

32-4082: Recombinant Human Cytokeratin 5

Alternative Name : Keratin 5,KRT5,EBS2,Epidermolysis Bullosa Simplex 2 Dowling-Meara/Kobner/Weber-Cockayne Types,Keratin 5 (Epidermolysis Bullosa Simplex,Dowling-Meara/Kobner/Weber-Cockayne Types),58 KDa Cytokeratin,Type-II Keratin Kb5,CK-5,K5,DDD1,CK5,DDD,

Description

Source : Escherichia Coli. KRT5 Human Recombinant produced in E.coli is a single, non-glycosylated polypeptide chain containing 613 amino acids (1-590) and having a molecular mass of 64.8 kDa.KRT5 is fused to a 23 amino acid His-tag at N-terminus. Cytokeratin 5 (KRT5) belongs to the keratin gene family. The type II cytokeratins are comprised of basic or neutral proteins which are arranged in pairs of heterotypic keratin chains coexpressed throughout differentiation of simple and stratified epithelial tissues. The type II cytokeratins are clustered in a region of chromosome 12q12-q13. The KRT5 type II cytokeratin is specifically expressed in the basal layer of the epidermis with family member KRT14. Mutations in these genes are linked with a complex of diseases termed epidermolysis bullosa simplex.

Product Info

Amount : 20 µg
Purification : Greater than 85% as determined by SDS-PAGE.
Content : The KRT5 solution (1mg/ml) contains 20mM Tris-HCl buffer (pH 8.0), 10% glycerol and 0.4M Urea.
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).Please avoid freeze thaw cycles.
Amino Acid : MGSSHHHHHH SSGLVPRGSH MGSMSRQSSV SFRSGGSRSF STASAITPSV SRTSFTSVSR
SGGGGGGGFG RVSLAGACGV GYGSRSLYN LGGSKRISIS TSGGSFRNRF GAGAGGGYGF
GGGAGSGFGF GGGAGGGFGL GGGAGFGGGF GPGFPVPCPP GGIQEVTVNQ SLLTPLNLQI
DPSIQRV RTE EREQIKTLNN KFASFDKVR FLEQQNKVLD TKWTLLQEQG TKTVRQNLEP LFEQYINNLR
RQLDSIVGER GRLDSELNRM QDLVEDFKNK YEDEINKRTT AENEVMLKK DVDAAYMNKV ELEAKVDALM
DEINFMKMFF DAELSQMOTH VS DTSVVL SM DNNRNLDLDS IIAEVKAQYE EIANRSRTEA ESWYQTKYEE
LQQTAGRHDG DLRNTKHEIS EMNRMIQRLR AEIDNVKKQC ANLQNAIADA EQRGELAL KD ARNKLAEELE
ALQKAKQDMA RLLREYQELM NTKLALDVEI ATYRKLEGE ECRLSGEGVG PVNISVVTSS VSSGYGSGSG
YGGGLGGGLG GGLGGGLAGG GSGSYSSSS GGVLGGGLS VGGSGFSASS GRGLGVGFGS
GGSSSSSVKF VSTTSSSRKS FKS.