

## 32-13214: EPHA2 Human, sf9

**Alternative Name :** EPHA2, ARCC2, CTPA, CTPP1, CTRCT6, ECK, EPHA2, sf9, EPH Receptor A2, sf9, Ephrin type-A receptor 2.

### Description

Source: Sf9, Baculovirus cells.

Sterile Filtered clear solution.

EPH Receptor A2 (EPHA2) is a member of the ephrin receptor subfamily of the protein-tyrosine kinase family. EPHA2 is a protein which binds ephrin-A ligands. EPH and EPH-related receptors are associated with mediating developmental events, particularly in the nervous system. Receptors in the EPH subfamily normally have a single kinase domain and an extracellular region containing a Cys-rich domain and 2 fibronectin type III repeats. The ephrin receptors are divided into two groups based on the similarity of their extracellular domain sequences and their affinities for binding ephrin-A and ephrin-B ligands. EPHA2 gene mutations are the cause of certain genetically-related cataract disorders.

EPHA2 Human Recombinant produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain containing 520 amino acids (27-537) and having a molecular mass of 57.3kDa (Molecular size on SDS-PAGE will appear at approximately 50-70kDa). EPHA2 is fused to 6 amino acid His-Tag at C-terminus and purified by proprietary chromatographic techniques.

### Product Info

**Amount :** 2 µg / 10 µg

**Purification :** Greater than 95.0% as determined by analysis by SDS-PAGE.

**Content :** EPHA2 protein solution (0.5mg/ml) containing Phosphate Buffered Saline (pH 7.4) and 10% glycerol.

**Storage condition :** Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Avoid multiple freeze-thaw cycles.

**Amino Acid :** ADPKEVVLLD FAAAGGELGW LTHPYGKGWD LMQNIMNDMP IYMYSVCNVM SGDQDNWLRT  
NWVYRGEAER IFIELKFTVR DCNSFPGGAS SCKETFNLYY AESDLGYGTN FQKRLFTKID TIAPDEITVS  
SDFEARHVKL NVEERSVGPL TRKGFYLAQ DIGACVALLS VRVYYKKCPE LLQGLAHFPE TIAGSDAPSL  
ATVAGTCVDH AVVPPGGEEP RMHCAVDGEW LVPIGQCLCQ AGYEKVEDAC QACSPGFFKF  
EASESPCLEC PEHTLPSPEG ATSCECEEGF FRAPQDPASM PCTRPPSAPH YLTAVGMGAK VELRWTPPQD  
SGGREDIVYS VTCEQCWPES GECGPCEASV RYSEPPHGLT RTSVTVSDLE PHMNYTFTVE ARNGVSGGLVT  
SRSFRTASVS INQTEPPKVR LEGRSTTSLV VSWSIPPPQ SRVWKYEVY RKKGDSNSYN VRRTEGFSVT  
LDDLAPDTTY LVQVQALTQE GQGAGSKVHE FQTLSPGSG NLAVHHHHHH.