

| UniprotKB ID | Entry name | organism | full name | oglnacscore | oglnacsites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extracellular region |
|--------------|-------------|--------------|--|-------------|-------------|-----------------------|----------|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| Q80U19 | DAAM2_MOUSE | Mus musculus | Disheveled-associated activator of morphogenesis 2 | 24.474581 | NaN | NaN | 33300544 | MALRRKSPHGLGFLCCFGGSDLPEI DLRDSHPLOYLEFSGPIPPEELNVR FAELVDELDTLTKNREAVFALPPEK KWQIYCSKRKEQEDPNKLATSWPEY YIDRINAMAAMQNYETEDEETDKR NQVVEDLKTALRTQPMRFVTRFIDL EGLTCLLNFLRGMDHTTCSRIHTS LIGCIKALMNNNSQGRAHVLAQPEAIS IIAQLRRTENSKTKVAVLEILGAVCLV PGGHKKVLOAMLHYQAYAAERTF QTLNLDRLSLGRYRDEVNLTAIM SFINAVLNAGAGEDNLEFRLHLRYE FLMLGIQPVIDKLRQHENAIDKHL DFFEMVRNEDDLELARRFDMVHID TKSASOMFELIHKKHKHTEAYPCLLS VLHHCLOMPYKRNGGYFQQWQLLD RILQQIVLQDERGVDPDLAPLENFNV KNIVNMLINENEVKQWRDQAEKFR KEHMELMSRLERKERECETKTLEK EEMMRTLNMKDKLARESQELROA RGQVAELVARHNESSTGPVSSPPPP GGPLTLSSRRTNDLPPPPPLPFDS CPPPPAPPLPPGGPPIPPAPPFCSSG PPPSHDPPSSNEAPLRKKRIPQPSHP LKSFNWVKLNEERVSGTVWNEIDD SQVFRILDLEDFEKMFSAQRHQAC MQEGPQREGRGNVRDGGAAASRPLPA VEASAHARTEKASRMVSAATGAKKEL GSTEDIYITSRKVKELSVIDGRRRAQN CILLSKLKSNDIEIRQAILRMDQE DLAKDMLEQLLKFIPEKSDIDLLEEH KHEIERMARADRFLEYMSRIDHYQQ RLQALFFKKKFQERLAEAKPKVEAIL LASRELTLSQRLKQMLEVVLAIGNF MNKGQGGGAYGFRVASLNKIADTKS SIDRNISLLHYLIMILEKHFDPDILNMP SELKHLSEAAKVNLAELEKEVSILRR GLRAVEVELEYQRHQARDPNDKFVP VMSDFITVSSFSFSELEDQLNEARD KFAKALTFGEQESKMQPDEFFGIF DTFLQAFLEARQDLEAMRRRKEEDE RRARMEFMLEQREKERWQRQRK VLAGGALEESGEFDDLVSALRSGEV FDKDLSKFKRNRKRPGSQVPEVTRE RAINRLNY | None | None | None | None | None | None | None | | |

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