Human EPHA2 Recombinant Protein



RPPB3453

Product Information Protein Information

Product SKU: Protein description:

RPPB3453 EPHA2 Human Recombinant produced in HEK cells is a single, glycosylated, polypeptide chain (Ala24-

Glu530) containing a total of 515 amino acids, having a calculated molecular mass of 56.9kDa. The EPHA2

Accession: protein is fused to a 2 aa C-terminal linker and a 6 aa C-terminal His tag.

P29317

Appearance:

Host: Filtered White lyophilized (freeze-dried) powder.

HEK 293.

Synonyms:

EPHA2, EPH Receptor A2, ECK, Tyrosine-Protein Kinase Receptor ECK, EC 2.7.10.1, CTRCT6, ARCC2, CTPP1, CTPA, Epithelial Cell Receptor Protein Tyrosine Kinase, Ephrin Type-A Receptor 2, Soluble EPHA2 Variant 1, Epithelial Cell Kinase, EC 2.7.10, EphA2.

Formulation:

EPHA2 was filtered ($0.4\mu m$) and lyophilized from 0.5mg/ml solution in phosphate buffered saline and 5% (w/v) trehalose.

Purity:

Greater than 95.0% as determined by SDS-PAGE.

Solubility:

It is recommended to add 200µl deionized water to prepare a working stock solution of approximately 0.5 mg/ml and let the lyophilized pellet dissolve completely. EPHA2 is not sterile! Please filter the product by an appropriate sterile filter before using it in the cell culture.

Stability:

Store lyophilized protein at -20°C. Aliquot the product after reconstitution to avoid repeated freezing/thawing cycles. Reconstituted protein can be stored at 4°C for a limited period of time; it does not show any change after two weeks at 4°C.

Amino Acid Sequence:

AQGKEVVLLD FAAAGGELGW LTHPYGKGWD LMQNIMNDMP IYMYSVCNVM SGDQDNWLRT NWVYRGEAER IFIELKFTVR DCNSFPGGAS SCKETFNLYY AESDLDYGTN FQKRLFTKID TIAPDEITVS SDFEARHVKL NVEERSVGPL TRKGFYLAFQ DIGACVALLS VRVYYKKCPE LLQGLAHFPE TIAGSDAPSL ATVAGTCVDH AVVPPGGEEP RMHCAVDGEW LVPIGQCLCQ AGYEKVEDAC QACSPGFFKF EASESPCLEC PEHTLPSPEG ATSCECEEGF FRAPQDPASM PCTRPPSAPH YLTAVGMGAK VELRWTPPQD SGGREDIVYS VTCEQCWPES GECGPCEASV RYSEPPHGLT RTSVTVSDLE PHMNYTFTVE ARNGVSGLVT SRSFRTASVS INQTEPPKVR LEGRSTTSLS VSWSIPPPQQ SRVWKYEVTY RKKGDSNSYN VRRTEGFSVT LDDLAPDTTY LVQVQALTQE GQGAGSKVHE FQTLSPEKLH HHHHH.