyperosmolarity generated at the site of tear breakup are far higher than in the tear meniscus [279]. There is evidence too, that tear hyperosmolarity is initiated as the tear film thins, and is amplified at the time of tear breakup [391]. Additionally, several of those inflammatory mediators, which have been demonstrated in the tears and ocular surface in DED, are known to be algesic compounds, including various prostanoids, cytokines and neurokinins. (see the TFOS DEWS II Pain and Sensation report [1224] for further details). The loss of lubrication between the globe and lids in DED has been suggested as a source of friction-related outcomes, including a reduction in tear volume in ADDE, loss of goblet cell gel mucin, degradation of glycocalyx mucin [218] and loss of the boundary lubricant, lubricin [82]. Filamentary keratitis is a particular source of pain, attributed to the drag of filaments on nociceptor endings at the base of the filament during blinking. A similar process may be responsible for symptoms of discomfort associated with LIPCOF [343]. The basis for pain associated with LWE is assumed to be due to hypersensitivity over the affected region of the lid wiper and the region of the keratopathy. In the healthy eye, this region of the lid margin has a mechanical sensitivity similar to that of the central cornea [250].

Thus tear hyperosmolarity is but one of the potential sources of discomfort in DED, another reason why the levels of tear osmolarity measured in DED patients with chronic pain may not always be significantly different from the osmolarity of asymptomatic patients [392].

Hypersensitivity (reduced threshold to stimulation) of corneal nerves in DED patients may also explain the occurrence of ocular discomfort at lower levels of tear osmolarity, due to exposure of corneal nerve endings with loss of the epithelial barrier [267,387,393].

Instillation of hyperosmolar drops within the ranges of osmolarity found in DED patients was shown to increase the sensitivity of the cold nociceptive neurons and to induce DED signs in a rat model. In this rat model, these nociceptors, which normally require more than 2 °C cooling, were activated by less than 1 °C cooling of the corneal surface when pretreated with hyperosmolar fluids [394]. This phenomenon may explain the cooling-induced discomfort and pain reported by DED patients. Upregulation of TRPM8 channels or control of the voltage-gated potassium channels (Kv1.1) may be involved in this process [395]. Both channels are well-established cooling sensors that can be regulated by a hyperosmolar stimulus [396].

Table 6
Sources of dry eye symptoms.

i. Visual Symptoms – (occurring in the interblink interval)
Tear film instability and breakup
Epithelial roughness in regions of tear breakup
ii. Symptoms of Discomfort
Tear Hyperosmolarity
General – affecting all tear compartments
Local – tear breakup related, local hotspots of hyperosmolarity
iii. Friction - Reduced lubrication - (Related to blinking and eye movements)
Low tear volume in ADDE
Loss of goblet cells; mucin
Loss of mature glycocalyx, loss of lubricin
Rough epithelium; punctate epithelial keratitis
Filamentary keratitis
SLK
LIPCOF - conjunctivochalasis
LWE
iv. Inflammatory mediators
Algesic mediators increasing sensory excitability
Prostanoids
Cytokines
Neurokinins
v. Neurosensory and central factors
Trigeminal hypersensitivity;
Neuropathic firing
Cognitive aspects of dry eye symptoms

ADDE, aqueous deficient dry eye; SLK, superior limbic keratoconjunctivitis; LIPCOF, lid parallel conjunctival folds; LWE, lid wiper epitheliopathy.

5.5. The ocular targets of dry eye disease

Clinical consequences at the ocular surface are independent of etiology. These consequences may include punctate epitheliopathy, filamentary keratitis, superior limbic keratitis, goblet cell loss, modification of the epithelial glycocalyx, LIPCOF, changes to Marx's line and MGD itself (Table 7). These are discussed below:

Table 7
Ocular targets of dry eye disease.

i. The Lacrimal Gland.
Inflammatory cell infiltration of ducts and acini
ii. The Meibomian Glands
Terminal duct obstruction; duct dilatation and gland loss
iii. The Cornea
Punctate epithelial keratopathy.
Filamentary keratitis
Superior Limbic Keratoconjunctivitis (SLK)
iv. The Conjunctiva.
a. General changes
Punctate epitheliopathy
Glycocalyx changes
Goblet cell loss
b. Bulbar changes
Lid parallel conjunctival folds (LIPCOF)
SLK
c. Tarsal changes
The lid margins changes
Marx's line migration
Lid wiper epitheliopathy.
v. Both Cornea and Conjunctiva.
Increased epithelial shedding
vi. Tear Film Instability
Early signs
Tear film breakup. Spot, dimple, line, area

5.5.1. The cornea

5.5.1.1. Punctate epitheliopathy and staining in dry eye disease. Evidence suggests that noxious influences at the ocular surface in DED lead to increased epithelial cell death (e.g. apoptosis) and increased epithelial shedding and turnover. It is likely that increased friction contributes to the increased shedding. No formal measurements of increased shedding or turnover have been made in DED and this would be of value.

Tabery has shown that punctate corneal epithelial staining in DED can be explained by the uptake of dye directly into individual epithelial cells and that fluorescein is taken up into the same cells that take up rose bengal [397,398]. Several studies suggest that staining cells on the cornea and conjunctiva have a defective glycocalyx, including a deficiency of MUC 16, [399–401] and in bullous keratopathy, too, superficial exfoliation and staining is associated with breaches in MUC16 [402]. Komuro et al. found, in patients with superior limbic keratoconjunctivitis, that areas of conjunctiva showing positive staining with rose bengal, had no galectin-3 expression, whereas in healthy regions showing rose bengal exclusion, galectin-3 was expressed normally [403].

The staining of individual, layer 1, corneal epithelial cells in DED states is thus attributed to the diffusion of dye across the defective glycocalyx of apoptotic cells, prior to shedding. The staining of small clusters of surface cells may have a similar explanation, but an additional possibility is that the dye enters the paracellular space around a cell that is about to be shed, across a defective tight junction and spreads into neighbouring cells across their plasma membranes, ie. by transmembrane spread [79]. Intercellular spread of dye between neighbouring cells via gap-junctions is less likely in the superficial epithelium since these are absent from layer 1 in the human cornea and connectivity is limited in the second layer [404]. An alternative view has been expressed [363].

5.5.1.2. The pattern of staining in dry eye. Epithelial staining of the exposed cornea and conjunctiva in DED has a characteristic horizontal, inter-palpebral distribution, which is of diagnostic value (Fig. 6). There has been a longstanding interest in its basis, particularly in relation to hyperosmolar hot-spots generated in the interblink interval. McMonnies [405], and others [406] have emphasized the role of partial blinking in extending the period of lower globe exposure to desiccating stress, emphasizing that the period of exposure will be a multiple of the number of partial blinks that occur in sequence. Partial blinking is common, both in normal subjects and in DED. A figure of up 22% was reported in one study of normal eyes [407] and may comprise from 20% to over 50% of all blinks [408–410]. Jansen et al. noted that in subjects engaged in tasks requiring a high degree of visual concentration, both the number of incomplete blinks and the interblink interval increased [290].

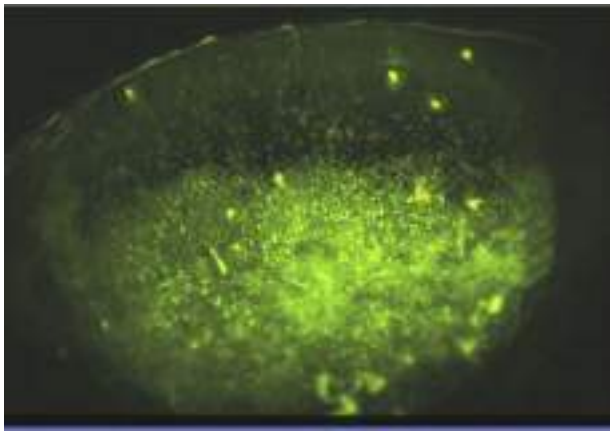

Fig. 6. Severe filamentary keratitis, with extensive corneal staining with fluorescein.

The region of meniscus-induced thinning (MIT), corresponding to the position of the 'black line' in the fluorescein-stained tear film, is also predicted to be a site of tear hyperosmolarity in the interblink interval [163,411]. However, the risk of hyperosmolar damage to the underlying corneal and conjunctival epithelium, due to MIT, is minimized by eye movements, especially in the vertical plane, which can distribute the effect over a wider area and reduce its damaging potential. Nonetheless, as McMonnies has observed, 'this may not be the case during reading, watching television, or similar activity, when up and down eye movements are limited and the zone of MIT maintains a more stable location on the ocular surface. Stability of its location is likely to be associated with an increased risk of hyperosmolarity-related epitheliopathy' [405]. This effect will be amplified in conditions where there is a severe restriction of eye movements, such as in progressive supranuclear palsy [412], progressive external ophthalmoplegia [413] and endocrine exophthalmos [414].

Additionally, it has been reported in normal subjects that, after short periods of up or downgaze, bands of MIT are imprinted onto the cornea which persist in the interblink interval. These may be accompanied by a secondary disruption of the tear film below [197]. These zones of thinning represent regions of potential hyperosmolar damage and hence a source of increased staining.

The above considerations summarize factors which might direct desiccating stress to the lower part of the exposed globe in any eye. These influences will be amplified in environmental conditions that increase desiccating stress and will be further so in DED states where early tear film breakup will determine the location of regional hotspots of hyperosmolarity. They appear to offer a reasonable explanation for the pattern and distribution of punctate epitheliopathy in DED.

5.5.1.3. Filamentary keratitis. Filamentary keratitis describes a condition of solitary or grouped filaments, usually no greater than 2 mm in length, which extend from the corneal epithelium into the tear film (Fig. 6). They seldom occur on the conjunctiva. It is associated with ocular surface disorders such as DED, SLK, viral

conjunctivitis, recurrent corneal erosion, neuroparalytic keratitis, post corneal transplantation, cataract surgery, ocular trauma and ptosis. In ptosis and SLK, filaments are commonly located under the upper lid; otherwise, for instance with severe aqueous deficiency, they have an interpalpebral location.

Corneal filaments stain particularly well with rose bengal and lissamine green. Using immunohistochemistry, Tanioka et al. showed that they have a twisted epithelial core, surrounded by secretory (MUC5AC) and membrane-associated (MUC16) mucins, inflammatory cells, and conjunctival epithelial cells, from which they concluded that filaments are spun out by an increased frictional action during blinking [415]. Frictional drag on the filaments during blinking results in severe, intractable, ocular pain and foreign body sensation [416]. Although filaments can be removed manually after instillation of a topical anesthetic, recurrence is not uncommon.

5.5.1.4. Superior limbic keratoconjunctivitis. SLK [417] is a bilateral, chronic inflammatory disease, affecting the upper bulbar conjunctiva, superior limbus and adjacent cornea. It can be a source of disabling discomfort. Typically, a patch of severe, perilimbal conjunctival hyperemia or inflammation, is accompanied by limbal thickening, punctate keratopathy, filamentary keratitis and a papillary reaction in the overlying superior tarsal conjunctiva. There may be a discrepancy between the level of pain experienced and the severity of clinical signs and the diagnosis may be missed if lissamine green staining is not carried out in the clinical workup of unexplained ocular discomfort. Staining with fluorescein is less apparent unless used with a suitable filter combination [74].

Histologically, squamous metaplasia, epithelial thickening with a decrease in the nuclear-cytoplasmic ratio, and a disappearance of goblet cells is reported in SLK [417]. Twenty five percent of SLK cases are associated with DED [418] and about 30% with thyroid disease [419], therefore it is important to examine hormone and autoantibody status. There is also an association with conjunctivochalasis affecting the upper bulbar conjunctiva [420,421].

In SLK, it is reported that the chronic inflammation may be related to the blink and to eye motion [418] and the association of SLK with upper bulbar conjunctivochalasis strongly supports recurrent frictional trauma as a trigger, particularly since surgery directed towards tightening the conjunctiva at this location is highly successful [421,422]. Similarly, in endocrine exophthalmos, an increase in the pressure of the upper lid against the globe may be invoked as the mechanism that precipitates SLK in thyroid disease with exophthalmos.

5.5.2. Conjunctiva

Although conjunctival goblet cell loss and a decrease in MUC5AC concentration in the tears are generally accepted as features of all forms of DED, reports of changes in the transmembrane mucins are less consistent [423]. This is in part due to differences in methodology, for instance in using immunohistochemistry to detect core mucin proteins on the one hand or the pattern of mucin glycosylation on the other. It is difficult to determine what level of glycan alteration is sufficient to disrupt the glycocalyx permeability barrier [154]. A loss of a mucin-like glycoprotein (probably MUC 16) from keratinized, surface conjunctival epithelial cells has been reported in SLK [424].

5.5.2.1. Modification of the epithelial glycocalyx mucins. There is evidence of an altered expression or glycosylation of transmembrane mucins in DED. In an immunohistochemical study, conjunctival mucosal epithelial membrane mucin expression was decreased in Sjögren syndrome [401]. More recently, Shimazaki-De et al. reported a reduced MUC16 mRNA expression in the conjunctiva in DED [425]. Similarly, surface immunoreactivity to MUC-1 appears to be reduced in Sjögren syndrome epithelium, suggesting disruption of normal epithelial differentiation [426], and Corrales et al. found significantly decreased mRNA expression of MUC1, MUC2, MUC4 and MUC5AC in ADDE patients [427].

In contrast, it was shown that the density of cells positive to KL6, a monoclonal antibody against a sialylated epitope of MUC1 was significantly increased in DED patients compared to normal [428]. Additionally, in Sjögren syndrome DED, mRNA, and proteins of both MUC16 and MUC1, were increased compared to normal subjects [429]. The basis for these conflicting findings needs to be resolved.

Gipson et al. [430] demonstrated an increase MUC1 and MUC16 mRNA and cell protein expression in impression cytology samples from postmenopausal women compared to normal. In contrast, Srinivasan et al. [431], found that MUC16 mRNA expression was significantly reduced in postmenopausal women with moderate to severe OSDI symptoms, while MUC1 mRNA expression remained unchanged, compared to non-symptomatic subjects.

In DED, some alterations of mucin glycosylation have been investigated. Garcher et al. showed a decrease of sialylated chains of mucins expressed in impression cytology samples from DED patients and CL users and in glaucoma patients treated with β-blockers [432]. In general, glycosyltransferases are the enzymes responsible for the initiation and elongation of glycan chains attached to the protein backbone. In mucins, the enzymatic addition of *N*-acetyl galactosamine (GalNAc) to serine and threonine residues by GalNAc-transferases (GalNAc-T) is the initial step in O-glycosylation. In ocular cicatricial pemphigoid (OCP), the conjunctival expression of GalNAc-transferases was increased in patients with early disease, which could play a role in maintaining epithelial wettability. Conversely, as might be predicted, expression was markedly reduced at the stage of conjunctival keratinization [433].

5.5.2.2. Goblet cell loss. Ralph [434] emphasized that conjunctival goblet cell loss is a feature of all forms of DED, and this has been confirmed in later reports, in Sjögren syndrome (SS), OCP, alkali burn, radiation keratitis, SLK, trachoma and after LASIK treatment [401,428,435–439]. In keeping with this, a decrease in MUC5AC staining has been shown by immuno-fluorescence in conjunctival impression samples from DED patients [440] and the expression of conjunctival, MUC5AC mRNA was also significantly decreased in SSDE [383,441], NSDE [427], and in patients with tear film instability [425]. MUC5AC mucin protein levels have also been reported to be decreased in human tear samples from patients with unspecified DED [384] and in patients with severe SSDE [383], and also in the mild DED of visual display terminal (VDT) users [442]. Versura et al., using an immunogold technique, demonstrated a decreased expression of sialic acid, *N*-acetyl-glucosamine and *N*-acetyl-galactosamine in the goblet cells of DED patients [443].

5.5.2.3. Lid Parallel Conjunctival Folds (LIPCOF). Lid parallel conjunctival folds (LIPCOF) are due to a redundancy of the bulbar conjunctiva and loss of adherence to the episclera that draws the conjunctiva into a series of folds, above the lower lid margin. It is likely that they result from the same general mechanism that leads to age-related bulbar conjunctival folds elsewhere on the ocular surface (conjunctivochalasis), which have a frictional relationship to blinking [343]. LIPCOF can be identified using slit-lamp biomicroscopy and white light, with the patient in primary gaze and measured at the lower lid margin at points directly below each nasal and temporal limbus [343]. Most recently, optical coherence tomography has also been used to quantify the degree of LIPCOF [444]. Using routine slit-lamp biomicroscopy, the number of conjunctival folds present above the inferior lid, are assessed relative to the height of the tear meniscus [445]. Of note, LIPCOF

disappear when the lower eyelid is retracted and then reappear after a few blinks when lid position is restored. LIPCOF are thought to result from inflammatory elastic fibre degradation, possibly involving MMPs [446], or from mechanical friction influencing lymphatic flow [447]. Their presence has good positive predictive value for DED [446,448,449].

5.5.3. The lids

5.5.3.1. Marx's line and the mucocutaneous junction. Marx's line is a vital dye staining pattern of the epithelium, located directly behind the mucocutaneous junction (MCJ) of the lid margin [36,76,104,341,450]. (Figs. 7 and 8) It can be demonstrated throughout life on the upper and lower lid margins, extending from the outer canthi to the punctal regions. In youth it is only a few cells wide, but it broadens with age [341] and, together with the MCJ takes an increasingly irregular course.

At the mucocutaneous junction, the epithelium changes from a hydrophilic, water–wettable, parakeratinized conjunctival epithelium [40] to the keratinized, hydrophobic epithelium of the lid margin skin. The tear meniscus overlies this hydrophilic epithelium and is pinned to the MCJ at this apex, marking its location. Knop prefers to regard the whole of this parakeratinised zone as the MCJ, stretching from the point at which skin keratinization ends, to the posterior border of the lid margin, or 'crest' (Fig. 9) [40].

It has been postulated that during the interblink interval differential effects of evaporation lead to a gradient of tear molarity with a hyperosmolar peak at the tip of the apex. This is suggested to encourage increased epithelial turnover immediately behind the MCJ, incomplete differentiation of the surface epithelial cells and an immature glycocalyx, which accounts for the uptake of stain referred to as Marx's line [163,451]. An argument against this solute gradient hypothesis is that hyperosmolarity should be cleared

Fig. 8. Marx's line of the upper lid of a young adult, stained with lissamine green. (courtesy of N Yokoi).

when the tears are refreshed at each blink. However, a number of recent reports suggest that there is not full apposition of the lid margins with every blink [322,343,452] and application of Navier-Stokes equations to tear dynamics at the meniscus, instead of lubrication theory, predicts a distinct absence of fluid flow and hence of convective mixing, at the meniscus apex, adjacent to the contact line [411]. This would tend to preserve evaporation-dependent hyperosmolarity at this site. Increased permeability at the site of Marx's line could permit the diffusion of proteins of at least 20Kd and provide a route for pro-inflammatory cytokines such as IL-1β, IFN-γ, TNF-α and MMPs to reach the terminal meibomian ducts.

Since IL-1β and IFNγ are able to induce the expression of cornified envelope precursor proteins in epithelial cells [453], their delivery over many years, could contribute to hyperkeratinization at this site, a key feature of MGD. This is supported by the findings of Yamaguchi et al. who reported an age-related, forward movement of the line which was positively correlated with both the

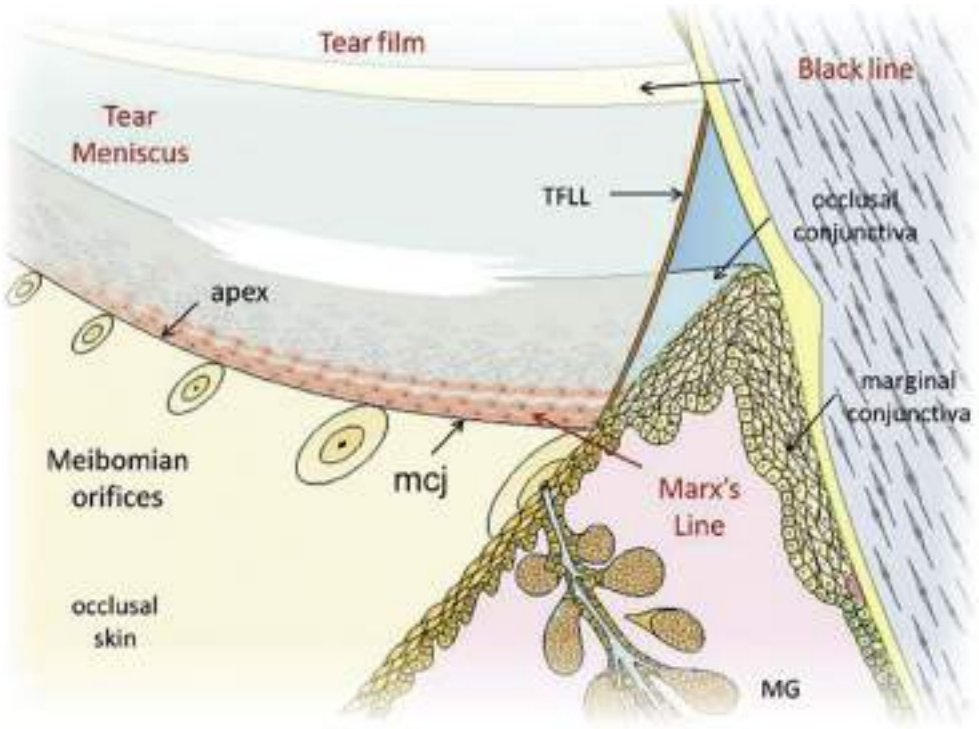

Fig. 7. Schematic view of the lower tear meniscus and lid margin. The meniscus overlies and wets both the occlusal part of the marginal mucosa and the adjoining surface, in contact with the globe. The peripheral apex of the meniscus is pinned at the mucocutaneous junction (MCJ) which forms the boundary between the stratified squamous keratinized epidermis of the lid margin skin and the stratified squamous, parakeratinized occlusal conjunctiva. It is located directly behind the meibomian gland orifices. The row of stainable epithelial cells that make up Marx's line lie under the apex of the tear meniscus, immediately behind the MCJ. (from Bron, A. J., et al. (2011). "A solute gradient in the tear meniscus. I. A hypothesis to explain Marx's line." Ocul Surf **9**(2): 70–91 - with permission) [163].

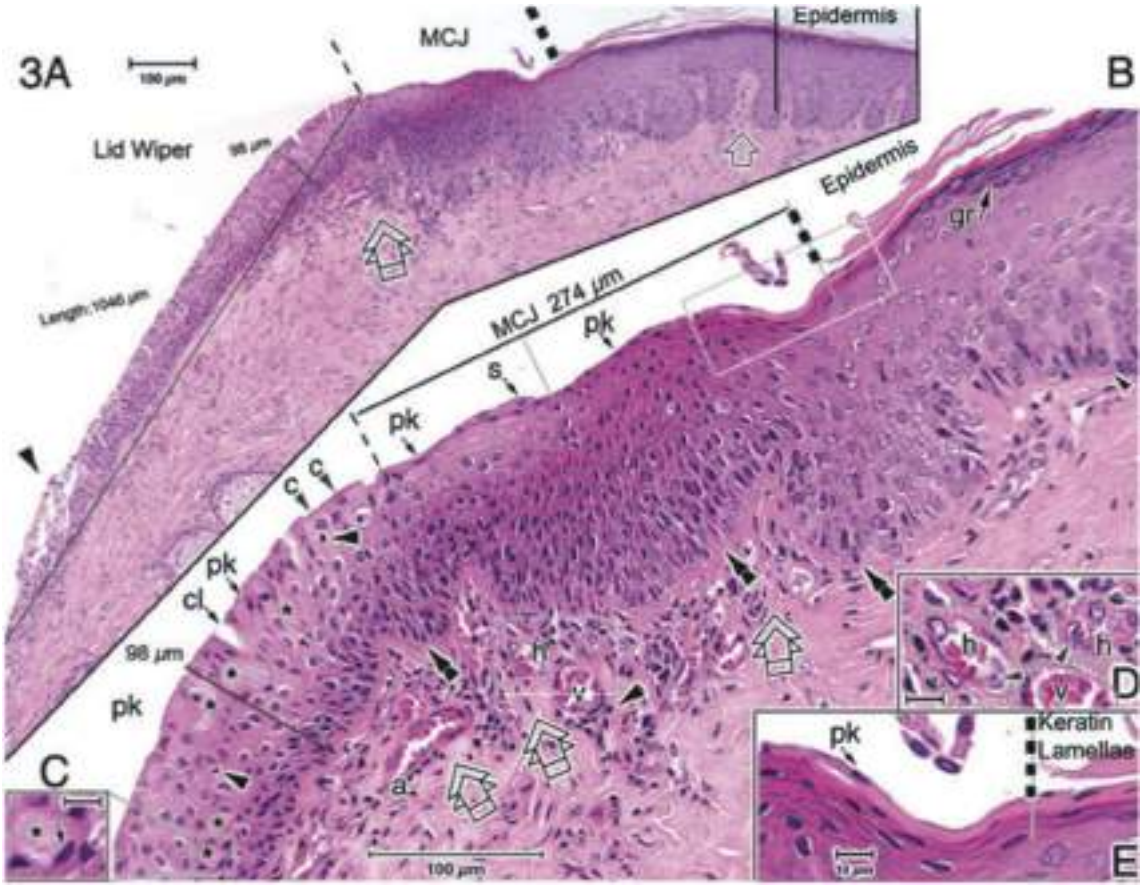

Fig. 9. According to Wolff (1946) [76] the marginal conjunctiva is a transitional zone between the skin and the conjunctiva proper, extending posteriorly for about 2 mm, from the mucocutaneous junction (MCJ) anteriorly, past the posterior lid margin and onto the tarsal plate, ending at the subtarsal fold. This H and E section passes through the posterior border of the upper lid margin in the mid-temporal region (A–E). In this figure, the MCJ is described by Knop et al. [36], as a *zone*, here, 274 μm wide, extending from the sharp edge of the cornified epidermis, to the 'crest' of the of mucosal epithelium, corresponding to the posterior lid margin. The Pathophysiology Subcommittee prefers to describe the MCJ as a line of junction between the epidermis and mucosal epithelium, marked clinically by the apex of the tear meniscus (See text and Fig. 7), in which case this stretch of transitional, mucosal epithelium is referred to as the occlusal zone of the marginal mucosa (Bron et al., 2011) [163]. In this section, Knop et al. [36] describe a continuous anterior zone (150 μm wide – B, grey line) of parakeratinised (pk) cells, followed by a zone of discontinuous pk cells interspersed with ordinary squamous (s) cells. On the tarsal plate, posterior to the crest (narrow dashed line in A, B), is the lid wiper region, which forms a thickened, cushion-like structure, is composed mainly of cuboidal cells, some columnar cells, and also goblet cells (asterisks in B), some of which reside in crypts. Here it reaches a maximal thickness of 98 μm and extends for a distance of about 1000 μm (A) to reach the sub-tarsal fold. Additional features include: a few intraepithelial lymphocytes (arrowheads in B), occasional small clefts (cl in B), vessels, including high endothelial venules (h), ordinary venules (v), and arterioles (a) underneath the MCJ, better seen in higher magnification (D, Scale bar: 100 lm (A,B); 10 lm (C–E). (gr in B = Granular layer). (from Knop, E., et al. (2011). "The lid wiper and muco-cutaneous junction anatomy of the human eyelid margins: an *in vivo* confocal and histological study." J Anat **218**(4): 449–461. – with permission) [40].

meibography scores and the quality of expressed meibum, implying an association with MGD [454].

In the presence of MGD, the line of stain may broaden and advance to involve the region of the meibomian orifices [454], or in the upper lid, with DED and contact lens wear, may broaden posteriorly to merge with LWE [344]. The lower lid may be similarly affected [346].

Of those agents known to be increased in concentration in dry eye tears, TNFα and neutrophil elastase, can cause shedding of glycocalyx mucins such as MUC 16 [241], and could increase the population of cells accessible to dyes, thereby broadening Marx's line. MMP9 causes proteolysis of tight junctional proteins such as zona occludens-1 and occludin [309,314,315] and could increase access to the paracellular compartment of the epithelium and the terminal meibomian ducts.

5.5.3.2. Lid-wiper epitheliopathy. LWE, the name given to the region of staining on the lid wiper epithelium, and considered to result from friction-related damage [324,342,344], has been demonstrated to affect both the upper and lower lid [346]. Although, for the upper lid it has usually been attributed to blinking, gaze movements also generate a relative movement between the lids and the globe and may be hypothesized to contribute frictional wear and tear to the LW region. Additionally, the initiation of a horizontal saccade is frequently accompanied by a blink, so that they are often combined in everyday viewing conditions [455].

The epitheliopathy can be demonstrated with rose bengal, lissamine green or fluorescein when it is seen as a narrow, irregular patch of staining of the lid-wiper region affecting either the upper and/or lower lid margin particularly in its central part. It is relevant that, during the blink, although the angular velocity of the upper lid margin is the same along the length of the lid, its linear velocity is highest at its centre, which traverses the full width of the lid aperture, while traversing least distance at its medial and temporal. Therefore the opportunity for frictional damage to the lid or globe is always greatest in the mid-zone of the palpebral aperture, relating more to the cornea than the bulbar conjunctiva. Because the lid-wiper zone makes a narrow band of contact as it traverses the palpebral aperture, the impact of shear stress will be concentrated more on the LW epithelium than the corneal epithelium or globe [324].

Korb et al. [324] compared the frequency of upper LWE in asymptomatic subjects without dry eyes, with that in a group of symptomatic DED patients, using sequential staining with a fluorescein/lissamine green combination. The epitheliopathy was graded on a scale of 0–3 using the horizontal lengths and average sagittal widths of the stained wiper. They found a LWE frequency of 16% in asymptomatic subjects, with 14% of Grade 1, 2%, Grade 2, and 0%, Grade 3. In symptomatic patients, 88% had LWE, with 22% Grade 1, 46% Grade 2, and 20% Grade 3. The overall prevalence of LWE was six times more in the DED group and the prevalence of LWE Grade 2 or greater was 16 times greater in DED patients than in controls ($P < 0.0001$).

In a study by Shiraishi et al. [346] the prevalence of lower-LWE was found to be significantly higher (39.5%) than upper-LWE in non-CL wearers (12.0%: $P < 0.001$) and the prevalence of both upper- and lower-LWE were significantly correlated with age ($P < 0.001$) but not sex or breakup time.

At first sight this is a surprising finding, since, while both lids are exposed to the frictional action of the globe during horizontal saccades, only the upper LW is exposed to extended friction during the blink, since the excursion of the lower lid during blinking is small. This puzzle was addressed in a further study, in which lid movement and globe displacement were followed during spontaneous blinking and lid pressure against the globe measured with a blepharo-tensiometer. The authors found no relationship between eyelid pressure and any grade of upper-LWE, but eyelid pressure in eyes with grade 3 lower-LWE (27.9 ± 2.8 mmHg) was significantly higher than with grade 0 lower-LWE (19.7 ± 1.3 mm Hg; $p < 0.05$). Also, lower eyelid pressure was significantly correlated with the length of horizontal movement of the lower eyelids during blinking ($p < 0.05$) and the degree of posterior movement of the eye globe ($p < 0.05$). The authors concluded that one cause for the development of lower-LWE was the application of a higher pressure from the lower eyelid.

It is possible, too, that another factor operates. The upper lid and globe move together in vertical gaze, but not synchronously – there is a small relative movement between them. In contrast, the lower lid moves only a small amount in vertical gaze, so that there is a rapid movement of the globe in relation to the lower lid wiper, a potential source of significant friction during reading and working at a computer.

6. Inflammatory responses in dry eye – innate and adaptive immunity

In general, immune processes are classified as innate or adaptive. Innate immune responses are considered to be fast and non-specific, whereas adaptive responses evolve over time, are specific and generate memory. These processes occur at the same time and cross-talk between each system is critical for the development of an effective response.

The immune responses of the ocular surface are not different from those at other mucosal surfaces [234,456,457]. The microenvironment of the ocular surface is constantly exposed to environmental challenges and maintains surveillance over desiccation, microorganisms, pollution and allergens and other noxious agents. Insults may be either acute or chronic and the immune system deals with them accordingly.

6.1. Innate immune responses in dry eye disease

6.1.1. Barriers and inflammatory signals

A critical component of the innate immune system is to provide a physical barrier between the eye and the external environment, for instance preventing the adherence of microorganisms and passage of toxins across the surface epithelia. Elements that accomplish this include the gel mucin of the tears, the glycocalyx, the epithelium itself and a stream of antimicrobial defense proteins including lactoferrin, lysozyme, lipocalin and trefoil peptides and surface molecules such as the defensins (α and β) [159,458–461]. However, the corneal and conjunctival epithelia are considered to be the “gate keepers” of the ocular surface [462].

This defense system can be hijacked by the hyperosmolar stress of DED, through the activation of MAPK that in turn activate the master regulator, NF_KB, production of IL-1 (chiefly) and of TNF-α. These have major downstream effects by inducing a cascade of other mediators and cellular signals that amplify the inflammatory immune response. IL-1 and TNF-α then up-regulate MMP-9 production by corneal epithelial cells, which is associated with disruption of the epithelial corneal barrier [316].

An aspect of the innate defense system involves activation of pattern recognition receptors (PRRs) such as the Toll-like receptors (TLRs) and the NOD-like receptor (NLR) that mediate cytosolic, inflammasome inflammation. They both participate in the inflammatory response of DED [463]. Stimulation of these receptors is associated with the up-regulation of IL-1, TNF-α and also of IL-6.

6.1.2. Recruitment signals and inflammatory cells

Experimentally, the expression of IL-1, TNF-α and IL-6 by ocular surface epithelia is critical to the inflammatory response of DED. A step in the amplification process is the generation of signals that recruit both innate and adaptive inflammatory cells to the site of inflammation. These signals may be soluble or membrane-bound and include chemokines and adhesion molecules [464]. In an experimental model of DED, induced by desiccating stress (DES) and scopolamine, the increased expression of inflammatory cytokines by the cornea and conjunctiva was greatly reduced in IL-1 receptor knockout mice [465].

Chemokines produced at the ocular surface during an inflammatory response (eg. CCL3, CCL4, CCL5, CXCL9, CXCL10, and CX3CL1, [306, 466–469] can bind macrophages, dendritic cells, neutrophils and activated T cells in which the respective chemokine receptors are up-regulated [470].

The other critical step in the homing of these inflammatory cells to the ocular surface is the expression of endothelial adhesion molecules [464] such as the intracellular adhesion molecule-1 (ICAM-1) which is expressed by conjunctival and corneal epithelium and by blood vessel endothelium in DED [471]. ICAM-1 is an adhesion molecule that binds to inflammatory cells expressing the ligand, integrin leukocyte functional antigen 1 (LFA-1), causing rolling, transmigration and activation at the site of inflammation and in lymphoid organs [464,472]. Such molecules, located at the surface of the eye, represent potentially accessible therapeutic targets. Lifitegrast, an ICAM inhibitor, has recently been approved by the United States Food and Drug Administration for the treatment of DED [473].

Three distinct cell types are involved in the innate inflammatory response, neutrophils, NK cells and monocyte/macrophages. The role of neutrophils in DED is an area of current investigation and the importance of NETs was alluded to earlier (see Section 4.8). However, in a DES model of DED, depletion of neutrophils led to increased $CD4^+$T cell activation and increased corneal staining, demonstrating that at some stage, neutrophils may play a protective role [474].

Recent studies of DED models suggest that NK cells may make a significant contribution in the pathogenesis of DED [105,475–477]. The recruitment or activation of resident ocular NK cells has been associated with the increased production of inflammatory cytokines that include IFN-γ, IL-6, IL-17 and IL-23, which stimulate macrophages, antigen-presenting cells (APCs) and auto-reactive T cells. NK cells may be an early source of IFN-γ that is responsible for the

activation and differentiation of Th1 T cells, induction of costimulatory signals by APCs and which, itself, is a key inflammatory cytokine causing conjunctival epithelial damage and goblet cell loss [475,478].

The infiltration of the conjunctiva by monocytes, that differentiate into tissue-associated macrophages, is a notable feature of murine DED. Indeed, infiltration by $CD11b^+$ (monocyte/macrophages) and $CD14^+$ macrophages correlates with disease progression in a mouse model of autoimmune lacrimal keratoconjunctivitis [479]. Monocytes can differentiate into two types of tissue macrophage; M1 cells are associated with pro-inflammatory responses whereas M2 cells are regulatory. DED has been shown to induce an M1 phenotype in a desiccating stress model [480].

6.1.3. The features of innate immunity

Other elements that are considered to be part of the innate immune system are the gamma/delta (γ/δ) T cells and the complement system. γ/δ T cells are frequently found in close proximity to epithelial cells, including the conjunctival epithelium [476]. γ/δ T cells can produce IL-17 [481] in the ocular surface but their specific role during DED remains unknown. Studies investigating the role of complement, in the ocular surface inflammation of DED, are limited to observations in animal models where nude mice receiving serum from mice with dry-eye, develop DED associated with the recruitment of inflammatory cells and cytokines by the activation of C3a/C5a and C3b/C5b and the formation of the membrane attack complex (MAC) [482]. These observations were also supported by the demonstration of C3b expression in the conjunctiva of diseased mice and the dampening of disease by neutralizing the complement pathway with the systemic administration of cobra venom [482].

6.2. Adaptive immune responses of the ocular surface

6.2.1. Initiation of adaptive immunity by antigen presentation

The presence of $CD4^+$ T cells at the ocular surface in DED and the successful treatment of ocular surface inflammation with topical cyclosporine suggested a potential role for adaptive immunity in DED [483]. Initiation of an adaptive immune response requires that antigens at the site of inflammation are processed and presented by professional APCs that migrate to regional lymphoid tissue to activate and expand antigen-specific effector T cells. Although the antigen or antigens that initiate this response in DED are not known, the expression of auto-antigens is hypothesized to be a key trigger to the inflammatory epitheliopathy in Sjögren syndrome. This is regarded as the basis for the production of autoantibodies to type 3 muscarinic acetylcholine receptor (anti-M3R Ab) and the Kallikrein family of proteins including Klk1 and Klk13 [482, 484–486] and the generation of autoreactive T cells [487].

Evidence for ocular surface antigen presentation as the initiating step in the adaptive immune response has come from the correlation between the accumulation of mature CD11c APCs, the activation of antigen-specific $CD4^+$ T cells in draining lymph nodes during desiccating stress, and the reduction of $CD4^+$ T cell infiltration in animals depleted of ocular surface macrophages and APCs [479]. As ocular surface tissue from inflammatory conditions is characterized by the up-regulation of MHC II and other stimulatory signals, the activation of circulating, primed T cells that are recruited to the cornea and conjunctiva of patients with DED is another plausible pathway of antigen presentation in the generation of local adaptive immune responses [471,482,488].

6.2.2. Lymphoid tissues and the ocular surface

Although the spleen is considered to be the main lymphoid tissue responsible for the immune-regulation of intraocular antigens, its role in the immunity of ocular surface inflammation is not considered to be dominant [457]. Also, the role of the thymus in the regulation of the ocular surface immune response is poorly understood. However, evidence from DED in animal models and patients with ocular graft-versus-host disease (GVHD) GVHD in which thymic damage is caused by conditioning before hematopoietic stem cell transplantation, suggests that central tolerance, regulated by the thymic environment, may be important in ocular surface immunity [489].

6.2.2.1. Conjunctiva-associated lymphoid tissue, or. As in other mucosae, such as the gut, the conjunctiva is equipped with local, stromal collections of lymphoid tissue that are involved in the induction of mucosal tolerance and the regulation of inflammation and immune defense at the ocular surface [490]. These foci constitute CALT, the local equivalent of the MALT collections in mucosae throughout the body [118]. They form part of the immune lymphoid circuit.

CALT foci have access to the epithelial surface, and germinal centre/follicle formation has been identified in response to local antigen exposure. Evidence for both homeostatic and pathological responses to protein, microbes and microbial products has been demonstrated in animal models and postulated to occur in humans [491–493].

6.3. Inflammation, the meibomian gland and dry eye

A striking feature of the human meibomian gland is its apparent resistance to inflammation and infection. For example, there is no peer-reviewed evidence of inflammation or infection in this tissue in obstructive MGD [36,494,495,976]. Further, exposure of human meibomian gland epithelial cells to a bacterial toxin (i.e. lipopolysaccharide [LPS]) does not induce the expression of pro-inflammatory gene ontologies, other than that associated with Toll-like receptor signaling [496]. In contrast, LPS stimulates a marked upregulation of genes linked to defense, cytokine and chemokine production, chemotaxis, Toll-like receptor signaling pathways and inflammatory and immune responses in immortalized human corneal and conjunctival epithelial cells [496]. It is possible that this apparent resistance to inflammation and infection within the human meibomian gland is due to the presence of innate anti-inflammatory and anti-infective factors.

In support of this hypothesis, the most highly expressed gene in the human meibomian gland encodes for leukocyte-associated immunoglobulin-like receptor-1 (LAIR-1) [505]. LAIR-1 is an inhibitory receptor that suppresses immune cell activation and reduces pro-inflammatory cytokine production [497,498]. Expression of the LAIR-1 gene is upregulated during human meibomian gland epithelial cell differentiation [43], as are those for uteroglobin (suppresses inflammation [1202]), phospholipase A2 (kills gram-positive bacteria and is a key bactericide in human tears [499]) and CCL28 (has antimicrobial activity against gram-positive and gram-negative bacteria [500]). Recently, investigators had also found that human meibomian gland epithelial cell lysates inhibit the growth rate of the Gram-negative bacteria, *Pseudomonas aeruginosa*, *in vitro* [501]. In addition, human MGD is associated with a significant increase in intraglandular transcripts for [a] S100 calcium binding proteins A8 and A9 (S100A8/9, also called calprotectin; in high concentrations this heterodimer has anti-inflammatory and anti-microbial functions and makes epithelial cells more resistant to bacterial invasion [502–504,1217]; [b] peptidase inhibitor 3, skin-derived (also called elafin [1203], inhibits bacterial infection [1202]); and [c] S100A7 (also called psoriasin, an antimicrobial peptide [1202]) [505].

These findings do not mean that human meibomian glands

cannot become inflamed or infected. A single meibomian gland, for instance, may develop a chalazion (i.e. inflammation associated with a blocked gland), that may become secondarily infected. Further, LPS can induce leukotriene B4 secretion from human meibomian gland epithelial cells [60], and isotretinoin can induce the expression of some inflammatory mediators in these cells [45]. However, neither inflammation nor infection is a characteristic of obstructive MGD, which affects multiple glands [36,506].

7. Research in animal models and cellular models *in vitro*

The use of animal models to study DED is a source of hypothesis generation, which enables pathological mechanisms to be examined in relation to clinical disease. The influence of risk factors such as age, sex and environment can also be explored and, in the case of Sjögren syndrome, the effect of immune dysregulation on immune tolerance. A general review animal models of DED is provided by Schrader et al. [507].

7.1. Animal models of Non-Sjögren dry eye

7.1.1. Overview

The Subcommittee has focused on the following two models: The desiccating environmental stress model (DES) involves exposure to a combination of low humidity and increased airflow with or without muscarinic blockade. The muscarinic receptor blockade model (SCP) involves systemic injection of scopolamine to suppress parasympathetic nervous system function and thereby inhibit lacrimal gland secretion.

There are acute and chronic DES models and there is great interest in recovery from damage after removal of the initiating cause because of its relevance to self-perpetuating disease.

7.1.2. The desiccating stress model

The desiccating stress or desiccating environmental stress (DES) model, first described by Dursun et al. [508] and later modified by several investigators [509,510] combines a high air flow, a low relative humidity and cholinergic blockade, to impair lacrimal gland secretion. It has become a standard DED model and has been used to study pathogenesis of DED, and potential therapies [511–515] The DES model recapitulates several features of DED, including corneal staining, conjunctival goblet cell loss, conjunctival infiltration with $CD4^+$ T cells, increased cytokines in tears, and apoptosis of ocular surface epithelium [378,427,516–518]. Interestingly, DES induces profound epithelial changes, with increased production of cytokines, chemokines and MMPs that precede the initation of the immune response [309,378,519], but a significant modulation of the immune system occurs (described below).

Another feature of DED is activation of MAPK that include extracellular signal regulated kinases, JNK and p38 MAPK. Increased levels of active, phosphorylated JNK1 and JNK2 in ocular surface epithelia treated with hypertonic saline *in vivo* and in cultured human corneal epithelial cells exposed to hyperosmolar media has been reported [309–311]. Furthermore, JNK2 but not JNK1, appears to mediate desiccation-induced corneal epithelial disease (by stimulating production of MMP-1, MMP-9, and cornified envelope precursors) since JNK2KO mice were resistant to DED-induced changes [520].

7.1.2.1. Initiation of dry eye by desiccating stress. Disruption of afferent and efferent immunoregulation of the ocular surface is recognised as a major process underlying DED inflammation [234,521]. Pro-inflammatory cytokines (IL-1, TNF-α and IL-6) and chemokines, released from stressed ocular surface epithelia cause epithelial damage and activate antigen-presenting cells (APCs) and NK cells [234,475]. In addition, activation of an innate, NK cell response not only damages target tissues but promotes APC maturation through IFN-γ [475,476,522]. These activated APCs on the ocular surface migrate to draining lymph nodes (DLNs) via newly-formed lymphatic vessels (facilitated by VEGF-C and VEGF-D) [512,523,524] and help to prime naïve T cells in the draining lymph nodes (DLNs), leading to the activation and expansion of IFN-γ-secreting $CD4^+$ T (Th1) and IL-17-secreting $CD4^+$ T (Th17) cells [476,479,525,526]. These unrestrained effector T cells home to the ocular surface via blood vessels under the influence of increased levels of local, ocular surface chemokines [516,527,528]. Increased levels of IL-17 and IFN-γ from activated T cells on the ocular surface lead, in addition, to disruption of the epithelial corneal barrier and to decreased conjunctival goblet cell density [516,523,529].

Although both $CD4^+$ and $CD8^+$ T cells take part in the adaptive immune response to antigens, $CD4^+$ T cells predominate at the ocular surface in chronic DED [487]. Naïve $CD4^+$ T cells in the lymphoid tissue differentiate into four functional phenotypes designated according to their major cytokine products. These are classified as Th1, Th2, Th17 and T regulatory (Treg) lymphocytes. Upon antigen presentation, the cytokine milieu present at the time of T cell activation is a major determinant of the final outcome of differentiation. The resolution of the adaptive immune response is mediated by the elimination of these effector $CD4^+$ T cells by activation-induced apoptosis at the site of inflammation, which results in the generation of memory, antigen-specific T cells characterized by the differential expression of surface markers including $CD45RB^+$, $CD44^+$ and $CD69^+$.

Th2 $CD4^+$ T cells have been associated with the development of allergic responses in the ocular surface and also have a role in maintaining homeostatic levels of conjunctival goblet cells [105]. Niederkorn and colleagues elegantly showed that adoptive transfer of CD4+T cells into immunodeficient mice, primed *in vivo* during DES, recapitulates the DED phenotype observed in donor mice [487]. Mice developed DED, with lacrimal gland infiltration, corneal staining, goblet cell loss, $CD4^+$T cell infiltration in the conjunctiva and production of cytokines and matrix metalloproteinases.(Niederkorn, Stern et al., 2006) (Fig. 10) [487].

Th1 $CD4^+$ T cells are the classic pathogenic T cell subset associated with the generation and progression of immune-related dry eye disease [478]. These populations of effector T cells are differentiated by the presence of IL-12 and are characterized by their production of IL-2 and IFN-γ at the site of inflammation. The production of IFN-γ by Th1 $CD4^+$ T cells is a major determinant of the pathological changes observed in the ocular surface of dry eye patients including epithelial cell death, loss of goblet cells and epithelial cells, and squamous cell metaplasia [516,530–532]. The recruitment of Th1 $CD4^+$ T cells to the ocular surface is regulated by their expression of LFA-1 and its interaction with ICAM expressed in the ocular tissues of patients with DED [472]. Moreover, their increased expression of CCR5 and CXCR3 makes them responsive to the chemoattractants CCL5 and CXCL10, which are also produced in the inflamed ocular surface in response to DES [467,527]. IFN- γ has been shown to antagonize IL-13 in the lung and the gut and this is also true on the ocular surface. As noted, IL-13 promotes goblet cell homeostasis in physiological conditions [105], while IFN-γ promotes goblet cell apoptosis [478,530]. IFN-γ knock-out mice are resistant to desiccating stress; however, when reconstituted with IFN-γ, they develop goblet cell loss similarly to wild-type mice [478]. Adoptive transfer of $CD4^+$ T cells from donor mice exposed to DES that received anti-IFN-γ were less pathogenic to immunodeficient mice recipients, yielding less corneal apoptosis and greater number of PAS^+ filled goblet cells [530]. Mice that received subconjunctival injections of anti-IFN-γ antibody showed decreased

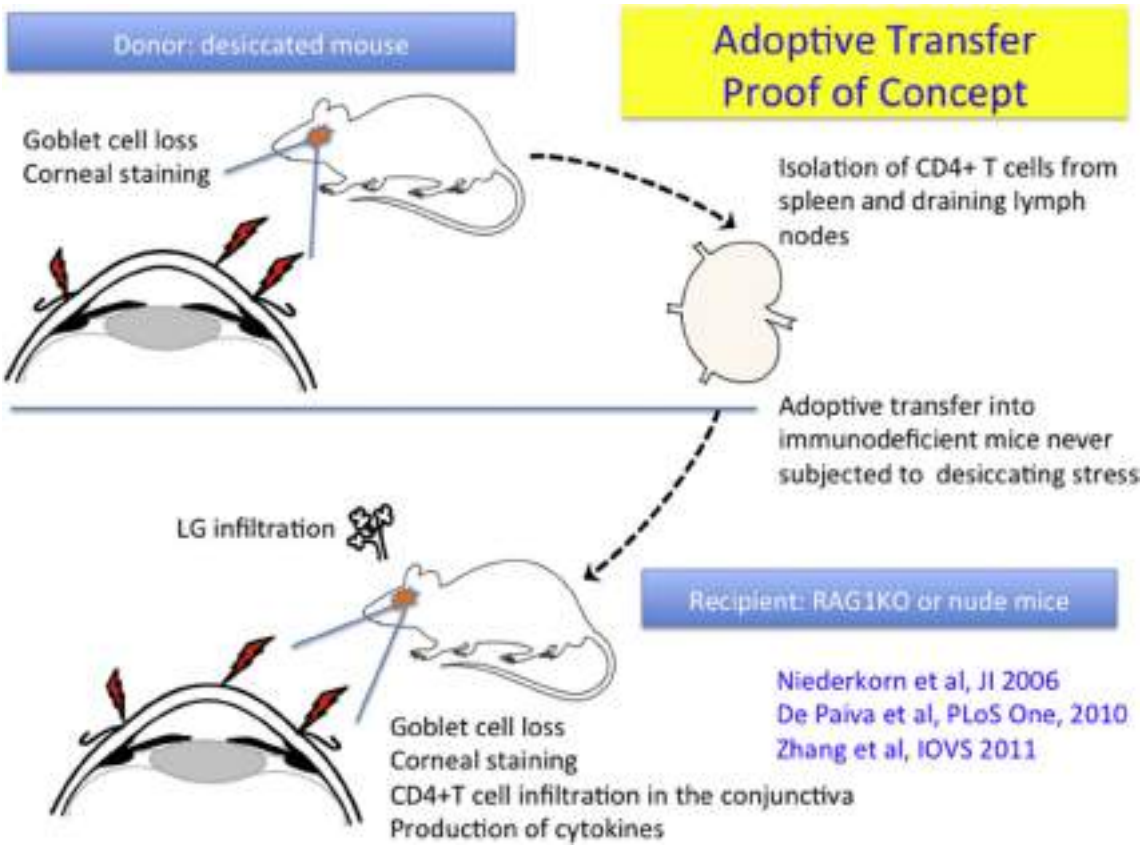

Fig. 10. Schematic depiction of adoptive transfer experiments. Mice that were subjected to desiccating stress (DES) had goblet cells loss, corneal staining and CD4+T cell infiltration in the conjunctiva. CD4+T cells were isolated from spleen and draining cervical lymph nodes using magnetic beads and adoptively transferred into immunodeficient mice that were never exposed to DES. Adoptively transferred recipients of $CD4^+$ T cells developed DED, with lacrimal gland (LG) infiltration, corneal staining, goblet cell loss, $CD4^+$T cell infiltration in the conjunctiva and production of cytokines and matrix metalloproteinases, recapitulating the DED phenotype observed in donor mice.

corneal and conjunctival apoptosis [530].

Th17 $CD4^+$ T cells are the prototype of auto-reactive T cells associated with chronic inflammatory diseases. The presence of IL-17 in the tear fluid of patients with DED and their localization to the ocular surface in animal models of DED induced by DES or auto-immune mechanisms, support the role of these cells in the progression of disease [458,516,526]. It has also been demonstrated *in vitro* that differentiation of naïve $CD4^+$ T cells into Th17 cells is possible, by co-culturing T cells with corneal epithelial cells that most likely are a source of IL-17 during ocular surface inflammation. Just as the recruitment of Th1 $CD4^+$ T cells is enhanced at the ocular surface in DED, CCL20 is expressed in the ocular surface of animals exposed to desiccating stress and Th17 T cells expressing CCR6 are potentially responsive to this signal and can be recruited to the ocular surface [516,527]. IL17 causes damage to the corneal epithelium directly and through the up-regulation of MMP9 and MMP3 and the inhibition of protection by Treg cells [516,529]. Both IL-17 and IFN- γ have been found to be elevated in tears and in conjunctival impression cytology of DED patients [235,441,458].

Tregs cells are characterized by the expression of $CD4^+CD25^{+hi}Foxp3^+$ markers and their role in the maintenance of peripheral tolerance has been found to be critical in immune responses to allo- and auto-antigens in non-ocular diseases. Studies of mice undergoing desiccating stress suggests a significant role of Tregs in regulating and dampening the inflammatory response. If adoptive transfer of Tregs is performed, a significant amelioration of DED inflammation is observed and this correlates with the regulation of "ocular specific $CD4^+$T cells" [533]. $CD8^+$ T cells can also function as regulatory cells, as depletion of $CD8^+$ T cells during the initiation phases of DES generated more pathogenic T cells. A functional (but not numeric) defect in Tregs has also been shown in DES experiments [529].

The role of B cells in the adaptive responses of the ocular surface in DED remains unclear. B cells and autoantibody production appear to be implicated in the systemic and ocular manifestations of Sjögren syndrome in patients and in animal models [482]. In contrast, their role in chronic DED in non-autoimmune patients is not obvious. However, in addition to the production of pathogenic autoantibodies, the role of professional APCs and activation of auto-reactive T cells cannot be underestimated [534].

7.1.2.1.1. Distinction between models based on desiccating stress alone and those induced by scopolamine or by a combination of the two. A majority of experimental evidence supporting the DED inflammation described above has been derived from a murine model of DED that combines DES with systemic muscarinic acetylcholine receptor inhibition using scopolamine [509,535]. DED increases tear evaporation through low humidity and high airflow, and SCP induces tear deficiency by antagonizing muscarinic activity in the lacrimal glands. Low humidity alone is capable of inducing DED, but the kinetics are delayed compared to DES [536]. A recent advance in our understanding of murine DED inflammation is that DES without muscarinic blockade and SCP induce DED through different primary mechanisms [537]. DES without muscarinic blockade induced greater conjunctival CD3 (+) T-cell infiltration, and higher Th17-cell activity and Treg dysfunction than SCP, while SCP reduced tear volume to a greater extent than DES. SCP

decreased Th17 activity and increased Th2 and Treg responses without influencing Th1 activity.

It should be noted that by inhibiting cholinergic activity, scopolamine also has a marked impact on the nature of inflammatory and ocular surface responses in DED. Scopolamine would interfere with the ability of the parasympathetic nervous system to respond to cytokines released during activation of the innate immune system, and to provide a negative feedback control of innate immune responses to restore homeostasis [1220]. Scopolamine would also prevent the ability of the cholinergic anti-inflammatory pathway to counteract abnormal chronic and hyper-activated inflammatory responses [538]. As additional considerations, cholinergic neurotransmitters are known to regulate meibomian gland epithelial cells [42] and goblet cells [539], but this modulatory activity would be suppressed by using scopolamine. Further, goblet cells appear to be dependent on constant neural input from the ocular surface [540], but this communication may be hampered by scopolamine. Overall, given that the immune system and ocular surface are physiologically important in the development and recovery from DED, scopolamine's removal of a major regulatory system (i.e. cholinergic pathway) limits the physiological relevance of this SCP model for understanding immunological and ocular surface processes in human DED.

7.1.2.2. Acute versus chronic evaporative dry eye models. The existing DED models are created in the acute setting [509,535], which raises questions as to how findings in these models relate to those in the clinical setting, where DED is generally encountered as a chronic disorder. A chronic, murine DED model has recently been developed, which seeks to address this question [541]. In brief, acute DED was first induced using the same DES method for 14 days, and affected mice were then transferred to a normal humidity environment and maintained for an additional 4 months without any DES or SCP challenge. DED severity peaked at the end of the DES, and after removal from the DES, the corneal epitheliopathy gradually regressed to lower levels, but never normalized. Further, the chronic phase was accompanied by Th17 responses at the ocular surface. These findings suggest that, following the induction of acute DED, corneal epitheliopathy and inflammation may persist into a long-term chronic phase, even without continued exposure to DES.

The mice selected for use in this chronic study had almost no corneal fluorescein staining at experimental inception [542]. In contrast, untreated mice typically seem to have higher and more variable staining [82,1213,1214], as do humans (see Section 4.14.1). It will be of interest to learn whether this chronic model is reproduced in mice with higher initial degrees of corneal fluorescein staining.

7.1.2.3. An age-related dry eye model. Another chronic model of DED is the aged C57BL/6 mouse, which also develops spontaneous DED and MGD [545]. Interestingly, female and male mice showed comparable goblet cell loss but greater corneal staining was observed in female mice. Adoptive transfer of aged $CD4^+$ T cells transferred the DED phenotype to RAG1KO mice, indicating that aging leads to generation of spontaneous autoreactive T cells [545]. The findings of spontaneously activated cells in aged mice of both Th1 and Th17 phenotype warrants further investigation.

7.1.2.4. Relevance of mouse models for human inflammatory diseases. Mouse models can be invaluable to help clarify the underlying physiological and pathological processes in many human conditions. Ideally, such understanding may be translated into therapies for various human diseases. However, potential treatments that have been discovered and validated in mouse models do not always translate successfully to human therapies. This is especially true for treatments targeting inflammatory pathways. Genomic responses to inflammatory challenges have demonstrated poor correlations between different mouse models in comparison with human responses [546]. Although some studies indicate that experimental findings with mice can be predictive of therapeutic success in humans [547,548], the fact remains that almost 150 clinical trials involving investigational anti-inflammatory therapies based on mouse data have failed [546], including several potential treatments for DED [549]. Some of these clinical trials were based on data from DES [514,550,1218] and botulinum toxin [551] mouse models. These results underscore the need to show whether a given mouse model mimics, or fails to mimic, a relevant human disease [546,552,1204].

7.2. Animal models of Sjögren Syndrome dry eye

7.2.1. Introduction

Sjögren syndrome is a chronic autoimmune disorder that affects exocrine glands, notably the lacrimal and salivary glands, causing DED and dry mouth in addition to affecting other organ systems. The clinical features of human disease are elaborated in a later section of this report.

Several animal models have been used to study the pathogenesis of Sjögren syndrome and have provided insights into the disorder, including its heterogeneity (Table 8). While animal models recapitulate one or more aspects of Sjögren syndrome, no perfect model exists. This section focuses on ocular events in autoimmune murine models.

7.2.2. Animal models of Sjögren Syndrome

A review of the animal model literature shows a dichotomy in reports, with rheumatological research focused on the salivary gland as the target organ and ophthalmological research favoring the lacrimal gland. With the exception of reports in MRL/lpr, NZB/NZW and NOD mice [553–556], there is relatively little comparative information about salivary and lacrimal pathology in the same animals. Ocular manifestations are also less widely studied in the rodent model. There is a need for more studies, embracing pathological changes in both organs which could identify common pathways of glandular destruction and delineate gland-specific differences. The temporal sequence of pathological events may be a guide to cellular and molecular mechanisms. It is still unclear as to what extent ocular surface manifestations in Sjögren syndrome are secondary to lacrimal or meibomian gland involvement or to the targeting of corneal and conjunctival autoantigens.

The influence of age, disease duration and sex is important in human Sjögren syndrome and age is one of the strongest risk factors for the DED [557–561]. Similarly, in animal models, the full Sjögren syndrome-phenotype often takes time to evolve [536,562,563] Two models illustrate this well. First, as noted, non-immune C57BL/6 mice spontaneously develop a lacrimal-dependent DED, starting at post-menopausal age (6–9 months), up to an elderly age of 24 months [545]. Second, the non-obese diabetic (NOD) mouse, variant strain ($NOD.B10.H2^b$) has a mild Sjögren syndrome phenotype at 10 weeks of age, but develops severe dacryoadenitis and DED at one year of age [564]. This suggests that a defined level of immune dysregulation is required to establish the histological Sjögren syndrome phenotype, dependent on such factors as the tissue accumulation of lymphocytes, loss of T regulatory cells and/or the generation of autoantibodies. Similar to human Sjögren syndrome, the presence of histologic lesions is considered one of the most important criteria for diagnosing Sjögren syndrome in the animal model [565].

The strong predeliction of Sjögren syndrome for women has

Table 8
Murine models of Sjögren syndrome.

Age of onset	Model	Sex predilection	Main mechanism	Main organs affected	References
0–3 Weeks	TGF-β KO	♀ = ♂	Gene deletion of TGF- β	Fatal systemic autoimmunity, inclusive of LG	(Shull et al., 1992, McCartney-Francis et al., 1997) [596,1112]
	Scurfy	♀ = ♂	Deletion in the forkhead domain of Foxp3	Fatal systemic autoimmunity, inclusive of LG	(Brunkow et al., 2001, Sharma et al., 2006) [1113,1114]
4 Weeks	CD25KO	♀ = ♂	Lack of T regs; autoreactive T cells	LG; SMG; colon; ocular surface	(Sharma et al., 2006, de Paiva et al., 2010, Pelegrino et al., 2012, Rahimy et al., 2010) [536,581,1114,1115]
	MRL.lpr	♀	Autoreactive T cells; disruptive Fas-Fas ligand system	LG; SMG; ocular surface	(Jabs and Prendergast, 1991a, Jabs and Prendergast, 1991b, Toda et al., 1999) [556,574,1116]
		♀	Defect in the negative selection of organ-specific T cells	LG; ocular surface	Yeh et al., 2009, Li et al., 2008, Chen et al., 2010) [589,590,1117]
8 Weeks	IL-12 Tg	♀ = ♂	Transgenic mice with Increased expression of IL-12 in thyroid gland	SMG	(Vosters et al., 2009) [1118]
	NFKbiz KO	♀ = ♂	Epithelial apoptosis prior to lymphocytic infiltration	LG	(Okuma et al., 2013) [602]
12 Weeks	C57BL/6.NOD-Aec1Aec2	♀?	Transfer of 2 auto-reactive loci from NOD to non-autoimmune C57BL/6; milder SS phenotype than parenteral NOD	LG; SMG; ocular surface	(Cha et al., 2002, Robinson et al., 1998, You et al., 2015, Bulosan et al., 2008, Cha et al., 2004) [593,1119–1122]
	TSP1 KO	?	Lack of autologous activation of TGF-β; autoreactive T cells	LG; ocular surface	(Turpie et al., 2009, Contreras-Ruiz et al., 2013, Gandhi et al., 2013) [563,599,1123]
	NOD	♀ (S); ♂ (D)	Inbred strain that develops autoreactive T cells; a defect in T regs is controversial	SMG; LG; pancreas	(Tsubota et al., 2001, Lieberman et al., 2015, da Costa et al., 2006, D'Alise et al., 2008, Skarstein et al., 1995) [1124-1128]
	NHE8 KO	♀ = ♂	NHE are a group of membrane proteins that exchange extracellular Na + for intracellular H+	LG; ocular surface	(Xu et al., 2015) [1129]
14 Weeks	DN TGFβRII	♀ = ♂	Autoreactive T cells due to disruptive TGF-β signaling under T cell promoter	LG; ocular surface	(de Paiva et al., 2011) [1130]
16 Weeks	NOD.B10.H2b	♂	Replacement of the NOD MHC I-Ag7Idd1 diabetes susceptibility locus by the MHC I-Ab locus; milder SS phenotype than parenteral NOD in young mice.	LG; SMG; ocular surface	Yoon et al., 2008, Yamachika et al., 1998, Robinson et al., 1998, Coursey et al., 2015) [564,618,1120,1131]
	OPN-Tg	♀?	Increased expression of osteopontin	SMG	(Husain-Krautter et al., 2015) [1132]
3 months	Tet-mev1 conditional knock-out	♂	Increased mitochondrial oxidative stress	LG; ocular surface	(Uchino et al., 2012) [770]
4 Months	Neurturin KO	♀ = ♂	Defective parasympathetic innervation of their lacrimal glands	LG; ocular surface	(Song et al., 2003) [1133]
	Act1.CD40 DKO	?	Deletion of negative regulator of B cell survival	SMG > LG; skin around eyes	(Qian et al., 2008) [612]
5 months	ArKO	Neither	Estrogen deficiency due to knock-out of aromatase (converts androgens to estrogens)	No inflammation in LG or meibomian gland; increased tear volume in male, not female, compared to wildtype controls	(Rahimi Darabad et al., 2014, Darabad et al., 2013) [691,692]
6 Months	NZB/NZW F1	♀	Hybrid inbred strain, autoreactive T cells	LG; ocular surface	(Kotzin and Palmer, 1988, Gilbard et al., 1987) [1134,1135]
	ArKO	♀	Estrogen deficiency due to knock-out of aromatase	SMG	(Iwasa et al., 2015, Ishimaru et al., 2003) [714,1136]
9 months	C57BL/6	♀ = ♂	Unknown; has accumulation of autoreactive T cells	LG; ocular surface	(McClellan et al., 2014) [1137]
11.5 months	SOD1 KO	♂	Knock-out of anti-oxidant defences (superoxide Dismutase)	LG; ocular surface, MGD	(Kojima et al., 2012) (Ibrahim et al., 2014) [1138,1139]
14 months	BAFF tg	♀ = ♂	Accumulation of B cells	SMG	Groom et al., 2002) [611]
12–17 months	ArKO	♀ = ♂	Estrogen deficiency due to knock-out of aromatase	SMG	(Shim et al., 2004) [1140]

Abbreviations: ♀ - Female, ♂ - Male, S – sialodenitis, D - dacryoadenitis G- Lacrimal Glands and SMG- Submandibular Glands.

been linked in large part to the sex-related differences in, and sex steroid actions on, the immune system. It is discussed in depth in the TFOS DEWS II Sex, Gender & Hormones report [1222]. Non-Sjögren DED (NSDE) is also more prevalent in women than in men [559,566,567]. Some intriguing observations have been made in animal models. In the NOD mouse, a model of Sjögren syndrome, the susceptibility of the lacrimal gland and salivary gland to inflammatory infiltration shows a strong sex bias with sialoadenitis developing in female mice and dacryoadenitis in male mice [568]. In contrast, as in humans, inflammation is significantly greater in lacrimal and salivary glands of female MRL/lpr mice, as compared to age-matched males [569]. Unfortunately, only limited information is available concerning the sex-related differences in lacrimal and salivary glands in animal models of Sjögren syndrome [555,556,569,570].

7.2.3. A consideration of specific models

Hallmarks of Sjögren syndrome include lymphocytic infiltration, production of autoantibodies and glandular loss secondary to epithelial apoptosis. It is still unclear whether lymphocytic infiltration precedes, or is necessary for, glandular apoptosis, and the relevant antigen(s) have not been identified. The following section groups different Sjögren syndrome models according to their potential relevance to the pathogenesis of human Sjögren syndrome. The grouping is somewhat arbitrary as many models could be included in more than one category.

7.2.3.1. Infiltrating autoreactive T cells. The presence of activated, autoreactive T cells within the lacrimal or salivary glands is a pathognomonic feature of human Sjögren syndrome and the focus score (the number of mononuclear cell infiltrates containing at least 50 inflammatory cells in a 4 mm^2 glandular section), in a minor salivary gland biopsy is an integral part of the current, international classification criteria for Sjögren syndrome [565,571]. Other components are the presence of serum antibodies, and subjective and objective evidence of DED and dry mouth. Several Sjögren syndrome models which exhibit glandular T cell infiltration can be included here, including the NOD, CD25KO, Scurfy, MRL/lpr, AIRE-KO, IL-12 transgenic (Tg), C57BL/6.NOD-Aec1Aec2 (Aec), NOD.B10.$H2^b$, and Osteopontin (OPN) Tg mice.

In MRL/lpr mice, a genetically-determined disruption of the Fas-Fas ligand system leads to tissue infiltration by lymphocytes, many of which are T cells [572,573,1205]. Interestingly, the genetic background in which the Fas mutation occurs, influences the phenotype and severity of dacryoadenitis and goblet cell loss [574–576]. The two most commonly used inbred laboratory strains of mouse show a distinct bias in their ability to mount an immune response: BALB/c mice and C57BL/6 are Th2 and Th1-skewed, respectively [577]. This may explain why goblet cell density is influenced by genetic background in the MRL/lpr mutation, with goblet cell density higher in the MRL/lpr.BALB/c mouse and lower in the MRL.lpr.B6 compared to respective wild-type controls [578,579].

In both CD25KO and the autoimmune regulator gene (AIRE) KO mice, disrupted immune tolerance leads to accelerated lacrimal gland destruction, with a severe phenotype [580–582]. CD25 is the IL-2 receptor α chain, the binding arm of the heterotrimeric IL-2 receptor [583–585]. It is expressed on T and B cells. In its absence, as in the CD25 knockout (CD25KO), cells are unable to respond to IL-2, no T regulatory cells are generated, spontaneous autoreactive T cells arise and these cells fail to undergo activation-induced cell death [586,587]. CD25KO mice develop age-dependent dacryoadenitis and systemic autoimmunity. This is also accompanied by ocular surface staining, goblet cell loss and the appearance of M3R antibodies [536,582].

Mice lacking the AIRE KO develop a $CD4^+$ T-cell-mediated autoimmune disease that targets multiple organs, including the lacrimal gland and ocular surface [588]. The AIRE KO mouse in a NOD background, shows severe squamous metaplasia and ocular surface staining which parallels the level of lacrimal gland infiltration [588,589] while the same mutation in a C57BL/6 background leads to significant goblet cell loss and $CD4^+$ T cell infiltration of the cornea and meibomian periglandular region, compared to wild-type controls [590].

While there is evidence for a role for $Th17^+$ cells in corneal barrier disruption and sialoadenitis [516,526,591], its role in dacryoadenitis is still controversial. Some of the autoimmune models that have been used to investigate dacryoadenitis have both $Th1^+$ and $Th17^+$ T-cells infiltrating the lacrimal gland, making it difficult to determine the individual contributions of Th subsets (TSP-1 KO, MRL/lpr, CD25KO, and Aec). Dacryoadenitis in CD25-IL-17DKO appeared earlier and was more extensive than in the CD25KO parental strain and it was accompanied by greater IFN-γ-receptor expression and caspase 3 levels [562], suggesting that IL-17A may have a minor role in counterbalancing IFN-γ. $Th\text{-}1^+$ cells have been implicated in colitis, experimental autoimmune uveitis and Sjögren syndrome [475,478,532,562,582,592]. Both NOD. IFN-γ KO, and NOD. IFN-γ receptor KO, mice have ameliorated sialoadenitis [593] and the same pattern is observed in CD25-IFN-γ DKO [536,562]. These findings indicate that a mixture of both Th1 and Th17 cells are involved in dacryoadenitis and that therapies targeting more than one subset may be beneficial in Sjögren syndrome.

7.2.3.2. Disruptive TGF-β signaling. TGF-β is a pleiotropic cytokine involved in epithelial differentiation, mitosis, cell motility, fibrosis and immune-regulation [308]. TGF-β is critical for the induction of $CD4^+Foxp3^+$ cells, the regulatory T cells involved in keeping other cells in check [594], but also of T helper (Th) 17 cells [595]. TGF-β null mice succumb to massive systemic autoimmunity, affecting both exocrine glands, shortly after birth, making it difficult to investigate the specific role of TGF-β in Sjögren syndrome [596–598]. Two other animal models with disruptive TGF-β signaling develop moderate Sjögren syndrome with aging: the thrombospodin-1 knock-out (TSP-1) KO and the dominant negative TGF-β receptor type II (DN TGFBRII). These mice develop dacryoadenitis and ocular surface manifestations that are accompanied by Th1 and Th17 responses [563,599]. TSP-1KO mice also have anti-SSA and anti-SSB serum antibodies [563]. The dual role of TGF-β in promoting Tregs (anti-inflammatory) and generating $Th17^+$ cells can be appreciated further by subjecting the DN TGFBRII and TSP1KO to DES, where Th17 cells are involved in corneal barrier disruption [529], Interestingly, both models show a paradoxical improvement of corneal staining compared to their own baseline prior to exposure to DES [105,600]. This effect was shown to be DC-mediated in the TSP-1KO mice [600]. Polymorphism in the thrombospondin gene was found to be associated with post-refractive surgery-related, chronic ocular surface inflammation in active duty U.S. Army soldiers [601]. Future studies are needed to delineate the specific role of TGF-β in Sjögren syndrome.

7.2.3.3. Glandular apoptosis. Glandular apoptosis is another hallmark of Sjögren syndrome and it is present ubiquitously in almost all Sjögren syndrome models. It is unclear whether it follows or precedes immune infiltration, since the initiating trigger for Sjögren syndrome is unknown. A recent manuscript reported that IκB-ζ-deficient, lacrimal epithelial cells exhibited enhanced apoptosis that preceded lymphocytic infiltration, demonstrating that epithelial cell death could be an initiating factor in Sjögren syndrome [602]. There is evidence suggesting that immune cells

participate in the disorganisation and apoptosis in exocrine glands. IFN-γ has been implicated in epithelial cell loss, inducing apoptosis in salivary gland cell lines [603,604]. As noted above, NOD.IFN-γ KO and NOD.IFN-γ receptor KO mice have a lower salivary gland focus score and caspase 3 activity compared to the NOD strain [593] and CD25-IFN-γ double KO mice have significantly lower caspase 3 levels and a lesser degree of dacryoadenitis compared to the parental $CD25^{+}$KO strain [536,562]. Cultured rat and human conjunctival goblet cells are exquisitely sensitive to IFN-γ and minute concentrations will induce apoptosis [605]. In another report, IFN-γ blocked carbachol-induced high molecular weight glycoconjugate secretion and reduced goblet cell proliferation [606]. The authors concluded that this could explain the goblet cell loss and mucin deficiency in DED. These studies indicate that the glandular epithelium can function as both an initiator and bystander target of infiltrating lymphocytes.

7.2.3.4. B cells and immunization models. SS is accompanied by polyclonal B cell activity and patients with Sjögren syndrome have an increased risk of lymphoma compared to the general population [607–609]. Increased serum autoantibodies (anti-SSA/Ro 52 kDa, anti-SSA/Ro 60 kDa, anti-SSB/La, rheumatoid factor, anti-α-fodrin, antimuscarinic receptor type 3 (M3R)), have been used as diagnostic criteria [565,610] but some SS patients are serum autoantibody negative.

BAFF (B-cell activating factor) is a member of the TNF superfamily and regulates B cell survival. BAFF Tg mice, mostly used as an SLE model when younger, develop leukocytic infiltration of submandibular glands with aging [611]. Act-1 is a negative regulator of BAFF and $CD40^{+}$. Act-1tg and Act1−/− mice develop both lacrimal gland and salivary gland infiltration by B and T cells (salivary gland > lacrimal gland) and have anti-SSA and anti-SSB antibodies [612]. NOD modified mice with impaired IgG1 secretion also have an ameliorated salivary phenotype (NOD.IL4 KO; NOD.B10.H2b.IL-4 KO; NOD. NOD.B10-H2b.C-Stat6 KO) [613,614]. Recent studies using M3R KO mice immunized with M3R peptides demonstrated that M3R autoreactive T cells can transfer sialodenitis to immunodeficient mice [615–617]. They also demonstrated that, similar to NOD and $CD25^{+}$KO models, IFN-γ is critical for inducing glandular apoptosis, since adoptive transfer recipients of M3R peptide immunized, M3R-IFN-γ DKO cells had neither a significant inflammation score nor showed apoptosis [616].

7.2.3.5. Effect of DES on autoimmune responses. The autoimmune response of mice to DES has been investigated in a few instances. Yoon and colleagues demonstrated increased conjunctival infiltration and corneal staining when 16 week-old $NOD.B10.H2^{b}$ mice were subjected to DES [618]. Upon removal of DES, $NOD.B10.H2^{b}$ mice had persistently lower tear production, goblet cell loss and increased $CD4^{+}$ T cells than C57BL/6 mice, indicating that DES in a genetically susceptible strain had prolonged effects [619]. In some other strains, such as the DN TGFBRII and TSP1KO, corneal staining and goblet cell numbers improved after DES [600]. The interaction between genetic susceptibility and DES deserves further study.

7.3. Animal models of meibomian gland dysfunction

Ideally, an animal model of human MGD will demonstrate the human MGD signs, as well as the tear film and ocular surface sequelae associated with MGD and EDE. The human MGD signs would include, among others, meibomian gland orifice obstruction and orifice metaplasia (a condition defined as an atypical growth and keratinization of duct epithelium [1219]), a reduced quality and altered lipid profile of meibum, cystic dilatation of the central duct, and acinar atrophy and loss [36,494,495,620–627,1206]. In particular, evidence of hyperkeratinization of the terminal meibomian gland duct is important, given that this is a predominant feature of human MGD [36,494,620–622,624,627]. In addition, MGD, and the resulting meibum insufficiency, promote tear film evaporation, hyperosmolarity and instability and ocular surface stress, and lead to increased friction, inflammation, eye damage (e.g. corneal squamous metaplasia, loss of corneal microvilli, glycocalyx disruption) and visual impairment [1,36,190,196,549,628–632,1207]. Meibum in human DED also contains cytokeratin-positive inclusions [183].

To date, a number of animal models of MGD have been identified or created that mimic, as least in part, human MGD. Monkey [633] and rabbit [39,634–636,1208] models that present hyperkeratinization of the meibomian gland terminal duct epithelium and meibomian gland orifice obstruction have been induced by polychlorinated biphenyl poisoning [633], systemic exposure to isotretinoin [1208] and the topical administration of epinephrine [39,634–636]. A common histopathological finding in these monkey and rabbit models is an abnormal dilatation of the ducts, which feature lumina filled with keratinized materials.

Similarly, rodent models of MGD have been discovered or developed. These are either natural, or have been generated by transgenic or knockout technologies, mutations, immunization, pharmaceutical treatment, exposure to desiccating stress, or alterations in nutrition (Table 9). The resulting strains may feature a variety of phenotypes, such as ductal hyperkeratinization, obstructed meibomian gland orifices, meibum and thickened ducts containing keratinized materials, and acinar cell atrophy, aplasia and loss (see Table 9 for references).

The following three strains show many of these aspects.

First, is a model induced in HR-1 hairless mice by feeding them a special diet with limited lipid content (HR-AD) [637]. This model was developed to facilitate understanding of MGD pathophysiology. After exposure to this diet for 4 weeks, mice develop hyperkeratinization of the meibomian gland ductal epithelium, loss of meibomian gland acini, and ultimately meibomian gland atrophy. Clinical examination of these mice reveals markedly plugged (i.e. obstructed) meibomian gland orifices, telangiectasia, and a toothpaste-like meibum compared with that of a normal eyelid. Of particular interest, topical azithromycin treatment in this mouse model significantly decreases the number of plugged orifices, the keratinization of meibomian gland ductal epithelium, the meibomian gland duct thickness, and the meibomian gland atrophy [637]. Azithromycin, in turn, is known to induce human meibomian gland epithelial cell differentiation [53,638–640], and is a very common therapy for human MGD [641].

A second model is induced by treatment with isotretinoin [642], a known and significant risk factor for the development of human MGD [643–652]. Treatment of rats for 3 months with isotretinoin led to keratinization and thickening of the meibomian gland ductal epithelium, a decrease in the quantity and size of acini, and many degenerated acini and acinar cell casts in the meibomian gland ducts. These isotretinoin-elicited effects could be inhibited by treatment with dehydroepiadrosterone, presumably, according to the investigators, through conversion to androgens [642]. Topical androgens, in turn, have been reported to be effective in the therapy of human MGD [653] (see TFOS DEWS II Sex, Gender, and Hormones report [1222]).

A third model involves interference with growth hormone (GH) action [57]. These include receptor (R) antagonist (A) transgenic mice (GHA) with reduced GH, as well as GHR KO mice with no GH activity. Many of the GHA and GHR KO meibomian glands present with hyperkeratinized and thickened meibomian gland ducts that contain cornified materials, secretory acini inserting into duct walls, and poorly differentiated acini. The GHR KO and GHA mice

Table 9
Rodent models for altered meibomian (MG) and/or sebaceous (SG) gland structure and/or function.

Condition	Glandular effect	Reference
Gene knockout		
Acyl-CoA:cholesterol acyltransferase-1	MG atrophy	(Yagyu et al., 2000) [660]
Autoimmune regulatory	T cell infiltration in MGs	(Yeh et al., 2009) [590]
Barx2	MG defects	(Tsau et al., 2011) [1141]
Blimp1	Enlarged MGs	(Horsley et al., 2006) [1142]
CCAAT-enhancer-binding proteins α and β	MG atrophy, reduced number of differentiated MG acinar cells	(House et al., 2010) [1143]
CD147	Lower number of MG acini, loss of lipid-filled meibocytes	(Mauris et al., 2015) [1144]
Cu, Zn-superoxide dismutase-1	Increased oxidative stress of the MG acinar epithelium	(Ibrahim et al., 2014) [1139]
Ectodysplasin-A	No MG	(Cui et al., 2005 Wang et al., 2016 Kuramoto et al., 2011) [672,1145,1146]
Ectodysplasin-A receptor	No MG	(Naito et al., 2002) [1147]
Growth hormone receptor	MGs feature hyperkeratinized and thickened ducts containing cornified materials, secretory acini inserting into duct walls, poorly differentiated acini, and reduced MG sizes	(Liu et al., 2016a) [57]
Krüppel-like family 5 (conditional disruption)	Malformed MG	(Kenchegowda et al., 2011) [1148]
Map3k1, Dkk2, c-Jun, Egfr, Shp2, Map3k1/jnk1, Map3k1/Rhoa (systemic or conditional knockouts)	MG hypoplasia	(Meng et al., 2014) [1149]
Melanocortin-5 receptor	Decreased production of sebaceous lipids	(Thiboutot et al., 2000) [1150]
Smad4	Ectopic row of hair follicles in place of MGs	(Huang et al., 2009) [1151]
Stearoyl-coenzyme A desaturase 1	No MG	(Miyazaki et al., 2001) [1152]
Stearoyl-coenzyme A desaturase loss of function (Scd3-Cre-induced, diphtheria chain A toxin-mediated depletion)	MGD-like ocular surface effects	(Dahlhoff et al., 2016) [1153]
Tumor necrosis factor receptor -associated factor 6	Modified MGs	(Naito et al., 2002) [1147]
Transgenic or gene overexpression		
Biglycan overexpression, under control of the keratocyte-specific keratocan promoter	MG aplasia	(Hayashi et al., 2005) [1154]
c-Myc overexpression	Enhanced sebum production	Zouboulis and Boschnakow, 2001) [1155]
Ectodysplasin receptor	Enlarged MGs	(Chang et al., 2009) [1156]
Ectodysplasin-A	SG hyperplasia	(Cui et al., 2003) [1157]
Growth hormone receptor antagonist	MGs feature hyperkeratinized and thickened ducts containing cornified materials, secretory acini inserting into duct walls, poorly differentiated acini, and reduced MG sizes	(Liu et al., 2016a) [57]
Human apolipoprotein C1	MG atrophy	(Jong et al., 1998) [1158]
K14-noggin	Formation of ectopic pilosebaceous units at the expense of MGs	(Plikus et al., 2004) [1159]
Kera-rtTA/tet-O-TGFα (ectopic stromal expression of TGF-α)	Abnormal MG morphogenesis	(Dong et al., 2015) [1160]
Keratin 5 − glucocorticoid receptor	No MG	(Cascallana et al., 2005) [1161]
Rat erbB2 overexpression in basal layer of mouse epidermis, under control of the bovine keratin 5 promoter	SG enlargement	(Kiguchi et al., 2000) [1162]
Rescued fatty acid transport protein 4 null	Abnormal MG development	(Lin et al., 2013) [1163]
Smad7 or parathormone-related protein overexpression	SG hyperplasia	(Zouboulis and Boschnakow, 2001) [1155]
Mutation		
"Rhino"	MG ductal hyperkeratinization, acinar cell loss and eventual atrophy	(Jester et al., 1988) [1164]
"Rough fur" (ruf)	SG hypertrophy	(Park et al., 2001) [1165]
ADAM metallopeptidase domain 17, also called tumor necrosis factor-α-converting enzyme	No MG	(Hassemer et al., 2013) [1166]
Downless locus	MG defects	(Majumder et al., 1998 Naito et al., 2002) [1147,1167]
Elongation Of Very Long Chain Fatty Acids gene	Protruding orifices and anatomical changes in MGs	(McMahon et al., 2014) [1168]
Protein phosphatase 1 regulatory subunit 13 like	No MG	(Toonen et al., 2012) [1169]
Stratifin (14-3-3σ)	MG atrophy in aged heterozygotes	(Lu et al., 2011) [1170]
Immunization		
Murine immunization with a human monoclonal anti-DNA antibody, bearing a major Id 16/6Id	Hypertrophic MGs	Chan et al., 1995) [1171]
Natural		
"Crinkled"	No MG	(Naito et al., 2002) [1147]
Age	MG atrophy	(Parfitt et al., 2013) [1172]
Pharmaceutical		
Isotretinoin	Keratinization and thickening of MG ductal epithelium, decreased number and size of MG acini, multiple degenerated MG acini	(Ibrahim et al., 2017) [642]
Environmental & pharmaceutical		
Desiccating stress and scopolamine	Increase in MG basal cell proliferation	(Suhalim et al., 2014) [656]
Nutrition		
HR-1 hairless mice fed special diet with limited lipid content	Hyperkeratinization of the MG ductal epithelium, toothpaste-like meibum, markedly plugged MG orifices, and loss and atrophy of MG acini,	(Miyake et al., 2016) [637]
n-3 fatty acid deficiency	Decreased MG meibum secretion	(Tanaka et al., 2015) [1173]

also have significantly smaller meibomian glands, as compared to wildtype controls [57]. Given that GH levels decline with aging, it is possible that this decrease contributes to the development of age-related MGD [1216].

Recently, Jester et al. have hypothesized that the primary target in MGD is the meibomian gland, as compared to duct hyperkeratinisation [654]. This hypothesis is based on studies in age-related [655] and evaporative stress mouse models [656]. They propose that key elements of MGD are glandular atrophy through a loss of meibocyte progenitors. Jester et al. also report that desiccating stress in mice results in a phase of acinar hyperproliferation, with a change in protein to lipid ratio causing an increase in lipid viscosity. According to this view, epithelial plugs within the gland ducts do not contain fully mature keratins [654]. Obata and colleagues have also found an age-related correlation between meibomian gland acinar epithelial cell loss and aging [495,624]. In contrast, other researchers have identified keratinization, meibomian gland orifice obstruction and metaplasia associated with human MGD during both aging [627] and in general [36,494,620–622,624]. Further, large quantities of nonlipid, protein-like inclusions that stain for cytokeratins have also been identified in abnormal meibum from DED patients [183]. These inclusions may possibly represent the keratinized materials that appear in the turbid meibum of elderly people [657].

Some of the mouse models listed in Table 9 may be also useful for studies on evaporative DED and corresponding ocular surface sequelae. Consistent with this proposal are the observations that MG absence in X-linked anhidrotic/hypohidrotic ectodermal dysplasia is associated with increased tear evaporation, scarce and shortened corneal microvilli (note: which would disrupt the glycocalyx [658]), corneal defects (e.g. neovascularization, keratinization, and squamous metaplasia), and ocular surface inflammation [1145,1215]. Further, MG atrophy in acyl-CoA:cholesterol acyltransferase-1 knockout mice is associated with corneal erosions [660].

Additional mouse models that display marked alterations in sebaceous gland structure and function (Table 9) may also serve as MGD models. However, studies have yet to be performed to examine these possibilities.

7.4. The microbiome of the ocular surface

There is evidence that the gut and ocular surface microbiome may influence the occurrence of DED. The ocular surface is constantly exposed to the environment but, compared to the lid margins, is a relatively sterile site, based on studies using conjunctival swabs [661,662]. The ocular surface microbiota are regulated by numerous antimicrobial factors produced by the lacrimal glands, goblet cells and conjunctiva, which are secreted into the tears, such as lactoferrin, lysosyme, defensins α and β and IgA [459–461,663]. Lately, there has been great interest in interactions between the host and the microbiota.

The term microbiota refers to the community of microbes that inhabit a particular site, and the microbiome refers to their collective genomes. Techniques used to assess the microbiome include traditional microbial culture and cultivation-independent techniques such as polymerase chain reaction (PCR) and 16S ribosomal DNA amplification and sequencing [664–666]. The literature has been unclear as to the presence of microbiota at the ocular surface [665,667,668], with some authors finding a stable presence, which may be modified by disease, while others indicate that microorganisms are present transiently, prior to their annihilation by ocular surface defence mechanisms. Recently a consensus has been reached that the ocular surface is a paucibacterial environment, but is not sterile [665,668].

The most common microbes cultured from the conjunctival surface, using traditional culture techniques, include Staphylococcus and *Propionibacterium acnes*, while more recent techniques indicate that there are many more genera to be found [664,665,668,669]. Swabs of the lid margins yield similar species, albeit at a higher count of colony-forming units [667].

Understanding the role of the microbiome in DED is important as this could provide a potential avenue for treatment. In a study by Graham et al., the bacterial population of the posterior lid margin and lower conjunctival sac of patients with and without DED were assessed using both conventional culture and 16S rDNA PCR [669]. A significantly greater number of bacteria were detected using the 16S rDNA PCR technique as compared to conventional culture which largely yielded coagulase negative staphylococci [669]. Interestingly, bacteria that are otherwise rarely associated with the ocular surface (*Rhodococcus erythropolis, Klebsiella oxytoca, and Erwinia species*) were identified in inflammatory DED, as well as at the normal ocular surface [669]. A significant difference in the mean bacterial count was found between the control group and the moderate to severe DED groups, a result supported by others [670]. Moreover, the authors found that a reduced goblet cell density was associated with greater bacterial presence [669]. Another study, comparing the ocular, oral and intestinal microbiome of controls and Sjögren syndrome patients showed that there was no difference in the ocular microbiome between the two groups [668]. Decreased diversity was noted in both in the oral and gut microbiome and specific changes in genera were observed. There was a relative decrease in the abundance of Bacteroides, Parabacteroides, Faecalibacterium, and Prevotella, with greater relative abundances of Pseudobutyrivibrio, Escherichia/Shigella, Blautia, and Streptococcus in Sjögren syndrome patients compared to controls. Furthermore, eye and systemic severity scores inversely correlated with microbial diversity [668].

Another study reported changes to the ocular surface microbiota that occur in the early stages of Sjögren syndrome-like disease in transpondin knock-out (TSP-1KO) mice, leading to the recommendation that TSP-1 derived peptides may be a means by which to reduce commensal flora and the resulting inflammation [671].

Tools to investigate the role of the microbiome in homeostasis and disease states involve the use of germ-free mice or the subjection of mice to a cocktail of antibiotics, either in drinking water or by oral gavage. Antibiotic treatment will induce changes in the bacterial community, leading to a dysbiotic state. Recently it was reported that mice subjected to DES, that drank oral antibiotics for 14 days prior to DES had greater goblet cell loss, greater T cell infiltration and worse corneal staining than mice that were subjected to the same protocol but drank normal water [668]. 16S sequencing of the stools of these mice indicated a decrease in Clostridium and an increase in Enterobacter, Escherichia/Shigella, and Pseudomonas after antibiotics + DES for 10 days.

A germ-free environment is very detrimental to ocular homeostasis in the mouse, as it predisposes to or worsens Sjögren syndrome-like disease in non-autoimmune and genetically predisposed mice, respectively [668,672]. Non-autoimmune C57BL/6 mice raised in germ-free conditions have Sjögren syndrome-like features, inclusive of dacryoadenitis and decreased EGF concentration in tears. This was accompanied by corneal staining, goblet cell loss and pathogenic $CD4^+$ T cell infiltration [672]. On the other hand, germ-free CD25KO mice have early onset of dacryoadenitis and greater numbers of $CD4^+IFN\text{-}\gamma^+$ cells infiltrating the lacrimal glands of RAG1KO recipients. These results suggest that signals provided by commensal bacteria and/or their metabolites are capable of modulating ocular health.

7.5. Cellular models of dry eye in vitro

Ocular surface cell cultures have been used to explore the roles of multiple factors and pathways involved in the pathophysiology and possible treatment of DED. Several such cultures have also been reported to serve as DED models *in vitro*.

Three DED models use the cornea. One model utilizes rabbit corneal cultures with experimental time frames of up to 21 days [673]. Studies with this model have used optical coherence tomography (OCT) to monitor the impact of DES, with a focus on changes in corneal layer thickness and in stromal scattering properties [673]. Another DED model *in vitro* utilizes human reconstructed corneal epithelium maintained in a controlled environmental setting (relative humidity <40% and 40 °C temperature) for 24 h and up to 72 h [674]. Culture conditions are controlled to mimic DED, and thereby permit identification of biomarkers that may be predictive of corneal damage and response to treatment. A third DED model uses human reconstructed corneal epithelium to assess the effects of severe osmotic stress and associated treatment on inflammatory pathway activity and barrier integrity [675].

A fourth DED model uses immortalized human meibomian gland epithelial cells [50]. This model involves exposure of these cells *in vitro* to isotretinoin [45], a well-known risk factor for the development of human MGD *in vivo* [643–652]. Exposure of human meibomian gland epithelial cells to isotretinoin: [a] alters the expression of thousands of genes, including an upregulation of genes for some inflammatory mediators (e.g. IL-8 and IL-1β), proteases (e.g. MMP-9), MAPK signaling, lytic vesicles, apoptosis and cell death, and suppresses genes linked to DNA replication, cell cycle, RNA transport and mitochondria; [b] increases the levels of pro-IL-1β, IL-1β and MMP-9 proteins; [c] decreases the signaling of the cell growth and survival mediator, phosphoinositide 3-kinase-protein kinase B; and [d] inhibits cell proliferation and induces cellular atrophy and death (e.g. by apoptosis, necrosis and/or autophagy) [45]. It is possible that these effects may be responsible for the acinar epithelial cell degeneration and atrophy, and reduced and abnormal secretions, that occur following isotretinoin induction of human MGD *in vivo* [643–652].

8. Human disease. Etiological classification of DED

It is still useful to discuss DED under two major headings, that of ADDE and EDE (Table 5).

9. Aqueous-deficient dry eye (ADDE)

ADDE is subdivided into Sjögren syndrome dry eye (SSDE) and non-Sjögren syndrome dry eye (NSDE).

9.1. Sjögren Syndrome and Sjögren Syndrome dry eye

9.1.1. Introduction

Sjögren syndrome is a chronic autoimmune disorder characterized by immune cell infiltration of exocrine glands (exocrinopathy or epitheliitis) and systemic complications due to autoantibody production, immune complex deposition and lymphocytic infiltration of many organs [676] (Table 10). The prevalence of primary Sjögren syndrome (pSS) in the USA has been estimated to be 0.6–1%, affecting between 0.4 million to 3.1 million adults [677]. However, this estimate is different than that of another study, which reported Sjögren syndrome afflicts less than 40,000 people in the USA [678]. More recent data indicate that the average annual incidence of pSS in a physician-diagnosed, population-based cohort in the USA is 5.8 per 100,000 [679], and that the prevalence of pSS in a geographically well-defined population in Omstead County, Minnesota, is between 2 and 10/10,000 inhabitants [680]. If translatable to the USA population as a whole, this estimate would indicate that between 65,000 and 326,000 people in the USA have pSS.

Table 10
Manifestations in primary Sjögren Syndrome.

Non-specific features
Musculoskeletal symptoms, Raynaud's phenomenon,
CNS – Symptoms of fatigue
Exocrine Epitheliitis (glandular)
Lacrimal and Salivary Glands–
Other glands - pancreas
Parenchymal Epitheliitis (extraglandular)
Bronchial, hepatic, renal - peri-epithelial lymphocytic infiltration
Endocrine Gland Involvement
Thyroid, adrenals, ovaries
Immunocomplex-mediated disease
Vasculitis - affecting small vessels of the skin, nerves, kidney as a result of B-cell hyperactivity)
Lymphoproliferative
B-cell lymphoma

From Ref. [697].

Sjogren syndrome occurs predominantly in women, with a female/male ratio of 9:1[557–559, 561] and it may lead to a very severe form of DED [681]. The disease may result from a range of aberrant immune responses to environmental and viral triggers occurring in genetically susceptible individuals. Hormonal environment is also important (see TFOS DEWS II Sex, Gender & Hormones report [1222]). It involves a loss of immune tolerance, the presentation of autoantigens and dysregulation of both the innate and adaptive immune systems [682,683]. The lacrimal and salivary glands are major targets of the epitheliitis, leading to gland destruction and the key symptoms of DED and dry mouth (sicca symptoms).

Historically, Sjögren syndrome was described as a condition in its own right, pSS or as part of a systemic autoimmune disorder (secondary Sjögren syndrome - sSS), such as rheumatoid arthritis, systemic lupus erythematosus (SLE) and Wegener's granulomatosis [684]. More recently, the American College of Rheumatology has recommended that the diagnosis of Sjögren syndrome should be given to *any* patient who fulfulls the diagnostic criteria of Sjögren syndrome [565] without distinguishing it as primary or secondary, recognizing them both to be a manifestation of immune dysregulation. The Subcommittee recognizes the value of this approach but the older terminology is retained here in relation to past literature.

Symptoms of DED and dry mouth are a major feature of Sjögren syndrome that are a result, at least in part, of infiltration of the salivary and lacrimal glands by T and B lymphocytes, dendritic cells (DCs), macrophages and other mononuclear cells, leading to tissue dysfunction or destruction [683]. In SSDE the lacrimal glands are considered to be primary targets of immune attack. This is less certain for the conjunctival epithelium and goblet cells, which are also involved clinically.

The signs and symptoms of SSDE are similar to those of NSDE. The ocular symptoms include blurred vision, grittiness and ocular discomfort and clinical signs include tear film instability, corneal and conjunctival staining, goblet cell loss and epithelial metaplasia [382,401,685,686]. However, the onset of SSDE is earlier and where populations of patients with either NSDE or SSDE are compared, SSDE patients are consistently younger and their disease more severe [610,687] suggesting a more rapid progression. There is also a greater risk of blindness with SSDE [688]. The higher frequency of severe MGD in patients with SSDE compared to NSDE contributes to its severity [375].

9.1.2. Hormonal influences

SS affects more women than men and its prevalence increases in

post-menopausal women [566,567,689]. Sex-related differences in the prevalence of DED have been linked, at least in part, to the effects of sex steroids (e.g. androgens and estrogens). These endocrine actions are detailed in the TFOS DEWS II Sex, Hormone & Gender report [1222]. In brief, sex steroids act on the meibomian gland, lacrimal gland, conjunctiva and cornea. Hormonal influences occur most likely after local, intracrine synthesis and appear to be mediated primarily through nuclear, and possibly membrane, receptors. Sex steroids impact multiple structural and functional aspects of the ocular surface and adnexa, including tissue architecture, gene expression, protein synthesis, immune activity, epithelial cell dynamics, aqueous secretion, meibum production, mucous output and tear film stability. For example, androgen deficiency has been linked to the development, and androgen administration to the treatment, of lacrimal gland inflammation (e.g. Sjögren syndrome), meibomian gland dysfunction (e.g. Sjögren syndrome and aging), corneal glycocalyx disruption, ocular surface damage, tear film instability, and aqueous-deficient and evaporative DED. In contrast, the precise role of estrogens in the physiology and pathophysiology of ocular surface and adnexal tissues is unclear and, in some cases, controversial. A foremost consideration is that a number of the sex steroid effects may be sex-specific (i.e. unique to males or females) [36,690–693]. Recognition of these sex-related differences and the determination of their underlying basis (e.g. sex steroid action) are extremely important. (A full discussion is provided in the TFOS DEWS II Sex, Hormones and Gender report [1222]).

9.1.3. Etiology: genetic susceptibility

Genetic susceptibility plays a role in the etiology of Sjögren syndrome. A number of associations have been made between pSS and gene loci or specific genes [694]. (Table 11) Increased risk of pSS has been associated with HLA II, IL-12A, BLK, STAT4, CXCR5 and IRF5 in a recent study of subjects of European descent, fulfilling the European-American consensus criteria [694]. Loci of interest are not identical in all geographic locations, indicating ethnic differences in susceptibility [695].

Some of the clinical and immunological similarities between pSS and SLE may have a genetic background. A number of Sjögren syndrome-associated gene polymorphisms including the MHC-II, STAT4, IRF5, BLK, and TNIP1 genes are shared with SLE and other autoimmune conditions. However, the genes CXCR5 and GTF2I, have been determined as risk factors only in Sjögren syndrome and conversely, many genes associated with of risk of SLE are not found in Sjögren syndrome [695].

Burbelo et al. [695] have proposed that the Sjögren syndrome-associated genes result in immune dysregulation via at least three pathways: 1. Activation of the IFN signaling pathway. 2. Activation of B-cell function antibody production and clearance pathways. 3. Activation of NFκB activity pathways.

A general prediction is that the possession of one or more such genetic risk factors influences affected individuals in terms of clinical manifestations, onset, severity and progression of the disease. Of interest is that none of the genes that confer risk are related to glandular physiology or sex. All polymorphisms occur in non-coding sequences, reflecting an epigenetic role that determines gene expression rather than gene product. All of the risk genes relate to the performance of the immune system.

Table 11
Sjögren syndrome-associated non-HLA genes.

Gene	Gene Function
STAT4	Transcription Factor
IRF5	Transcription Factor
IL12A	Cytokine
BLK	B cell kinase
CXCR5	Chemokine
TNIP	NFκB signaling
GTF2I	Transcription Factor
TNFAIP3	NFκB signaling

From Refs. [694,1174].

9.1.4. Etiology: viral infection

The etiology of Sjögren syndrome remains unclear and involves multiple factors. One of the theories of the onset of Sjögren syndrome relates to viral infection. Indeed, various associations between viral infection and DED have been reported, including Hep B, HLTV1, HIV and Epstein Barr virus (EBV). Importantly, it has been suggested that the generation of tertiary or ectopic lymphoid structures (TLS) in response to viral infection may provide a site for autoantibody production in genetically disposed individuals [696].

9.1.5. The inflammatory process in Sjögren syndrome

Our understanding of the destructive inflammatory process that occurs in the lacrimal glands of Sjögren syndrome patients is partly inferred from the study of labial minor salivary gland biopsies. The typical pathologic lesion in the minor salivary glands consists of clusters of round cell infiltrates whose composition depends on lesion severity. $CD4^+$ T cells predominate in milder lesions and $CD8^+$ T cells and B cells in more severe lesions [697]. The distribution of other infiltrating immune cells also correlates with the degree of inflammation, with macrophages increasing and interdigitating dendritic cells decreasing with increasing severity [698]. Patients with Sjögren syndrome have been categorized at diagnosis into distinct groups according to whether the predominating immune response is T or B cell mediated [699] and whether the T cell response is mainly of Th1, Th2 or Th17 type. According to Moutsopoulos [697], Th1 responses are the most common, Th2 cytokines predominate in mild lesions and Th17 reactivity correlates with greater lesion severity.

Of particular concern in Sjögren syndrome is the formation of germinal centres, which are predictive of a higher risk of lymphoma [700,701].

9.1.5.1. T cells. T cells, which play a major role in Sjögren syndrome inflammation, can be divided into several subsets according to the cytokines that they make. Th1 cells produce IFN-γ and IL-18; Th17 cells produce IL-17 and IL21 and Th2 cells secrete IL-4, IL-5 and IL-13. Historically, Sjögren syndrome has been identified as a Th-1-dependent autoimmune disease, with increased concentrations of IFN-γ in tears, conjunctiva, saliva, lacrimal and salivary gland and blood [235,478,522,582,702]. Moreover, a Th1/Th2 imbalance, with high levels of IFN-γ in blood, salivary glands, tears or conjunctiva correlates with a more severe phenotype, which may help to differentiate Sjögren syndrome-determined aqueous tear deficiency from a non-Sjögren syndrome aqueous tear deficiency [532,703].

Recently, Th17 cells have emerged as players in the pathogenesis of Sjögren syndrome and the interaction between Th1 and Th17 cells is starting to be elucidated. There is evidence for the presence of IL-17 in fluids such as the tears, saliva, serum and synovial fluid and in tissue lesions themselves in patients with Sjögren syndrome [704–706]. Data from animal models have shown a pro-inflammatory role for IL-17 in sialoadenitis, while its specific role in lacrimal gland inflammation is still under debate [536,562,581,582,591,707,708].

9.1.5.2. Epithelial cells. A contributor to glandular inflammation is the activation of acinar and ductal epithelial cells to perform

immune functions and act as non-professional APCs whereby they mediate the recruitment and activation of almost all types of immune cells that drive the activation and differentiation of T and B cells. The factors which trigger epithelial activation are not known, but it has been suggested that latent viral infection (see above) or other intrinsic factors are responsible for their activation, in the context of an appropriate genetic and environmental background [709]. Activated salivary gland epithelial cells express a range of immunomodulatory molecules implicated in innate and acquired immune responses. They can also present autoantigens released from exosomal vesicles [710] or apoptotic bodies [711]. They therefore play an important role in initiating and perpetuating the local autoimmune process in the salivary glands, in Sjögren syndrome. A key feature of the process is that while infiltrating lymphocytes remain activated, the activated glandular epithelial cells undergo apoptotic cell death [712]. It is yet to be determined whether lacrimal gland epithelial cells play an analogous role.

9.1.5.3. B cells. B cell hyperactivity is now recognised as a central element of Sjögren syndrome, underscoring the loss of immune tolerance. It is manifested by hypergammaglobulinaemia, cryoglobulinaemia and the production of multiple autoantibodies, directed, for instance, against α-fodrin, the M_3 muscarinic receptor, and the ribonucleoprotein components Ro52 and Ro60 (anti-Ro/SSA) and La (anti-La/SSB). The latter are included among the classification criteria for Sjögren syndrome and correlate with early disease onset, parotid gland enlargement, extraglandular manifestations and lymphocytic glandular infiltration [683].

B cells fulfill other functions besides producing autoantibodies, by acting as APCs and secreting cytokines that can sustain the immune response [713].

9.1.5.4. Dendritic cells. Dendritic cells help orchestrate the immune response. There is evidence of cross-talk between dendritic cells and epithelial cells. Epithelial cells secrete inflammatory cytokines that can activate dendritic cells and T-cells and these in turn can further activate the epithelium. For example, IFN-γ-stimulated expression of MHC-II and HLA-DR, a ligand for the T-cell receptor, by epithelial cells, is well documented in the literature [714]. Expression of HLA-DR by both epithelium and DCs has been noted previously and recently used as an endpoint in clinical trials for DED [381,715–717].

9.1.5.5. Autoantibodies. Circulating autoantibodies in Sjögren syndrome contribute to its pathophysiology and can be of diagnostic importance [718]. Autoantibodies directed against Ro/SSA and La/SSB autoantigens are one of the recommended diagnostic tests for Sjögren syndrome [684,719].

Similarly, autoantibodies against the M_3 muscarinic receptor can be found in a subset of Sjögren syndrome patients, and have been considered to be pathogenic [720]. Some studies showed that these autoantibodies have agonistic (ie., stimulating) activity whereas others showed that they have antagonistic (ie., inhibitory) activity [721–724], although the difference may be methodological. The prevalence of these antibodies in the sera of Sjögren syndrome patients varies considerably, which questions their usefulness for diagnostic or prognostic purposes [720].

9.1.6. The lacrimal gland in Sjögren syndrome

The loss of aqueous tear flow in Sjögren syndrome is a result of inflammatory cell infiltration of the lacrimal glands which leads to acinar and duct destruction. Infiltrating lymphocytes, epithelial, endothelial and neural cells are all potential sources of inflammatory cytokines and other mediators that are responsible for lacrimal tissue damage. Additionally, inflammatory changes within the gland may lead to a decrease in lacrimal secretion by reason of damage to secretomotor innervation, or inhibition of neurotransmitter release or action by cytokines or antibodies [725]. The lacrimal glands in Sjögren syndrome are heavily infiltrated with mononuclear cells, the majority of which are T lymphocytes, with a lesser number of B cells and plasma cells (Fig. 11) [726]. These T cells express the activation marker, IL-2R, and contain cytotoxic granules such as granzyme A [727,728]. The degree of lymphocyte infiltration of the lacrimal glands correlates well with lacrimal secretion. Poor reflex tear secretion on nasal stimulation correlates with the presence of Sjögren syndrome autoantibodies and with both lacrimal and salivary lymphocyte infiltration in DED patients [729].

According to earlier reports, as in salivary gland biopsies, $CD4^+$ T cells predominate over $CD8^+$ cells in the lacrimal gland infiltrate, while B cells make up the smallest numbers.

Because of the constraints on using lacrimal gland biopsies for investigative purposes there would be great value in setting up prospective, post-mortem studies of lacrimal pathology in well-characterized Sjögren syndrome cases and NSDE, to further our understanding of natural history and to identify the likely time points for therapeutic intervention.

9.1.7. The conjunctiva in Sjögren syndrome

It is not known if the conjunctiva is a primary target of inflammation in Sjögren syndrome or whether changes in it are secondary to lacrimal gland inflammation and the onset of DED. Much of what is known about the pathologic events in the conjunctiva of patients

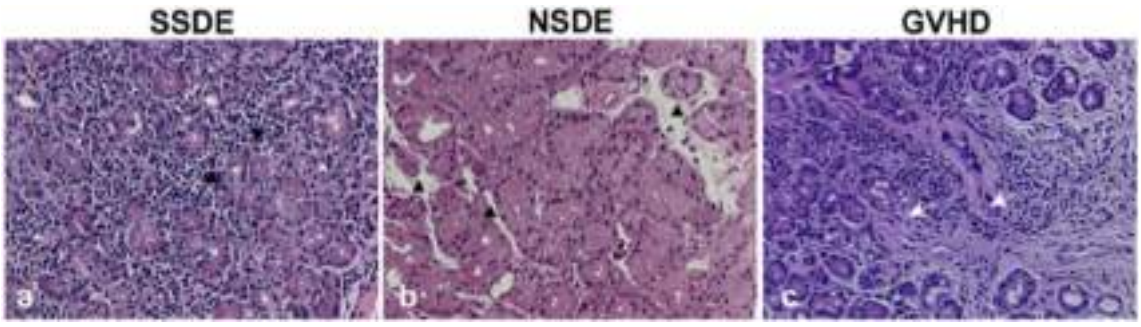

Fig. 11. Histopathology of the lacrimal gland in various forms of DED. (a) In SSDE there is marked intralobular lymphocytic infiltration (▲) and fibrosis is not apparent. (b) In a patient with NSDE inflammatory cell infiltration is limited and again, fibrosis is scarcely detected (▲). The acini maintain an almost normal structure. (c) By contrast, in a patient with chronic graft-vs-host disease (GVHD) there is marked interstitial periductal fibrosis (arrows) in addition to the lymphocytic infiltration. The periphery of the lobules is irregularly replaced by fibrotic tissue. (Courtesy of Y. Ogawa).