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32-13259: HLA-DRB1 Human, Sf9

DRB1, HLA DRB1, HLA-DR1B, HLA-DRB1, MHC class II antigen DRB1 16, DR-16, DR16, Human

Alternative Name: Leucocyte Antigen DRB1, MHC Class II HLA-DR-Beta Cell Surface Glycoprotein, MHC Class II HLA-

DRw10-Beta.

Description

Source: Sf9, Baculovirus cells. Sterile Filtered colorless solution.

Major Histocompatibility Complex Class II DR Beta 1 also known as HLA-DRB1 ia a member of the HLA class II beta chain paralogs. Molecule class II is a heterodimer consisting of an alpha (DRA) and a beta chain (DRB), both anchored in the membrane. HLA-DRB1 takes an essential part in the immune system by presenting peptides derived from extracellular proteins. Class II molecules are expressed in antigen presenting cells (APC: B lymphocytes, dendritic cells, macrophages). Furthermore, the beta chain is approximately 26- 28 kDa. It is encoded by 6 exons. While exon one encodes the leader peptide; exons 2 and 3 encode the two extracellular domains; exon 4 encodes the transmembrane domain; and exon 5 encodes the cytoplasmic tail.

HLA-DRB1 produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain containing 207 amino acids (30-227a.a.) and having a molecular mass of 24.0kDa. (Molecular size on SDS-PAGE will appear at approximately 28-40kDa). HLA-DRB1 is expressed with a 9 amino acid His tag at C-Terminus and purified by proprietary chromatographic techniques.

Product Info

Amount: $1 \mu g / 5 \mu g$

Purification: Greater than 85.0% as determined by SDS-PAGE.

HLA-DRB1 protein solution (0.25mg/ml) contains Phosphate Buffered Saline (pH 7.4) and 30% Content:

glycerol.

Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods Storage condition:

of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA).

Avoid multiple freeze-thaw cycles.

ADPGDTRPRF LWQPKRECHF FNGTERVRFL DRYFYNQEES VRFDSDVGEF RAVTELGRPD Amino Acid:

AEYWNSQKDI LEQARAAVDT YCRHNYGVVE SFTVQRRVQP KVTVYPSKTQ PLQHHNLLVC SVSGFYPGSI

EVRWFLNGQE EKAGMVSTGL IQNGDWTFQT LVMLETVPRS GEVYTCQVEH PSVTSPLTVE

WRARSESAQS KHHHHHH