

Product datasheet for DA3516X

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VEGF-E Orf Virus Protein

Product data:

Product Type: Recombinant Proteins

Description: VEGF-E orf virus recombinant protein, 20 μg

Species: Orf Virus

Expression Host: E. coli

Predicted MW: 35 kDa

Purity: >90% pure by SDS-PAGE and visualised by silver stain.

Buffer: Presentation State: Purified

State: Lyophilized

Buffer System: PBS, pH 7.4 without stabilizers.

Biological: The ED50 for stimulation of 3H-thymidine incorporation and cell proliferation by

human umbilical vein endothelial cells for VEGF-E has been determined to be in the range of

5-20 ng/ml.

Specific: 2 x 10e5 units/mg

Endotoxin: < 0.1 ng per µg of VEGF-E

Reconstitution Method: Restore in PBS or medium containing at least 0.1% human or bovine serum albumin to a

concentration not lower than 50 µg/ml.

(VEGF-E is soluble in water and most aqueous buffers).

Preparation: Lyophilized

Protein Description: Recombinant ov-VEGF-E (Orf virus). A DNA sequence encoding the mature variant of ovVEGF-E

isolate D1701 (Dehio et al., 1999; GenBank accession No. AF106020) was expressed in E. coli as a 132 amino acid residue fusion protein with an N-terminal His-tag sequence and a thrombin cleavage site. Recombinant VEGF-E homodimer was dimerized *in vitro* and has a

predicted mass of approximately 35 kDa.

Note: Centrifuge vial before opening!

Storage: Store lyophilized VEGF-E at -20°C to -70°C.

Reconstituted VEGF-E should be stored in working aliquots at -20°C.

Avoid repeated freezing and thawing.

Stability: Shelf life: six months from despatch.





Summary:

Based on sequence similarity to VEGF-A, a gene encoding a VEGF homologue has recently been discovered in the genome of Orf virus (OV) (Lyttle et al., 1994). Different isolates of Orf virus show significant amino acid sequence similarity to VEGF-A and described as a viral virulence factor that appears to be derived from captured host genes. All eight cysteine residues of the central cysteine knot motif characteristic of members of the VEGF family are conserved among other residues in the VEGF-E proteins (Dehio et al., 1999; Wise et al., 1999). Alignment of all mammalian VEGF sequences indicated that VEGF-E is distinct from the previously described VEGFs but most closely related to VEGF-A. Like VEGF-A, VEGF-E was found to bind with high affinity to VEGF receptor-2 (KDR) resulting in receptor autophosphorylation, whilst in contrast to VEGF-A, VEGF-E can not bind to VEGF receptor-1 (Flt-1). Furthermore VEGF-E can also not bind to VEGF receptor-3 (FLT-4). Therefore VEGF-E is a potent angiogenic factor selectively binding to VEGF receptor -2/KDR.