

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasmic reticulum | golgi apparatus | plasm memb |
|--------------|-------------|--------------|------------------------------|--------------|---------------|--------------------------|---|--|---------------|---------------|---------|---------|--------------|-----------------------|-----------------|------------|
| Q9H3P2 | NELFA_HUMAN | Homo sapiens | Negative elongation factor A | 18.844331 | T452 | T157;S225;S233;T277;S363 | 37340703;28510447;35138101;38665916;32119511;29485866;34019948;39302247 | MASMRSDTGLWLHNKLGATDEL WAPPSIASLLTAAVIDNIRLCFHGLS SAVKLKLLGLTLHLPRRTVDEMKGAL LMEIQLASLDSDPWVLMVADILKSF PDTGSLNLEEEQPNVQDILGELR EKVGECEASAMLPLECYLNKNALT TLAGPLTPPVKHFQLKRKPKSATLRA ELLQKSTETAQQLKRSAGVPFHAKG RGLLRKMDTTTPLKQIPKQAPFRSPT APSVFSPGTGNRTPIPPSRTLRRKERG VKLLDISELDMVGAGREAKRRRRTL DAEVVEKPAKEETVVENATPDYAAG LVSTQKLGSLNNEPALPSTSYLPSTP SVVPASSYIPSSETPPAPSSREASRPP EEPSAPSPTLPAQFKORAPMYNSGL SPATPTPAAPTSPLTPTTPPAVAPTTQ TPPVAMVAPQTQAPAQQPKKNLSL TREQMFAAQEMFKTANKVTRPEKA LILGFMAGSRENPCQEQQDVIQIKLS EHTEDLPKADGQGSTTMLVDTVFE MNYATGQWTRFKKYKPMTNVS | True | False | 4.434 | 5.0 | 0.982 | 0.806 | False | 0.754 |