

| 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 |
|--------------|------------|--------------|--|--------------|---------------|--------------------------|-------------------|--|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| Q8N9B5 | JMY_HUMAN | Homo sapiens | Junction-mediating and -regulatory protein | 18.419428 | S832 | S115;S121;S713;S888;S974 | 30379171;23301498 | MSFALEETLES DWVAVRPHVFDERE KHKFVFI VAWNEIEGKFAITCHNRTA QRQRSGSREQAGARGGAEAGGAAS DGSRGPGSPAGRGRPEATASATLVR SPGPRSSAWAEGGSPRSTRSLLGD PRLRSPGSKGAESRLRSPVRAKPIPG QKTSEADDAAGAAAAARAPPREAQ VSSVRIVSASGTVSEEIEVLEMVKED EAPLALSDAEQPPATELESPAEECS WAGLFSFQDLRAVHQQLCSVNSQL EPCLPVFPPEPSGMWTVLFGGAPEM TEQEIDTLCYQLQVYLGHLDTCCGW KILSQVLFTETDDPEEYYESLSELRO KGYEVLQRARKRIOELLDKHKNTE SMVELLDLYQMEDEAYSSLAEATTE LYQYLLQPFRRDMRELAMLRROQIKI SMENDYLGPRRIESLQKEDADWQR KAHMAVLSIQDLTVKYFEITAKAKA VYDRMRADQKKFGKASWAAAAERM EKLOYAVSKETLQMMRAKEICLQQR KHALKEEMQSLRGGTEAIARLDQLE ADYYDLQLQLYEVQFEILKCELLLT AQLESIKRLISEKRDEVVYDYTESM EAMLEKEEMAASAYLQREELQKLQ QKARQLEARRGRVSAKKSYL RNKKE ICIAKHNEKIQRRTRIEDEYRTHHTV QLKREKLHDEERKSAWVSQERQR TLDRLRTFKQRYPGQVILKSTRRLRLA HARRKGAASPVLQEDHCDSLPSVLQ VEEKTEEVGEGRVKRGPSQTTEPQS LVQLEDTSLTQLEATSLPLSGVTSEL PPTISLPLNNNLEPCSVTINPLPSPL PPTPPPPPPPPPPPPPLPAKDSGP ETLEKDLPRKEGNEKRIPKSASAPSA HLFDSSQLVSARKKLRKTAEGLQRR RVSSPMDEVLASLKRGSFHLKKVEQ RTLPPFPDEDDSNILAQIRKGVKLG KVQKDVRESFTLLPDTDPLTRSIHE ALRRIKEASPESEDEEALPCTDWE N | True | False | 2.521 | 4.745 | 1.345 | 1.278 | 1.255 | 2.436 | 1.192 |