

| UniprotKB ID | Entry name   | organism                | full name             | oglcnaScore | oglcna sites | phosphorylation sites | PMIDS    | sequence  | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extracellular region |
|--------------|--------------|-------------------------|-----------------------|-------------|--------------|-----------------------|----------|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| V5B9A3       | V5B9A3_TRYCR | Trypanosoma cruzi Dm28c | Coatomer subunit beta | 19.946178   | NaN          | NaN                   | 30984116 | MNVGSNQEVACTLLVGF DGSSVNT<br>KDLKAALKEGDINARAEALETMIRL<br>HLNGEPQNHMIMSVIKYITPLEDHLI<br>KKLVLYFWEVVEKTDKNGKLLSEMI<br>LICSFLREDLLHPNEYVVRGLTLRFLC<br>KVKEKELIEPLISSVVQNLTHRVTYV<br>RRSAVAHVHAIYKRFDPDLSDAPELV<br>EKFIGDENDVSARRNAFDMLECAP<br>ERAVRFLTGFRESTNMADAGAAFO<br>MSVDFARQMIRANPYDKAKYVAIL<br>FNVLQSKNPAVRYQCASTLLSLSSP<br>TAIRQAALTFIDLLKTHSDNSVRLIVV<br>DQLDGMRRMRFGDILQESLLDVMSV<br>LANSTTEICKKVVTLAVELVSNKNVE<br>VFLQAIKELLRSQSEEDVLDKSSIQ<br>EYRQLLIRAIHTAVIRQPQMAATVLP<br>MMMDYICDSSNGSQEVISFIREVLQ<br>VQPLL RAGTLKQLLEVFPMTSPGV<br>VRTVLWLF GAHVSSADEVLGVLALL<br>ANSLKPLPLTPPVPPAASVDGDANG<br>AESPVMRAVTTVREDGTYVTSYTTVP<br>TGTMAA VAAVTDTTEDAGVVPAGLR<br>LLITKGDYFVASALASTLSKLLIRLFT<br>SYGSVDTAVKSKAQDDALMLLHEII<br>RYGTAPDAACAMDEDSHEHIRLSIIT<br>IANPQSPFLQVFDVDSYKALNAVES<br>VVGTA VGETLPNDWSKAKEPTAFCS<br>IDTPHFTQLAQGKDAFLEATDDL<br>SAIANASIDKTEEFKRLKLEKTIPLSGF<br>CDPVYCEASVTVHQFDVSDWYLVN<br>KTSNLMQELTIELASLGGMKLCERP<br>QVHSLPPHGSLKLR TALKVSSSTETG<br>VIYASVLYDAPNGERCCVILNDIHVD<br>IMDYIKPASCLTTEFREKWTGTFDWE<br>NTIAVNTDKTDLRDYVEFVMKETNM<br>QLLEPYQDVDEEELQGLSPTDEDGG<br>YGYVSCSLYARTVFGEDALANVSIER<br>DAQGRISGVVIRSNTO SIAYGIGEK<br>LSLLDKGGIK | None          | None          | None    | None    | None          | None                  | None            |                 |                      |

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