

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
089032	SPD2A_MOUSE	Mus musculus	SH3 and PX domain-containing protein 2A	13.597734	NaN	T256;S405;S420;S546;S566;S592;S643;T728;S764;S766;S812;T822;S993;S1007;S1008;S1029	28528544;25153642	MLAYCVQDATVVDVEKRRSPSKHYV YIINVTWSDSTSQTIYRRYSKFFDLQ MQLLDKFPPIEGGQKDPKQRIIPFLPG KILFRRSHIRDVAVKRLKPIDEYCRAL VRLPPHISQCDEVFRFFEARPEDVN PPKEDYGSSKRKSVWLSSWAESPCK DVTGADTNAEPMILEQYVVVSNYKK QENSELSLQAGEVVDVIEKNESGW WVSTSEEQGWVPATYLEAQNTR DDSDINTSKTGEVSKRRKAHLRRLD RRWTLGGMVNRQHSREEKYVTVQP YTSQSKDEIGFEKGVTVVIRKNLEG WWYIRYLGKEGWAPASYLKKAKDDL PTRKKNLAGPVEIIGNIMEISNLLNK KASGDKEAPAEGEGSEAPITKKEISL PILCNASNGSALAIPERTTSKLAQGS PAVARIAPQRAQISSPNLRTTRPPRR ESSLGFQLPKPEPPSVEVEYYTIAEF QSCISDGISFRGGQKAEVIDKNSSG WWYVQIGEKEGWAPASYIDKRKKPN LSRRTSTLTRPKVPPPAPPSKPKEAE ENPVGACESQGSPLKVKEYEPEYDV PAFGFDSEPEMNEEPSGDRGSGDK HPAQPRRISPASSLQRAHFVKVGESSE DVALEEETIYENEGFRPYTEDTLRAR GSSGSDSPGSSSLSLAVKNSPKSD SPKSSLLKLAEKNAQAELGKNQS NISFSSSVTISTTCSSSSSSSSLSKNN GDLKPRSASDAGIRDTPKVGTKKDP DVKAGLASCARAKPSVRPKPVLNRA ESQSQEKMDISSLRRLRPTGQLRG GLKGSRSSESELPPQMASEGSRRGS ADIPLTATTPPCVPPKKEWEGQGATY VTCSAYQKVQDSEISFPEGAEVHVLE KAESGWYVRFGELEGWAPSHYLV AEENQQPD TASKEGDTGKSSQNEG KSDSLEKIEKRVQALNTVNQSKRAT PPIPSKPPGGFGKTSGTVAVKMRNG VRQVAVRPQSVFVSPPKDNNLSCA LRRNESLTATDSLGRVRRNSSFSTA RSAAAEAKGRLAERAASQGSSEPLLP TQRKGIPVSPVRPKPIEKSQFIHNNL KDVIYISIADYEGDEETAGFQEGVSM EVLEKNPNGWWYQCILDEVKPFKG WVPSNYLEKKN