

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
Q9D415	DLGP1_MOUSE	Mus musculus	Disks large-associated protein 1	33.732466	T525;T526;T531;S534	S169;S362;S365;S368;S372;S389;S418;S421;S425;S428;S437;S509;S516;S578;T579;S581;S605;T606;S608;S611;S947	19458039;34678516;22517741;35822049;22826440;22645316	MKGLSGSRSHHHGITCEAACDSLSH HSDHKPYLLSPVDHHPADHPYYTQR NSFQAEQCVGPFSDPLASSTFPRRHY TSQQELKDEALVPRTLATKANRLPT NLLDQFERQLPLSRDGYHTLQYKRT AVEHRSDSPGRIRHLVHSVQKLFK SHSLEGPSKGSVNGGKASPDESQTL RYGKRKSKERRSEKARSNASNAS PTSPSWWSDDNLDGDMCLYHTPS GVMTMGRCPDRSASQYFMEAYNTI SEQAVKASRSNNDIKCSTCANLPVT LDAPLLKKSASWSTLTVSRAREVYQ KASVNMDQAMVKSEACQQRSCQY LQVPQDEWSGYTPRGKDDEIPCRRM RSGSYIKAMGDEDSGSDTSPKPSP KVAARRESYLKATQPSLTELTLTKIS NEHSPKLQIRSHSYLRAVSEVSINRS LDSLDPAGLLTSPKFRSRNESYMRA MSTISQVSEMEVNGQFESVCESVFS ELESQAVEALDPLPGCFMRMRSHSY VRAIEKGCSDDECVSLRSSPPRRT TTVRTIQSSTGVIKLSSAVEVSSCITT YKKTPPPVPRTTTKPFISITAQSSTE SAQDAYMDGQGQRGDMISQSGLSN STESLDSMKALTAIEAANAQIHGPA SQHMGSNAAAVTTTTIATVTTEDR KKDFKKNRCLSIGIQVDDAEPEKRM AESKTSNKFQSVGVQVEEEKCFRRF TRNSVTTAVQADLDFHDNLENSLE SIEDNSCPGPMARQFSRDASTVSI QSGSNHYHACAADDDFDTFDPSIL PPDPWIDSITEDPLEAVQRSVCHRD GHWFLKLLQAERDRMEGWCKLME REERENLNPEDILGKIRTAVGSAQLL MAQKFYQFRELCEENLNPNAHPRP TSQDLAGFWDMLQLSIENISMKFDE LHQLKANNWKQMDPLDKKERRAPP PVPKKPAKGPAPLIRERSLESSORQE ARKRLMAAKRAASVRQNSATESAES IEIYIPEAQTRL

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