	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrace region
0	O94762	RECQ5_HUMAN	Homo sapiens	ATP-dependent DNA helicase Q5	20.148807	T456;T898	S488;S491;S727;S815;T839	30379171;38253038	MSSHHTTFPFDPERRVRSTLKKVFG FDSFKTPLQESATMAVVKGNKDVFV CMPTGAGKSLCYQLPALLAKGITIVV SPLIALIQDQVDHLLTLKVRVSSLNS KLSAQERKELLADLEREKPQTKILYI TPEMAASSSFQPTLNSLVSRHLLSYL VVDEAHCVSQWGHDFRPDYLRLGA LRSRLGHAPCVALTATATPQVQEDV FAALHLKKPVAIFKTPCFRANLFYDV QFKELISDPYGNLKDFCLKALGQEA DKGLSGCGIVYCRTREACEQLAIELS CRGVNAKAYHAGLKASERTLVQND WMEEKVPVIVATISFGMGVDKANVR FVAHWNIAKSMAGYYQESGRAGRD GKPSWCRLYYSRNDRDQVSFLIRKE VAKLQEKRGNKASDKATIMAFDALV TFCEELGCRHAAIAKYFGDALPACAK GCDHCQNPTAVRRRLEALERSSSW SKTCIGPSQGNGFDPELYEGGRKGY GDFSRYDEGSGGSGDEGRDEAHKR EWNLFYQKQMQLRKGKDPKIEEFV PPDENCPLKEASSRRIPRLTVKAREH CLRLLEEALSSNRQSTRTADEADLR AKAVELEHETFRNAKVANLYKASVL KKVADIHRASKDGQPYDMGGSAKSC SAQAEPPEPNEYDIPPASHVYSLKPK RVGAGFPKGSCPFQTATELMETTRI REQAPQPERGGEHEPPSRPCGLLDE DGSEPLPGPRGEVPGGSAHYGGPSP EKKAKSSSGGSSLAKGRASKKQQLL ATAAHKDSQSIARFFCRRVESPALLA SAPEAEGACPSCEGVQGPPMAPEKY TGEEDGAGGHSPAPPQTEECLRERP STCPPRDQGTPEVQPTPAKDTWKGK RPRSQQENPESQPQKRPRPSAKPSV VAEVKGSVSASEQGTLNPTAQDPFQ LSAPGVSLKEAANVVVKCLTPFYKEG KFASKELFKGFARHLSHLLTQKTSP GRSVKEEAQNLIRHFFHGRARCESE ADWHGLCGPQR	None	None	None	None	None	None	None	None	None