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Lieferung & Zahlungsart

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Zuschläge

- Mindermengenzuschlag
- Trockeneiszuschlag
- Gefahrgutzuschlag
- Expressversand

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Datasheet

NOG (Human) Recombinant Protein

Catalog Number: P8974

Regulation Status: For research use only (RUO)

Product Description: Human NOG (Q13253, 28 a.a. - 232 a.a.) partial-length recombinant protein with His tag at C-Terminus expressed in HEK293 cell.

Sequence:

QHYLHIRPAPSDNLPLVLDLIEHPDPIFDPKEKDLNETLL
RSLGHHYDPGFMATSPPEDRPGGGGAAGGAEDLA
ELDQLLRQRPSGAMPSEIKGLEFSEGLAQGKKQRLSK
KLRRKLMWLWSQTFPCVLYAWNDLGSRFWPRYVK
VGSCFSKRSCSVPEGMVCKPSKSVHLTVLRWRCQR
GGQRCGWIPQIYPIIIECKCSCHHHHHH.

Host: Human

Theoretical MW (kDa): 23.8

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

Form: Liquid

Preparation Method: HEK 293T cell expression system

Purity: > 90% by SDS PAGE

Storage Buffer: 50mM MES (pH 6.5) and 30% glycerol.

Storage Instruction: Store at -20°C. Aliquot the product after reconstitution to avoid repeated freezing/thawing cycles.

Entrez GeneID: 9241

Gene Symbol: NOG

Gene Alias: SYM1, SYNS1

Gene Summary: The secreted polypeptide, encoded by this gene, binds and inactivates members of the transforming growth factor-beta (TGF-beta) superfamily signaling proteins, such as bone morphogenetic protein-4 (BMP4). By diffusing through extracellular matrices more efficiently than members of the TGF-beta

superfamily, this protein may have a principal role in creating morphogenic gradients. The protein appears to have pleiotropic effect, both early in development as well as in later stages. It was originally isolated from *Xenopus* based on its ability to restore normal dorsal-ventral body axis in embryos that had been artificially ventralized by UV treatment. The results of the mouse knockout of the ortholog suggest that it is involved in numerous developmental processes, such as neural tube fusion and joint formation. Recently, several dominant human NOG mutations in unrelated families with proximal symphalangism (SYM1) and multiple synostoses syndrome (SYNS1) were identified; both SYM1 and SYNS1 have multiple joint fusion as their principal feature, and map to the same region (17q22) as this gene. All of these mutations altered evolutionarily conserved amino acid residues. The amino acid sequence of this human gene is highly homologous to that of *Xenopus*, rat and mouse. [provided by RefSeq]