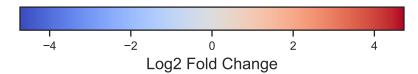
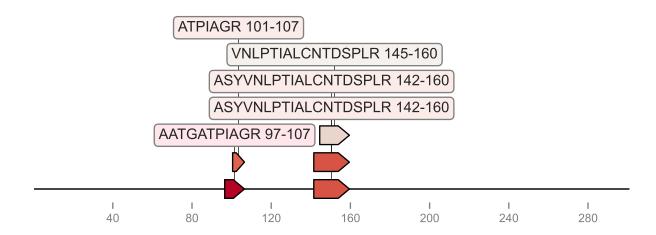


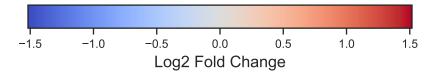


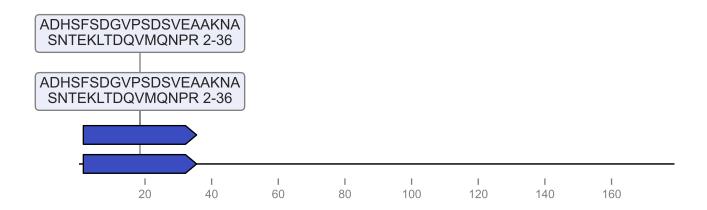
## tein SA {ECO:0000256|HAMAP-Rule:MF\_03016}

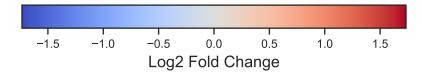


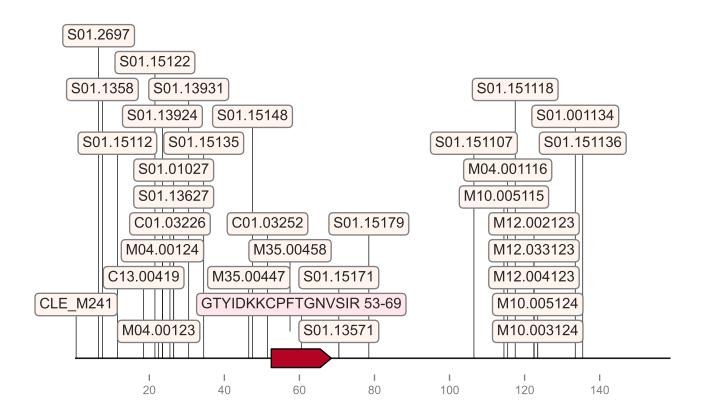


# B|Ensembl:ENSP00000405912.1, ECO:0000313|Ensembl:ENSP00000488479.1}



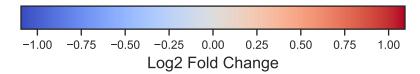


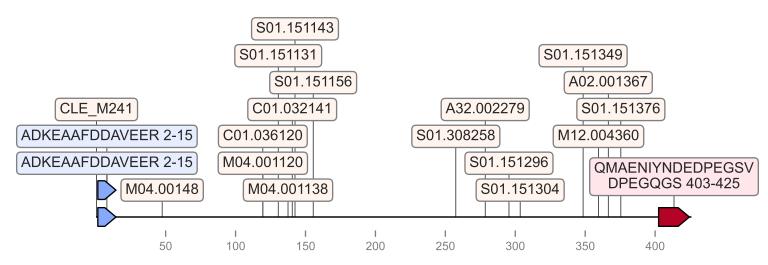


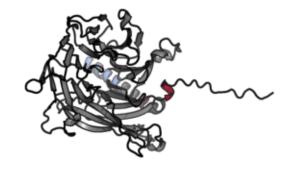




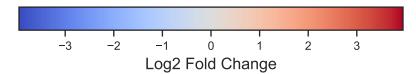
#### stone-binding protein RBBP4

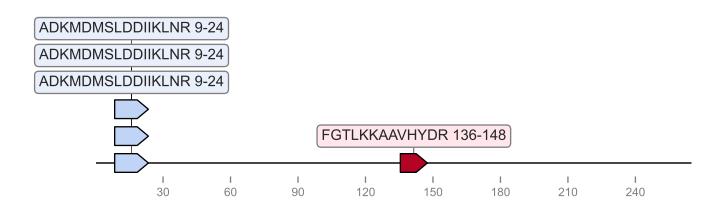




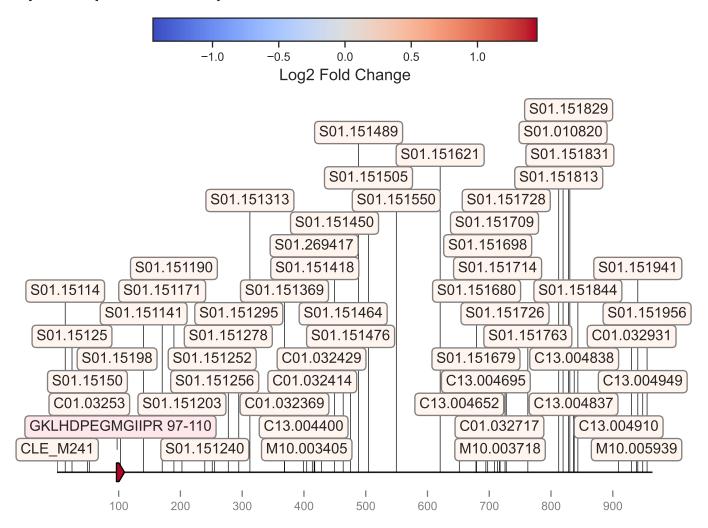


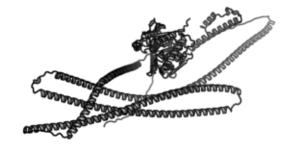
# r {ECO:0000313|Ensembl:ENSP00000421592.2}



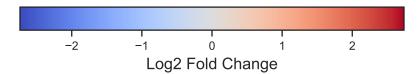


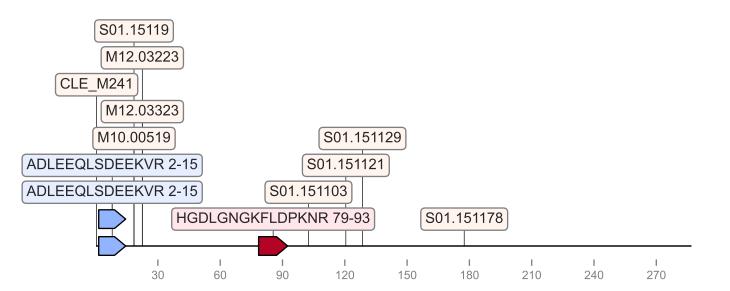
#### n-1 heavy chain {ECO:0000305}

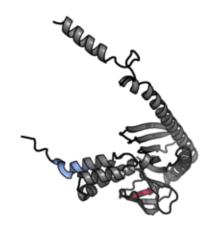


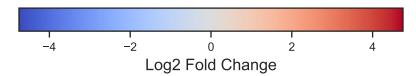


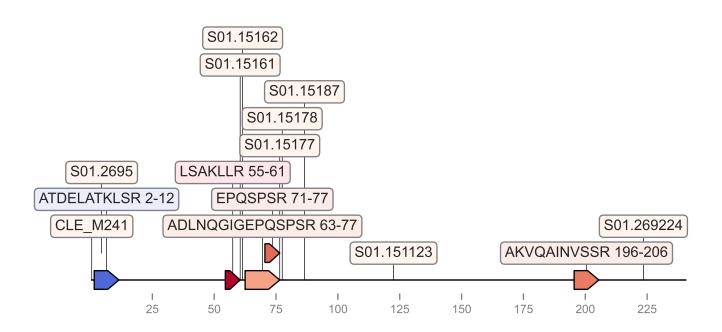
### n-capping protein subunit alpha-2





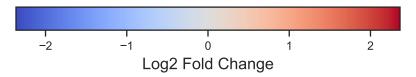


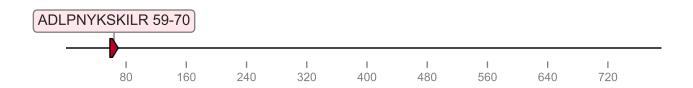






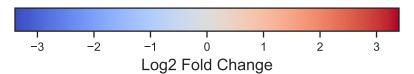
# ar cap-binding protein subunit 1

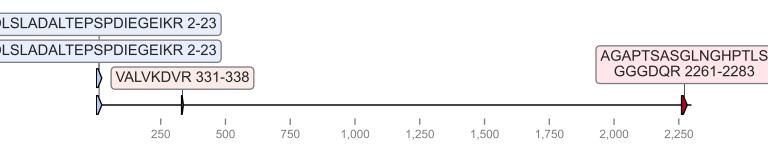




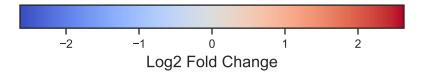


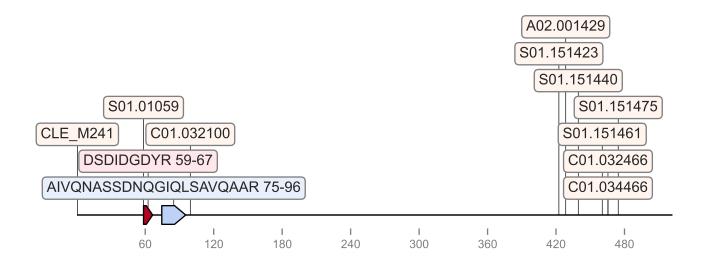
d protein {ECO:0000256|RuleBase:RU000686}





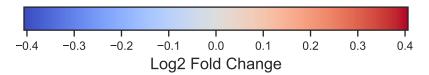
## -Importin subunit alpha-3

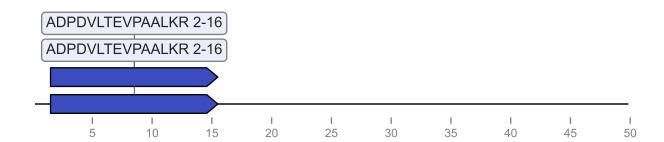




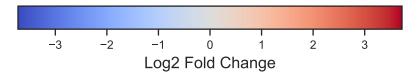


# subunit 1 {ECO:0000313|Ensembl:ENSP00000417342.1}



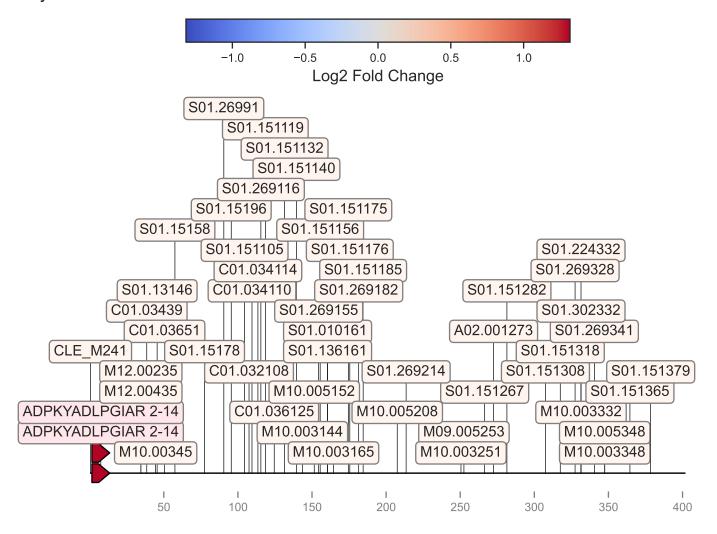


# up) {ECO:0000313|Ensembl:ENSP00000484849.2}

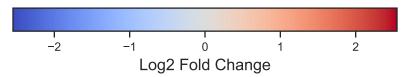


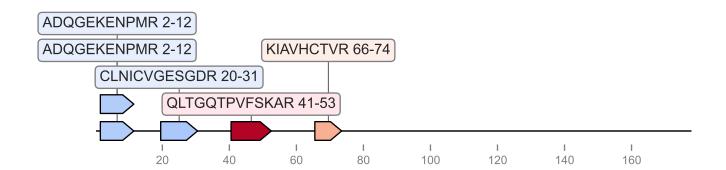
SIGNAI								
	<u> </u>	ı						
6	12	18	24	30	36	42	48	54

#### N2-Dynactin subunit 2

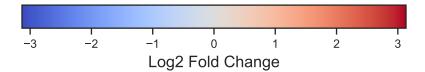


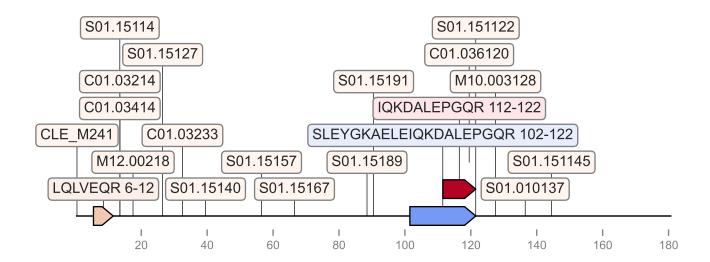




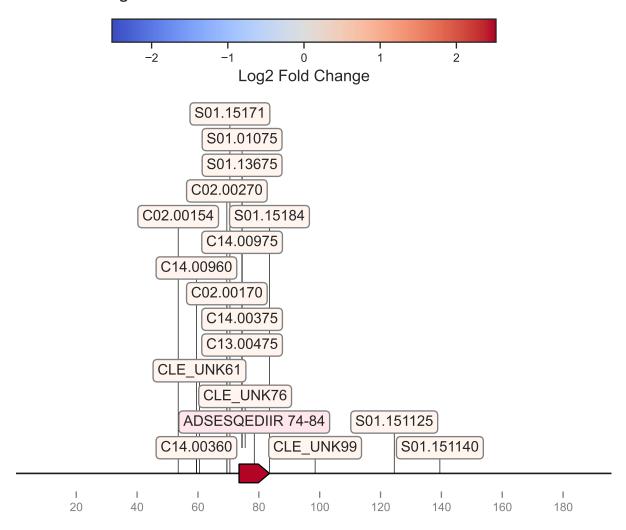


### phoribosyltransferase {ECO:0000305}

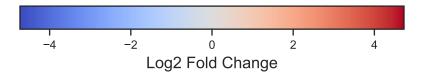


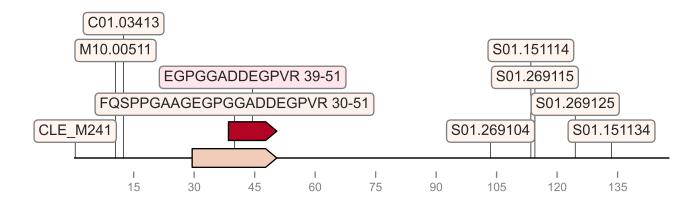


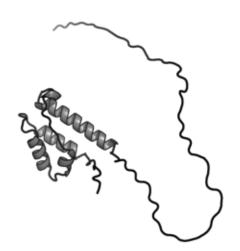




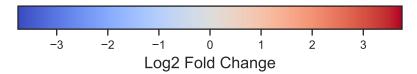


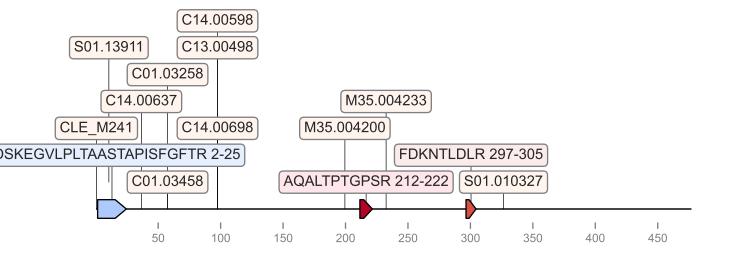


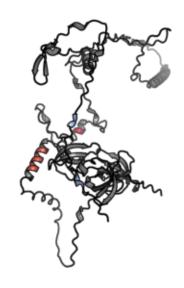


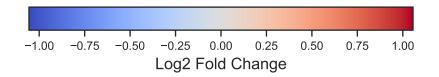


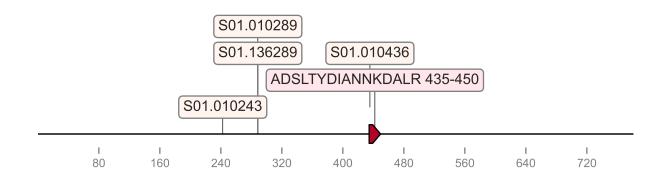
### nain and KOW motifs-containing protein

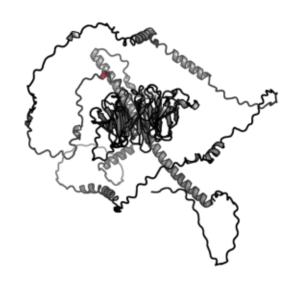




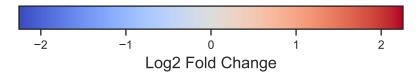


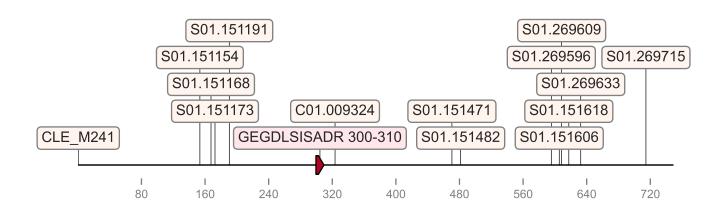






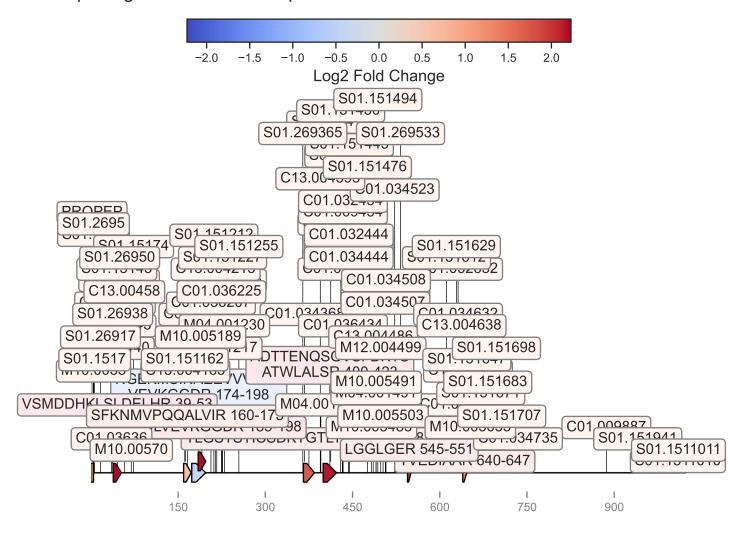
### eine--tRNA ligase, cytoplasmic





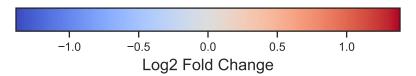


#### sium-transporting ATPase subunit alpha-1



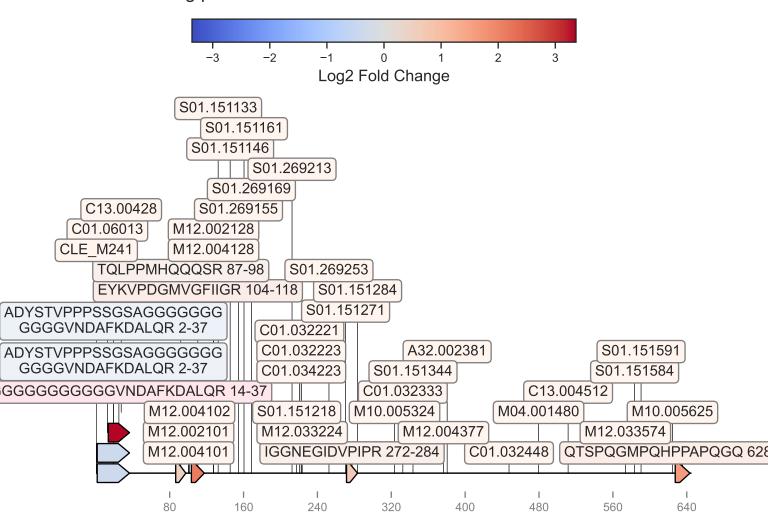


# ex subunit 1 {ECO:0000313|Ensembl:ENSP00000428697.1}



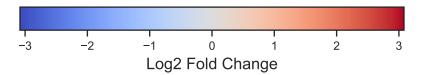


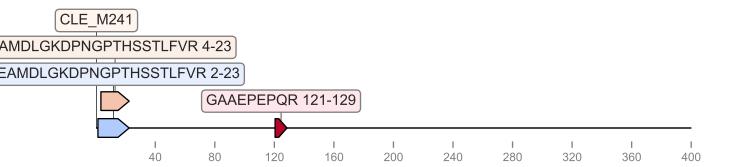
#### stream element-binding protein 1





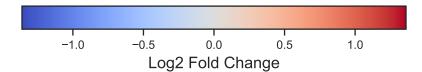
## peat-binding factor 2-interacting protein 1

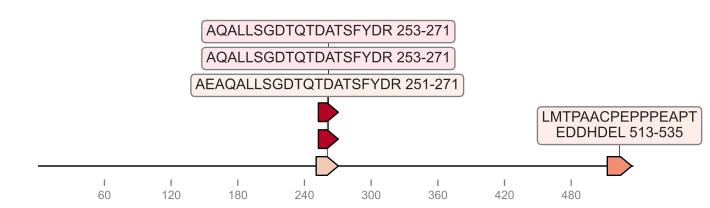




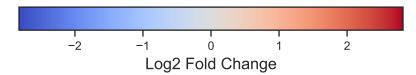


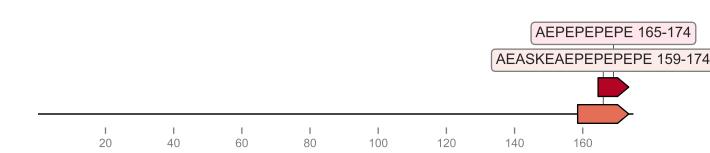
## init beta {ECO:0000256|ARBA:ARBA00022387}



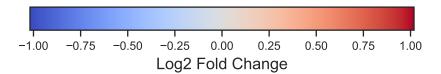


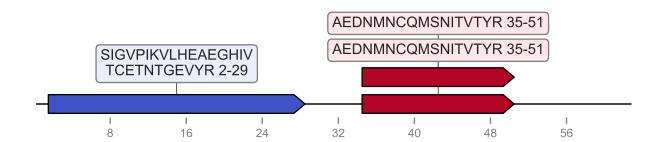
# ECO:0000313|Ensembl:ENSP00000457732.1}

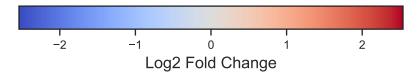


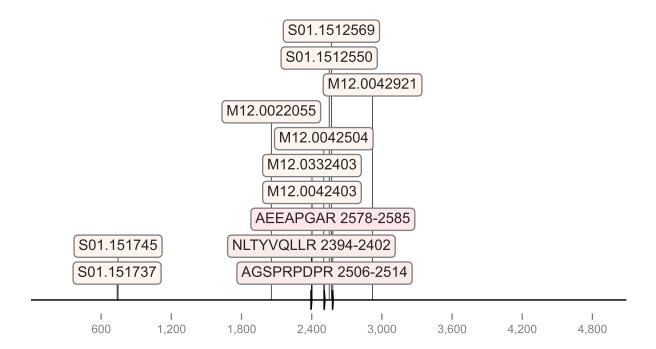


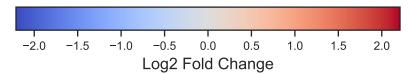
## D:0000256|ARBA:ARBA00020160, ECO:0000256|RuleBase:RU365050}

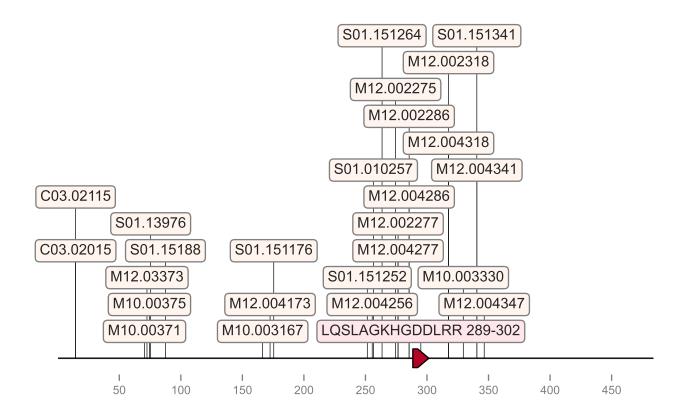


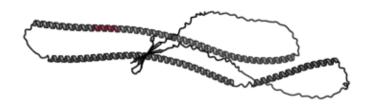




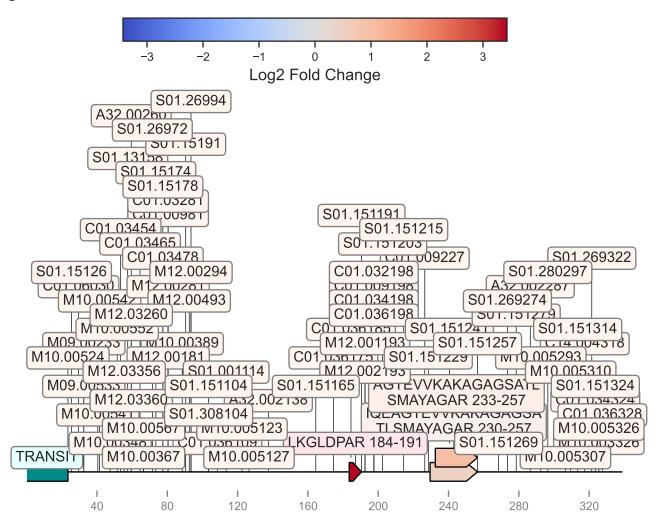


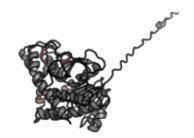




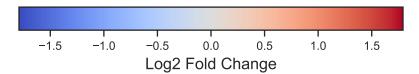


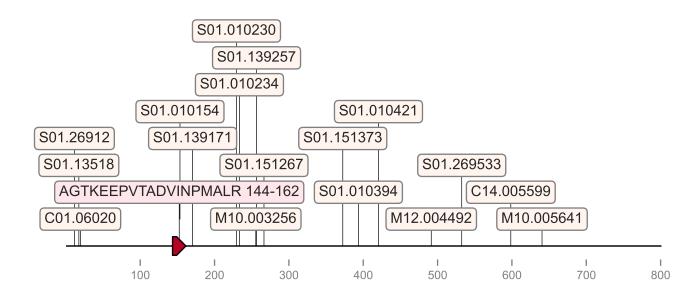
#### e dehydrogenase, mitochondrial





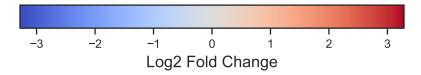
### J5 tri-snRNP-associated protein 1

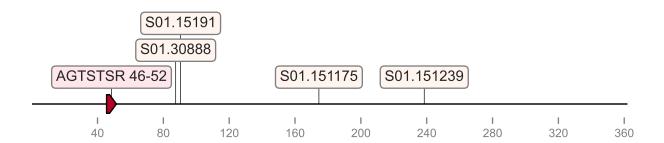






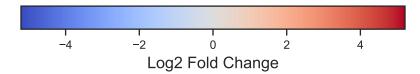
# import receptor subunit TOM40 homolog

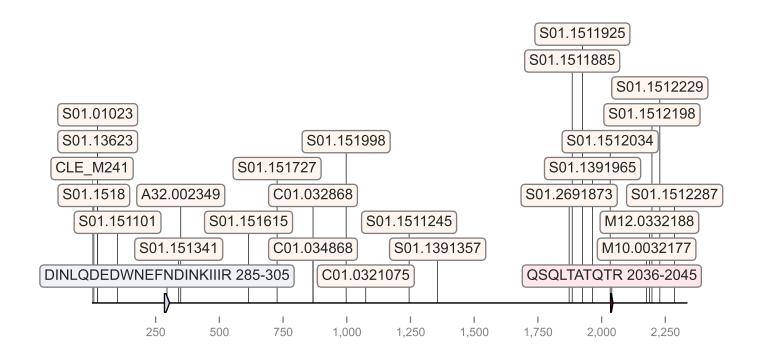


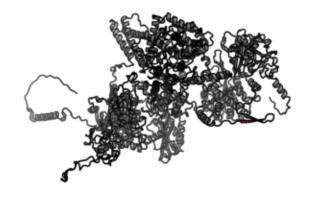




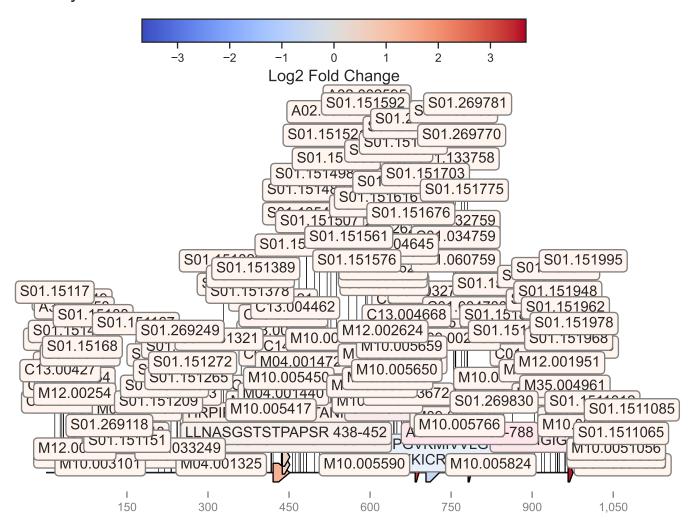
### RNA-processing-splicing factor 8





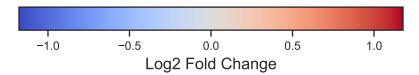


#### Y-ATP-citrate synthase



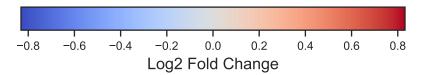


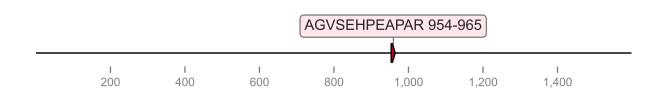
# alpha subcomplex subunit 5 {ECO:0000256|ARBA:ARBA00016385}

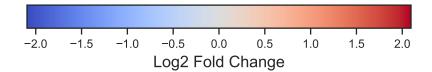


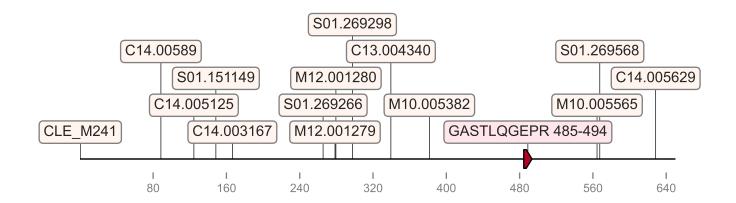


hange factor 5 {ECO:0000313|Ensembl:ENSP00000494816.1}



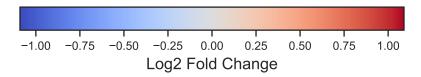


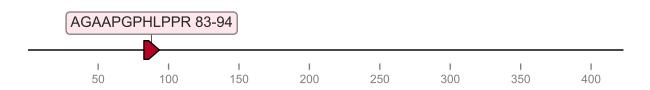






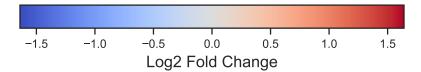
## uitin-conjugating enzyme E2 Q1

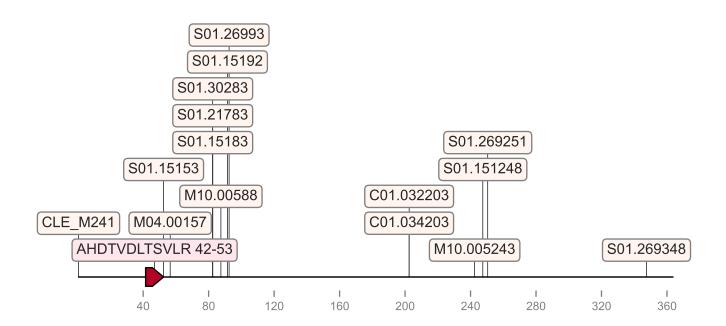


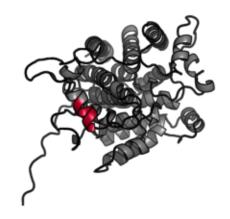




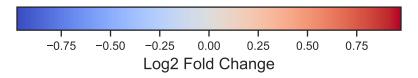
### osylhydrolase ARH3 {ECO:0000305}

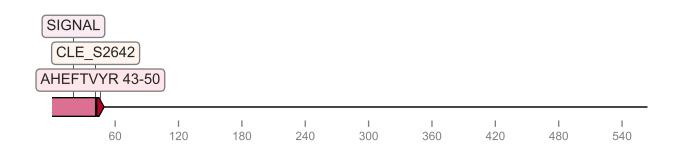


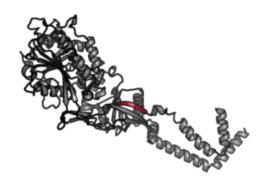




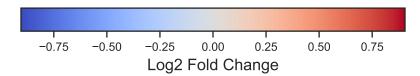
## plex subunit NCLN {ECO:0000305}

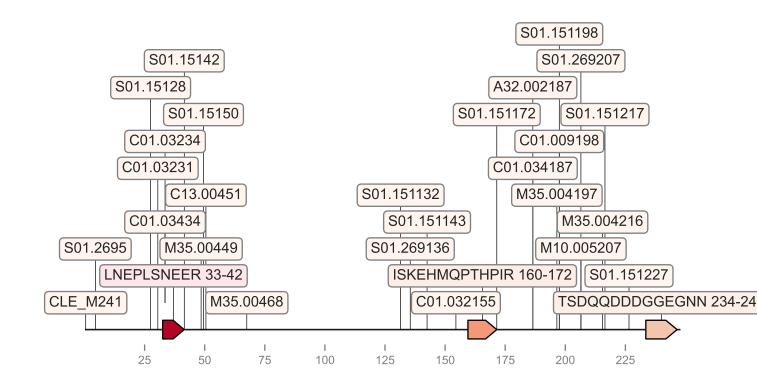






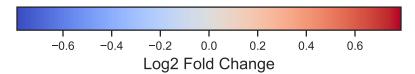
#### G-14-3-3 protein gamma

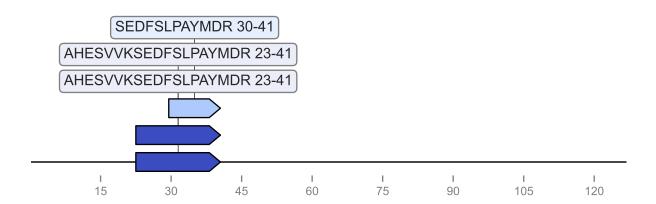




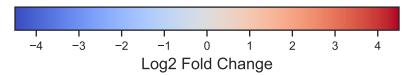


## e subunit 4 {ECO:0000256|RuleBase:RU367145}



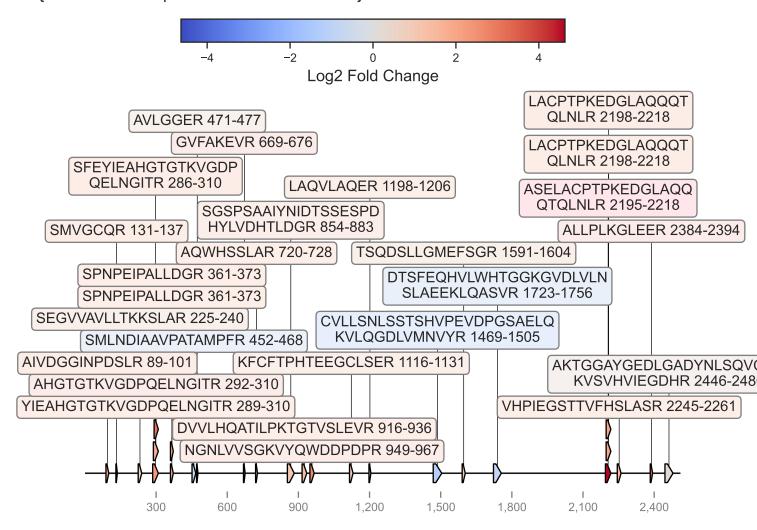


# ubunit Aalpha {ECO:0000313|Ensembl:ENSP00000391905.3}

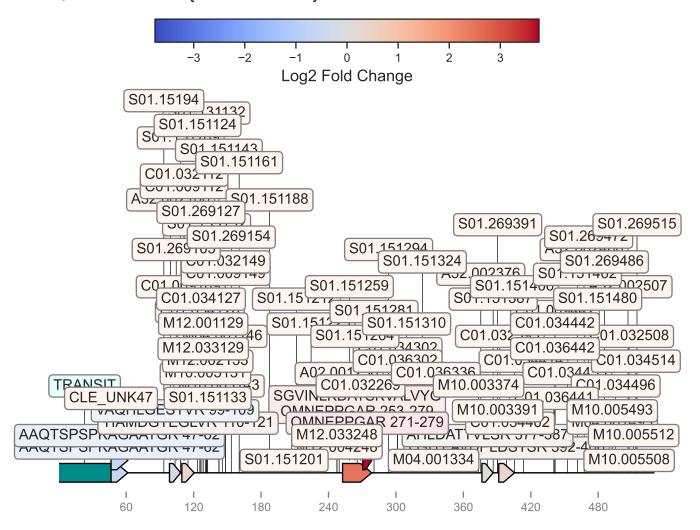


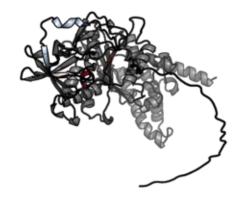


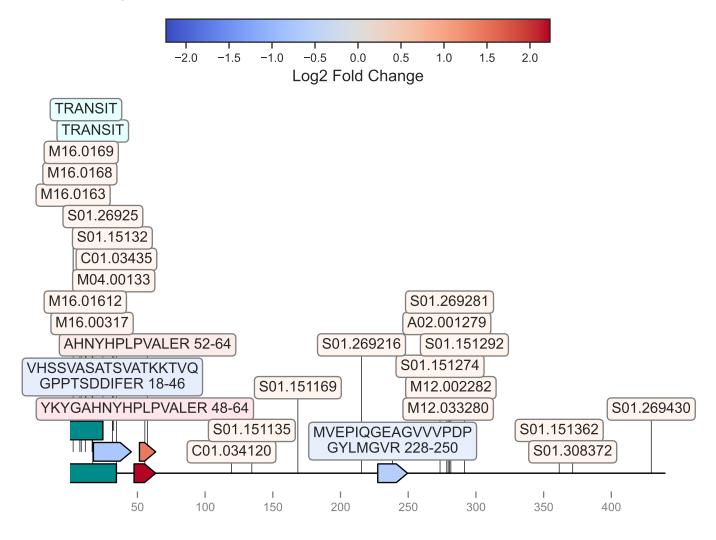
thase {ECO:0000256|ARBA:ARBA00018769}



#### subunit beta, mitochondrial {ECO:0000305}

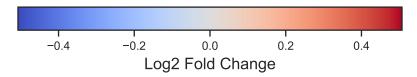


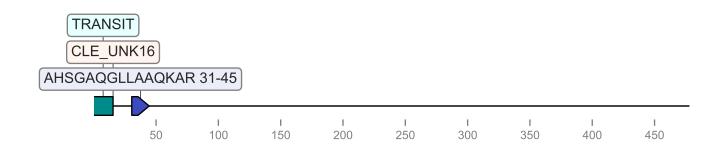






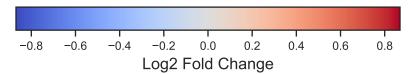
## sine--tRNA ligase, mitochondrial

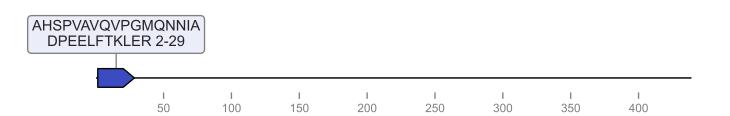




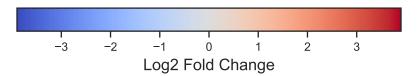


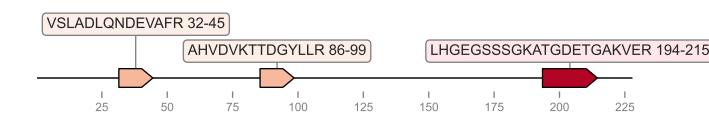
## 26 {ECO:0000313|Ensembl:ENSP00000418753.1}



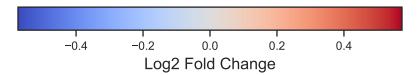


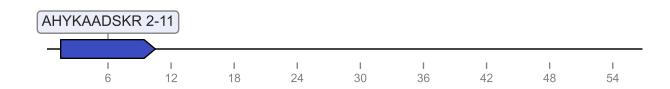
n S3a {ECO:0000256|HAMAP-Rule:MF\_03122}



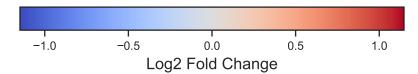


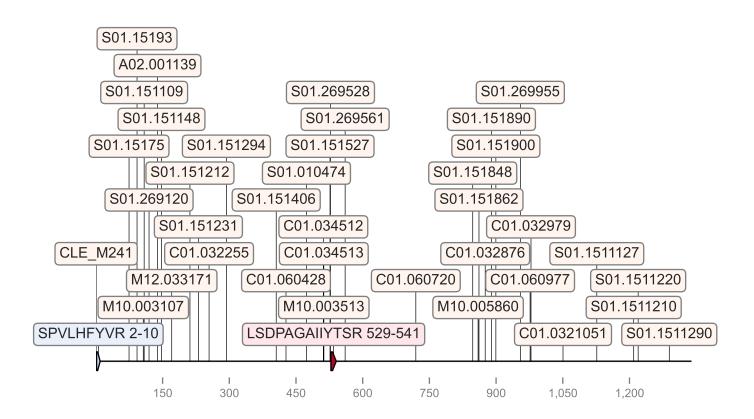
## ein {ECO:0000313|Ensembl:ENSP00000485717.1}

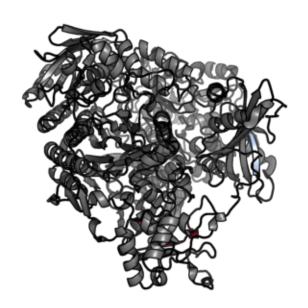


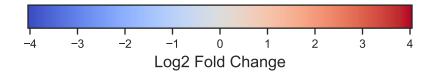


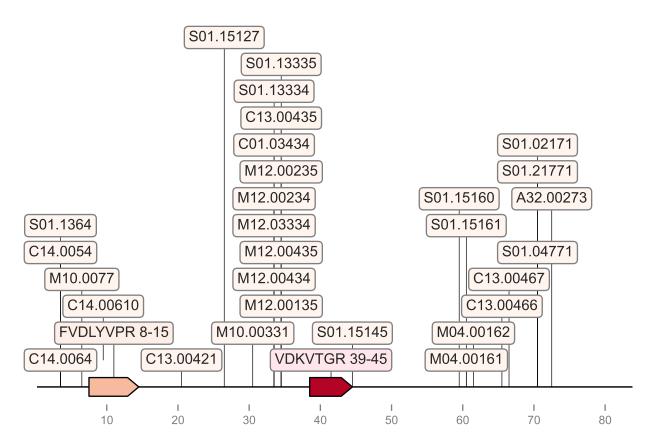
#### bosylformylglycinamidine synthase













### Isoform 2 of Protein SET

