

HLA-G Antibody

Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP51989

Product Information

Application WB **Primary Accession** P17693

Reactivity Human, Mouse, Rat

Host Rabbit Clonality Polyclonal Calculated MW 38224

Additional Information

Gene ID 3135

Other Names HLA class I histocompatibility antigen, alpha chain G, HLA G antigen, MHC

class I antigen G, HLA-G, HLA-60, HLAG

Dilution WB~~1:1000

Format 0.01M PBS, pH 7.2, 0.09% (W/V) Sodium azide, Glycerol 50%

Store at -20 °C. Stable for 12 months from date of receipt **Storage**

Protein Information

HLA-G {ECO:0000303|PubMed:1570318, ECO:0000312|HGNC:HGNC:4964} Name

Function [Isoform 1]: Non-classical major histocompatibility class Ib molecule

involved in immune regulatory processes at the maternal-fetal interface (PubMed:<u>19304799</u>, PubMed:<u>23184984</u>, PubMed:<u>29262349</u>). In complex with B2M/beta-2 microglobulin binds a limited repertoire of nonamer self-peptides derived from intracellular proteins including histones and ribosomal proteins (PubMed:7584149, PubMed:8805247). Peptide-bound HLA-G-B2M complex acts as a ligand for inhibitory/activating KIR2DL4, LILRB1 and LILRB2 receptors on uterine immune cells to promote fetal development while maintaining maternal- fetal tolerance (PubMed: 16366734, PubMed: 19304799,

PubMed: <u>20448110</u>, PubMed: <u>23184984</u>, PubMed: <u>27859042</u>,

PubMed: 29262349). Upon interaction with KIR2DL4 and LILRB1 receptors on decidual NK cells, it triggers NK cell senescence-associated secretory phenotype as a molecular switch to promote vascular remodeling and fetal growth in early pregnancy (PubMed: 16366734, PubMed: 19304799, PubMed: 23184984, PubMed: 29262349). Through interaction with KIR2DL4 receptor on decidual macrophages induces pro-inflammatory cytokine

production mainly associated with tissue remodeling (PubMed: 19304799). Through interaction with LILRB2 receptor triggers differentiation of type 1

regulatory T cells and myeloid-derived suppressor cells, both of which actively maintain maternal-fetal tolerance (PubMed:20448110, PubMed:27859042). May play a role in balancing tolerance and antiviral-immunity at maternal-fetal interface by keeping in check the effector functions of NK, CD8+ T cells and B cells (PubMed:10190900, PubMed:11290782, PubMed:24453251). Reprograms B cells toward an immune suppressive phenotype via LILRB1 (PubMed:24453251). May induce immune activation/suppression via intercellular membrane transfer (trogocytosis), likely enabling interaction with KIR2DL4, which resides mostly in endosomes (PubMed:20179272, PubMed:26460007). Through interaction with the inhibitory receptor CD160 on endothelial cells may control angiogenesis in immune privileged sites (PubMed:16809620).

Cellular Location

[Isoform 1]: Cell membrane; Single-pass type I membrane protein. Endoplasmic reticulum membrane. Early endosome membrane [Isoform 2]: Cell membrane; Single-pass type I membrane protein [Isoform 4]: Cell membrane; Single-pass type I membrane protein [Isoform 6]: Secreted Cell projection, filopodium membrane. Note=HLA-G trogocytosis from extravillous trophoblast's filopodia occurs in the majority of decidual NK cells.

Tissue Location

Expressed in adult eye (PubMed:1570318). Expressed in immune cell subsets including monocytes, myeloid and plasmacytoid dendritic cells and regulatory T cells (Tr1)(at protein level) (PubMed:20448110). Secreted by follicular dendritic cell and follicular helper T cells (PubMed:24453251) [Isoform 7]: Expressed in placenta, amniotic membrane, skin, cord blood and peripheral blood mononuclear cells

Background

Involved in the presentation of foreign antigens to the immune system. Plays a role in maternal tolerance of the fetus by mediating protection from the deleterious effects of natural killer cells, cytotoxic T-lymphocytes, macrophages and mononuclear cells.

References

Shukla H.,et al.Nucleic Acids Res. 18:2189-2189(1990). Geraghty D.E.,et al.Proc. Natl. Acad. Sci. U.S.A. 84:9145-9149(1987). Ishitani A.,et al.Submitted (APR-1992) to the EMBL/GenBank/DDBJ databases. Hampe A.,et al.DNA Seq. 10:263-299(1999). Shiina S.,et al.Submitted (SEP-1999) to the EMBL/GenBank/DDBJ databases.

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