

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|-----------|--------------|---------------|---|----------|--|
| Q8R0W0 | EPIPL_MOUSE | Mus musculus | Epiplakin | 16.794757 | NaN | T33;T1551;S2430;S2508;S3220;S3460;S3538;S3735;S3975;S4053;S4765;S5005;S5083;S5795;S6035;S6113;S6310 | 37453647 | MNGQAPPHDVVVANGTEKFIVPKIK KNQLGASTPSRPQAEALPTTARSIA GVYVEASGQTQSIYAAIKQGLLPTGL GLTLLEAQAATGGLVDLAQGGQLLPV SEALRRGLVGLLELKEKLLAAERAVTG YDPYGG EKLSLFQAIKKEVVDRTL WRLLAQLATGGLVDPTQGVQVAPE LACQGGLLDKETWLSLVESEPSMGT PGFSDPNTLEQLPYSVLLGRCVQDP SSGLPLLPLKTTFFHTLAGAASASMLL EAGVLNEEMVRDLQEGMLVSDVG TRPEVRRYLEGTGGLAGVLLPGGH KKSFFQATVEHLVSKGIALQLLEAQA ATRTL VHP TTGQRLWVEEAVKAGLV GPELHEQLLVAEQAVTGYDPPFSSS RIPVFQAMKKGLVDQPLALRLDDAQ LATGG LICPARRFRLPLEAALRFGL DEETRQRLSQAMGFSDPTTHDRLGY EQLLALSVDTPETGLAFLPLPGMSH ANEPQGPTFIDHCTRQALSKATTSIS VGRYQGRPVS L WELLFSESVPVKKR AMLAQRHQEGALSVEELAAELKNIV EQAAATAKVTFAGLRDTPVTPGELLKA EIINQDLFEQLERGQTS AQDVGSLDS VQRYLQGTGSIAGLLL PDSQERLSIY EARSKGLLRPGTALILLEAQAATGFII DPKENKRYSV E EALRAGVIGPDVYA KLLSAEHAVTGYTDPYSGEQISL FQA MQRDLIVRDHGIRLLEAQIATGGVID PVHSHRVPVDVAYQRGYFDQILNSIL LDPSDDTKGFFDPNTHENLTYLQLL ERCVHDS ETGLHLLPLSSTRPQLVD SSTRQAFQKLLSVKYGRFRGQRVS AWELVNSEYFTEDRRRQLLQRYRQR KITLEQVTQLLEKEMRRWTDITLPAL QGQVTAYQLLEAHIINQELLDQVLTG TISPDALLQVGDVHRYLRGSGTVGG VLLKPSNQRISLYQAMKQKLLPPSTA LALLEAQAATGTITDPCSMETLSVDE AVCRGVVGA E VYGK LKRAEHSITGY RDPFSGKKVSLFRAMKKGLVPVEQA TRLLEAQVSTGGVVDPTTHLHLPMP VAVQRGCIDREME AALS RSPETFTP DGRGHTSYAQLLEHCLQDKASGLHL LPLTEDAPNVPTDTQIQETLQASAGT EDGLSLWDL LTSCHFTEE QRRGYLE DVKVGKISVPQLQNTVRSWVHSAKL LARARITVPGPRGEVPATWLRDAGII TQETLEALA QGMQSPDEVAKQPTVK |

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