

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q16849	PTPRN_HUMAN	Homo sapiens	Receptor-type tyrosine-protein phosphatase-like N	22.903221	S449;S454;T457	S308	38253038	MRRPRRPGGLGGSGGLRLLCLLLL SSRPGGCSAVSAHGCLFDRRLCSHL EVCIQDGLFGQCQVGVGQARPLLQV TSPVLQRLQGVLRQLMSQGLSWHD DLTQYVISQEMERIPRLRPPPEPRFRD RSGLAPKRPGPAGELLQDIPTGSAP AAQHRLPQPPVGGKGGAGASSLSPL QAELLPPLEHLLPPQPPHPSLSYE PALLQPYLFHQFGRDGSRVSEGS GMVSVGPKAEAPALFRTASKGIF GDHPGHSYGDLPGPSAQLFQDSGL LYLAQELPAPSRARVPRLPEQGSSSR AEDSPEGYEKEGLDRGEKPAFPAV QPDAALQRLAAVLAGYGVLRQLTP EQLSTLLTLLQLLPKGAGRNPGGVV NVGADIKKTMEGPVEGRDTAELPAR TSPMPGHPTASPTSSEVQVPSVPS SEPPKAARPPVTPVLEKKSPLGQSQ PTVAGQPSARPAEEYGYIVTDQKPL SLAAGVKLEILAEHVMSSGSFINI SVVGPALTFRIRHNEQNLSLADVTD QAGLVKSELEAQTGLQILQTVGVGQR EEAAAVLPQTAHSTSPMRSVLLTLV ALAGVAGLLVALAVLCVRQHARQQ DKERLAALGPEGAHGDITTFEYQDLC RQHMATKSLFNRAEGPPEPSRVSSV SSQFSDAAQASPSHSSSTPSWCCEP AQANMDISTGHMILAYMEDHLRNR DRLAKEWQALCAYQAEPTCATAQ GEGNIKKNRHPDFLPYDHARIKLV ESSPSRSDYINASPIEHDPMPAYIA TQGPLSHTIADFWQMVWESGCTVIV MLTPLVEDGVKQCDRYWPDEGASL YHVYEVNLYSEHIWCEDFLVRSFYI KNVQTOETRTLTOFHFLSWPAEGTP ASTRPLLDFRRKVNKCYRGRSCHIIV HCSDGAGRTGTYILIDMVLNRMAKG VKEIDIAATLEHVRDQRPGLVRSKDQ FEFALTAVAEVNAIKALPO	True	True	2.476	5.0	1.698	1.744	4.281	4.532	2.247

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