

Produktinformation



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Datasheet

SHH (Human) Recombinant Protein (Q01)

Catalog Number: H00006469-Q01

Regulation Status: For research use only (RUO)

Product Description: Human SHH partial ORF (NP_000184, 181 a.a. - 280 a.a.) recombinant protein with GST-tag at N-terminal.

Sequence:

IHCSVKAENSVAAKSGGCFPGSATVHLEQGGTKLVKD LSPGDRVLAADDQGRLLYSDFLTFLDRDDGAKKVFYVI ETREPRERLLLTAAHLLFVAPHNDS

Host: Wheat Germ (in vitro)

Theoretical MW (kDa): 36.74

Applications: AP, Array, ELISA, WB-Re (See our web site product page for detailed applications information)

Protocols: See our web site at http://www.abnova.com/support/protocols.asp or product page for detailed protocols

Preparation Method: *in vitro* wheat germ expression system

Purification: Glutathione Sepharose 4 Fast Flow

Storage Buffer: 50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.

Storage Instruction: Store at -80 °C. Aliquot to avoid repeated freezing and thawing.

Entrez GenelD: 6469

Gene Symbol: SHH

Gene Alias: HHG1, HLP3, HPE3, MCOPCB5, SMMCI, TPT, TPTPS

Gene Summary: This gene encodes a protein that is instrumental in patterning the early embryo. It has been implicated as the key inductive signal in patterning of the ventral neural tube, the anterior-posterior limb axis, and the ventral somites. Of three human proteins showing sequence and functional similarity to the sonic hedgehog protein of Drosophila, this protein is the most similar. The protein is made as a precursor that is autocatalytically cleaved; the N-terminal portion is soluble and contains the signalling activity while the C-terminal portion is involved in precursor processing. More importantly, the C-terminal product covalently attaches a cholesterol moiety to the N-terminal product, restricting the N-terminal product to the cell surface and preventing it from freely diffusing throughout the developing embryo. Defects in this protein or in its signalling pathway are a cause of holoprosencephaly (HPE), a disorder in which the developing forebrain fails to correctly separate into right and left hemispheres. HPE is manifested by facial deformities. It is also thought that mutations in this gene or in its signalling pathway may be responsible for VACTERL syndrome, which is characterized by vertebral defects, anal atresia, tracheoesophageal fistula with esophageal atresia, radial and renal dysplasia, cardiac anomalies, and limb abnormalities. Additionally, mutations in a long range enhancer located approximately 1 megabase upstream of this gene disrupt limb patterning and can result in preaxial polydactyly. [provided by RefSeq]