

## 32-6666: AOC3 Human

**Alternative Name :** VAP-1, AOC3, HPAO, VAP1, Membrane primary amine oxidase, Copper amine oxidase, HPAO, Semicarbazidesensitive amine oxidase, SSAO, Vascular adhesion protein 1.

### Description

Source: Sf9, Baculovirus cells.

Sterile Filtered colorless solution.

Amine Oxidase Copper Containing 3, also referred to AOC3, a copper amine oxidase with a topaquinone cofactor. AOC3 is a cell adhesion protein which participates in recirculation & extravasation of lymphocyte by mediating the binding of lymphocytes to peripheral lymph node vascular endothelial cells in an L-selectin independent fashion. Amine Oxidase Copper Containing 3 acts in adipogenesis. The protein catalyzes the oxidative deamination of small primary amines such as methylamine, benzylamine & aminoacetone in a reaction that produces an aldehyde, ammonia and H<sub>2</sub>O<sub>2</sub>.

AOC3 Human produced in Sf9 Baculovirus cells is a single, glycosylated polypeptide chain containing 746 amino acids (27-763 aa) and having a molecular mass of 82.8kDa. AOC3 is fused to a 9 amino acid His tag at C-terminus and purified by proprietary chromatographic techniques.

### Product Info

**Amount :** 2 µg / 10 µg

**Purification :** Greater than 90.0% as determined by SDS-PAGE.

**Content :** The AOC3 solution (0.25 mg/ml) contains 10% Glycerol and Phosphate-Buffered Saline (pH 7.4).

**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). Avoid multiple freeze-thaw cycles.

**Amino Acid :** ADPGRGGDGG EPSQLPHCPS VPSAQPWTH PGQSQFADL SRELTAVMR FLTQRLGPG  
VDAAQARPSD NCVFSVELQL PPKAAALAH DRGSPPARE ALAIVFFGRQ PQPNVSELVV GPLPHPSYMR  
DVTVERHGGP LPYHRRPVLF QEYLDIDQMI FNRELQASG LLHHCCFYKH RGRNLVTMTT APRGLQSGDR  
ATWFGLYYNI SGAGFFLHHV GLELLVNHKA LDPARWTIQK VFYQGRYYDS LAQLEAQFEA GLVNVVLIPI  
NGTGGSWSLK SPVPPGPAPP LQFYPQGRF SVQGSRVASS LWTFSFGLGA FSGPRIFDVR FQGERLVYEI  
SLQEALAIYG GNSPAAMTTR YVDGGFGMGK YTTPLTRGVD CPYLATYVDW HFLLESQAPK TIRDAFCVFE  
QNQGLPLRRH HSDLYSHYFG GLAETVLVVR SMSTLLNYDY VWDTVFHPSG AIEIRFYATG YISSAFLFGA  
TGKYGNQVSE HTLGTVHTHS AHFKVDLVA GLENWVWAED MVFVPMVAVPW SPEHQLQRLQ  
VTRKLEEMEE QAAFLVGSAT PRYLYLASNH SNKWGHPRGY RIQMLSFAGE PLPQNSSMAR GFSWERYQLA  
VTQRKEEES SSSVFNQNDP WAPTVDVDFSD INNETIAGKD LVAVVTAGFL HIPHAEDIPN TVTVGNGVGF  
FLRPYNFFDE DPSFYSADSI YFRGDQDAGA CEVNPLACLQ QAAACAPDLP AFSHGGFSSH HHHHHH