

Distinct NF-kB activation pathways engaged by T-cell receptor and co-receptor CD28 on T-cells

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The transcription factor nuclear factor-kB (NF-kB) is critical for the induction of inflammatory responses in T-cells, survival and differentiation. Antigen receptor (TCR) and co-receptor CD28 are the central regulators of NF-kB activation in T-cells. Progress in understanding NF-kB activation in T-cells has occurred over the years with the identification of individual adapters such as ADAP and GRB-2 and enzymes such as PKC-0 that regulate NF-kB. However, little is known whether the engagement of distinct modules by the TCR and CD28 account for the cooperative effects of the two receptors in activating NF-kB. In this review, we discuss recent advances in our understanding of NF-kB regulation by TCR and CD28.

Keywords: Adaptors; ADAP; VAV-1; GRB-2; NF-kB

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Introduction

Nuclear factor-kB (NF-kB) includes a family of transcription factors that act as dimers and regulate genes involved in the inflammatory and immune responses, growth, development and differentiation. NF-kB family is comprised of five members: RelA (p65), RelB and c-Rel, and the precursor proteins NF-κB1 (p105) and NF-κB2 (p100), which are processed into p50 and p52, respectively [1]. Activation signals from antigen receptors, inflammatory cytokines and infections all lead to the processing of p105 and p100 and their dimerization with RelA component. Complex of NF-kB dimers bind to the kB sites on the promoters or enhances of various target genes leading to the activation or repression of a specific signaling pathway. Examples include the IL-2 promoter that concurrently binds to transcription factors NF-kB, NFAT and AP1 leading to the regulation of its expression during an inflammatory response [2]. Defects in the of NF-kB pathway is linked to the immune

disregulation such as inflammatory disorders, autoimmune diseases as well as cancer [3,4]. The NF-kB pathway therefore is tightly regulated at multiple checkpoints, and various receptors including CD28 are believed to use individual signalling components for its regulation. In resting T cells NF-kB dimers remain bound to inhibitors of κB (IκB) molecules. Stimulation by antigen receptors and co-receptors induces IKK (IkB Kinase) complex activation which is responsible for IkB phosphorylation leading its degradation via ubiquitin pathway. IKK consists of three subunits, two active kinases IKKβ catalytically IKKα, [5]. The main regulatory/structural subunit, NEMO consequence of IkB degradation is the liberation of NF-kB dimers from cytoplasmic retention pool and subsequent translocation to the nucleus. However, a second phase of NF-kB regulation occurs in the nucleus which includes post-translational modifications of NF-kB subunits to fine tune their activity on promoters and enhancers of multiple genes [1,6]. These regulatory steps are common to all cell http://www.smartscitech.com/index.php/ics

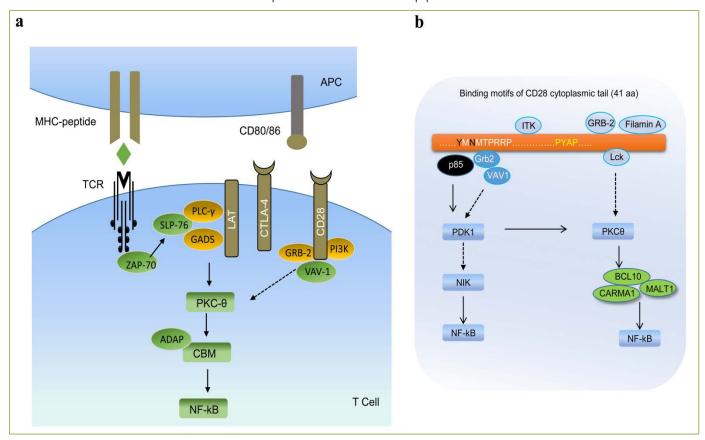


Figure 1. NF-kB activation in T-cells: a) upon engagement of the antigen receptor (TCR) by peptide-MHC, proximal signaling events activate tyrosine kinase, zeta-chain-associated protein kinase 70 (ZAP-70). ZAP-70 phosphorylates SLP-76 and LAT thereby recruiting several proteins resulting in the formation of a signalosome complex responsible for the activation of several downstream targets including protein kinase C theta (PKC-θ). PCK-θ is essential intermediate that couples TCR signals to NF-kB via its association with I kappa B kinase (IKK) and activation of Carma1-Bcl10-Malt1 (CBM) complex. IKK complex phosphorylates I kappa Bs (IkBs), leading to their polyubiquitination, release of NF-kB dimers and their translocation into the nucleus. ADAP is a scaffold protein that regulates CBM complex assembly upon TCR ligation and is required for TCR mediated NF-kB signaling but not CD28 pathway [23,25]. CD28 co-stimulation dependent NF-kB activation, on the other hand, is dependent on the binding of GRB-2 to its cytoplasmic tail. b) GRB-2 binds to the YMNF motif in the cytoplasmic tail of CD28, and a point mutation (N-Q) that selectively disrupts this binding, abrogates CD28 driven NF-kB activation. VAV1 is required for CD28 NF-kB activation likely via its association with IKKα and GRB-2^[23,39]. CD28 engagement also facilitates activation of PDK1 and its binding to the PKC-θ to activate NF-kB via classical pathway. CD28 ligation has been shown to activate alternate-like NF-kB pathway by recruiting p52/RelA dimers in a Vav1 dependent pathway, however, precise mechanism remains unclear ^[45]. In this situation, NIK, which is required for the alternate pathway is likely to be activated via PI3K-PDK1-AKT1 pathway upon CD28 engagement. PKC-θ also associates with CD28 in the immunological synapse via Lck, and regulates its localization and downstream signaling ^[46]

types including immune cells. However, NF-kB activation in T-cells is orchestrated by various unique signalling receptors and their potential crosstalk in the regulation of NF-kB has been unexplored. Proximal events upstream of IKK complex especially role of adapters in T-cells and NF-kB activation will be main subject of this review briefly touching on the other aspects of NF-kB regulation in T-cells.

Adapter molecules in T-cell activation

TCR and CD28 generate digital biochemical signals which are amplified and converted into effector functions in a series of well-defined signaling pathways ^[7-10]. One of the earliest events of antigen ligation is the activation of src and syk family kinases such as Lck and ZAP-70 respectively,

phosphorylating various enzymes and adapter proteins. Adapters are a unique group of proteins that have well defined structural domains but lack enzymatic activity ^[9,11]. Instead, they function as scaffolds bringing other proteins into the proximity of each other, nucleating and bridging multi-molecular signaling complexes ^[8,12]. Adapters can also induce conformational changes in their binding partners, potentially capable of regulating their activity and function ^[11,12]. Adapter scaffolds can regulate T-cell function in a positive or negative manner dependent upon the signaling complex and pathway. The importance of adapters in immune cell function and development has been underscored by *in vivo* and cell-line approaches, leading to important insights of their role in the transmission and integration of early signaling events generated by antigen receptor (TCR)

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to downstream effector functions. Several adapter for these functions are exclusively expressed in the hematopoietic cells, while others are more widely expressed^[12]. Characterization of key positive adapters in T-cells has helped in our understanding of T-cell activation mechanisms and regulation. Key positive adapters include:SH2-DOMAIN CONTAINING LEUKOCYTE-SPECIFIC PHOSPHOPROTEIN (SLP-76), mice null for SLP-76 have severe disruption in thymic T-cell development and IL-2 production [13-15]; LINKER FOR T-CELL ACTIVATION (LAT) is a transmembrane adapter with multiple cytoplasmic tyrosine phosphorylation/binding sites, an important signaling node to nucleate multi-protein complex of GADS, SLP-76, PLC-gamma, ITK and GRB-2. LAT or SLP-76 null mice displayed similar phenotype and a complete block in thymopoiesis at the pro-T3 stage [16]. In addition to this, Jurkat T-cell mutant lines lacking either SLP-76 or LAT expression showed a vital requirement of both molecules in TCR-induced PLC₇1 phosphorylation. extracellular signal-regulated kinase (ERK) activation, Ca²⁺ influx and IL-2 promoter activity [13,17,18]: GRB-2 RELATED ADAPTER DOWNSTREAM OF SHC (GADS) is a SH2and SH3-domain-containing adaptor protein and plays crucial role in TCR-mediated signalling by linking LAT with SLP-76 adapter, thereby coupling membrane-proximal events to downstream signaling pathways. GADS-null mice revealed impaired T-cell development, with specific defects in both positive and negative selection of thymocytes [19]. GADS also associates with the serine/threonine kinase hematopoietic progenitor kinase-1 (HPK1) which has been implicated in the activation of the JNK pathway ^[20]. Adapters such as ADAP and SKAP1 with critical roles in the regulation of T-cell adhesion, viral transmission, NF-kB regulation is described in detail elsewhere [10,23,24,25].

Negative regulators of T-cell activation include PHOSPHOPROTEIN ASSOCIATED WITH GEMS (PAG), a transmembrane adapter protein that binds Csk (c-terminal Src kinase) which phosphorylates and inactivates src kinases (e.g. Lck) in resting cells ^[21]. Other adapters that negatively regulate T-cell function is SH2-INTERACTING TRANSMEMBRANE ADAPTOR PROTEIN (SIT) by binding to Csk and protein tyrosine phosphatase 2 (SHP-2) enzymes ^[22].

Role of adapters in NF-kB activation

Generation of knock-out mice deficient for individual signalling proteins, together with biochemical functional studies, have identified several adapter proteins required for the co-stimulation induced NF-kB activation^[26]. These initial reports also showed that co-stimulation induced NF-κB activation was dependent on the initial tyrosine

phosphorylation cascade that engages adapters multiprotein complexes containing several proteins required for NF-κB activation such as Src homology domain-containing leukocyte phosphoprotein 76 (SLP-76), growth factor receptor bound protein-2 (GRB-2), GRB-2 related adapter downstream of Shc (GADS), adhesion and degranulation promoting adapter protein (ADAP) and proteins with enzymatic activity as well as adapter-like functions, such as phospholipase C (PLC) γ and the exchange factor Vav1 [27]. While Vav1 contributes to Rac-dependent reorganization of the actin cytoskeleton, activated PLCy1 activates protein kinase C (PKC) to generate diacylglycerol (DAG) and IP3. Adapter SLP-76 has no intrinsic enzymatic activity but its expression is required for the activation of NF-kB as shown in Jurkat cells J14 deficient in SLP-76 [28]. A role for SLP-76 in vivo has been less certain given that there are fewer mature T-cells in the periphery due to a severe block in thymic development at double negative (DN) stage [14,15]. Disruption in the thymic T-cell development as a consequence of defective NF-kB could not be ruled out. GRB-2 on the other hand has been demonstrated to directly associate with CD28 cytoplasmic tail [29,30], and its role in the activation of NF-kB pathway has been examined previously in Jurkat cells [31], showing its expression is required for the CD28 linked NF-kB pathway. In our recent study, we demonstrated that GRB-2 is an essential component of CD28 pathway and is vital for achieving full NF-kB activation while intriguingly, it did not participate directly in the TCR driven pathway ^[23]. Another study claimed a role for GADS in this pathway ^[32], however, we have consistently observed stronger association of GRB-2 but not GADS with CD28 [23]. The point of CD28/TCR convergence of NF-kB activation remains unknown.

CD28 and NF-kB activation

Signals from co-receptors, particularly CD28 appear to cooperate with primary TCR signaling for optimal T-cell activation [33]. NF-kB pathway is no exception, and CD28 is a crucial component to achieve full immune response as demonstrated by CD28 KO cells that have diminished IL-2 production and activation. However, despite the importance of CD28 in potentiating TCR activation of T-cells (i.e. co-stimulation), increasing evidence has shown that its ligation alone can induce signaling events in T-cells [34,35]. This is further supported by studies on the use of mitogenic CD28 antibody which can induce proliferation and cytokine burst in the absence of TCR ligation [36]. We previously showed that GEF VAV1 binding to CD28 involves the intermediate binding of another adaptor GRB-2 [29,30]. Other investigators have shown that the loss of either GRB-2 or GADS binding to CD28 can abrogate NK-κB activation in Jurkat T-cells [37]. In addition, Tuosto et al. have reported

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that non-mitogenic anti-CD28 can deliver a unique signal leading to the recruitment of p52/Rel-A complexes on Bcl-xL promoter $^{[38]}$. They showed that CD28 can co-operate with VAV-1 to activate NF- κB in a pathway involving Rac-1 and mitogen-activated kinase kinase 1 $^{[35,39]}$.

On the other hand, Ca²⁺-independent PKC subfamily member PKC- θ predominantly expressed in T-cells has important and non-redundant roles in T-cell activation particularly via regulation of NF-kB ^[40,41]. The activity of PKC- θ is essentially regulated by its membrane localization and conformational changes ^[42]. Recently, a conserved proline-rich motif in the V3 domain of PKC- θ was found to be required for association with CD28 and its immunological synapse localization and downstream effector functions ^[43]. Thus, CD28 directly engages proximal pathways leading to the NF-kB activation in T-cells.

New insights into the regulation of NF-kB by CD28 and TCR

Despite the progress made in understanding NF-kB regulation in T-cells, studies dissecting the individual components of TCR and CD28-mediated NF-κB activation in primary T-cells have been lacking. In our recent paper, we addressed this issue by using primary T-cells from various knock-out ($Cd28^{-/-}$, $adap^{-/-}$) and knock-in (i.e. Cd28 Y-170F) mice in conjunction with transfected Jurkat T-cells and showed that the TCR and CD28 use distinct pathways for the activation of the NF-kB pathway in T-cells [23]. CD28 engaged GRB-2 via YMNM motif in its cytoplasmic tail, which was required for NF-kB induction. Using Y170F knock-in mice, NF-kB activation was significantly dampened, so was the case when endogenous GRB-2 was depleted. CD28 induced NF-kB pathway was further delineated by showing Vav1 as an important component of CD28/GRB-2 pathway. Vav1 expression significantly up-regulated NF-kB, and its depletion abrogated NF-kB activation in response to CD28 engagement. Surprisingly, CD28 KO primary T-cells had normal NF-kB response when engaged by anti-TCR antibodies. Further, CD28 activated NF-kB pathway was fully functional in ADAP deficient primary cells but defective TCR pathway. In both cases (ADAP or CD28 deficient primary cells), synergy in NF-kB co-ligation of CD3 and CD28 was lost. The independent nature of CD3 and CD28 pathways in NF-kB activation was also supported by results from LAT deficient cells showing normal CD28 activation, but no activation via TCR/CD3 pathway. Our findings provide evidence that the CD28 and TCR pathways regulate NF-kB activity via different signaling modules of GRB-2/VAV1 and LAT/ADAP respectively.

Concluding remarks

In lymphocytes, NF-kB controls expression of diverse set of genes involved in the productive immune response, division and growth. NF-kB has been topic of intense study for several decades, and recent years have witness, complex interplay of signaling pathways that shape the spatial and timely outcome of NF-kB activation [44]. To successfully interfere NF-kB pathway in clinical settings, receptors as well as co-receptor (stimulatory/inhibitory) engaging NF-kB pathway entail further understanding. In this direction, we have uncovered that CD28 and CD3 use unique signaling modules of GRB-2/Vav1 and LAT/ADAP to achieve full activation of NF-kB in T-cells respectively.

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