

DESCRIPTION

Source	Mouse myeloma cell line, NS0-derived		
	Human JAM-A (Ser28 - Ala242) Accession # Q9Y624	IEGRMD	Human IgG ₁ (Pro100 - Lys330)
	N-terminus		C-terminus

N-terminal Sequence Analysis	Ser28
Structure / Form	Disulfide-linked homodimer
Predicted Molecular Mass	50 kDa (monomer)

SPECIFICATIONS

SDS-PAGE	60-65 kDa, reducing conditions
Activity	Bioassay data are not available.
Endotoxin Level	<0.01 EU per 1 µg of the protein by the LAL method.
Purity	>90%, by SDS-PAGE under reducing conditions and visualized by silver stain.
Formulation	Lyophilized from a 0.2 µm filtered solution in PBS. See Certificate of Analysis for details.

PREPARATION AND STORAGE

Reconstitution	Reconstitute at 100 µg/mL in sterile PBS.
Shipping	The product is shipped at ambient temperature. Upon receipt, store it immediately at the temperature recommended below.
Stability & Storage	<p>Use a manual defrost freezer and avoid repeated freeze-thaw cycles.</p> <ul style="list-style-type: none"> ● 12 months from date of receipt, -20 to -70 °C as supplied. ● 1 month, 2 to 8 °C under sterile conditions after reconstitution. ● 3 months, -20 to -70 °C under sterile conditions after reconstitution.

BACKGROUND

The family of junctional adhesion molecules (JAM), comprising at least three members, are type I transmembrane receptors belonging to the immunoglobulin (Ig) superfamily (1, 2). These proteins are localized in the tight junctions between endothelial or epithelial cells. Some family members are also found on blood leukocytes and platelets. Human JAM-A, also known as platelet adhesion molecule 1 (PAM-1) and platelet F11 receptor (3), is predominantly expressed at intercellular junctions of both epithelial cells and endothelial cells (1 - 4). It is also expressed on circulating blood cells including neutrophils, monocytes, platelets, erythrocytes and lymphocytes (5). Human JAM-A cDNA predicts a 299 amino acid (aa) residue precursor protein with a putative 27 aa signal peptide, a 210 aa extracellular region containing two Ig-like V-subset domains, a 24 aa transmembrane domain and a 38 aa cytoplasmic domain. The human and mouse proteins share approximately 67% aa sequence homology. Human JAM-A also shares approximately 35% and 32% aa sequence homology with human JAM-B and JAM-C, respectively. JAM-A exhibits homophilic interactions to regulate tight junction assembly and modulate paracellular permeability. This homophilic interaction also mediates platelet aggregation and adhesion to endothelial cells and may play a role in thrombosis (3). JAM-A binds heterotypically with the β2 integrin lymphocyte function-associated antigen-1 (LFA-1). This JAM-A-LFA-1 interaction is involved in leukocyte adhesion and transmigration (6). JAM-A has also been shown to bind reovirus attachment protein sigma-1 to permit reovirus infection and signal virus-induced apoptosis (7).

References:

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3. Sobocka, M.B. *et al.* (2000) *Blood* **95**:2600.
4. Martin-Padura, I. *et al.* (1998) *J. Cell Biol.* **142**:117.
5. Williams, L.A. *et al.* (1999) *Mol. Immunol.* **36**:1175.
6. Ostermann, G. *et al.* (2002) *Nature Immunol.* **3**:151.
7. Barton, E.S. *et al.* (2001) *Cell* **104**:441.