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Authors

Dinasarapu, Ashok Reddy
Chandrasekhar, Anjana
Hajishengallis, George
[et al.](#)

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Integrin beta-2

Ashok Reddy Dinasarapu¹, Anjana Chandrasekhar¹, George Hajishengallis², Shankar Subramaniam³

Integrins are heterodimeric transmembrane (TM) glycoproteins containing one each of α and β subunit, which are held together by non-covalent forces. Integrin $\beta 2$ (CD18) is the β subunit for four heterodimers: $\alpha D\beta 2$, $\alpha X\beta 2$, $\alpha M\beta 2$ and $\alpha L\beta 2$. Integrin $\beta 2$ family plays an essential role in leukocyte recruitment and activation during inflammation. Structurally, while most part of the $\alpha\beta$ dimer is extracellular, both the subunits traverse the plasma membrane and terminate as short cytoplasmic domains. Each heterodimeric integrin exists on the cell surface mainly in an inactive (bent) form until they receive stimulating signals from other receptors (*via* inside-out signaling), and the end result of integrin activation is a shift in integrin conformation from a bent to an extended one. The binding of cytoplasmic proteins to α - and/or β -subunit carboxy-terminal tails is an essential part of the activation process, as these interactions stabilize the extended integrin conformation and provide connections to the cytoskeleton. The binding of extracellular ligand to the extended form of integrin (*via* outside-in signaling) triggers a large variety of signal transduction events that modulate cell behaviors such as adhesion, proliferation, survival or apoptosis, shape, polarity, motility, and differentiation, mostly through effects on the cytoskeleton. The receptors $\alpha M\beta 2$ (Complement Receptor type 3, CR3) and $\alpha X\beta 2$ (Complement Receptor type 4, CR4) are regarded to be the most important mediators for complement-driven phagocytosis.

KEYWORDS

CD18; Cell surface adhesion glycoprotein LFA-1/CR3/P150,959 beta subunit precursor); Cell surface adhesion glycoproteins LFA-1/CR3/p150,95 subunit beta; Complement receptor C3 beta-subunit; Complement receptor C3 subunit beta; Integrin beta chain, beta 2; Integrin beta-2; Integrin, beta 2 (complement component 3 receptor 3 and 4 subunit); ITGB2; LAD; LCAMB; Leukocyte cell adhesion molecule CD18; Leukocyte-associated antigens CD18/11A, CD18/11B, CD18/11C; LFA-1; MAC-1; MF17; MFI7

IDENTIFIERS

Molecule Page ID:A004263, Species:Human, NCBI Gene ID: 3689, Protein Accession:NP_001120963.1, Gene Symbol:ITGB2

PROTEIN FUNCTION

Inflammation which occurs due to infection or tissue injury, controls a cascade of cellular and microvascular reactions that allow the removal of pathogens or cell debris, and finally give rise to wound healing, repair and homeostasis. The process of the inflammation includes recruitment (migration) of free-flowing immune cells such as polymorphonuclear neutrophils (PMN) and monocytes/macrophages to the site of infection (Simon and Green 2005; Nourshargh *et al.* 2005). The essential steps during leukocyte recruitment includes tethering and rolling, activation, firm adhesion, intraluminal crawling, and extravasation. Firm adhesion and crawling are largely mediated by $\beta 2$ -integrins (Kolaczowska and Kubes 2013, Hajishengallis and Chavakis 2013). Signaling *via* adhesion molecules of the $\beta 2$ integrin family plays an essential role for immune cell recruitment and activation during inflammation. An important function of these recruited leukocytes is the phagocytosis of complement opsonized particles mediated by integrin $\alpha M\beta 2$ (Anderson and Springer 1987). Therefore, integrin $\beta 2$ -mediated leukocyte migration contributes crucially to the performance of the immune defense system.

Integrin structure: Integrins are noncovalently associated $\alpha\beta$ heterodimeric cell surface glycoproteins. The known 18 α and

8 β subunits in humans generate 24 different heterodimeric receptors, each of which exhibits distinct ligand-binding specificities and tissue distribution (Takada *et al.* 2007; Hynes 2002). Both the α and β subunits are type I membrane proteins (single-pass transmembrane (TM) proteins, which have their N-terminus exposed to the extracellular or luminal space), with a large extracellular ligand-binding region (a.k.a. ectodomain) and generally a short cytoplasmic tail that binds multiple cytoskeletal and adaptor/signaling proteins that regulate the affinity of integrin for extracellular ligands (Hynes 2002; Suzuki and Naitoh 1990; Anthis and Campbell 2011). The integrin heterodimers adopt a shape that resembles a large “head” on two “legs,” with the head containing the sites for ligand binding and subunit association (Campbell and Humphries 2011). The extracellular region of the α subunit is composed of a β -propeller fold with seven blades (W1-W7), a Thigh domain, and two Calf domains (Calf-1 and Calf-2) (Xiong *et al.* 2001; Zhu *et al.* 2009; Xie *et al.* 2010). Further, an I-(inserted or αA) domain (in β -propeller) is present in nine of the α subunits (Lee *et al.* 1995). The extracellular region of the β subunit is composed of a plexin-semaphorin-integrin (PSI) domain, an I-like (or βA) domain, a hybrid domain, four integrin-epidermal growth factor (I-EGF1 to I-EGF4) folds and a β tail domain (βTD) (Xiong *et al.* 2001; Zhu *et al.* 2009; Xie *et al.* 2010; Tan *et al.* 2001; Shi *et al.* 2007; Shi *et al.* 2005; Zhu *et al.* 2008). The I-like domain is inserted into hybrid domain, which in turn is inserted into PSI domain. The two α -helical TM domains of a resting integrin adopt a ridge-in-groove packing (Zhu *et al.* 2009; Lau *et al.* 2009) and the association of the TM domains is specific (Vararattanavech *et al.* 2008). The cytoplasmic tails of the β subunits (other than $\beta 4$ and $\beta 8$) contain one or two highly conserved NxxY/F motifs (x represents other amino acids) that can recognize a wide variety of signaling and cytoskeletal proteins (e.g. adaptor molecules such as ILK, DAB1, Dok-1 and FHL2) that connects integrins to the actin cytoskeleton or activate a range of signaling pathways. In contrast, apart from having a highly conserved juxtamembrane GFFKR motif, α cytoplasmic tails are divergent in their lengths and sequences.

¹Department of Bioengineering, University of California, San Diego, CA 92093, US. ²School of Dental Medicine, University of Pennsylvania, PA 19104, US. ³Department of Bioengineering, University of California at San Diego, CA 92093, US.

Correspondence should be addressed to Ashok Reddy Dinasarapu: adinasarapu@ucsd.edu

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Integrin $\beta 2$ family genes and selectivity: The human CD18 gene, a.k.a ITGB2, is located on chromosome 21q22.3 and encodes a 95-kDa glycoprotein, Integrin $\beta 2$ (Kishimoto *et al.* 1987). The human CD11 genes such as ITGAL, ITGAM, ITGAX and ITGAD are located on chromosome 16p11.2 and encode glycoproteins αL (CD11a, 180kDa), αM (CD11b, 160kDa), αX (CD11c, 150kDa) and αD (CD11d, 145 kDa), respectively (Tan 2012; Fu *et al.* 2012; Luo *et al.* 2007). Integrin $\beta 2$, exclusively expressed on leukocytes, forms heterodimers with the above four α subunits and these heterodimers are signal transducer receptors involved in phagocytosis, degranulation and cell adhesions. Even though $\beta 2$ integrin is common to all these heterodimers, differences in divergent α tails confer structural variations between these integrins. For example, $\alpha L\beta 2$ and $\alpha M\beta 2$ integrins show distinct chemokine-induced activation kinetics (Weber *et al.* 1999), sites for the docking of specific cytosolic molecules such as selective recruitment of the Src kinase Hck to $\alpha M\beta 2$ but not $\alpha L\beta 2$ (Tang *et al.* 2006), and specific association of CD45 cytoplasmic domain with αL (Geng *et al.* 2005). See 'Interactions with Ligands and Other Proteins' section for further details.

Leukocyte migration/adhesion: The movement of leukocytes from the bloodstream to the tissue occurs in several distinct steps as explained above. The $\beta 2$ integrin family of adhesion molecules plays a central role in firm adhesion and subsequent crawling on the endothelium, during which leukocytes seek an appropriate site for diapedesis through endothelial junctions (Grönholm *et al.* 2011; Gahmberg *et al.* 1999). See 'Interactions with Ligands and Other Proteins' section for further details.

Phagocytosis: Phagocytosis is a physiological process by which specialized cells (e.g. macrophages) recognize, bind and internalize materials such as cell debris, microbes, necrotic/apoptotic cells through the use of phagocytic receptors such as Fc γ receptors (utilizes membrane pseudopods), scavenger receptors (mediates binding to modified lipoprotein particles) or integrins (utilizes membrane ruffle mechanism). Integrin activation through bidirectional (inside-out and outside-in) signaling leads to the interaction between particle and integrin which results in an actin-driven uptake of the particle. Activated integrins link actin dynamics to extracellular components that involves cytoskeletal remodeling and cell-shape changes during phagocytosis. However, integrin signaling is also exploited by a variety of pathogens for entry into host cells (Dupuy and Caron 2008). See 'Interactions with Ligands and Other Proteins' section for further details.

REGULATION OF ACTIVITY

Integrins lack enzymatic (intrinsic) activity and the interactions between the membrane proximal regions of α and β are crucial for maintaining integrins in resting state (Chua *et al.* 2011). Integrins use classical bidirectional (a.k.a inside-out and outside-in) signaling and non-classical signaling processes (integrin clustering and membrane ruffling) to integrate the intracellular and extracellular environments (Lim and Hotchin 2012). Inside-out signaling refers to intracellular signaling events that result in a higher-affinity state of the ectodomain of integrin for its cognate ligands. Regulatory events that mediate inside-out signaling converge on the cytoplasmic tails of the α and β chains, which transduce signals to their ectodomains (Dustin *et al.* 2004).

Intracellular signaling pathways, which regulate the

interactions of integrins with their ligands, affect a wide variety of biological functions. Integrin activation is usually initiated by integrin β subunit cytoplasmic tail (Calderwood *et al.* 1999) through the recruitment of cytosolic proteins and many of these interactions are modulated by tail phosphorylation (Gahmberg *et al.* 2009; Fagerholm *et al.* 2004; Liu *et al.* 2000). Signaling molecules implicated in inside-out signaling through $\alpha L\beta 2$ include talin, Vav1, PKD1, several adaptor proteins (SLP-76, ADAP, and SKAP-55), the Ras family GTPase Rap1, and two of its effectors, RAPL and RIAM (Ménasché *et al.* 2007). Apart from talin, kindlin-3 was shown to bind to, and activate Integrin $\beta 2$ and that a direct interaction of kindlin with the β subunit cytoplasmic tail is required, but not sufficient, for integrin activation (Moser *et al.* 2009). Integrin-linked kinase (ILK) interacts with the cytoplasmic domains of integrin $\beta 2$ (also $\beta 1$ and $\beta 3$) (Hannigan *et al.* 1997; Hannigan *et al.* 1996; Delcomenne *et al.* 1998) which acts as a proximal receptor kinase that regulates integrin-mediated signal transduction. Spleen tyrosine kinase (Syk) is constitutively associated with the cytoplasmic tail of $\beta 2$ integrin (Willeke *et al.* 2003; Woodside *et al.* 2002). Syk is known to be phosphorylated and activated upon $\beta 2$ integrin mediated adhesion (Mócsai *et al.* 2002; Willeke *et al.* 2003). Syk and Zap-70 (Zeta-chain-associated protein kinase) are non-receptor cytoplasmic tyrosine kinases with two Src homology (SH) $_2$ - domains, a kinase domain and two interdomains (A and B). Syk and Zap-70 transmit signals from the immune receptors (B-Cell receptor and T-Cell receptor), CD74, Fc Receptor and integrins (Mócsai *et al.* 2002; Turner *et al.* 2000). The inside-out activation leads to an increase in the binding affinity of integrin ectodomains for their extracellular ligands (known as 'outside-in' activation) (Calderwood *et al.* 1999; Tadokoro *et al.* 2003; Li *et al.* 2007; Wegener *et al.* 2007; Lim *et al.* 2007; García-Alvarez *et al.* 2003; Calderwood 2004). Outside-in signaling is analogous to signaling by conventional receptors and is defined as stimulation of intracellular signaling pathways as a consequence of ligation of $\alpha L\beta 2$ with any of its extracellular ligands, such as intracellular adhesion molecule 1 (ICAM-1). Guanine nucleotide exchange factors Cytohesin-1 and Cytohesin-3, activated by PI(3,4,5)P $_3$, bind $\beta 2$ integrin which leads to an increase cell adhesion through an affinity-independent processes, such as integrin clustering, rather than integrin activation (Calderwood 2004). Cytohesin-1 interacts with the cytoplasmic domains of the integrin β -chain common to all $\beta 2$ integrins such as $\alpha L\beta 2$ and $\alpha M\beta 2$ and regulates cell adhesion (Geiger *et al.* 2000; Hyduk and Cybulsky 2011; El Azreq *et al.* 2011).

$\alpha M\beta 2$ also mediates events (classified as non-classical) such as integrin clustering and membrane ruffling in a ligand independent fashion in macrophages following treatment with phorbol 12-myristate 13-acetate (PMA) or lipopolysaccharide (LPS) (Patel and Harrison 2008; Williams and Ridley 2000). The bacterial endotoxin LPS is a potent stimulator of monocyte/macrophage activation and induces adhesion of monocytes while PMA is used in monocyte differentiation. The integrin clustering (in phagocytic function) occurs through the cytoplasmic tails which is different from the extracellular clustering (promotes differentiation to macrophage) of monocyte integrins. The association of Rack1 to integrin $\beta 2$ (coimmunoprecipitated with $\alpha L\beta 2$) in vivo (Liliental and Chang 1998) requires a treatment with PMA which promotes cell spreading and adhesion. These findings suggest that Rack1 may link protein kinase C directly to integrin $\beta 2$ and participate in the regulation of integrin functions (Liliental and Chang 1998).

Bacteria derived fMLP (N-formyl-Met-Leu-Phe, also known as fMLF) induces chemotactic migration and it activates α M β 2 or α L β 2 in human neutrophils through vasodilator stimulated protein (VASP) (Deevi *et al.* 2010) and cytohesin-1 (El Azreq *et al.* 2011). Both VASP and cytohesin-1 function as 'negative regulators' of inside-out function of α M β 2 (El Azreq *et al.* 2011). fMLP, activates Rap1 and inside-out signaling of β 2 integrins (Deevi *et al.* 2010), triggers phosphorylation of VASP on S239 and, thereby, controls membrane recruitment of C3G (a guanine nucleotide exchange factor for Rap1), which is required for activation of Rap1 and antibacterial (β 2 integrin-dependent) functions of neutrophils.

INTERACTIONS

Integrins are heterodimeric ($\alpha\beta$) type I membrane receptors which have their N-terminus exposed to the extracellular space with a large extracellular ligand-binding region and a short cytoplasmic tail that binds multiple cytoskeletal adaptor or signaling proteins that regulate the affinity of integrin for extracellular ligands. The adhesion of integrins to the extracellular matrix is regulated by binding of the cytoskeletal protein talin to the cytoplasmic tail of the β -integrin subunit. Activation is initiated by tail separation and propagation of conformational changes to the outside of the cell. Rap1, a small GTPase, controls activation of integrin (α M β 2) in a talin-dependent manner (Lim *et al.* 2010). Ligand interaction with activated β 2 integrins takes place *via* an inserted I-domain in the α subunit (Shimaoka *et al.* 2003; Shimaoka *et al.* 2005). Integrin α L I domains interact with β 2 in the following orders of affinity: ICAM-1 > ICAM-2 > ICAM-3 (Guermontprez *et al.* 2001). Leukocyte integrins α L β 2, α M β 2 and α X β 2 act as collagen receptors and the α domains favor different collagen subtypes and also differ in their requirements for activation (Lahti *et al.* 2013).

α L β 2 (CD11a/CD18; Leucocyte Function-associated Antigen-1, LFA-1): α L β 2, a leukocyte-restricted integrin, is essential for the adhesion, migration, proliferation of leukocytes, immune synapse formation, and NK cell cytotoxicity (Kinashi 2007; Smith *et al.* 2007; Bryceson *et al.* 2006; Bhunia *et al.* 2009). Ectopic expression of talin head domain induces α L β 2 activation possibly via association of talin head domain with the membrane proximal NPXF motif in the β 2 tail (Li *et al.* 2007; Kim *et al.* 2003). Another actin-binding protein -actinin binds to the membrane proximal sequence of the β 2 tail of α L β 2 (Pavalko and LaRoche 1993; Stanley *et al.* 2008). Interestingly, the binding of filamin to triplet Thr motif of the β 2 tail has an inhibitory effect on α L β 2-mediated T cell adhesion (Takala *et al.* 2008) (Bhunia *et al.* 2009). RAPL (regulator of adhesion and cell polarization enriched in lymphoid tissues) associates with Rap1-GTP, and the activating effect of this complex on α L β 2 requires the membrane proximal Lys1097 and Lys1099 in the α L tail (Tohyama *et al.* 2003). Collectively, a multifaceted (positive and negative) regulatory network of molecules at the cytoplasmic face of the α L β 2 allows fine-tuning of α L β 2 activity in cells under different contexts such as physiological conditions, and in different regions of a polarized and migrating cell (Chua *et al.* 2013). Studies in mice have led to identification of developmental endothelial locus-1 (Del-1) as an endogenous antagonist of LFA-1 (Choi *et al.* 2008) and it inhibits transmigration to inflamed tissues (Eskan *et al.* 2012).

Integrin α L β 2 interacts with ICAM1-4. ICAM-1 is an inducible molecule that is up-regulated by inflammatory cytokines on endothelium, leukocytes, and multiple other cell

types, whereas ICAM-3 is constitutively expressed on leukocytes and absent from endothelium and most other cell types under normal conditions (Springer 1990; Fawcett *et al.* 1992). ICAM-1/ α L β 2 interaction is essential for T-cell activation as well as for migration of T-cells to target tissues (Anderson and Siahaan 2003). CD47, also called Integrin Associated Protein (IAP), has been demonstrated to associate with β 2 integrins. The interaction between Jurkat T-cell β 2 integrins and CD47 were detected by fluorescence lifetime imaging microscopy (Azcutia *et al.* 2013) and that CD47 is necessary for induction of α L β 2 high affinity conformations that bind to their ligand ICAM-1. ICAM-1, as a member of super-IgG family, consists of five IgG-like domains (D1–D5) and binds to α M β 2 via D3 domain (Diamond *et al.* 1991) and α L β 2 via D1 domain (Staunton *et al.* 1991), respectively. Cytohesin-1 interacts with the intracellular portion of the integrin β 2 chain (Kolanus *et al.* 1996). Colocalization of CD82 antigen or Cytohesin-1 with α L β 2 at an adhesion foci results in enhanced interaction between α L β 2 and ICAM-1 during T cell-T cell and T cell-APC interactions (Shibagaki *et al.* 1999; Kolanus *et al.* 1996). Except α L β 2, all other β 2 Integrins binds to Fibrin.

α M β 2 (CD11b/CD18; Complement Receptor type 3, CR3; Macrophage-1 antigen, Mac-1; the iC3b receptor): α M β 2, a leukocyte restricted integrin, mediates leukocyte migration, adhesion, phagocytosis, degranulation and the maintenance of immune tolerance. The receptor α M β 2 is regarded to be the most important mediator for complement-driven phagocytosis. Signaling via α M β 2 predominantly occur, in polymorphonuclear neutrophils (PMN), upon ligand binding and may have a unique role in neutrophil migration (Walzog *et al.* 1996; Yan *et al.* 1997; Ross and Lambris 1982). Integrin α M β 2 binds ligands such as intercellular adhesion molecule -1 (ICAM-1) on inflamed endothelial cells, the complement C3 (fragments such as iC3b), fibrinogen and fibrin, collagens and coagulation factor X (Plow *et al.* 2000; Walzog *et al.* 1995). Being expressed on phagocytes, it interacts with iC3b opsonized pathogen (Bajic *et al.* 2013). Complement C3 deposition on the bacterial surface and α M β 2 on the macrophage surface play important roles in the uptake of the highly virulent *Francisella tularensis* subsp. *Tularensis*, an infectious facultative intracellular pathogen (Schulert and Allen 2006; Clemens *et al.* 2005; Clay *et al.* 2008). Complement receptors, particularly α M β 2, have long been postulated to allow for safe passage for intracellular pathogens (Wright and Silverstein 1983). There is an increasing evidence for signaling crosstalk between complement receptors and TLRs (Hajishengallis and Lambris 2010; Hajishengallis and Lambris 2011; Ivashkiv 2009). For example, TLR2 is able to trans-activate α M β 2 through inside-out signaling including the activation of Rac1, PI3K and cytohesin-1 (Harokopakis *et al.* 2005; Sendide *et al.* 2005). Integrin β 2 signaling can also negatively regulate TLR responses (Ivashkiv 2009; Wang *et al.* 2010). Specifically, α M β 2 can inhibit TLR4 signaling by promoting the degradation of MyD88 and TRIF (Han *et al.* 2010).

Integrin α M β 2's function is dependent on the activation of outside-in and inside-out two way signals (Abram and Lowell 2009). Signaling via α M β 2 plays an important role in regulating production of interleukin-12 (IL-12), a key mediator of cell-mediated immunity (Marth and Kelsall 1997). In addition, engagement of α M β 2 has been shown to down-regulate IL-12 production (Marth and Kelsall 1997) and avoid initiation of the oxidative burst in macrophages following phagocytosis of apoptotic cells (Kim *et al.* 2005). Known key players during

inside-out activation of $\alpha\text{M}\beta 2$ include Rap1, talin1 and CamKII. CamKII phosphorylation of S756, allows Rap1 and talin to be recruited to $\beta 2$ and consequently activate $\alpha\text{M}\beta 2$ (Lim *et al.* 2011). Ceramide, a constituent of atherogenic lipoproteins, binds with CD14 (membrane anchored) and induces clustering of CD14 with co-receptors in lipid rafts (Ceramide recruits $\alpha\text{M}\beta 2$ and CD36 to the proximity of CD14). CD14 lacks a transmembrane signaling domain and signals through TLR4 or TLR2 and plays a major role in the inflammatory response of monocytes to LPS (Pfeiffer *et al.* 2001).

Integrin $\alpha\text{M}\beta 2$ is a known ligand of RAGE (Advanced glycosylation end product-specific receptor) protein. RAGE and $\alpha\text{M}\beta 2$ have been shown by Ma *et al.* 2012 to interact with C1q, both individually ($\alpha\text{M}\beta 2$ /RAGE and RAGE/C1q complexes) and together as a complex ($\alpha\text{M}\beta 2$ /RAGE/C1q complex) (Ma *et al.* 2012). The outcome of C1q interaction with these proteins, is enhanced phagocytosis. The tri-complex of $\alpha\text{M}\beta 2$ /RAGE/C1q shows more efficient phagocytosis than C1q/RAGE or RAGE/ $\alpha\text{M}\beta 2$. RIAM (Rap1-interacting adaptor molecule), in contrast to the previous study (Lim *et al.* 2010), regulates the recruitment of talin (via Rap1) to $\alpha\text{M}\beta 2$ in complement-mediated phagocytosis in human myeloid cell lines (HL-60 and THP-1) and macrophages derived from primary monocytes (Lee *et al.* 2009; Medraño-Fernandez *et al.* 2013).

Integrin $\alpha\text{M}\beta 2$ interacts with fimbriae of *Porphyromonas gingivalis* (*P. gingivalis*) (Hajishengallis *et al.* 2007). *P. gingivalis* (Harokopakis *et al.* 2005) and *Mycobacterium bovis* BCG (Sendide *et al.* 2005) can activate $\alpha\text{M}\beta 2$ through inside-out signaling via TLR2 to facilitate bacterial uptake. CyaA (*Bordetella pertussis*) uses the $\alpha\text{M}\beta 2$ as a cell receptor and CyaA intoxication leads to increased intracellular cAMP level and cell death (Guermontprez *et al.* 2001). RrgA on pneumococcal pilus 1 promotes nonopsonic $\alpha\text{M}\beta 2$ -dependent uptake of *S. pneumoniae* by murine and human macrophages. RrgA- $\alpha\text{M}\beta 2$ -mediated phagocytosis promotes systemic pneumococcal spread from local sites (Orrskog *et al.* 2012). Complement iC3b covalently bound to the gonococcus serves as a primary ligand for $\alpha\text{M}\beta 2$ adherence. However, gonococcal porin and pili also bound to the I-domain of $\alpha\text{M}\beta 2$ in a non-opsonic manner. $\alpha\text{M}\beta 2$ -mediated endocytosis serves as a primary mechanism by which *N. gonorrhoeae* elicits membrane ruffling and cellular invasion of primary, human, cervical epithelial cells and this data suggest that gonococcal adherence to $\alpha\text{M}\beta 2$ occurs in a co-operative manner, which requires gonococcal iC3b-opsonization, porin and pilus (Edwards *et al.* 2002; Jones *et al.* 2008). CD14 cooperates with $\alpha\text{M}\beta 2$ to mediate phagocytosis of *Borrelia burgdorferi*. Complement enhances phagocytosis of *B. burgdorferi* in a C3-dependent manner (Hawley *et al.* 2013). αM interacts with leukocidin A/B (LukAB), which is produced by *S. aureus* upon encountering neutrophils and is both necessary and sufficient for *S. aureus* to kill human neutrophils, macrophages and dendritic cells (DuMont *et al.* 2011, DuMont *et al.* 2013a). The α subunit of the $\alpha\text{M}\beta 2$ integrin acts as a cellular receptor for LukAB (DuMont *et al.* 2013b).

$\alpha\text{X}\beta 2$ (CD11c/CD18; p150,95; Complement Receptor type 4, CR4): Integrin $\alpha\text{X}\beta 2$ is a receptor for iC3b, C3dg, and C3d fragments of complement C3 (Myones *et al.* 1988, Vik and Fearon 1985; Chen *et al.* 2012; Micklem and Sim 1985) and was shown to bind with apparently equal affinity (Vik and

Fearon 1985). Integrin $\alpha\text{X}\beta 2$ also shares some functional properties with $\alpha\text{M}\beta 2$ as an adhesion surface molecule. Both $\alpha\text{M}\beta 2$ (Wright and Jong 1986) and $\alpha\text{X}\beta 2$ bind bacterial LPS and β -glucans and promote phagocytosis of unopsonized bacteria and yeast. A large number of intracellular proteins have been found to interact with the cytosolic tails (CTs) of this integrin linking $\alpha\text{X}\beta 2$ to the cytoskeleton (Chua *et al.* 2012).

$\alpha\text{D}\beta 2$ (CD11d/CD18): Integrin $\alpha\text{D}\beta 2$ is a multiligand macrophage receptor with recognition specificity identical to that of the major myeloid cell-specific integrin $\alpha\text{M}\beta 2$. Integrin $\alpha\text{D}\beta 2$ is capable of supporting cell adhesion to various extra cellular matrix (ECM) proteins, including fibronectin, vitronectin, fibrinogen, CCN1 (Cyr61) and others. $\alpha\text{D}\beta 2$, selectively binds ICAM-3 and VCAM-1 and does not appear to bind ICAM-1 (Van der Vieren *et al.* 1995; Van der Vieren *et al.* 1999; Grayson *et al.* 1998). The αD I-domain is responsible for the binding function and that the mechanism whereby αD I-domain recognizes its ligands is similar to that utilized by $\alpha\text{M}\beta 2$.

CMAP, a complement database, documents the biochemical methods used to identify these interactions (Yang *et al.* 2013).

PHENOTYPES

Leukocyte emigration, from the bloodstream to tissue to sites of inflammation, is a dynamic process and involve multiple steps in an adhesion cascade. Various adhesion molecules are expressed on both resting and stimulated endothelial cells and leukocytes (Nagendran *et al.* 2012; Muller 2003). Leukocyte adhesion and tethering defects involve $\beta 2$ integrins and selectin ligands (Bunting *et al.* 2002). Selectins are found on both leukocytes and endothelial cells and primarily mediate cellular margination and rolling. Defects in a number of these adhesion molecules result in recognized clinical syndromes called Leukocyte Adhesion Deficiency (LAD) syndrome in which leukocytes (particularly neutrophils) cannot leave the vasculature to migrate normally into tissues under conditions of inflammation or infection. Affected individuals display blood neutrophilia, suffer from recurrent infections, and invariably develop aggressive periodontitis leading to premature loss of primary and permanent teeth (Bowen *et al.* 1982; Anderson and Springer 1987; Arnaout 1990; Shaw *et al.* 2001; Wright *et al.* 1995; Etzioni 1999).

LAD I, in which the $\beta 2$ -integrin family is deficient or defective.

LAD II, in which the fucosylated carbohydrate ligands for selectins are absent.

LAD III, in which the activation of β integrins ($\beta 1$, $\beta 2$, and $\beta 3$) are defective (Karaköse *et al.* 2010; Plow *et al.* 2009; Jurk *et al.* 2010). LAD III is mainly due to mutations in fermitin family member 3 (FERMT3, aka KIND3). All LAD III patients have premature stop codons or nonsense mutations in both alleles of their FERMT3 gene (Malinin *et al.* 2009; Manevich-Mendelson *et al.* 2009; Kuijpers *et al.* 2009; Svensson *et al.* 2009; Kuijpers *et al.* 1997). Kindlin-3 is a cytoplasmic protein that acts cooperatively with talin-1 in activating $\beta 1$, $\beta 2$, and $\beta 3$ integrins. LAD III is characterized by bleeding disorders and defective recruitment of leukocytes into sites of infection.

MAJOR SITES OF EXPRESSION

$\alpha\text{L}\beta 2$ (CD11a/CD18, LFA-1): Integrin $\alpha\text{L}\beta 2$ is the only integrin expressed on all leukocyte lineages.

$\alpha\text{M}\beta 2$ (CD11b/CD18, CR3; Mac-1): Expressed on

polymorphonuclear leukocytes (mainly, neutrophils), mononuclear phagocytes (dendritic cells, monocytes and macrophages), lymphocytes (mainly, natural killer (NK) and $\gamma\delta$ T-cells) and microglia.

$\alpha X\beta 2$ (CD11c/CD18, CR4): Expressed on mononuclear phagocytes (dendritic cells, monocytes and macrophages), polymorphonuclear leukocytes (mainly, neutrophils), activated B lymphocytes and natural killer (NK) cells.

$\alpha D\beta 2$ (CD11d/CD18): Expressed on macrophages and eosinophils.

SPLICE VARIANTS

Integrin $\beta 2$ is a 95-kDa glycoprotein, encoded by the ITGB2 gene and is located on chromosome 21q22.3 (Kishimoto *et al.* 1987). Human ITGB2 spans approximately 40 kb of DNA and contains 16 exons (Weitzman *et al.* 1991). Two transcript variants encoding the same protein have been identified.

REGULATION OF CONCENTRATION

The expression of the leukocyte integrins is cell-specific and is coordinately regulated during leukocyte differentiation through transcriptional and post-transcriptional mechanisms (Miller *et al.* 1986; Noti *et al.* 2001; Noti and Reinemann 1995; Back *et al.* 1992). The promoters for the CD11a-d (Pahl *et al.* 1992; Nueda *et al.* 1993; López-Rodríguez *et al.* 1995; Cornwell *et al.* 1993; Noti *et al.* 1992; López-Cabrera *et al.* 1993; Agura *et al.* 1992; Hickstein *et al.* 1992) and CD18 (Rosmarin *et al.* 1992; Agura *et al.* 1992) genes lack classical TATA boxes but instead appear to be controlled by initiator elements positioned within 100 bp of their ATG translational start codons. Cis elements are found within 500 bp upstream of the ATG site, some of which control cell-specific expression.

ANTIBODIES

Monoclonal antibodies (mAbs) directed against the CD18 ($\beta 2$): blocking IB4 (Bednar *et al.* 1996) and an activating KIM-127. KIM127 is a widely used mAb that recognizes a $\beta 2$ subunit epitope (on epidermal growth factor (EGF)-like domain 2) that is cryptic on bent $\alpha L\beta 2$, but exposed when the integrin extends (Beglova *et al.* 2002; Kamata *et al.* 2002; Chen *et al.* 2010). Efalizumab is a monoclonal antibody, which is specific for $\alpha L\beta 2$ to treat psoriasis. Anti-integrin $\beta 2$ mAb MEM-48 is available from Sigma.

Table 1: Functional States

STATE DESCRIPTION	LOCATION	REFERENCES
$\beta 2$ (CD18)	plasma membrane	
$\beta 2$ /DAB1	integrin complex	Calderwood DA <i>et al.</i> 2003
$\beta 2$ /FHL2	integrin complex	Wixler V <i>et al.</i> 2000
$\beta 2$ /DOK1	integrin complex	Calderwood DA <i>et al.</i> 2003
$\beta 2$ /PKC	integrin complex	Fagerholm S <i>et al.</i> 2002
$\beta 2$ /ILK	integrin complex	Hannigan GE <i>et al.</i> 1997; Delcommenne M <i>et al.</i> 1998
$\beta 2$ /(Syk Zap-70)	integrin complex	Willeke T <i>et al.</i> 2003; Woodside DG <i>et al.</i> 2002; Miura Y <i>et al.</i> 2000
$\beta 2$ /Hsp40 (<i>F. tularensis</i>)	integrin complex	Dyer MD <i>et al.</i>
$\alpha D\beta 2$ (CD11d/CD18)	alphaD-beta2 integrin complex	Van der Vieren M <i>et al.</i> 1995
$\alpha D\beta 2$ /VCAM-1	alphaD-beta2 integrin complex	Grayson MH <i>et al.</i> 1998; Van der Vieren M <i>et al.</i> 1999
$\alpha D\beta 2$ /ICAM-3	alphaD-beta2 integrin complex	Van der Vieren M <i>et al.</i> 1995
$\alpha D\beta 2$ /Fibrinogen	alphaD-beta2 integrin complex	Yakubenko VP <i>et al.</i> 2006
$\alpha M\beta 2$ (CR3; CD18/11b)	alphaM-beta2 integrin complex	Arnaout MA <i>et al.</i> 1988; Sándor N <i>et al.</i> 2013
$\alpha M\beta 2$ /CD14	alphaM-beta2 integrin complex	Pfeiffer A <i>et al.</i> 2001; Ross GD <i>et al.</i> ; Zarewych DM <i>et al.</i> 1996
$\alpha M\beta 2$ /CD23	alphaM-beta2 integrin complex	Ross GD <i>et al.</i> ; Lecoanet-Henchoz S <i>et al.</i> 1995
$\alpha M\beta 2$ /CD59	alphaM-beta2 integrin complex	Ross GD <i>et al.</i>
$\alpha M\beta 2$ /Collagen	alphaM-beta2 integrin complex	Ross GD <i>et al.</i>
$\alpha M\beta 2$ /Fibrinogen	alphaM-beta2 integrin complex	Diamond MS <i>et al.</i> 1993; Ross GD <i>et al.</i>
$\alpha M\beta 2$ /ELANE	alphaM-beta2 integrin complex	Cai TQ and Wright SD 1996
$\alpha M\beta 2$ /Heparan sulfate	alphaM-beta2 integrin complex	Ross GD <i>et al.</i>
$\alpha M\beta 2$ /fH	alphaM-beta2 integrin complex	DiScipio RG <i>et al.</i> 1998; Ross GD <i>et al.</i>
$\alpha M\beta 2$ /FX	alphaM-beta2 integrin complex	Altieri DC and Edgington TS 1988; Ross GD <i>et al.</i>
$\alpha M\beta 2$ /β-glucan	alphaM-beta2 integrin complex	Ross GD <i>et al.</i>
$\alpha M\beta 2$ /LN-8	alphaM-beta2 integrin complex	Wondimu Z <i>et al.</i> 2004
$\alpha M\beta 2$ /GPIba	alphaM-beta2 integrin complex	Josefsson EC <i>et al.</i> 2005
$\alpha M\beta 2$ /uPAR-GPI	alphaM-beta2 integrin complex	Pliyev BK <i>et al.</i> 2010; Ross GD <i>et al.</i> ; Xue W <i>et al.</i> 1994
$\alpha M\beta 2$ /uPAR-GPI/uPA	alphaM-beta2 integrin complex	
$\alpha M\beta 2$ /FcyRIIa (CR3/CD32)	alphaM-beta2 integrin complex	Annenkov A <i>et al.</i> 1996; Ross GD <i>et al.</i>
$\alpha M\beta 2$ /FcyRIIIB (CR3/CD16)	alphaM-beta2 integrin complex	Ross GD <i>et al.</i> ; Preynat-Seauve O <i>et al.</i> 2004; Sehgal G <i>et al.</i> 1993; Zhou M <i>et al.</i> 1993
$\alpha M\beta 2$ /HP	alphaM-beta2 integrin complex	El-Ghmati SM <i>et al.</i> 2002
$\alpha M\beta 2$ /PR-3	alphaM-beta2 integrin complex	David A <i>et al.</i> 2003
$\alpha M\beta 2$ /iC3b	alphaM-beta2 integrin complex	Gordon DL <i>et al.</i> 1987
$\alpha M\beta 2$ /ICAM[1,2,4]	alphaM-beta2 integrin complex	Diamond MS <i>et al.</i> 1990; Hermand P <i>et al.</i> 2000; Ross GD <i>et al.</i>
$\alpha M\beta 2$ /Talin-1	alphaM-beta2 integrin complex	Lim J <i>et al.</i> 2007
$\alpha M\beta 2$ /Kindlin-3	alphaM-beta2 integrin complex	
$\alpha M\beta 2$ /FUT4 (CR3/CD15)	alphaM-beta2 integrin complex	Skubitz KM and Snook RW 1987
$\alpha M\beta 2$ /RAGE	alphaM-beta2 integrin complex	Ma W <i>et al.</i>
$\alpha M\beta 2$ /RAGE/C1q	alphaM-beta2 integrin complex	Ma W <i>et al.</i>
$\alpha M\beta 2$ /CYR61(CCN1)	alphaM-beta2 integrin complex	Schober JM <i>et al.</i> 2002; Schober JM <i>et al.</i> 2003
$\alpha M\beta 2$ /CCN2	alphaM-beta2 integrin complex	Schober JM <i>et al.</i> 2002
$\alpha M\beta 2$ /MPO	alphaM-beta2 integrin complex	El Kebir D <i>et al.</i> 2008; Johansson MW <i>et al.</i> 1997; Lau D <i>et al.</i> 2005
$\alpha M\beta 2$ /PLG	alphaM-beta2 integrin complex	Lishko VK <i>et al.</i> 2004
$\alpha M\beta 2$ /CyaA (<i>B. pertussis</i>)	alphaM-beta2 integrin complex	Guermonprez P <i>et al.</i> 2001
$\alpha M\beta 2$ /App1 (<i>C. neoformans</i>)	alphaM-beta2 integrin complex	Stano P <i>et al.</i> 2009
$\alpha M\beta 2$ /RrgA (<i>S. pneumoniae</i>)	alphaM-beta2 integrin complex	Orrskog S <i>et al.</i>
$\alpha M\beta 2$ /LPS (<i>E. coli</i>)	alphaM-beta2 integrin complex	Van Strijp JA <i>et al.</i> 1993; Ross GD <i>et al.</i> ; Wright SD and Jong MT 1986
$\alpha X\beta 2$ (CR4, CD11c/18)	alphaX-beta2 integrin complex	Shelley CS <i>et al.</i> 2002; Lecoanet-Henchoz S <i>et al.</i> 1995
$\alpha X\beta 2$ /CD23	alphaX-beta2 integrin complex	Lecoanet-Henchoz S <i>et al.</i> 1995
$\alpha X\beta 2$ /FUT4 (CR4/CD15)	alphaX-beta2 integrin complex	Skubitz KM and Snook RW 1987
$\alpha X\beta 2$ /Fibrinogen	alphaX-beta2 integrin complex	
$\alpha X\beta 2$ /iC3b	alphaX-beta2 integrin complex	Micklem KJ and Sim RB 1985; Chen X <i>et al.</i> 2012
$\alpha X\beta 2$ /ICAM-1	alphaX-beta2 integrin complex	
$\alpha X\beta 2$ /LPS (<i>E. coli</i>)	alphaX-beta2 integrin complex	Ingalls RR and Golenbock DT 1995
$\alpha L\beta 2$ (LFA-1, CD11a/18)	alphaL-beta2 integrin complex	
$\alpha L\beta 2$ /CD45	alphaL-beta2 integrin complex	Geng X <i>et al.</i> 2005
$\alpha L\beta 2$ /CD47	alphaL-beta2 integrin complex	Azcutia V <i>et al.</i> 2013

α L β 2/CD82	alphaL-beta2 integrin complex	Shibagaki N <i>et al.</i> 1999
α L β 2/Cytohesin-1	alphaL-beta2 integrin complex	Kolanus W <i>et al.</i> 1996; Geiger C <i>et al.</i> 2000
α L β 2/ICAM[1-4]	alphaL-beta2 integrin complex	Edwards CP <i>et al.</i> 1998; Hermand P <i>et al.</i> 2000; Huang C and Springer TA 1995; Li N <i>et al.</i> 2013; Mizuno T <i>et al.</i> 1997; Shimaoka M <i>et al.</i> 2001
α L β 2/RanBPM	alphaL-beta2 integrin complex	Denti S <i>et al.</i> 2004
α L β 2/DNAM-1	alphaL-beta2 integrin complex	Shibuya K <i>et al.</i> 1999
α L β 2/JAB1	alphaL-beta2 integrin complex	Bianchi E <i>et al.</i> 2000; Kinoshita SM <i>et al.</i>
α L β 2/FUT4	alphaL-beta2 integrin complex	Skubitz KM and Snook RW 1987
α L β 2/ESM-1	alphaL-beta2 integrin complex	Bécharde D <i>et al.</i> 2001
α L β 2/Rack1	alphaL-beta2 integrin complex	Liliental J and Chang DD 1998
α L β 2/VacA (H. pylori)	alphaL-beta2 integrin complex	Cover TL <i>et al.</i> 2008; Sewald X <i>et al.</i> 2008

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SUPPLEMENTARY

Supplementary information is available online.

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This molecule exists in 66 states , has 66 transitions between these states and has 0 enzyme functions.(Please zoom in the pdf file to view details.)

