

## Positive and negative adaptors in T-cell signalling

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### OVERVIEW OF T-CELL SIGNALLING

T-cell receptor (TCR) engagement results in the initiation of intracellular signalling cascades that lead to the initiation, amplification and/or inhibition of specific functions. These functions include the production of different cytokines and chemokines, the induction of anergy, the differentiation of T-cell subsets (i.e. Th1/Th2/TRegs) and the generation of cytotoxic T cells (CTLs). T-cell signalling is initiated by the Src kinase p56lck and its ability to interact with the coreceptors CD4 and CD8.<sup>1,2</sup> The concordant binding of the antigen-receptor complex and the coreceptors to major histocompatibility complex (MHC) antigens brings p56lck to the TCR $\zeta$ /CD3 chains, leading to the phosphorylation of immunoreceptor-based tyrosine activation motifs (ITAMs) as defined by the sequence YXX(L/I)X<sub>6–8</sub>YXX(L/I).<sup>3,4</sup> The phosphorylated ITAMs then recruit the syk family tyrosine kinase zeta associated protein-70 (ZAP-70), or the related protein SYK, via tandem binding to Scr homology 2 (SH2) domains.<sup>3,4</sup> Phosphorylation leads to the activation of phospholipase C $\gamma$  (PLC $\gamma$ 1) which in turn hydrolyses phosphatidylinositol 4,5-bisphosphate (PIP2) to diacylglycerol (DAG) and phosphoinositol 1,4,5-trisphosphate (IP3). This is mediated by the combined action of adaptors and the Tec kinases, inducible T-cell kinase (ITK) and resting lymphocyte kinase (RLK).<sup>5</sup> DAG activates protein kinase C (PKC), while IP3 induces the release of intracellular Ca<sup>2+</sup> for activation of the serine/threonine phosphatase calcineurin. Calcineurin in turn dephosphorylates the transcription factor nuclear factor in activated T cells (NFAT), allowing its entry into the nucleus.<sup>6</sup> Proximal events also activate small GTP-binding proteins p21ras, p21rac and Cdc42, leading to the remodelling of the cytoskeleton and the activation of mitogen-activated protein kinases. These include the extracellular signal regulated kinase (ERKs), c-Jun N-terminal kinase (JNK) and p38, which in turn activate a variety of transcription factors such as NF $\kappa$ B and activator protein-1 (AP-1). It is the orchestration of this array of intracellular

pathways that leads to multiple functions attributed to T cells in the immune response.

### ADAPTOR PROTEINS IN T CELLS

Recent advances in T-cell signalling have been made with the identification of haematopoietic specific adaptor proteins, or molecular scaffolds.<sup>7–9</sup> These adaptors lack enzymatic activity, and instead possess an array of binding sites and modules that bind to other proteins. A single protein therefore has the potential to regulate multiple discrete events and group proteins in the integration of signalling pathways. Adaptor molecules can function as both positive and negative regulators of T-cell function. Examples of adaptors with a positive effect include growth factor receptor-bound protein-2 (GRB-2), the linker for activation of T cell (LAT), GRB-2-related adaptor downstream of Shc (GADS) and the SH2-domain-containing leukocyte protein of 76 kDa (SLP-76), while others such as the phosphoprotein associated with (glycosphingolipid enriched microdomains) GEMs (PAG), the SH2-interacting transmembrane adaptor protein (SIT) and downstream of tyrosine kinases (DOK) have a negative regulatory function. In addition, a number of other adaptors such as the adhesion and degranulation promoting adaptor protein (ADAP), the Src kinase-associated phosphoprotein of 55 kDa (SKAP-55) and the Wiskott–Aldrich syndrome protein (WASP) have regulatory effects on the cytoskeleton, on adhesion and on the ability of T cells to form conjugates with antigen-presenting cells (APCs).

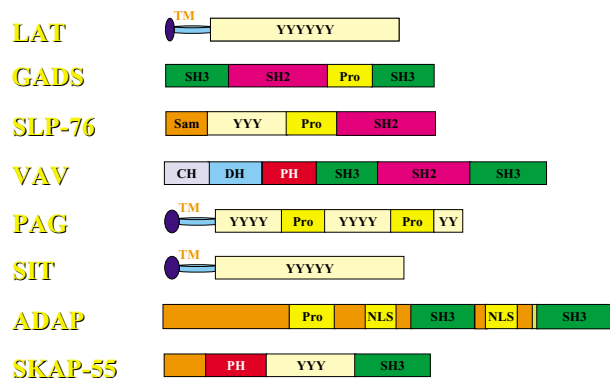
### POSITIVE REGULATORS OF T-CELL SIGNALLING

#### Linker for activation of T cell (LAT)

LAT is the first in an array of adaptors that is needed for the successful transmission of signals from the TCR/CD3 complex (Fig. 1). It is a transmembrane adaptor with a short extracellular region and a tyrosine-rich cytoplasmic tail.<sup>10,11</sup> The nature of the binding sites on LAT and the properties of LAT-deficient T cells have provided insights into the manner in which the adaptor is connected to signalling. There are binding sites for PLC $\gamma$ 1, GRB-2 and GADS. Tyrosine residue 132 binds the SH2 domain of PLC $\gamma$ 1, residues at 171, 191 and 226 bind GRB-2, and residues 171 and 191 bind to GADS.<sup>12</sup> PLC $\gamma$ 1 binding could couple LAT to Ca<sup>2+</sup> mobilization, GRB-2 binding to the activation of ERKs, and GADS to the downstream adaptor

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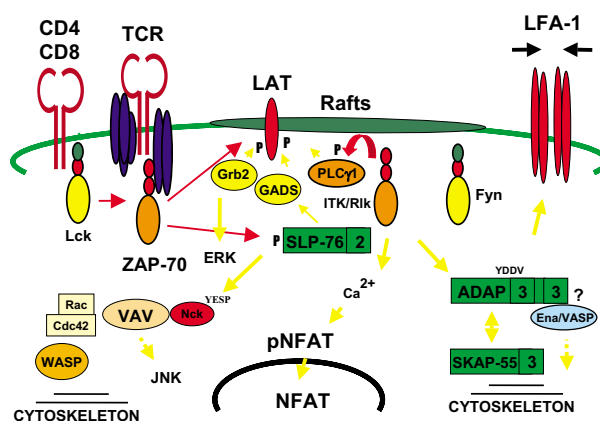


**Figure 1.** Structure and functional domains of adaptor proteins. The structures of immune cell adaptors discussed in this review are shown, including linker for activation of T cell (LAT), SH2-domain-containing leukocyte protein of 76 kDa (SLP-76), Grb2-related adaptor downstream of Shc (GADS), phosphoprotein associated with GEMS (PAG), SH2-interacting transmembrane adaptor protein (SIT), adhesion and degranulation promoting adaptor protein (ADAP), Vav and Src kinase-associated phosphoprotein of 55 kDa (SKAP-55). Adaptor proteins, by definition, lack enzymatic activity or transcriptional domains, but instead express a variety of modular domains or binding sites, which mediate interactions with other proteins. In this way they act as scaffolds, facilitating the formation of multiprotein complexes and the integration of diverse signalling cascades. YYY, tyrosine phosphorylation sites; Pro, proline-rich region; Sam, XXX; SH2, Src homology 2 domain; SH3, Src homology 3 domain; PH, pleckstrin-homology domain; NLS, nuclear localization sequence; DH, Dbl-homology domain; CH, calponin-homology domain.

**SLP-76.** LAT-deficient Jurkat cells are defective in tyrosine phosphorylation of several proteins, including PLC $\gamma$ 1, Ca<sup>2+</sup> mobilization and NFAT gene transcription.<sup>13,14</sup> LAT-deficient mice also show a block at the pre-TCR stage of thymic differentiation,<sup>15</sup> while knock-in mice expressing the LAT-PLC $\gamma$ 1 binding mutant have confirmed the importance of this interaction for PLC $\gamma$ 1 activation, calcium mobilization and NFAT function.<sup>16</sup> Interestingly, the loss of binding did not affect activation of the ERK pathway in normal cells, although an ERK activation defect has been observed in Jurkat cells.<sup>12,17,18</sup> LAT function requires acylation, which is needed for localization to lipid rafts (otherwise known as GEMs or microdomains).<sup>13,14</sup> These microdomains are enriched in relatively ordered cholesterol-rich domains that possess key proteins needed for signalling.<sup>19</sup> Other LAT-related functions include cytoskeletal-dependent spreading<sup>20</sup> and Th1/Th2 differentiation, where deficient mice show defects in Th1 development.<sup>21</sup>

#### Grb2-related adaptor downstream of Shc (GADS)

GADS is a member of the family of SH2- and SH3-domain-containing adaptor proteins. GADS plays a critical role in TCR-mediated signalling by linking LAT with another T-cell adaptor protein, SLP-76, coupling membrane-proximal events to downstream signalling pathways. The GADS protein consists of a central SH2 domain flanked by a proline-rich region and two SH3 domains (Fig. 1).<sup>22</sup> Following TCR stimulation, GADS binds to phosphorylated LAT via its SH2 domain, whilst the



**Figure 2.** Role of adaptors in T-cell signalling. Ligation of the T-cell receptor (TCR) leads to the activation of Src family tyrosine kinases (PTKs), such as Lck, which phosphorylate key tyrosine residues within immunoreceptor-based tyrosine activation motifs (ITAMs) on CD3 and TCR $\zeta$  chains. This results in the phosphorylation and activation of the syk family PTK zeta associated protein-70 (ZAP-70), which subsequently phosphorylates the linker for activation of T cell (LAT) and SH2-domain-containing leukocyte protein of 76 kDa (SLP-76), allowing them to interact with SH2-containing proteins. SLP-76 and its associated molecules are recruited to LAT in the plasma membrane via an interaction with the Grb2-related adaptor downstream of Shc (GADS). This leads to the SLP-76-dependent activation of phospholipase C $\gamma$  (PLC $\gamma$ 1), which results in intracellular Ca<sup>2+</sup> mobilization and protein kinase C (PKC) activation. Phosphorylation of SLP-76 allows SH2-mediated binding to the adaptors Vav and Nck, leading to cytoskeletal reorganization. SLP-76 also binds to adhesion and degranulation promoting adaptor protein (ADAP) [previously termed Fyn T-binding protein (FYB)/SLP-76-associated protein (SLAP)] in a pathway that involves Fyn-T and binding to Src kinase-associated phosphoprotein of 55 kDa (SKAP-55) and SKAP-55R.

SH3 domains bind to polyproline residues (residues 224–244) on SLP-76, promoting the formation of a complex between SLP-76 and LAT (Fig. 2). In this way, GADS allows SLP-76 and its associated proteins to be recruited to the membrane. Analysis of GADS-deficient mice revealed impaired T-cell development, with specific defects in both positive and negative selection of thymocytes.<sup>23</sup> Expression library screening has revealed that, in addition to a connection with SLP-76, GADS associates with the serine/threonine kinase haematopoietic progenitor kinase-1 (HPK1) which has been implicated in the activation of the JNK pathway.<sup>22,24,25</sup>

#### SH2-domain-containing leukocyte protein of 76 kDa (SLP-76)

SLP-76 is another multidomain adaptor protein, linked to LAT-GADS, which plays an important role in the activation of PLC $\gamma$ 1 and other downstream signalling pathways. SLP-76 has several distinct domains: an N-terminal SAM domain, an acidic region with key tyrosine motifs (YESP, YESP and YEPP), a central proline-rich region and a carboxy-terminal SH2 domain<sup>26</sup> (Fig. 1). As in the case of LAT and GADS, studies involving SLP-76-deficient T cells and SLP-76 knock-out and mutant mice have provided insights into the role of this

adaptor in T-cell development and function.<sup>27,28</sup> Phosphorylation of PLC $\gamma$ 1 in SLP-76-deficient Jurkat T cells is severely impaired, resulting in defective Ca<sup>2+</sup> mobilization and inositol 1,4,5-triphosphate and interleukin (IL)-2 production.<sup>27</sup> Interestingly, this loss of phosphorylation is more selective than that seen in LAT-deficient T cells, where a reduction in the phosphorylation of multiple substrates has been observed. This is consistent with the notion that SLP-76 operates downstream of LAT. The mechanism by which SLP-76 is coupled to PLC $\gamma$  phosphorylation is unclear, but involves Tec family kinases such as ITKs and RLKs. Both kinases phosphorylate SLP-76, but ITK has also been shown to bind to the adaptor.<sup>29,30</sup> Studies involving SLP-76<sup>-/-</sup> mice have revealed that SLP-76 is required for pre-TCR signalling, with a developmental block at the double negative (DN) stage.<sup>31,32</sup> Examination of mice expressing different mutations of SLP-76 have demonstrated a differential requirement for specific domains of SLP-76 in the regulation of T-cell development, proliferation and effector function, providing an insight into the specific molecular mechanisms underlying SLP-76 function.<sup>30</sup>

SLP-76 has also been implicated in the regulation of cytoskeletal changes in activated T cells though its association with the adaptor proteins Vav, Nck and ADAP [previously termed Fyn T-binding protein (FYB)/SLP-76-associated protein (SLAP)].<sup>7-9</sup> TCR ligation is accompanied by the remodelling of the actin cytoskeleton, an event needed for signalling. The N-terminus of SLP-76 has two key tyrosine residues which directly associate with the SH2 domain of Vav.<sup>33-35</sup> Phosphorylation of these residues and binding is regulated by ZAP-70 and SYK.<sup>35,36</sup> SLP-76 can cooperate with Vav in the up-regulation of IL-2 gene transcription,<sup>33,34</sup> although Vav-SLP-76 binding *per se* is not essential.<sup>35</sup> Vav is a multidomain protein comprising a calponin homology (CH) domain followed by an acidic (Ac) motif, a DBL-homology (DH) domain, a pleckstrin-homology (PH) domain, a zinc finger (ZF)-like domain, two SH3 domains and an SH2 domain (Fig. 1). The presence of so many domains within Vav suggests that it may serve to interact with, or bring together, many signal transduction pathways. The DH domain is central to Vav function as it possesses guanine nucleotide exchange (GEF) activity for members of the Rho/Rac family of GTPases, which include the proteins Rac and Cdc42. GTP-bound Cdc42 subsequently associates with WASP. This protein, when activated by Cdc42 and PIP<sub>2</sub>, can stimulate actin polymerization through the Arp2/3 complex.<sup>37</sup> SLP-76 also binds the adaptor Nck, another WASP-binding protein, via its YESP domain,<sup>38,39</sup> and a model has been proposed whereby SLP-76 acts as a scaffold bringing Nck and WASP into proximity with Vav and Cdc42-GTP.<sup>40</sup> Together, these findings suggest that SLP-76 is connected to the regulation of the actin cytoskeleton, although evidence directly demonstrating this link has yet to be published.

### NEGATIVE REGULATORS OF T-CELL SIGNALLING

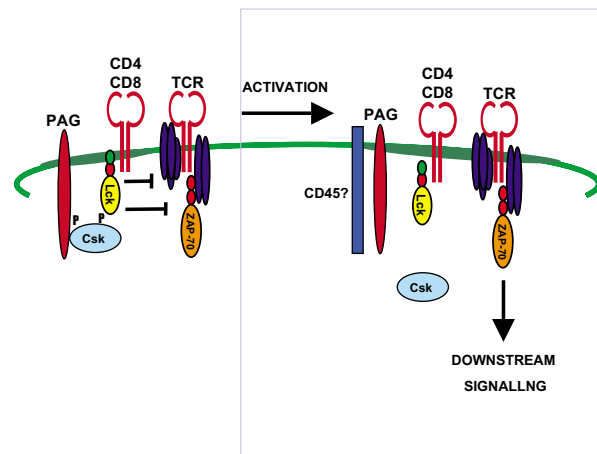
Adaptors were originally described as molecules involved in the positive regulation of T-cell signalling events. However, there are increasingly more negative regulatory adaptor proteins being described, which are important in the control of T-cell

receptor signalling. Such proteins are important in terminating TCR signalling and preventing the inappropriate activation of T cells. Two recently identified transmembrane adaptors with negative regulatory function, PAG and SIT, are discussed in more depth below.

### Phosphoprotein associated with GEMS (PAG)

PAG, also known as Csk-binding protein (Cbp), is a transmembrane adaptor protein found in lipid rafts.<sup>41,42</sup> PAG comprises a short 16-amino acid extracellular domain and a 397-amino acid cytoplasmic domain containing a dicysteine motif (CSSC), a palmitoylation site for targeting PAG to GEMS<sup>43</sup> and 10 tyrosine residues (Fig. 1).<sup>41,42</sup> In resting T cells, PAG is constitutively tyrosine-phosphorylated and binds the major negative regulator of Src kinases, the tyrosine kinase c-terminal Src kinase (Csk) (Fig. 3). Csk phosphorylates the C-terminal tyrosine of Src kinases such as Lck, causing it to bind to its internal SH2 domain, leading to kinase domain inactivation. Csk SH2 domain binding to at least one of the tyrosine-based signalling motifs in the cytoplasmic tail of PAG leads to its localization to the rafts and activation.<sup>44-46</sup> Once in the rafts, Csk inhibits the activity of Src family kinases. Following TCR stimulation, PAG becomes transiently dephosphorylated by the phosphatase CD45, which results in the release of Csk from the GEMS, allowing Src kinases to initiate downstream signalling events<sup>47</sup> (Fig. 3). In this way, the PAG-Csk complex transmits negative regulatory signals, helping to keep resting T cells in a quiescent state.

PAG has also been shown to bind the cytoplasmic adaptor protein EBP50 [ezrin/radixin/moesin (ERM)-binding phosphoprotein of 50 kDa], which is also known as NHERF (Na<sup>+</sup>/H<sup>+</sup> exchange regulatory factor). The EBP50 protein interacts with ERM family proteins which are known to bind the actin cytoskeleton, and the PAG-EBP50-ERM complex has been shown to regulate the process of synapse formation and activation in T cells.<sup>48,49</sup>



**Figure 3.** Negative regulation of T-cell signalling by phosphoprotein associated with GEMS (PAG). PAG is constitutively tyrosine-phosphorylated in resting T cells, and associates with Csk, a negative regulator of Src kinases, via its SH2 domain. Following T-cell receptor (TCR) engagement, PAG is dephosphorylated by a PTPase, most likely CD45. This leads to the dissociation of Csk, which relieves Src kinases of inhibition, enabling them to transmit signals downstream.

### SH2-interacting transmembrane adaptor protein (SIT)

SIT is a glycosylated transmembrane adaptor protein, expressed by lymphocytes, which is involved in the negative regulation of antigen receptor signalling.<sup>50–52</sup> This adaptor has an extracellular domain of 18 amino acids, a 20-amino acid transmembrane region and a 136-amino acid cytoplasmic tail containing five potential sites of tyrosine phosphorylation, which mediate binding to the Grb2 and Src family kinases (Fig. 1).<sup>50,52</sup> Amongst these five phosphorylation sites is an immunoreceptor tyrosine-based inhibition motif (ITIM) which mediates the association of SIT with the tyrosine phosphatase SH2 domain-containing protein tyrosine phosphatase 2 (SHP-2).<sup>50</sup> Mutation of the tyrosine residue within this ITIM almost completely abrogates the interaction between SIT and SHP-2. However, surprisingly, this mutation has no effect on SIT-mediated inhibition of NFAT activity, indicating that the interaction between SIT and SHP-2 is not required for this effect.<sup>50</sup> SIT has also been shown to interact with Grb-2 via two consensus YxN motifs, but the mutation of both of these binding sites has no effect on the inhibitory function of SIT.<sup>52</sup> Coprecipitation experiments suggest that Csk could represent the effector molecule which mediates the negative regulatory function of SIT on TCR-mediated signalling.<sup>52</sup> However, further studies are required to formally demonstrate this and to elucidate the precise mechanism by which SIT mediates inhibition of TCR signalling events.

### ADAPTOR PROTEINS AND THE CYTOSKELETON

Modification of the actin cytoskeleton has been shown to be important in antigen recognition and T-cell activation. TCR-mediated cytoskeletal changes not only are required for cell motility, but are important in the regulation of cellular adhesion. Recently, there has been much attention focused on the signalling pathways involved in TCR-mediated regulation of the cytoskeleton. Indeed, several adaptors have now been described which have been implicated in the regulation of adhesion and cytoskeletal changes in activated T cells. Two such adaptor proteins, ADAP and SKAP-55, are discussed below.

#### Adhesion and degranulation promoting adaptor protein (ADAP)

Upon activation, the SH2 domain of SLP-76 interacts with the adaptor ADAP.<sup>53,54</sup> It is expressed as two isoforms of 120 and 130 kDa (ADAP-120/130), with the 130-kDa isoform being preferentially expressed in peripheral T cells.<sup>55</sup> ADAP contains several proline-rich regions and an SH3 domain which mediate binding to SKAP-55 (see below), 16 putative tyrosine phosphorylation sites which allow binding to SH2 domains of SLP-76 and Fyn, two putative nuclear localization sequences (NLS), and an Ena (enabled)/VASP (vasodilator-stimulated phosphoprotein) homology 1 domain binding site (EVH1) (E/DFPPPPXD/E)<sup>7,53,56–58</sup> (Fig. 1). Mapping studies have identified the Y595 and Y651-YDDV motifs as binding sites for SLP-76, while the Y625-YDGI motif binds the SH2 domain of the

Src family kinase Fyn.<sup>56,57</sup> Following TCR stimulation, Fyn phosphorylates these signalling motifs and regulates binding of SLP-76 and Fyn to ADAP (Fig. 2).<sup>53,54,57,59</sup> Tyrosine phosphorylation of ADAP following TCR ligation is diminished in Fyn-deficient T cells, supporting a link between this kinase and the ADAP signalling pathway.<sup>53</sup>

The role of ADAP as a positive or negative regulator of T-cell function was initially uncertain. Transfection studies with ADAP have yielded conflicting results on the role of this adaptor in the regulation of T-cell activation. One study found that ADAP attenuated IL-2 production,<sup>53</sup> whereas another report suggested a negative regulatory role for this protein.<sup>54</sup> However, cotransfection of ADAP with Fyn and SLP-76 was shown to up-regulate TCR-driven IL-2 transcription.<sup>56,57</sup> Furthermore, ADAP-deficient peripheral T cells exhibit defects in proliferation, cytokine production and lymphocyte functional antigen-1 (LFA-1) clustering following TCR stimulation.<sup>60,61</sup> Similarly, ADAP was shown to enhance  $\beta$ 1 integrin clustering in mast/basophilic cells; however, clustering of the high-affinity immunoglobulin E (IgE) receptor *Fc $\epsilon$ RI* was unaffected.<sup>62,63</sup> The effect of ADAP on integrin clustering has been reported to facilitate stromal cell-derived factor 1 alpha (SDF1 $\alpha$ )-mediated T-cell migration<sup>64</sup> and enhance conjugate formation between T cells and antigen-presenting cells (APCs).<sup>65</sup> Together, these findings suggest that ADAP is important in coupling TCR-mediated actin cytoskeletal changes to the activation of integrin function, but exactly how this is achieved is unclear. One possible mechanism by which ADAP may regulate integrin adhesion is through an association with proteins of the Ena/VASP family. These proteins are important in the regulation of actin dynamics, controlling processes such as fibroblast migration, axon guidance, and T-cell polarization, and in the actin-based motility of the intracellular pathogen *Listeria monocytogenes*.<sup>66</sup> ADAP was recently shown to co-localize with Ena/VASP proteins and WASP, Vav and F-actin at the interface between T cells and anti-CD3-coated beads.<sup>58</sup> In activated T cells, ADAP associates with Ena/VASP family proteins and is found within multiprotein complexes containing WASP, Nck, and SLP-76. Inhibition of binding between ADAP and Ena/VASP proteins or between WASP and the Arp2/3 complex inhibits TCR-mediated actin rearrangement, suggesting that these interactions are important in linking TCR signalling to cytoskeletal remodelling.<sup>58</sup> Clearly, further studies are required to determine the precise role of ADAP in the regulation of the actin cytoskeleton.

#### Src kinase-associated phosphoprotein of 55 kDa (SKAP-55)

ADAP constitutively associates with another adaptor protein, SKAP-55 or Scap1 (NCBI assignment).<sup>25,67</sup> SKAP-55 is a T-cell-specific adaptor protein which comprises a unique N-terminal region followed by a pleckstrin-homology (PH) domain and a carboxy-terminal SH3 domain (Fig. 1). The SH3 domain of SKAP-55 associates with the proline-rich region of ADAP<sup>25,68</sup> and, at the same time, the tyrosine-based RKxxYxxY motif in SKAP-55 mediates ADAP SH3 domain binding.<sup>69</sup> Significant progress has been made in identifying the role of this adaptor in T-cell function. A recent study demon-

strated that SKAP-55 potently increases T-cell/APC conjugate formation and enhances integrin-mediated adhesion.<sup>65</sup> SKAP-55 co-localizes with F-actin at the T-cell/APC synapse, enhances cellular adhesion via binding of LFA-1 to fibronectin and intercellular adhesion molecule-1 (ICAM-1), and promotes clustering of LFA-1.<sup>65</sup> SKAP-55-induced integrin clustering, in turn, mediates high-avidity integrin binding. T-cell/APC conjugate formation induces the translocation of SKAP-55 to lipid rafts, a process which is regulated by LFA-1 and TCR engagement. The translocation of SKAP-55 to the membrane brings it into close proximity with other signalling components which include the kinase Fyn, which has the capacity to phosphorylate SKAP-55, linking this protein to downstream signalling events such as the enhancement of ERK activity.<sup>69</sup> During T-cell/APC interactions, engaged receptors and their associated tyrosine kinases assemble at the area of cell contact into spatially organized supramolecular activation clusters (SMACs), which comprise an inner central SMAC (cSMAC) enriched with TCR, CD2 and CD28 surrounded by a peripheral SMAC (pSMAC) enriched with LFA-1.<sup>70,71</sup> Adhesive interactions play a central role in the formation of SMACs, and since ADAP and SKAP-55 have been shown to enhance LFA-1-mediated adhesion during T-cell/APC interactions, it is likely that these adaptors control the formation of SMACs. However, future studies will help to clarify the role of these adaptors in the formation of the immunological synapse.

#### FUTURE DIRECTIONS IN ADAPTOR PROTEIN RESEARCH

The recent identification of adaptor proteins has greatly contributed to our understanding of the way in which membrane proximal signalling events lead to the initiation and integration of downstream signalling pathways in T cells. By acting as scaffolds for the assembly of proteins, these proteins act as integrators of T-cell signalling. Although many adaptor proteins have been characterized, the biological functions of a host of others need to be elucidated. Understanding the functions of the different signalling domains of adaptors will require some complex studies in which adaptors with defined mutations in their signalling domains are introduced back into genetically modified mice or deficient cell lines. Further to this, information on the assembly, spatial dynamics and intracellular location of adaptor complexes will help to elucidate the precise mechanisms by which adaptors integrate diverse signalling networks. New imaging technologies, coupled with the development of new fluorescent labels and sensors and the use of more sophisticated computer software for image acquisition and analysis, may provide insights into the dynamics of adaptor proteins and their interactions with other signalling components in living cells.<sup>72-74</sup> Hopefully, such studies will lead to a better understanding of TCR signalling in the future.

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