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# Signal Transduction During Activation and Inhibition of Natural Killer Cells

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Natural killer (NK) cells provide an early defense against intracellular pathogens, such as parasites and viruses, and protect against spontaneous tumors. The NK cell response is not only cytotoxicity, which is mediated through polarized release of cytolytic granules towards target cells, but also secretion of cytokines and chemokines. In many cases, the potent release of cytokines, such as interferon (IFN)- $\gamma$  and tumor necrosis factor (TNF)- $\alpha$ , is sufficient for the effector function of NK cells, in the absence of perforin-dependent killing of other cells (Strowig *et al.*, 2008). Removal of tumor cells by NK cells occurs not only through a perforin-dependent pathway, but also by expression of the Fas ligand and 'tumour necrosis factor (TNF)-related apoptosis-inducing ligand' (TRAIL) (Smyth et al., 2002). NK cells belong to the innate immune system, but they share some features of adaptive immunity, and are integrated in the immune system as a whole, through reciprocal regulation with dendritic cells, macrophages, and other cell types (Moretta et al., 2008). For example, NK cells can promote a T helper type 1 (Th1) response through their ability to prime dendritic cells and to produce interferon (IFN)-y. Priming in vivo is required for NK cells to become responsive and this step involves specific interactions with other cells types. There are four distinct phenotypic subsets of NK cells: CD16<sup>+</sup> CD56<sup>dim</sup> NK cells in peripheral blood, CD16<sup>-</sup> CD56<sup>bright</sup> NK cells that are preferentially recruited into tissues through expression of L-selectin (CD62L) (Cooper et al., 2001), mucosal NK cells that produce IL-22 (Vivier et al., 2009), and uterine NK cells that may promote proper vascularization during early pregnancy (Ashkar et al., 2000; Hanna et al., 2006; Rajagopalan et al., 2006).

In the absence of a receptor repertoire generated by somatic DNA recombination and mutation, NK cells must rely on innate, germ-line encoded receptors to distinguish healthy cells from those that should be disposed of. To achieve this task, NK cells use a large number of receptors, each one with unique specificity for ligands and signaling properties. Some of the ligands are expressed broadly on many cell types, while others are expressed predominantly by hemapoietic cells. The expression of several ligands of NK cell activation receptors is induced as a result of infection, stress, or cell transformation. Some of the NK cell receptors have inhibitory functions and serve to protect healthy cells by recognizing major histocompatibility complex (MHC) class I molecules.

The focus of this Unit is on signaling by receptors that activate NK cell effector function, and those that inhibit NK cell responses. How NK cells respond to soluble stimulators, such as cytokines and chemokines, is fairly well understood and will not be reviewed here. However, it should be noted that the response of NK cells to cytokines and chemokines during viral infections is highly regulated by positive and negative signaling through different signal transducers and activators of transcription (STAT) molecules (Lee *et al.*, 2007; Nguyen *et al.*, 2002). To a great extent, it is generally assumed that signaling components in NK cells participate in signal transduction in a way similar to pathways

established in other cells. However, it is important to keep in mind that it is not always the case. As much as possible, work done with NK cells will be highlighted here.

# NK cell activation

NK cell cytotoxicity is a highly regulated process. It involves NK cell adhesion to target cells and the interaction between activating NK cell receptors and their respective ligands on the target cell surface. This induces intracellular signaling pathways resulting in polarization and release of cytotoxic granules towards the attached target. Various experimental systems have been used to study signaling after receptor stimulation. In this respect, it is worth noting that crosslinking with Abs is no substitute for the receptor–ligand interactions that occur upon contact of NK cells with other cells. Likewise, NK cell lines may not faithfully reproduce the signal transduction that occurs in primary NK cells.

#### Activation Receptors in Natural Cytotoxicity

In contrast to T or B cells, NK cells do not possess a single activating receptor that dominates their regulation. Instead, they possess a large array of receptors, which can act in synergy to regulate NK cell effector functions (Bryceson *et al.*, 2006a; Lanier, 2005). These receptors recognize ligands on infected or transformed cells (Bottino *et al.*, 2005; Gasser and Raulet, 2006) and thereby enable NK cells to detect and eliminate these cells. Interestingly, freshly isolated resting human NK cells can only be activated by triggering two or more activating receptors in combination (Bryceson *et al.*, 2006b). The only exception seems to be the low affinity Fc receptor Fc  $\gamma$ RIIIa (CD16), which is sufficient by itself to induce activation of resting NK cells. Therefore, all stimulatory receptors for natural cytotoxicity should be considered as co-activating. In the following we will discuss the receptor proximal signaling pathways of the different co-activating NK cell receptors.

Natural Cytotoxicity Receptors (NCR)-The NCR NKp30, NKp44, and NKp46 were among the first identified activating receptors in NK cells (Cantoni et al., 1999; Pende et al., 1999;). While NKp46 expression is specific for NK cells, NKp30 and NKp44 can also be found on subsets of T cells. NCR are immunoglobulin-like type I transmembrane glycoproteins that are expressed as monomers and contain a positively charged amino acid in their transmembrane domain. This charge facilitates the association of the NCR with signaling partner chains containing a negative charge in their transmembrane region. NKp46 and NKp30 associate with disulfide-linked homodimers or heterodimers of the TCR  $\zeta$  and/or the FcR  $\gamma$  chains, whereas NKp44 associates with the disulfide-linked homodimer DAP12. All these signaling partners contain an immunoreceptor-based activation motif (ITAM) in their cytoplasmic tail. While several viral ligands for the NCR have been identified (Arnon et al., 2006), the cellular ligands for these receptors remain poorly defined (Byrd et al., 2007). The intracellular protein BAT3 has been described as an NKp30 ligand (Pogge von Strandmann et al., 2007). How BAT3 is exposed on the surface of target cells is unknown. NCR-mediated NK cell activation is critically dependent on the ITAM motif of the associated partner chains and is very similar to the extensively studied signal transduction of the T cell receptor or B cell receptor. Briefly, the most important signaling steps are as follows (Figure 1). NCR engagement induces Src-family kinase-mediated tyrosine phosphorylation of the ITAM sequence in the associated partner chains. This induces binding of tyrosine kinases Syk and ZAP70 to the ITAM. These kinases phosphorylate transmembrane adapter molecules such as LAT and NTAL leading to the association, phosphorylation and activation of several signaling complexes, including phosphatidylinositol-3-OH kinase (PI3K), phospholipase C (PLC-y1, PLC-y2) and Vav-1, 2, 3. There are differences between human and mouse NK cells and between different ITAM-coupled

receptors as to the importance of the different PLC-γ and Vav isoforms (Billadeau *et al.*, 1998; Cella *et al.*, 2004; Tassi *et al.*, 2005; Ting *et al.*, 1992; Upshaw *et al.*, 2005).

**NKG2D**—The C-type lectin-like receptor NKG2D is a type II transmembrane glycoprotein which is expressed as a disulfide-linked homodimer on the surface of NK cells,  $\gamma\delta$  T cells and CD8<sup>+</sup> T cells (Bauer et al., 1999). NKG2D carries an arginine residue in the central portion of its transmembrane region, which mediates the association with the signaling partner chain DAP10 (Wu et al., 1999). DAP10 is also a disulfide-linked homodimer. One homodimer of NKG2D associates with two homodimers of DAP10, thereby forming a hexameric complex (Garrity et al., 2005). NKG2D can bind several ligands such as the human MICA/B and ULBP1, 2, 3, or 4, and the mouse Mult1, H60, and the Rae-1 family. The expression of these ligands is up-regulated on infected, stressed, or transformed cells (Gasser et al., 2005; Joncker and Raulet, 2008). This makes NKG2D an important coactivating receptor for the recognition of tumors (Guerra et al., 2008). Signaling of human NKG2D is critically dependent of DAP10, while a mouse-specific splice variant of NKG2D can also associate with DAP12 (Diefenbach et al., 2002). The short cytoplasmic tail of DAP10 contains a tyrosine based signaling motif (YINM), which can be phosphorylated by Src-family kinases and bind either the p85 subunit of PI3K or the adapter molecule Grb2 (Figure 1). Binding of both molecules is essential for NKG2D-mediated Ca<sup>2+</sup> flux and cytotoxicity (Upshaw et al., 2006). PI3K activity is important for the production of phosphatidyl-inositol-3,4,5-trisphosphate (PIP3), facilitating the plasma membrane recruitment of pleckstrin homology domain containing proteins such as PLC- $\gamma 1$ , PLC- $\gamma 2$ , Vav1 and Tec-family kinases. Grb2 recruits Vav1 and is important for the phosphorylation and activation of SLP76 and PLC- $\gamma 2$ . Vav1 is important for the induction of actin reorganization and the polarization of the microtubule organizing center (MTOC) toward target cells (Cella et al., 2004; Graham et al., 2006). NKG2D signaling does not require Syk or ZAP70 kinases, and is independent of the transmembrane adapters LAT and NTAL (Billadeau et al., 2003; Chiesa et al., 2006; Zompi et al., 2003).

2B4, NTB-A, and CRACC-2B4, NTB-A and CRACC are members of the SLAMrelated receptors (Claus et al., 2008). The expression of these monomeric immunoglobulinlike type I transmembrane glycoproteins is not limited to NK cells and can be found on many other immune cells. While 2B4 binds to CD48, which is widely expressed on cells of hematopoietic origin, NTB-A and CRACC are homophilic. The intracellular domain of these receptors contains immunoreceptor tyrosine-based switch motifs (ITSM), which can be phosphorylated by Src-family kinases upon receptor cross-linking (Figure 1). The phosphorylated ITSM then recruits small SH2-domain containing adapters such as SAP (SH2D1A) (Sayos et al., 1998), EAT-2 and ERT (which is only expressed in mice) (Eissmann et al., 2005; Eissmann and Watzl, 2006; Roncagalli et al., 2005). SAP then recruits the Src-family kinase Fyn by an untypical SH2-SH3 domain interaction (Chan et al., 2003; Latour et al., 2003). This interaction is crucial for 2B4-mediated NK cell activation. Mutations in SAP are the cause for a genetic disorder called X-linked lymphoproliferative disease (XLP). These patients show an uncontrollable and fatal lymphoproliferation in response to EBV infection (Nichols et al., 2005). 2B4 and NTB-A-mediated NK cell activation is defective in XLP patients (Benoit et al., 2000; Bottino et al., 2001; Nakajima et al., 2000; Parolini et al., 2000; Tangye et al., 2000), demonstrating the importance of SAP for the function of these receptors. Similarly, 2B4-mediated NK cell activation is also defective in Fyn-deficient mice (Bloch-Queyrat et al., 2005). CRACC signaling is independent of SAP and seems to rely on the adapter EAT-2 (Bouchon et al., 2001; Cruz-Munoz et al., 2009). Another adapter protein that binds to phosphorylated 2B4 is 3BP2 (Saborit-Villarroya et al., 2005). Phosphorylated 3BP2 interacts with the signaling molecules Vav1, LAT, and PLC-γ. Stimulation of 2B4 leads to phosphorylation of LAT,

Vav1, PLC-γ1, c-Cbl, Grb2 and SHIP (Bottino *et al.*, 2000; Chen *et al.*, 2004; Watzl *et al.*, 2000).

Under certain circumstances 2B4, NTB-A and CRACC can also inhibit NK cell functions (Veillette, 2006). In humans inhibitory signaling of these receptors has been reported in rare cases of SAP deficiency, such as XLP patients (Parolini et al., 2000; Bottino et al., 2001), or in immature NK cells, which express low levels of SAP (Sivori et al., 2002). However, inhibitory signaling of 2B4 in mice has been reported. A recent report has suggested that the amount of 2B4 expression, the strength of cross-linking, and the expression level of SAP determine if 2B4 is activating or inhibitory (Chlewicki et al., 2008). High levels of 2B4 expression, heavy cross-linking, and low SAP expression would favor inhibitory signaling of the receptor. The basis for this inhibitory signaling is not completely understood. Phosphorylated 2B4 can recruit the phosphatases SHP-1, SHP-2, and SHIP and the inhibitory kinase Csk (Eissmann et al., 2005). SAP competes with these inhibitory molecules for binding to the receptor, which could explain the negative signaling in the absence of SAP. EAT-2 and ERT can mediate inhibitory signaling by 2B4 when bound to the receptor (Roncagalli et al., 2005). However, as EAT-2 can induce positive signals when bound to NTB-A (Eissmann and Watzl, 2006) or CRACC (Cruz-Munoz et al., 2009), the basis for its inhibitory signaling when bound to 2B4 is controversial.

**Other Receptors**—Other co-activating NK cell receptors include DNAM-1, NKp80, and CD2. Not much is known about the signal transduction of these molecules. DNAM-1 (CD226, also known as PTA-1) is an immunoglobulin-like type I transmembrane glycoprotein that is expressed on NK cells, T cells and monocytes (Shibuya *et al.*, 1996). The ligands for DNAM-1 are poliovirus receptor (CD155) and Nectin-2 (CD112), both of which are expressed on epithelial and endothelial cells and are over-expressed on certain tumors (Bottino *et al.*, 2003; Tahara-Hanaoka *et al.*, 2004; Iguchi-Manaka *et al.*, 2008). The cytoplasmic tail of DNAM-1 contains three tyrosine residues that can be phosphorylated by Src-family kinases upon receptor engagement and recruit actin-binding proteins (Ralston *et al.*, 2004; Shibuya *et al.*, 2003). Interestingly, DNAM-1-mediated NK cell activation is dependent on its association with lymphocytes function-associated antigen 1 (LFA-1) and DNAM-1 signaling is therefore involved in promoting cellular adhesion (Shibuya *et al.*, 2003).

The C-type lectin-like receptor NKp80 (also known as KLRF1) is a type II transmembrane glycoprotein that is expressed as a disulfide-linked homodimer on the surface of NK cells and a subset of CD8<sup>+</sup> T cells (Kuttruff *et al.*, 2009; Vitale *et al.*, 2001). The ligand for NKp80 is the activation-induced C-type lectin (AICL) (Welte *et al.*, 2006), which is selectively expressed on myeloid cells, such as monocytes and macrophages, and the NKp80-AICL interaction seems to contribute to reciprocal activation of NK and myeloid cells at sites of inflammation. NKp80 carries two tyrosine-based signaling motifs in its cytoplasmic tail, but nothing is known so far about the signal transduction of this receptor.

CD2 is a member of the immunoglobulin super-family and is expressed as a type I transmembrane glycoprotein on T cells and NK cells. The ligands for CD2 are CD58 (LFA-3) and, to a lower extent, CD48 in humans, but CD48 is the only ligand in mice (Davis *et al.*, 1998). The cytoplasmic domain of CD2 contains a proline-rich sequence that interacts with the Src-family kinase Lck (Bell *et al.*, 1996) and several adapter molecules (Li *et al.*, 1998). However, the exact signaling mechanism of CD2 is still unclear. Signaling through CD2 has been shown to enhance cellular adhesion and CD2 is linked to the cytoskeleton.

Several members of the killer cell immunoglobulin-like receptors (KIR) in humans, some of the Ly49 receptors in mice, and the CD94/NKG2C receptor complex carry a negative charge in their transmembrane region through which they associate with the ITAM-containing partner chain DAP12 (Lanier *et al.*, 1998; Lanier *et al.*, 1998; Smith *et al.*, 1998). Except for Ly49H and Ly49P in mice, which confer resistance to mouse cytomegalovirus (Arase *et al.*, 2002; Kielczewska *et al.*, 2009; Smith *et al.*, 2002), the function of activating isoforms in the KIR, Ly49, and CD94 receptor families remains unknown.

## Fcy Receptor Signaling

The CD56<sup>dim</sup> subset of NK cells expresses the Fc $\gamma$  receptor CD16, through which NK cells mount antibody-dependent cellular cytotoxicity (ADCC). CD16 associates with disulfidelinked homodimers or heterodimers of the TCR  $\zeta$  and/or the FcR  $\gamma$  chains (Lanier *et al.*, 1991). The signaling of CD16 follows therefore the classical ITAM pathway of Src-family kinase-mediated tyrosine phosphorylation, Syk association leading to PI3K, Vav1 and PLC- $\gamma$ 1 and PLC- $\gamma$ 2 activation (Billadeau *et al.*, 1998; Ting *et al.*, 1992). Signaling for ADCC involves a PI3K-dependent activation of the ADP-ribosylation factor (Arf)-6, which couples CD16 to the production of phosphatidylinositol-4,5-bisphosphate (PIP2) by a phosphatidylinositol-4-phosphate 5-kinase (PI5K) and the activation of phospholipase D (Galandrini *et al.*, 2005). PI5K activity in primary human NK cells is required for degranulation but not for granule polarization (Micucci *et al.*, 2008). In contrast to other ITAM coupled NK cell receptors, triggering of CD16 is by itself sufficient to activate cytotoxicity and cytokine production in resting human NK cells (Bryceson *et al.*, 2006b). So far it is unknown which part of the CD16 signaling cascade is responsible for this unique ability of this receptor.

## Signaling by Integrins

NK cells express several types of integrins, including LFA-1 ( $\alpha$ L\beta2, CD11a/CD18) and VLA-4 ( $\alpha$ 4 $\beta$ 1, CD49d/CD29). Integrin  $\alpha$ 4 $\beta$ 1 binds to the vascular cell adhesion molecule (VCAM)-1 (CD106), an Ig-like molecule expressed on activated endothelial cells, and to the extracellular matrix component fibronectin. LFA-1 binds to ICAM-1, ICAM-2, and ICAM-3, which are expressed on endothelium and upregulated during inflammation. Interaction of integrins with their ligand on other cells is a complex process, as it requires an 'inside-out' signal from other receptors that promotes a change in the integrin from a bent, low affinity conformation to extended forms, which are of intermediate or high affinity. The strength of adhesion is determined by the integrin affinity and avidity for the ligand, and by the interaction of the integrin cytoplasmic domains with the cytoskeleton and signaling molecules. The  $\beta 1$  and  $\beta 2$  integrin cytoplasmic tails bind to talin, a protein that links them to the actin cytoskeleton and is essential for 'inside-out' activation, even when such signals originate from different receptors and signaling pathways (Cantor et al., 2008). Inside-out signaling, which has been studied mainly following stimulation of the T-cell receptor (TCR), requires the scaffolding proteins 'adhesion and degranulation-promoting adapter protein' (ADAP, formerly known as SLAP-130 or Fyb) and 'Src kinase-associated phosphoprotein of 55kDa' (SKAP55) (Mor et al., 2007; Wang and Rudd, 2008). ADAP is required, independently of LFA-1, for polarization of the MTOC and recruitment of dynein in T cells (Combs et al., 2006). The molecule cytohesin is a guanine-nucleotide exchange factor for Arf-6, and promotes inside-out signaling through binding to the membraneproximal part of the  $\beta$ 2 cytoplasmic tail (Geiger *et al.*, 2000). The small GTPase Rap1 and its effector 'regulator for cell adhesion and polarization enriched in lymphoid tissues' (RAPL) are key elements of inside-out signaling. RAPL binds directly to the aL subunit of LFA-1. Furthermore, WAVE2 regulates inside-out signaling by recruiting vinculin and talin to the immunological synapse and by promoting Rap1 activation through a CrkL-C3G pathway (Nolz et al., 2007; Nolz et al., 2008). The current model (Figure 2) proposes that a

complex RAPL–Rap1GTP–RIAM interacts with talin, and that RIAM provides a link to the ADAP/SKAP-55 complex (Cantor *et al.*, 2008; Menasche *et al.*, 2007; Mor *et al.*, 2007; Wang and Rudd, 2008). C3G and RIAM are guanine exchange factors for Rap1. However, ADAP is dispensable for NK cell function, including adhesion, cytotoxicity, and cytokine production (Fostel *et al.*, 2006). This could either reflect the fact that an inside-out signal is not essential for LFA-1 signaling in NK cells, or be due to a redundancy of ADAP with another signaling component.

The importance of  $\beta 2$  (CD18) integrins is underscored by the severe infections suffered by individuals with homozygous mutations in CD18, a disease known as leukocyte adhesion deficiency type I (LAD-I). Recently, the genetic basis for the even more severe LAD-III disease has been mapped to the gene encoding Kindlin-3 (Hidalgo and Frenette, 2009). Kindlin-3, which is expressed in hematopoietic cells, binds to the cytoplasmic tail of  $\beta 1$ ,  $\beta 2$ , and  $\beta 3$  integrins and promotes inside-out signaling. Other proteins that link  $\beta 1$  and  $\beta 2$  integrin directly to the cytoskeleton include filamin and  $\alpha$ -actinin (Cantor *et al.*, 2008). Conformational changes in integrins, which are required to achieve high affinity binding to ligands, are facilitated by force, as it occurs during adhesion under flow (Alon and Dustin, 2007). Linkage of integrins to the underlying cellular cytoskeleton is required for their response to shear forces and the ensuing switch to higher affinity states.

**β1 Integrins**—Protein kinase D1 (PKD1) promotes clustering and activation of integrin  $\beta$ 1 in T cells through membrane recruitment of Rap1 (Burbach *et al.*, 2007). The analysis of 'outside-in' signaling by  $\beta$ 1 integrins is complicated due to a number of *cis* interactions with tetraspannin proteins, such as CD9, CD63, and CD81, which recruit the signaling enzymes PI4K, PKCa and PKC $\beta$ II (Berditchevski, 2001). The cytoplasmic tail of  $\beta$ 1 integrin binds to integrin-linked kinase (ILK), which phosphorylates myosin light chain and Akt (Wu and Dedhar, 2001). Through a direct association of  $\alpha$ 4 with paxillin,  $\alpha$ 4 $\beta$ 1 integrin provides an 'inside-out' signal to the  $\beta$ 2 integrin LFA-1 (Cantor et al., 2008).

Engagement of  $\beta$ 1 integrin specifically on NK cells results in phosphorylation of tyrosine kinase Pyk2 and the cytoskeleton protein paxillin, and formation of a phospho-Pyk2– paxillin complex (Gismondi *et al.*, 1997).  $\beta$ 1 integrin crosslinking induces formation of a Shc-Pyk2-Grb2 complex and leads to Erk-dependent secretion of interferon- $\gamma$  (Mainiero *et al.*, 1998). Furthermore, NK cells secrete IL-8 in response to  $\beta$ 1 integrin binding to fibronectin through a signaling pathway that involves activation of the small GTPase Rac1 by Vav, and of the Rac1 effector Pak1, which leads to activation of the p38 MAPK (Mainiero *et al.*, 2000).

**β2 Integrins**—Some of the known interactions of LFA-1 with signaling components are depicted in Figure 2. Most of them are derived from experiments performed with T cells. It is therefore important to keep in mind that LFA-1 has unique signaling properties in NK cells, which are not shared with LFA-1 in T cells. First, in resting NK cells, inside-out signaling is not strictly required for binding of LFA-1 to ICAM-1 (Barber and Long, 2003). Slow and signaling-dependent binding of resting NK cells to ICAM-1 implies that LFA-1 can transmit its own inside-out signal to other LFA-1 molecules. However, co-engagement of NK cell activation receptors 2B4 and CD2 results in stronger inside-out signals (Barber and Long, 2003). Second, binding of LFA-1 to ICAM-1, either attached to beads or expressed on insect cells, is sufficient to promote polarization of cytolytic granules (Barber *et al.*, 2004; Bryceson *et al.*, 2005). This contrasts with signaling in T cells, where LFA-1 cannot signal on its own. Nevertheless, LFA-1 engagement in T cells does enhance TCR-dependent polarization of cytolytic granules (Anikeeva *et al.*, 2005). In addition, Mg<sup>2+</sup>-induced binding of the high affinity form of LFA-1 on T cells to ICAM-1 stimulates F-actin reorganization independently of TCR signals (Porter *et al.*, 2002).

As first described in platelets with  $\beta$ 3 integrin, which activates MAP kinase signaling through Vav, integrins can signal independently of actin polymerization (Miranti *et al.*, 1998). In NK cells, engagement of LFA-1 by ICAM-1 on insect cells was sufficient to induce actin-independent activation of the Vav1-Rac1 pathway (Riteau *et al.*, 2003). Therefore, LFA-1 has the unique ability to provide early activation signals to NK cells. These early LFA-1-dependent signals are sensitive to inhibition by MHC class I-specific inhibitory receptors (Barber *et al.*, 2004). Consistent with its ability to signal independently of actin polymerization, LFA-1 engagement on NK cells induces phosphorylation of the Wiskott-Aldrich syndrome protein (WASp), a known regulator of the actin cytoskeleton (Gismondi *et al.*, 2004), and phosphorylation of Pyk2 and Pyk2 association with paxillin (Gismondi *et al.*, 2003).

The tyrosine kinase Syk binds directly to integrin  $\beta$  chains through its SH2 domain but in a phospho-tyrosine independent way (Woodside *et al.*, 2002). Signaling by  $\beta$ 2 and  $\beta$ 3 integrins in neutrophils, platelets, and myeloid cells requires Syk and an ITAM-containing transmembrane adapter molecule, which can be either DAP12 or the FcR  $\gamma$  chain (Jakus *et al.*, 2007). Similarly, calcium signaling through the  $\beta$ 2 cytoplasmic tail of LFA-1 in T cells required the TCR, a requirement that could be fulfilled by the TCR  $\zeta$  chain alone (Sirim *et al.*, 2001). It remains to be seen whether these unusual signaling properties contribute to the unique ability of LFA-1 to trigger granule polarization in primary NK cells.

#### Key Events in NK cell Activation

NK cell activation uses highly redundant signaling pathways originating from many different surface receptors (Figure 1). Additionally, while several critical and related signaling molecules are expressed selectively in B-cells (e.g. Syk and SLP-65) and T-cells (e.g. ZAP70 and SLP-76), NK cells express all of these proteins. Therefore, it has been difficult to identify critical signaling pathways using genetic approaches (Tassi *et al.*, 2006). Furthermore, even the genetic deletion of both Syk and ZAP70 does not significantly impair NK cell development and cytotoxic function. NK cell activity is also largely intact in mice lacking all three ITAM-containing signaling subunits DAP12, TCR  $\zeta$  and FcR  $\gamma$  (Chiesa *et al.*, 2006; Colucci *et al.*, 2002). However, a few molecules and signaling pathways have emerged as essential for NK cell activation.

**Src-family Kinases**—Basically, all activating NK cell receptors depend for their phosphorylation and function on the activity of Src-family kinases. This makes this family of kinases essential for NK cell functions. NK cells express many different members of this family, including Lck, Fyn, Src, Lyn, Yes and Fgr, and the activity of these kinases is highly redundant. Therefore, no single Src-family kinase is essential for NK cell function (Mason *et al.*, 2006; van Oers *et al.*, 1996). However, chemical inhibition of Src-family kinases efficiently blocks NK cell activation mediated by ITAM and non-ITAM coupled receptors. Several of the Src-family kinases are enriched in membrane microdomains, also referred to as lipid rafts, through the posttranslational addition of a lipid anchor (Cherukuri *et al.*, 2001).

The initial step in co-activating NK cell receptor signaling seems to be receptor clustering induced by ligand binding. Receptor cross-linking either by ligands, some of which are preclustered on target cells (Eleme *et al.*, 2004), or plate-bound antibodies is sufficient to induce signal transduction by all co-activating NK cell receptors. Whether ligand-induced conformational change occurs in these receptors is unknown. Receptor clustering induces a concentration of co-activating receptors such as 2B4 or NKG2D in membrane microdomains that are enriched in Src-family kinases, and where receptor phosphorylation is facilitated **PLC-\gamma2**—Exocytosis of lytic granules in NK cells is strongly dependent on PLC- $\gamma$  phosphorylation and subsequent mobilization of Ca<sup>2+</sup> from the endoplasmic reticulum. While the two isoforms of PLC- $\gamma$  (PLC- $\gamma$ 1 and PLC- $\gamma$ 2) may be redundant in ITAM signaling pathways, signaling by the NKG2D/DAP10 receptor has an exclusive requirement for PLC- $\gamma$ 2 (Upshaw *et al.*, 2005). PLC- $\gamma$ 2 is essential for NK cell cytotoxicity (Caraux *et al.*, 2006; Tassi *et al.*, 2005) and over-expression of PLC- $\gamma$ 1 can only partially rescue the cytotoxicity defect in PLC- $\gamma$ 2 deficient NK cells (Regunathan *et al.*, 2006).

**Vav**—Vav proteins are essential components for NK cell activation pathways. NK cells express all three Vav isoforms (Vav1, Vav2 and Vav3), all of which have guanine nucleotide exchange factor (GEF) activity and thereby activate small GTPase proteins of the Rho family, primarily Rac1. These GTPases are important mediators of actin cytoskeleton reorganization, which is essential for the clustering of co-activating receptors and the polarization of the lytic granules toward the target cell (Figure 3). Rac1 also activates a PAK1–MEK–Erk pathway, which is important for granule polarization and release (Jiang *et al.*, 2000). Vav1 is required for DAP10-mediated signaling pathways, whereas Vav2 and Vav3 are essential for the function of ITAM-coupled receptors (Cella *et al.*, 2004; Chan *et al.*, 2001; Colucci *et al.*, 2001). Absence of all three Vav isoforms severely compromises NK cell cytotoxicity (Cella *et al.*, 2004). The central role of Vav in the activation of NK cell responses is highlighted by the fact that it is selectively inactivated during inhibition by MHC class I-specific receptors as a preferred substrate of the ITIM-bound tyrosine phosphatase SHP-1 (Figure 3).

**Granule Polarization and Degranulation**—Killing of target cells by cytotoxic lymphocytes is a complex process (Fischer *et al.*, 2007; Orange, 2008; Stinchcombe and Griffiths, 2007). The effector cell has to reorient its cytoskeleton, including the MTOC, and move cytolytic granules towards the target cell. Granules have to make contact with the plasma membrane where they will eventually fuse and release their content into the synaptic cleft. Studies of NK cells from patients with genetically defined immuno-deficiencies have identified the Wiskott-Aldrich syndrome protein (WASp) as an essential component of actin-dependent formation of the NK cell immune synapse (Gismondi *et al.*, 2004; Orange *et al.*, 2002a).

Several signals can induce granule polarization in NK cells. Granule polarization in a human NK cell line during contact with sensitive target cells required activation of the MAPK Erk through a PI3K-Rac1-MEK pathway (Jiang et al., 2000). Pharmacological inhibition of another MAP kinase, JNK, suggested that it is required for polarization (Li et al., 2008). A role for tyrosine kinase Pyk2 in polarization is supported by experiments with a dominantnegative mutant of Pyk2, which blocked MTOC and paxillin movement to the synapse, and NK cell cytotoxicity (Sancho et al., 2000). There may be a direct link between cytolytic granules and the actin cytoskeleton during polarization. Granule polarization in NK cells is dependent on the WASp-interacting protein (WIP) (Krzewski et al., 2008). A key connection between microtubules and actin during granule polarization may be provided by the Cdc42-interacting protein-4 (CIP4), a molecule known to bind directly to tubulin. CIP4 is associated with WASp at the MTOC and is required for MTOC polarization to the NK cell immune synapse (Banerjee et al., 2007). Myosin IIA, which is known as a regulator of actin dynamics, is required for a late step in the degranulation pathway, downstream of granule polarization in the cell line NKL (Andzelm et al., 2007). A feature unique to human NK cells is that LFA-1 binding to ICAM-1 is sufficient to induce granule polarization

(Barber *et al.*, 2004; Bryceson *et al.*, 2005), which is not the case in T cells. Furthermore, in mouse NK cells that were derived from embryonic stem cells polarization required coengagement of LFA-1 and NKG2D, and was dependent on talin (Mace *et al.*, 2009).

As shown in T cells, the Rab27a GTPase effector protein Munc13-4, which is defective in patients with familial hemophagocytic lymphohistiocytosis 3 (FHL3) (Fischer *et al.*, 2007), is required for NK cell degranulation (Marcenaro *et al.*, 2006). FHL4 is a disease caused by mutations in the 'soluble N-ethylmaleimide-sensitive factor' (SNF) attachment protein receptor' (SNARE) syntaxin 11. The defect could be traced to a lack of degranulation by NK cells (Arneson *et al.*, 2007; Bryceson *et al.*, 2007). Expressed mutations of the 'NF- $\kappa$ B essential modifier' (NEMO) result in a syndrome known as hypohidrotic ectodermal dysplasia with immune deficiency (HED-ID). NK cells from HED-ID patients were defective in natural cytotoxicity but not ADCC (Orange *et al.*, 2002b). How activation of the transcription factor NF- $\kappa$ B by NEMO contributes to NK cell cytotoxicity is still unknown.

# INHIBITION

It was appreciated long ago that the activity of NK cells was under stringent negative regulation. Specifically, normal expression of MHC class I molecules protects cells from lysis by NK cells (Ljunggren and Kärre, 1985). These early findings were given a mechanistic basis when NK cell inhibitory receptors specific for MHC class I were identified in mouse and man (Ciccone et al., 1992; Karlhofer et al., 1992). As knowledge about these inhibitory receptors advanced, it became clear that they were the prototypes of a very large group of receptors with similar signaling properties (Long, 2008). The definition of a sequence motif, now known as the 'immunoreceptor tyrosine-based inhibition motif' (ITIM), for binding of the tyrosine phosphatase SHP-1 to human inhibitory 'killer cell Iglike receptors' (KIR) (Burshtyn et al., 1996) led to the identification of many other ITIMcontaining receptors with inhibitory properties in various types of cells (Long, 2008). Although the ITIM is evolutionary ancient, conserved from bony fish to primates, some of the ITIM-containing receptor families have evolved rapidly. For example, the inhibitory receptors for classical MHC class I molecules in mice, the Ly49 family, are completely different from their counterparts in humans, which are members of the KIR family. Despite the structural differences between KIR and the lectin-like Ly49 and CD94-NKG2A, signaling through the conserved ITIM is remarkably similar.

#### Inhibitory Receptors

There are many ITIM-containing receptors on different cell types and they bind to a large variety of ligands (Long, 2008). NK cells alone express several of these ITIM-containing receptors, some of which bind to non-MHC ligands. The human KIR family (also known as CD158) includes inhibitory receptors for HLA-B and HLA-C. In the mouse, members of the Ly49 family bind to H-2 class I molecules with varying degrees of specificity. The ITIMcontaining CD94-NKG2A (also known as KLRC1 and CD159) heterodimeric receptor is conserved in both species but the ligands are not. Human CD94-NKG2A binds to the nonclassical MHC molecule HLA-E, whereas the ligand in the mouse is Qa1. LILR (also known as ILT, LIR, and CD85) is the third family of ITIM-containing receptors on human NK cells that bind to MHC class I, and they display permissive binding to several different MHC class I molecules. In addition to these MHC-specific receptors, NK cells express several other ITIM-containing receptors (Long, 2008): NKR-P1 (CD161) binds to human LLT1 and to mouse Ocil/Clrb molecules; Siglec-7 (CD328) binds to sialic acids; LAIR-1 (CD305) binds to collagen; KLRG1 binds to cadherins; CEACAM-1 (also known as BGP and CD66a) binds to itself; PILRa binds to CD99; and finally CD300a (also known as IRp60), which has no known ligand. Considering how many cells express ligands for any one of

these inhibitory receptors, one wonders how much inhibition has to be overcome during NK cell activation by contact with target cells.

Individual receptors within the KIR, Ly49, and CD94-NKG2 families are expressed randomly among NK cells. Thus, each individual NK cell expresses its own small repertoire of inhibitory receptors. Furthermore, some of these receptors may have a self MHC class I ligand, but others may not. Therefore, within most individuals, there are NK cells that have no inhibitory receptor for self, and others that express several. To ensure proper balance between activation and inhibition for every NK cell, NK cells undergo a form of education, which is determined by the MHC class I repertoire of the host (Anfossi *et al.*, 2006; Gasser and Raulet, 2006; Yokoyama and Kim, 2006). The more MHC class I detected by inhibitory receptors on any given NK cell, the more intrinsically responsive that NK cell will be. Conversely, NK cells with no inhibitory receptor for self MHC will be hyporesponsive. The education mechanism is not known yet. It could be a form of anergy provoked by continuous stimulation due to weak inhibition (Gasser and Raulet, 2006). Alternatively, inhibitory receptors may deliver an instructive signal that permits acquisition of responsiveness (Yokoyama and Kim, 2006).

# **Inhibitory Signals**

The first step in the inhibitory signaling pathway is phosphorylation of ITIM sequences in the inhibitory receptor. How this is achieved is still unknown. The model proposing that ITIMs become phosphorylated in *trans* by tyrosine kinases that are involved in activation pathways is difficult to reconcile with evidence that inhibition occurs at a very early step, independently of actin polymerization (Figure 3). Once phosphorylated, ITIMs recruit mainly the tyrosine phosphatases SHP-1 and SHP-2. Relative selectivity for SHP-1 or SHP-2 among different ITIM-containing receptors has been observed (Long, 2008). Furthermore, some of the ITIM-containing receptors bind additional molecules through their phosphorylated ITIMs: LAIR-1 and human LILR bind the inhibitory 'C-terminal Src kinase' Csk; Siglecs bind SOCS3; KLRG1 binds SHIP; CD300 binds PI3K; and KIR binds  $\beta$ arrestin 2 (Yu *et al.*, 2008). Deficiency in  $\beta$ -arrestin 2 results in diminished inhibition through KIR. By an unknown mechanism, which involves binding to phosphorylated ITIM,  $\beta$ -arrestin 2 facilitates SHP-1 and SHP-2 recruitment to inhibitory KIR (Figure 3).

During inhibition of NK cells by an ITIM-containing receptor, most of the tyrosinephosphorylated proteins detectable during activation have greatly diminished phosphorylation. Rather than being a consequence of fairly 'permissive' dephosphorylation of many signaling components by the ITIM-bound SHP, it is due to inhibition of a very proximal signaling step, which is independent of actin polymerization, and occurs through dephosphorylation of Vav1 by SHP-1 (Figure 3). Given the central role of Vav1 in promoting Rac1-dependent actin cytoskeleton rearrangement, synapse formation, and clustering of receptors, this inhibitory mechanism prevents tyrosine phosphorylation of activation receptors even before it occurs (Endt *et al.*, 2007; Watzl and Long, 2003). The important concept here is not so much that Vav1 *per se* is selectively targeted for dephosphorylation, but rather that inhibition does rely on selection of a key component in a signaling pathway. This opens the possibility of multiple, finely tuned inhibitory mechanisms, which could be controlled by the engagement of different inhibitory receptors in different cellular contexts. It would be interesting to identify SHP substrates during inhibition by various ITIM-containing receptors in different cells.

The early, proximal, and actin-independent signals that lead to Vav1 activation during NK cell contact with other cells are not well characterized. However, engagement of integrins alone is sufficient to induce Vav1 phosphorylation (Gismondi *et al.*, 2003; Riteau *et al.*, 2003). This would be consistent with the model (Figure 3), since LFA-1-dependent NK cell

activation by ICAM-1 is blocked by co-engagement of KIR with its cognate ligand HLA-C (Barber *et al.*, 2004).

Inhibition of natural cytotoxicity by KIR and by CD94-NKG2A involves a second component in the negative signaling pathway besides Vav1 dephosphorylation by SHP-1 (Figure 3). During incubation of NK cells with target cells that express an MHC class I ligand for the inhibitory receptor, the small adapter Crk becomes phosphorylated and forms a complex with the tyrosine kinase Abl (Peterson and Long, 2008). Crk phosphorylation by Abl, and Abl binding to Crk, are steps known to dissociate Crk from the scaffold protein Cbl and from the Rap1-specific GEF C3G. The Cbl–Crk–C3G complex, as well as the CAS–Crk–C3G and the paxillin–Crk–C3G complexes, that form during NK cell activation may contribute to the strength of LFA-1-mediated adhesion through activation of Rap1 (Figure 2). Phosphorylated Vav1 is also bound to phosphorylated Cbl during activation (Peterson and Long, 2008). During inhibition, the Cbl–Crk–C3G complex is not simply absent, due to a position downstream of the point of inhibition, but it is actively disassembled through recruitment of a tyrosine kinase and specific phosphorylation of an adapter protein (Peterson and Long, 2008). The ability of ITIM-containing receptors to activate a specific tyrosine phosphorylation event suggests additional signaling functions of this large receptor family.

This Unit has described many positive and negative signaling pathways that act in parallel and in synergy to regulate NK cell responses. To achieve some understanding of these different signaling pathways it is necessary to study each one in isolation, and a few in controlled combinations, using reductionist approaches. However, a challenge for the future is to study and to understand the complex integration and interaction of these different signals in NK cells.

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# Figure 11.9B.1.

Complexity in Activation Pathways. The redundant signaling pathways emanating from ITAM-, DAP10-, and ITSM-coupled receptors are shown. NKG2D/DAP10 and the ITSM-coupled receptor 2B4 synergize for activation of primary resting NK cells.



### Figure 11.9B.2.

LFA-1 Signaling. For the sake of clarity, Src-family kinases, which are involved in many of the signaling steps, have not been included in the diagram. Most of the information shown is derived from data obtained with T cells. LFA-1 signaling in NK cells may be different in some respects. For example, inside-out signals for LFA-1 are not a stringent requirement in NK cells, and LFA-1 signaling in NK cells is sufficient to induce granule polarization. The detailed pathways leading to actin reorganization and to granule polarization are not known.

Watzl and Long



### Figure 11.9B.3.

Model of Inhibitory Signaling in NK Cells. Binding of  $\beta$ -arrestin 2 to phosphorylated ITIM enhances the recruitment of SHP-1 and SHP-2 to phosphorylated ITIMs. The dephosphorylation of Vav1 by SHP-1 blocks its GEF activity and activation of Rac1. Phosphorylation of the adapter Crk by Abl, which is induced by inhibitory receptor engagement, results in disassembly of the Cbl–Crk–C3G complex. Inhibition is dominant over activation because it occurs upstream of the actin-dependent clustering and phosphorylation of co-activation receptors.