

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrace region
Q99N57	RAF1_MOUSE	Mus musculus	RAF proto-oncogene serine/threonine-protein kinase	30.704947	NaN	S29;S43;S233;S252;S259;T268;T269;S289;S296;S301;S338;S339;T340;T341;S471;T491;S494;S497;S499;S621;S642	40128797	MEHIQGAWKTI SNGFGLKDAVFDGSSCISPTIVQQFGYQRRASDDGKLTDSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVRGLQPECCAVFRLQEHKGGKARLDWNTDAASLIGEELQVDFLDHVPLTTHNFARKTFLKLAFCDICQKFLLNGFRCQTCGYKFHEHCSTKVPTMCVDWNSNIRQLLFPNSTVGDSGVPAPPSFPMRRMRESVSRMPASSQHRYPHAFHTFNTSSPSSEGLSQRQRSTSTPNVHMVSTTLHVDSRMIEDAIRSHSESASPSALSSSPNNLSPTGWSQPKTPVPAQRERAPGSGTQEKNKIRPRQRDSSYYWEIEASEVMLSTRIGSGSFGTVYKKGWHGDVAVKILKVVDPPTPEQLQAFRNEVAVLRKTRHVNILLFMGYMTKDNLAIVTQWCEGSSLYKHLHVQETKQMFQLIDIARQTAQGM DYLHAKNIIHRDMKSN NIFLHEGLTVKIGDFLATVKSRWSGSQQVEQPTGSVLWMAPEVIRMQDDNPFQSDVYSYGIVLYELMAGELPYAHINN RDQIIFMVGRGYASPDLSRLYKNCPKAMKRLVADCVKKVKEERLFPQILSSIELLQHS LPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF	True	False	5.0	3.0	3.0	False	4.0	4.0	False