

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O88307	SORL_MOUSE	Mus musculus	Sortilin-related receptor	13.680566	NaN	S114;S2207	22645316	MATRSSRRESRLPFLFALVALLPRGALGGGWTQRLHGGPAPLPQDRGFFVVOGDPRDLRLGTHGDAPGASPAARKPLRTRRSAALQPQPIQVYGQVSLNDSHNQMVVHWAGEKSNVIVALARDSLALARPKSSDVVVSVDYGKSFSEKLNFGVGNNEAVISQFYHSPADNKR YIFVDAYAQYLWITFDFCSTIHGFSIPFRAADLLLHASKASNLLLGFD RSHPNKQLWKSDDFGQTWIMI QEHVKSFSWGIDPYDQPNAIYIERHEPFGFSTVLRSTDFQSRNQEVILEEVRDFQLRDKYMFATKVVHLPGSQQSSVQLWVSFGRKPMRAAQFVTKHPINEYYIADAAEDQVFVCVSHSNSTNLYISEAELKFSLSLENVLYYSPGGAGSDTLVRYFANEPFADFHRVEGLQGVYIATLIN GSMNEENMRSVITFDKGGTWEFLQAPAFTGYGEKINCELSQGCSLHLAQRLSOLLNLQLRRMPILSKESAPGLIIATGSVGKNLASKTNVYISSSAGARWREALPGPHYTTWGDHGGIIMAI AQGETNELKYSTNEGETWKTFFVSEKPVFVYGLLTPGKSTVFTIFGSNKESVHSWLILQVNATDALGVPCTENDYKLWSPSDERGN ECLLGHKTVFKRRTPHATCFNGEDFDRPVVVSNCSTREDEYECDFGFKMSEDL SLEVCVPDPEFSGKPYSPVPCPVGSSYRRTRGYRKISGDTCSGGDVEARLEGE LVPCLAEENEFILYAMRKS IYRYDLASGATEQLPLSGLRAAVALDFDYERNCLYWSDLA LDTIQRLCLNGSTGQEVINSGLETV EALAFEPLSQLLYWVDAGFKKIEVANPDGDFRLTIVNSSVLDRPRALVLPQEGVMFWDWGD LKPGIYRSYMDGSAAYRLVSEDKWPNGISVDSQWIY WTDAYLDCIERITFSGQQRSVILDSL PHPYAI AVFKNEIYWDDWSQLSIFRA SKHSRSQVEILASQLTGLMDMKVFY KGKNAGSNACVPQPCSL LCLPKANN SKSCRCPEGVASSVLP SGDLMCDCPQGYQRKNNTCVKEENTCLRNQYRC SNGNCINSIWWCDFDND CGDMSDERNCPTTVCDADTQFRCQESGTCIPL SYKCDLEDDCGDNSDESHCEMHQCRSDEFNCSSGMCIRSSWVCDGDND CRDWSDEANCTAIYHTCEASNFC HNGHCIPQRWACDGDADCQDGSDE

DPVSCEKKCNGFHCPCNGTCIPSSKH
CDGLRDCPDGSDEQHCEPFCTRFM
DFVCKNRQQCLFHSMVCDGIVQCR
DGSDEDAAFAGCSQDPEFHKECDE
FGFQCQNGVCISLIWKCDGMDDCG
DYSDEANCENPTEAPNCSRYFQFHC
ENGHCIPNRWKCDRENDCGDWS
EKDCGDSHVLPSPTPGPSTCLPNYF
HCSSGACVMGTWVCDGYRDCADGS
DEEACPSLANSTAASTPTQFGQCDR
FEFECHQPKKIPNWKRCDBGHQDC
QDGQDEANCPTHSTLTCTSREFKCE
DGEACIVLSERCDFGLDCSDESDEK
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LTWMRPKKMPASCVYNVYRVVG
ESIWKLETHSNKTSTVLKVLKPDTT
YQVKVQVHCLNKVHNTNDFVTLRT
PEGLPDAPRNLQLSLNSEEEGVILG
HWAPPVHTHGLIREYIVEYSRSGSK
MWASQRAASNSTEIKNLLLNALYTV
RVAAVTSRGIGNWSDSKSITTIGKV
IQAPNIHIDSYDENSLSFTLTMDGDI
KVNGYVNLFWSFDAHKQEKKTL
FRGGSALSHRVSNLTAHTSYEISAW
AKTDLGDSPLAFEHILTRGSSPPAPS
LKAKAINQTAVECIWTGPKNVYGFIF
YATSFLDLYRNPKSVTTLHNKTIV
SKDEQYLFLVRVLPYQGPSSDYVVV
KMIPDSRLPPRHLHAVHIGKTSALIK
WESPYDSPDQDLFYAIAVKDLIRKTD
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LKIITENDHVLLFWKSLALKEKQFNE
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FFKVSNLKMGHNYFTVQARCLFGS
QICGEPVLLYDELSSGADA AVIQAA
RSTDVAVVVPILFLILLSLGVGFAIL
YTKHRRQLQSSFAFANSHYSSRLGS
AIFSSGDDLGEDDEDAPMITGFSDD
VPMVIA