

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|--|--------------|-------------------------------|-----------------------|-------------------------------------|---|
| Q4LDE5 | SVEP1_HUMAN | Homo sapiens | Sushi, von Willebrand factor type A, EGF and pentraxin domain-containing protein 1 | 24.466594 | T1224;T1262;S2088;S2090;S2092 | NaN | 35254053;38665916;28657654;35289036 | MWPRLAFCCWGLALVSGWATFQQ MSPSRNFSFRLFPETAPGAPGSIPAP PAPGDEAAGSRVERLQAFRRRVRL LRELSELELVFLVDDSSSVGEVNF RSELMFVRKLLSDFPVVPTATRVAIV TFSSKNYVPRVDYISTRARRQHKCA LLLQEIPAIYRGGGTYTKGAFQAAA QILLHARENSTKVFLITDGYNNGGD PRPIAASLRDSGVEIFTFGIWQGNIR ELNDMASTPKEEHCCYLLHSFEEFEA LARRALHEDLPSGSFIQDDMVHCSY LCDEGKDCCDRMGSKCGTHTGHF ECICEKGYGKGLQYECTACPSGTYK PEGSPGGISSCPCDENHTSPPGST SPEDCVCREGYRASGQTCELVHCPA LKPPENGYFIQNTCNNHFNAACGVR CHPGFDLVGSSIILCLPNGLWSGSES YCRVRTCPHLRQPKHGHISCSTREM LYKTTCLVACDEGYRLEGSCKLTCQ GNSQWDGPEPRCVERHCSTFQMPK DVIISPHNCGKQPAKFGTICVSCRQ GFILSGVKEMLRCTTSGKWNVGVQA AVCKDVEAPQINCPKDIEAKTLEQQ DSANVTWQIPTAKDNSGEKVSVHV HPAFTPPYLFPIGDVAIVYTATDLSG NQASCIFHIKVIDAEPVIDWCRSPP PVQVSEKVHAASWDEPQFSDNSGA ELVITRSHTQGDLPQGETIVQYTAT DPSGNNRTCDIHIVIKGSPCEIPFTPV NGDFICTPDNTGVNCTLTCLLEGYDF TEGSTDKYYCAYEDGVWKPTYTTEW PDCAKKRFANHGFKSFEMFYKAAR CDDTDLMKKFSEAFETTLGKMVPSF CSDAEDIDCRLEENLTKKYCLEYNY DYENGFAIGPGGWGAANRLDYSYDD FLDTVQETATSIGNAKSSRIKRSAPL SDYKIKLIFNITASVPLPDERNDTLE WENQORLLQTLTITNKLKRTLNDK PMYSFQLASEIILIADSNLETKKASP FCRPGSVLRGRMCVNCPLGTYYNLE HFTCESCRIGSYQDEEGQLECKLCP SGMYTEYIHSRNISDCKAQCKQGTY SYSGLETCECPLGTYQPKFGSRSC SCPENTSTVKGAVNISACGVPCPE GKFSRSGLMPCHPCPRDYYQPNAG KAFCLACPFYGTTPFAGSRSITECSS FSSTFSAAEESVPPASLGHIKKRHE ISSQVFHECFNPNCHNSGTCQQLGR GYVCLCPLGYTGLKCETDIDECPLP CLNNGVCKDLVGEFICECPSGYTGQ RCEENINECSSPCLNKGICVDGVA GYRCTCVKGFVGLHCETEVEQCQS NPCLNNAVCEQVGGFLCKCPPGFL GTRCGKNVDECLSQPCKNGATCKD GANSFRCLCAAGFTGSHCELNINEC QSNPCRQATCVDELNSYSCKQP |

GFSGKRCETEQSTGFNLD FEVSGIY
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