

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extract region
P28028	BRAF_MOUSE	Mus musculus	Serine/threonine-protein kinase B-raf	24.474581	NaN	S135;S316;S348;T356;T379;S382;T384;S431;S432;S714;S735;T738	33300544	MAALSGGGSSSGGGGGGGGGGG GDGGGGAEEQQALFNGDMEPEAGA GAAASSAADPAIPEEVVNIKQMIKLT QEHIEALLDKFGGEHNPPSIYLEAYE EYTSKLDALQREQLLESVLFQTPT DASRNNPKSPQKPIVRVFLPNKQRT VVPARCGVTVRDSLKKALMMRGLIP ECCAVYRIQDGEKKPIGWDTDISWL TGEELHVEVLENVPLTTHNFVRKTF FTLAFCDFCRKLLFQGFRCQTCGYK FHQRCSTEVPLMCVNYDQLDLLFVS KFFEHHVPQEEASFPETALPSGSSS APPSDSTGPOILTSPPSPKSIPIPPF RPADEDRNQFQGRDRSSAPNVHI NTIEPVNIDDLIRDQFRGDDGGSTTG LSATPPASLPGSLTNVKALQKSPGPQ RERKSSSSSEDRSRMKTIGRRDS SDDWEIPDGQITVGQRIGSGSFGTVY KKGWHGDVAVKMLNVTAPTPQQLQ AFKNEVGVLKTRHVNILLFMGYST KPOLAIVTQWCEGSSLYHLLHIETK FEMIKLIDIARQTAQGM DYLHAKSII HRDLKSNNIFLHEDLTVKIGDFGLA TVKSRWGGSHQFEQLSGSILWMAP EVIRMQDKNPYSFQSDVYAFGIVLYE LMTGQLPYSNINNRDQIIFMVGRGY LSPDLSKVRSNCPKAMKRLMAECLK KKRDERPLFPQILASIELLARS LPKIH RSASEPSLNRAGFQTEDFSLYACASP KTPIQAGGYGGFPVH	True	True	4.65	4.559	5.0	2.456	1.983	5.0	2.967