

Review

CD8: Adhesion Molecule, Co-Receptor and Immuno-Modulator

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CD8 is a cell surface glycoprotein found in cytotoxic T lymphocytes, which are important components in cellular immunity, esp. in the immune response to cancer and chronic infections. There are two forms of CD8, either as an $\alpha\alpha$ homodimer or $\alpha\beta$ heterodimer. It acts as an “assistant” or co-receptor in the function of cytotoxic T cells where specific immunity is mediated by interaction of specific T cell receptor ($\alpha\beta$ TCR) and its ligand peptide major histocompatibility complex (pMHC). CD8 also binds to pMHC but away from the interface of pMHC and TCR contact, thereof no influence on the specificity of this interaction. If the TCR and CD8 bind to the same pMHC at the same time, CD8 is defined as a co-receptor, functioning through its signalling *via* its cytoplasmic tyrosine phosphorylation pathway; if CD8 binds to pMHC independently of the TCR, it is defined as an adhesion molecule. At present, the co-receptor function theory is dominated in the field. Recent study has also shown that murine CD8 $\alpha\alpha$ binds to TL antigen, an MHC homologue, therefore acts as an immuno-modulator. In this review, we discuss these current understandings of the three aspects of the CD8 functions and their structural basis. *Cellular & Molecular Immunology*. 2004;1(2):81-88.

Key Words: CD8, T cell, adhesion, co-receptor, immuno-modulator

Introduction

As science progresses, and the body of information concerning the functions and mechanisms of biological pathways invariably increases through experiment and discussion, our knowledge of the nature of the proteins involved typically broadens to encapsulate the new facts. With this always being the case in science, we often find that the more we know, the more there is to learn. One such example of these is CD8, a cell surface glycoprotein expressed on all cytotoxic T lymphocytes (CTLs), or killer cells.

CD8 was first discovered as a cell surface marker in mice, useful only to distinguish between cytotoxic T cells and the two subsets of CD4 helper T (Th) cells (Th1 and Th2) (1, 2). At this time, these cell surface markers were termed lyt-1 (expressed on helper T cells) or lyt-2/3 (expressed on cytotoxic T cells) (3). Later experiments revealed human homologues of these markers (leu-1, 2a & 2b) (4) suggesting an important functional role due to their evolutionary conservation. This function, due to the ability of CD8 to bind to the same peptide/major histocompatibility complex (pMHC) as its corresponding T cell

receptor (TCR), was first thought to be that of an adhesion molecule (5), increasing the avidity and/or stability of the TCR/pMHC interaction (3, 6). This idea was later revised due to the extremely low binding affinity of CD8/pMHC and the failure to detect any specific enhancement to TCR/pMHC binding (7). Presently, CD8 is described most often as a co-receptor to TCR function because it binds to the same pMHC as the TCR (8). As a co-receptor, it is thought to have an important role in the signalling *via* p56^{lck} to induce TCR activation through the CD3 ζ chain tyrosine phosphorylation pathway (9, 10). There is also evidence to suggest that CD8 is involved in the recruitment of lipid rafts (11), which interact with the TCR/CD3 complex, also possible important in TCR signalling and activation (12, 13). Although CD8 is primarily recognized as a co-receptor, and the evidence is very much in favour of this being its role, there has recently become available data, which expands the function of CD8 further to that of an immuno-modulator. Leishman et al. carried out experiments to show that intestinal intraepithelial T lymphocytes (iIELs) express CD8 that has the ability to bind to a non-classical MHC class I-like molecule, TL antigen (TL), independently of TCR/CD3 but still signal the TCR through p56^{lck} type pathways (14). These data provide evidence that CD8 can modulate the immunological response of the TCR without being classed as a co-receptor or adhesion molecule by definition.

This review will expand upon the issues of CD8 as an adhesion molecule, a co-receptor (Figure 1) and an immuno-modulator (Figure 2). A more detailed account of the structure and function of CD8 (in comparison with CD4) as well as an account of its evolutionary and cross-species dynamics are described. Lastly, possible future research will be considered to uncover more about its function and the potential for manipulation of this molecule for use in therapy.

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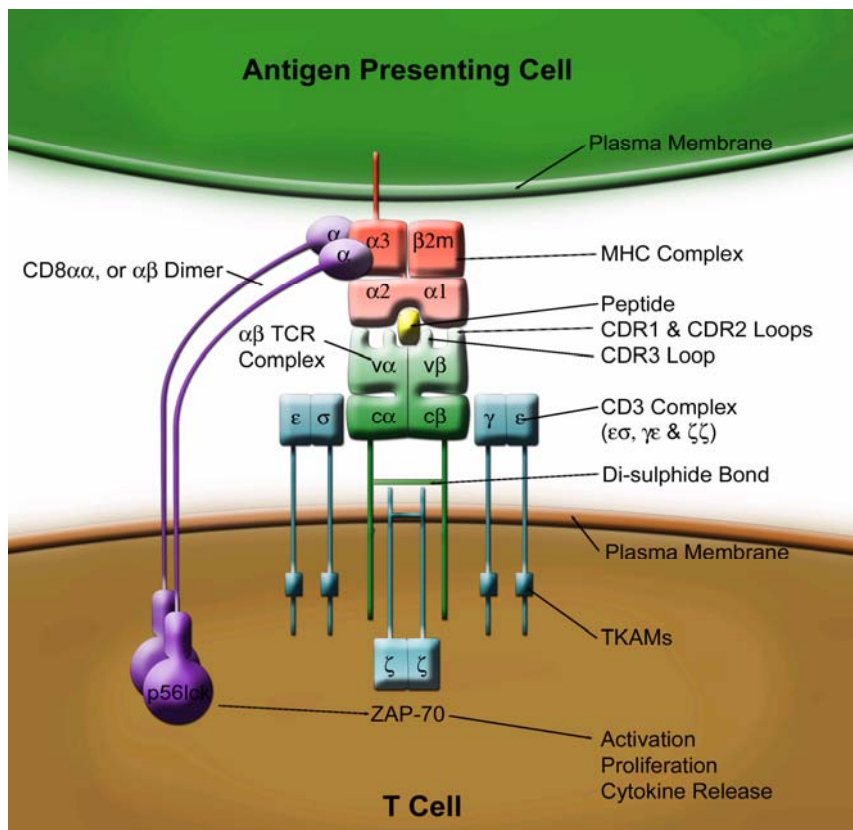


Figure 1. Schematic of the interaction between TCR, pMHC, CD8 $\alpha\alpha$ co-receptor and the CD3 complex (it is thought that this complex forms a V-shape in reality (reviewed in Gao et al.)). pMHC I (red), anchored through its transmembrane domain to its antigen presenting cell (green), presents short peptides (yellow) to a T cell receptor (TCR in green, $\alpha\beta$ TCR heterodimer) bound to a T cytotoxic cell (brown) via their transmembrane domains. The TCR recognizes the pMHC I via CDR1, 2 and 3 loops from its variable α and β regions ($v\alpha$ and $v\beta$). CD8 $\alpha\alpha$ (purple) anchored to the T cell via its transmembrane domain, binds to the $\alpha 3$ domain of pMHC independently of the dominant TCR/pMHC interaction. This is thought to be of importance in activation signals transduced via CD8 associated p56^{lck} tyrosine kinase to TCR/CD3 $\zeta\zeta$ mediated T cell activation. CD3 molecules contain tyrosine kinase activation motifs (TKAMs), CD3 $\epsilon\sigma$ and $\gamma\epsilon$ have one each, $\zeta\zeta$ possesses three. These TKAMs are thought to be important in acting as part of p56^{lck} to ZAP-70 kinase pathways and TCR signalling.

Comparative view of the structure and function of CD8 $\alpha\alpha$, CD8 $\alpha\beta$ and CD4

CD8 forms dimers in its native state; these dimers can be of two sorts. The first is the $\alpha\alpha$ homodimer form, encoded by the *leu-2a* gene in human (*lyt-2* in mice). The second is the $\alpha\beta$ heterodimer encoded by *leu-2a* and *leu-2b* (*lyt-3* in mice) genes (3, 15). The structural and functional differences between these two forms are considered.

Structure and function of CD8 $\alpha\alpha$

The α chain of CD8 is physically larger than the β chain at 34-37 kD. It is comprised of an extracellular 122 amino acids N-terminal Ig-like domain with a 48 amino acids stalk region. A transmembrane domain separates this from the cytoplasmic region, which is comprised of 28 amino acids (15) and contains two vicinal cysteine motifs responsible for src kinase p56^{lck} interaction via a zinc chelate complex (16). Two α subunits, covalently linked by a disulphide bond, form the homodimer CD8 structure (17).

The expression of CD8 $\alpha\alpha$ is present on $\alpha\beta$ T cells (although CD8 $\alpha\beta$ is predominant) and exclusively on NK cells and $\gamma\delta$ T cells (18). The binding affinity (Kd) of CD8 $\alpha\alpha$ /pMHC analyzed by surface plasmon resonance (SPR) using BiacoreTM is around 0.2 mM with kinetics of around 2 to 3 orders of magnitude faster than that of TCR/pMHC binding (7). The heterologous form of CD8 (CD8 $\alpha\beta$) shares these characteristics. The fast kinetics is thought to be important in ensuring that the binding of

pMHC is dominated by the TCR allowing immune recognition and response (19). The main function of the α chain has previously been determined as important for interaction with p56^{lck} through its cytoplasmic domain, possibly resulting in a signalling cascade and T cell activation (20). The homologous form of CD8, however, has been shown to be a less effective T cell activator than its heterologous counterpart (21, 22). More recent data have alluded to a role as an immuno-modulator due to the ability of two α chains to bind in a cross-link fashion with non-classical MHC molecule (TL antigen) independently of TCR/CD3 complex (only shown in murine systems) (14). Evidence has also provided a role for the α chain in the recruitment of lipid rafts, via palmitoylation events, important in TCR signalling (23).

Structure and function of CD8 $\alpha\beta$

The CD8 β chain is 32 kD and its amino acid sequence shares homology with around 20% of that of the α chain (24). Both the stalk region of the β chain (10-13 amino acids shorter than the α chain (15)) and the cytoplasmic region are shorter than that of the α chain (22). There is no p56^{lck} binding domain in the cytoplasmic region of the β chain, therefore it seems to have no involvement in signalling. Combined with the longer α chain, a recent proposed orientation by Kern et al. could confer greater thermodynamic stability to this form of CD8, explaining to some degree, its greater ability to induce $\alpha\beta$ T cell activation (15). These conformational differences have been the main area of study in alluding to the differences observed in the two forms of CD8.

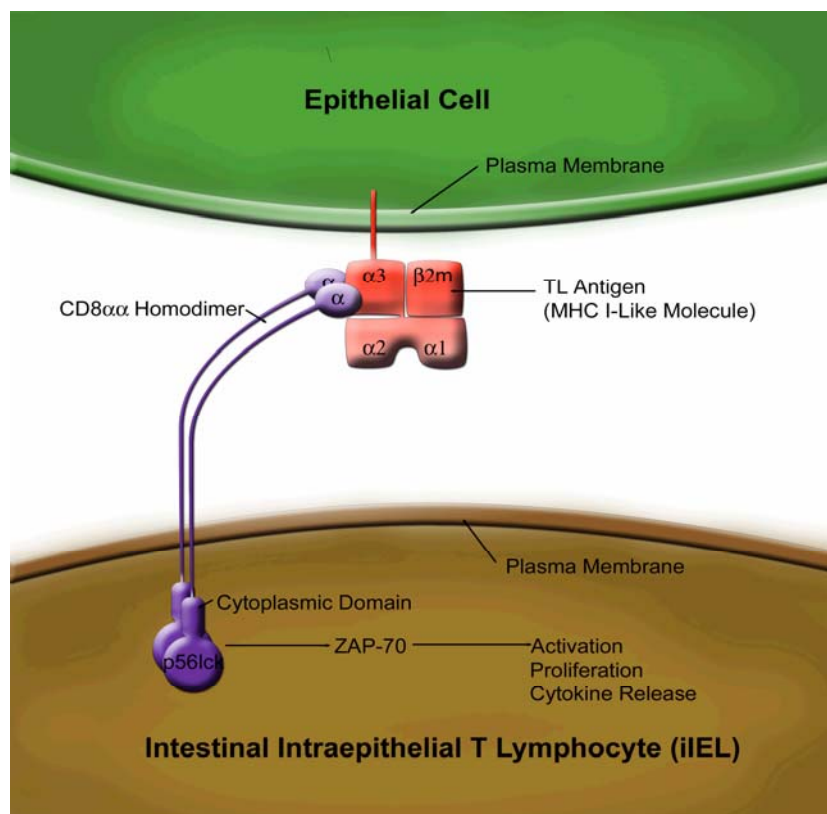


Figure 2. Schematic of CD8 $\alpha\alpha$ (purple) bound to TL antigen (MHC I-like molecule in red). CD8 $\alpha\alpha$, bound to an intestinal intraepithelial T lymphocyte (iIEL in brown) *via* its transmembrane domain can bind to TL independently of TCR. TL, bound to an epithelial cell (green) *via* its transmembrane domain, interacts with CD8 $\alpha\alpha$ *via* its $\alpha 3$ domain. This interaction causes CD8 $\alpha\alpha$ associated p56^{lck} activation and subsequent initiation of ZAP-70 pathways. This can mediate iIEL responses in the absence of TCR/pMHC binding. In this way CD8 $\alpha\alpha$ acts as an immuno-modulator.

Although the $\alpha\beta$ form shares much of its binding affinity characteristics with CD8 $\alpha\alpha$, data have shown T cell activation to be more effective when it is expressed (25, 26). A recent report, using a chimeric form of CD8 α with the stalk region replaced by the CD8 β region, has implicated the stalk region in mediating this difference (27). This is most likely due to specifically regulated *O*-linked glycosylation modifications that are important during binding to pMHC, which only occurs in the β chain (19, 27). This function is also of importance when considering negative and positive selection events during thymocyte maturation, due to the restriction in TCR activating peptide repertoire these glycosylations would invoke. Increased T cell production of interleukin-2, a cytokine important for CTL maturation and activation (28), is another characteristic associated with CD8 β expression only (29). Expression of high affinity CD8 $\alpha\beta$ /pMHC is thought to be crucial during positive selection when binding to self-pMHC is required, and this interaction becomes much weaker during negative selection allowing the TCR to become dominant to ensure self-reactive T cells are destroyed (30, 31). As with the α chain, the β chain is palmitoylated, leading to lipid raft recruitment thought to be important in signalling pathways (11, 16, 31-33).

Structure and function of CD4

CD4, unlike CD8, is expressed exclusively on T helper, rather than T cytotoxic cells and interacts with MHC class II (pMHC II) rather than MHC class I (pMHC I) molecules (8). Also unlike CD8 species, it classically forms only a monomeric rather than a dimeric molecule on the cell

surface (34), although new evidence has highlighted the possible role of dimeric and oligomeric forms of functional CD4 (35). The monomeric form of CD4 comprises of four Ig-like domains (D1-D4), a single transmembrane domain followed by a short cytoplasmic tail (36, 37). Crystal structures of CD4 have led to the identification of short linker regions between D2, D3 and D4 allowing for a degree of CD4 flexibility (34, 36, 38), unlike the rigidity retained by CD8 molecule. The N-terminal Ig-like V domain interacts with the $\alpha 2$ and $\beta 2$ proximal domains of the pMHC II molecule (34). In this way, it binds in a dissimilar orientation to CD8, which forms a “clamp” like bond to pMHC. Although key differences exist in the binding modes of CD4 and CD8, both bind to pMHC independently of the TCR/pMHC and do not play a physical role in the dominant peptide interaction.

The main consensus for the function of CD4, is that of a co-receptor involved in TCR signalling, but not critical for TCR/pMHC binding (34, 39, 40). p56^{lck} interaction and subsequently TCR activation *via* phosphorylation events involving the CD3 ζ chain occurs *via* the N-terminus of the cytoplasmic tail of CD4 *via* a zinc clasp (41, 42). More in function than in structure, CD4 and CD8 are much alike.

Evolutionary facts and functional diversity of CD8 in different species

CD8 has been studied in a number of model organisms including mice, human and fish. By comparing the function and identifying gene homology, a broader picture of a conserved role (if any) for CD8 can be identified.

As previously discussed, CD8 was first discovered as a cell surface marker in mice, used to identify different populations of T cells. As described by Ledbetter et al., *lyt-1* (CD4) is expressed only on helper T cells, whereas *lyt-2* (CD8 α) and *lyt-3* (CD8 β) are expressed exclusively on cytotoxic T cells (4, 15). Human homologues to these genes were later described by Ratnoffsky et al., *leu-3* (*lyt-1*), *leu-2a* (*lyt-2*) and *leu-2b* (*lyt-3*) (3). It is commonly accepted that CD8 homologues exist in all mammals due to the evolutionary conservation of the genes and the similarities that exist in the workings of the immune systems of these organisms. Not only mammalian species share CD8 co-receptor homology, birds and fish also share many immunological genes, which confer similar functions. Chickens are one of the more widely studied avian species sharing specific sequences in genes that encode for both CD4 (CT4) and CD8 (CT8)-like proteins (43). Similar findings have been described in rainbow trout, using a degenerate primer based PCR protocol to identify CD8 α -like sequences. This study indicates that CD8 genes have been conserved for over 400 million years and have evolved little in vertebrate systems (44).

The genes may well be conserved for CD8, but the immunology and physiology of different species such as human, mice, birds and fish are diverse. Is then the function of CD8 co-receptor also conserved, or does it have a multitude of roles as we consider different organisms?

One well-documented fact about human CD8 (hCD8) is its role as a co-receptor and its ability to bind to pMHC "away" from the direct contact interface of TCR/pMHC I acting as a T cell activator *via* p56^{lck} mediated pathways (8, 39, 41). When compared to murine CD8 (mCD8), we find almost identical functionality. Some key differences, however, include the affinity of mCD8, which is in the region of four times stronger than that of hCD8 with more overall contact, although this does not seem to endow mCD8 with any greater functionality (32), although some evidence exists implicating mCD8 with a greater role for TCR/pMHC stability than hCD8 (45). Studies comparing the crystal structures of hCD8 and mCD8 to pMHC I, have confirmed that, although some structural differences in binding orientation do exist between the two species, overall functional homology is maintained, i.e. neither mCD8 nor hCD8 has any contact with the peptide of the TCR/pMHC complex (46). Both bind to the $\alpha 3$ domain of the pMHC molecule inducing a conformational twist either away (mCD8) or toward (hCD8) the T cell to allow binding to occur (there has been some discussion on whether this shift mediates a similar change to the TCR/pMHC interaction, e.g. Gao and Jakobsen, 2000). There are also some differences in the pattern of hydrogen bonding between the two species. In human, three hydrogen bonds are formed between the T cell proximal CD8 $\alpha\alpha$ subunit and the MHC $\alpha 2$ domain, in addition to the hydrogen-bonding between CD8 $\alpha\alpha$ and pMHC heavy chain $\alpha 3$ domain, light chain $\beta 2m$ (17). In the murine model hydrogen bonds are exclusively formed between the heavy chain $\alpha 3$ domain and light chain $\beta 2m$ of the pMHC and N-terminal amino acids in the CD8 α chain (47). Recent data, using murine CD8 $\alpha\alpha$ on intestinal intra-epithelial T lymphocytes have suggested an immuno-

modulator type role for CD8 (14, 48). Whether this is also true for hCD8 has not yet been explored, although this will be discussed in more depth later. The function of CD8 seems to be of a similar nature in avian models as well. Experimental data indicate that CD8 in avian models also interact with a tyrosine kinase molecule sharing homology to the mammalian p56^{lck} (49) mimicking the hCD8 in TCR signalling and activation. As we move away from mammalian and avian models and consider fish, we do observe more of a difference in CD8, both structurally and functionally. Studies involving rainbow trout confirm that the cytoplasmic tail of the CD8 α molecule in this species lacks the consensus sequence required for interaction with a p56^{lck}-like molecule (44). Whether it interacts with other signalling molecules is not yet known, although it does contain a highly charged proximal portion of the consensus motif, allowing this possibility to be the likely target for future work (44).

CD8 functions: adhesion molecule versus co-receptor

CD8 was first discovered as a cell surface marker important for the initial discovery that a heterologous population of lymphocytes existed as part of the immune system (50). Later, this information was expanded somewhat describing not only different populations according to cell surface marker, but also in terms of immunological function (2, 51, 52). This indicated that specific roles existed for these cell surface markers in immune functions. Experiments using antibodies against the markers provided evidence that T cell activation is reliant on their functions (51, 53, 54). Definitively, CD8 can be classed as a co-receptor (Figure 1). This statement is validated due to the nature of CD8 to bind to the same pMHC as its TCR, without being directly involved in peptide recognition (17).

Adhesion molecule

CD8 has been shown to attach to the $\alpha 3$ domain of pMHC I in an antibody/antigen type conformation with (in human complex) or without (in murine complex) hydrogen bonding between the $\alpha 2$ and $\beta 2m$ domains (17, 46). For the CD8 $\alpha\alpha$ /pMHC I complex structure, it has been determined that binding is asymmetric with one chain contributing around 75% of the interaction. What percentage of the binding the β chain commands is still under scrutiny (17, 21). Mutagenesis data indicate that residues 223-229 of the MHC $\alpha 3$ chain are of particular importance in the CD8 interaction (55). These residues form a loop, which remains flexible pre-CD8 binding, but becomes locked between CDR1-like and CDR3-like loops within the extracellular Ig-like domain of both CD8 α chains (17). Residues Gln 115, Asp 112 and Glu 128 have also been shown as crucial factors in the MHC $\alpha 2$ domain (by further mutagenesis work) for successful CD8 binding (47). It was first thought that this CD8 (and CD4)/pMHC complex was needed solely to add stability to the TCR/pMHC complex, endowing additional affinity of the T cells to pMHC on antigen presenting cells (5, 6, 56). This theory was later expanded upon giving rise to the idea of CD8 and TCR

signalling *via* tyrosine kinase p56^{lck} pathways (6, 42, 56, 57). Presently, the idea of CD8 as an adhesion molecule is unlikely due to SPR analysis indicating a significantly lower K_d (binding affinity) when compared to other adhesion molecules such as CD2 and CD48 (58, 59). Also, no conformational changes to the TCR/pMHC complex occur with/without CD8 binding; therefore, CD8 does not induce an energetically favourable shift in binding properties. Therefore the most commonly accepted role for CD8 is that of a TCR signalling molecule as a co-receptor.

TCR signalling

The most important function of CD8 when considering $\alpha\beta$ TCR is its involvement in activation pathways. The CD8 α tail region is associated with tyrosine kinase p56^{lck} *via* two vicinal cysteines that interact through a zinc chelate complex (22, 57, 60). Upon binding with the $\alpha 3$, $\alpha 2$ and β_2 m of pMHC, activation pathways are thought to initiate creating a signalling cascade originating from CD8 α linked p56^{lck} and leading to ZAP-70 activation on the TCR/CD3 complex (61). CD3 is a signal transducer with three subunits. CD3 $\delta\epsilon$ and $\gamma\epsilon$ heterodimers containing a single tyrosine kinase activation motif, and a disulphide linked $\zeta\zeta$ homodimer with three tyrosine kinase activation motifs are responsible for transmitting the signal from the CD8 α chain to the TCR (47, 62). Knockout studies including the CD3 subunits have confirmed the data (63-67).

CD8 β chain is not directly involved in p56^{lck} signalling as there is no cytoplasmic tail sequence with a tyrosine phosphorylation site, so it would not be unexpected to find the $\alpha\beta$ heterodimer as a poorer TCR activator than the homodimer. This, however, is not the case. It has been experimentally proven the heterodimeric form of CD8 is a more potent activator than the homodimeric form (25). It has been suggested that the unique structure of CD8 β stabilizes the interaction between CD8 α and p56^{lck} allowing more efficient transduction (29, 68). Perhaps more interesting is the role of the β chain in the creating of lipid rafts. The CD8 β cytoplasmic tail is palmitoylated at a membrane-proximal cysteine partitioning CD8 into lipid rafts, or detergent-insoluble membranes (11). These rafts are made up of ordered microdomains, enriched with sphingolipids and cholesterol (11-13, 69, 70). Molecules such as phosphatases (CD45) are excluded allowing uninhibited TCR signalling to take place. Other important molecules such as lck and the linker for activation of T cells (LAT) have been shown to associate increasingly with CD8 β during raft recruitment, molecules vital to T cell activation (71, 72). These events inevitably lead to lck phosphorylation, mobilization of intracellular calcium and ZAP-70/CD3 activation (16). Moreover, it has been postulated that CD8 β can form a functional link with CD3 δ , which in turn is part of the TCR/CD3 complex, creating an important bond between TCR and co-receptor (16). How this interaction is formed and how it functions are not yet known, but knockout mice lacking either CD8 β or CD3 δ have been shown to suffer from extremely poor TCR mediated activation (73). There is also evidence to suggest that CD8/pMHC binding is enhanced by the β chain due to the presence of unique glycans on the stalk region that

modulate the distal binding surface of the CD8 $\alpha\beta$ globular head domain and therefore, its link to pMHC (74).

Immuno-modulator

Recent evidence has implicated a new role for CD8 $\alpha\alpha$ in murine intestinal intraepithelial T lymphocytes (iIELs) that reside in the epithelial tissue layer barrier, which shields internal tissues from the external environment (75). This role is that of an immuno-modulator, rather than a co-receptor or adhesion molecule (Figure 2). The distinction between these two functions is in the ability of CD8 $\alpha\alpha$ to bind to an MHC I-like thymus leukemia antigen molecule (TL) independently of the TCR (14).

The expression of the monomeric form of CD8 is unusual on $\alpha\beta$ TCR expressing T cells because of the increased efficiency observed in TCR mediated T cell activation in cells expressing the dimeric form (21). iIELs may mature in a more specific environment compared to other T cells, in this way they may only be exposed to peptides expressed by intestinal epithelial cells, which in turn could influence activation pathways in the periphery (76). Lieshmann et al. found that TL binds very strongly to CD8 $\alpha\alpha$ due to cross-linkage interaction involving both CD8 chains (76). The crystal structure of this complex has shown that one CD8 $\alpha\alpha$ homodimer binds to one TL molecule (77). The affinity of binding between CD8 $\alpha\beta$ and TL was not of the same affinity and deemed insufficient to operate independently of TCR/TL interaction (14). Binding data have indicated that the complementary binding loops of CD8 α interact differently with the β_2 m domain of TL and the conventional MHC is perhaps in part responsible for this difference (78). In this model, the TCR does not interact with TL, rather, the high affinity bond between CD8 and TL, and the subsequent p56^{lck} signalling pathways are sufficient to induce or suppress immune response (48). Whether the ability of CD8 $\alpha\alpha$ to act as an immuno-modulator can be transferred to NK cells and $\gamma\delta$ T cells, where it is also commonly expressed, is still a matter for scrutiny. Recent data concerning the functions of TL have indicated that both $\alpha\beta$ and $\gamma\delta$ TCR recognize it independently of antigen (79). Also, due to the self-reactive nature of CTL that bind TL and the low affinity of this interaction, it has been postulated that CD8 $\alpha\alpha$ may be required for activation in this unique situation (48). Therefore, the role of CD8 as an immuno-modulator may only occur in this isolated environment and may not contribute towards a broader function for CD8 $\alpha\alpha$ expression.

Conclusion and future perspectives

Although evidence exists for a new role for CD8 as an immuno-modulator, this function seems to be isolated to a unique subpopulation of T cells in the murine system. Therefore, the present experimental data point strongly to the major role of co-receptor for CD8 important in TCR signalling and T cell activation. With this in mind, one of the major benefits of CD8 mediated research may come in the form of therapeutic drugs for regulation of immune

responses for conditions such as cancer, allergy and autoimmune diseases such as type I diabetes. It has already been shown that blocking CD8 with anti-CD8 antibody, soluble CD8, or synthetic peptides drastically reduces immune responsiveness (reviewed in Gao et al., 2002). Refinement in the control of CD8 could, therefore be useful in immune suppression for organ transplantation patients.

There are still important discoveries to come in the future for CD8 based research. The structure of CD8 $\alpha\beta$ has not yet been fully described, and a broader understanding of the function of CD8 $\alpha\alpha$ as an immuno-modulator and in other potential roles will, no doubt, be useful. Although many models have been proposed, we still lack the triple complex crystal structure of CD8/pMHC/TCR needed to understand the physical orientation of CD8 in relation to TCR and CD3. These discoveries and others will be key for our continuing increase in knowledge of antigen recognition and TCR signalling and will help to define what there is still to learn.

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