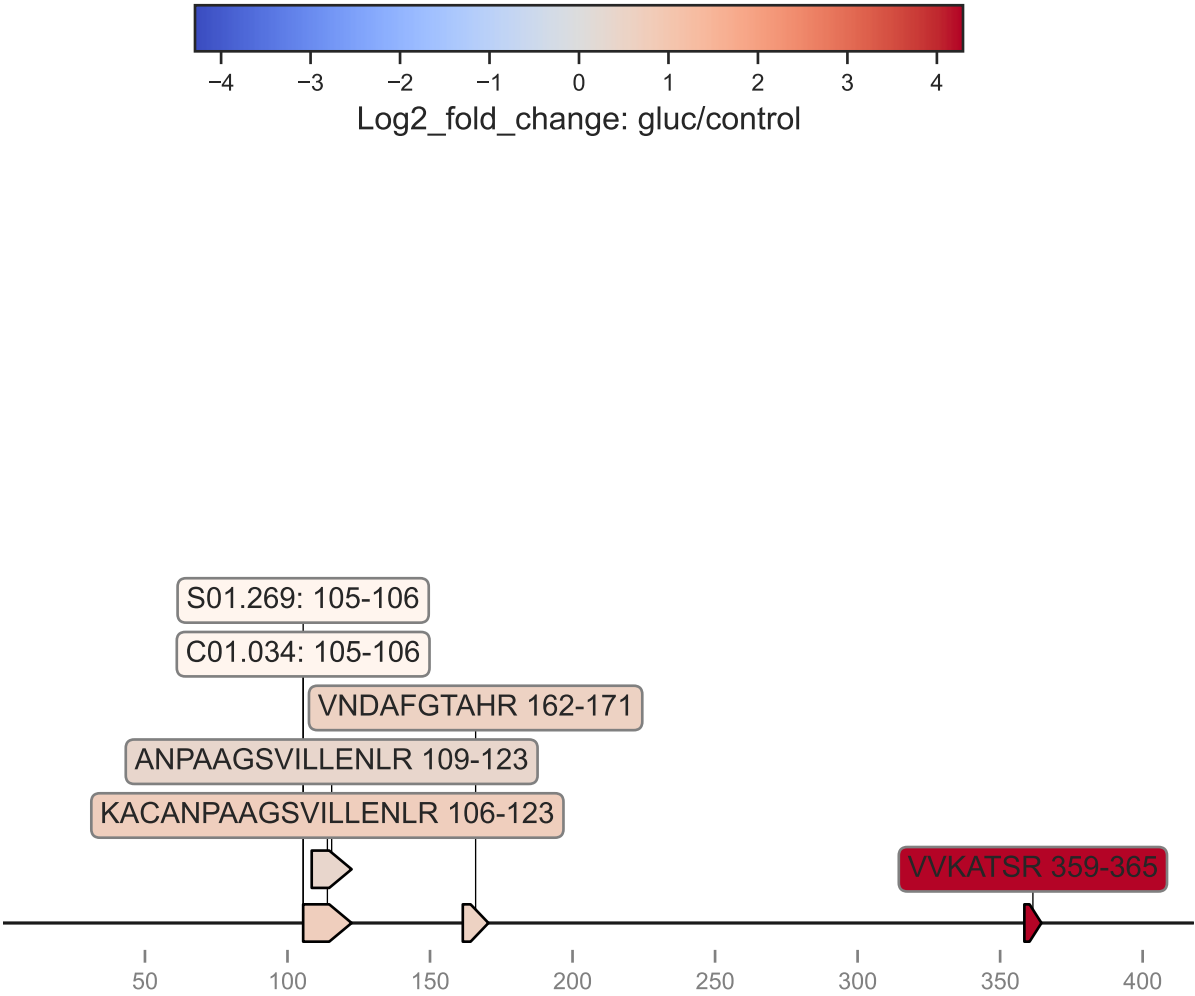
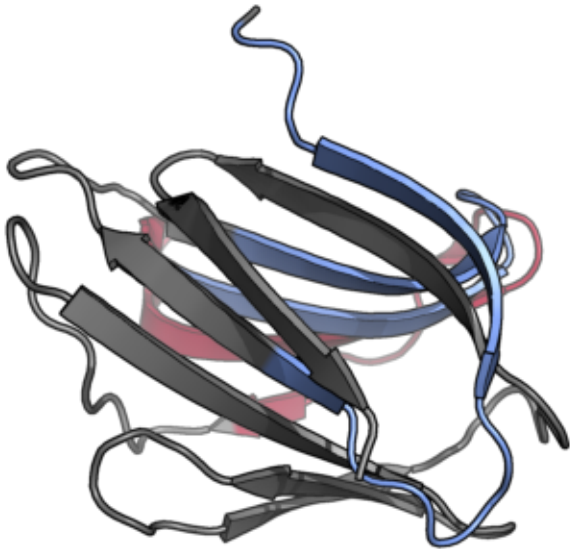
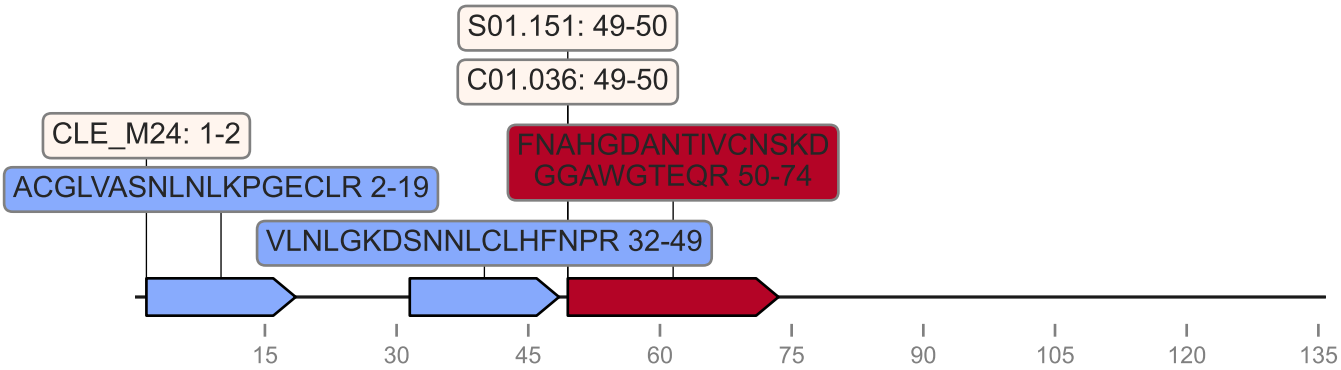
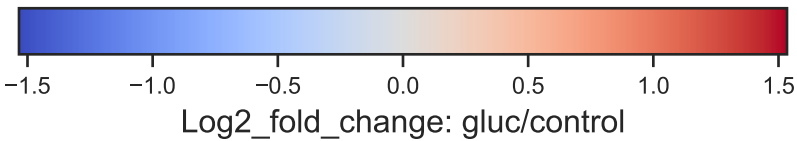


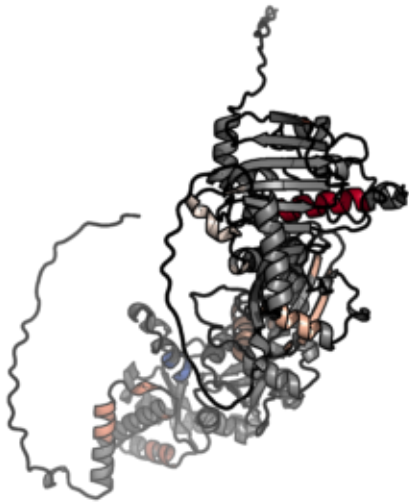
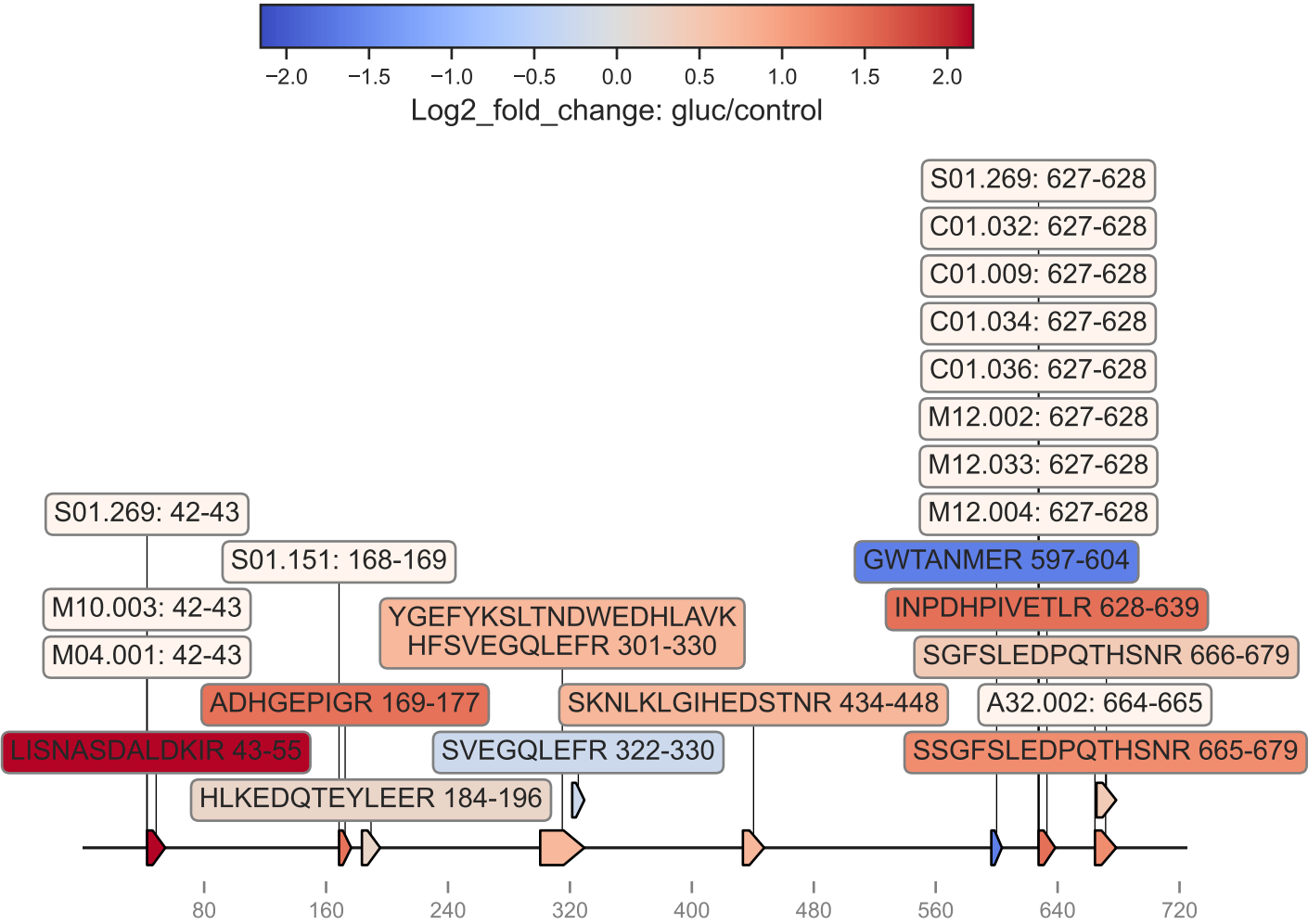
P00558-PGK1-Phosphoglycerate kinase 1



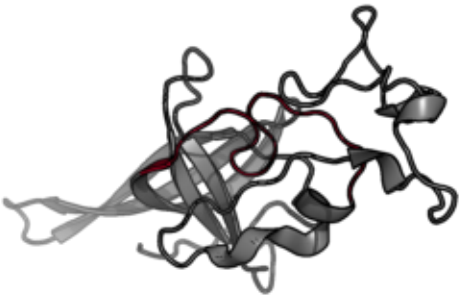
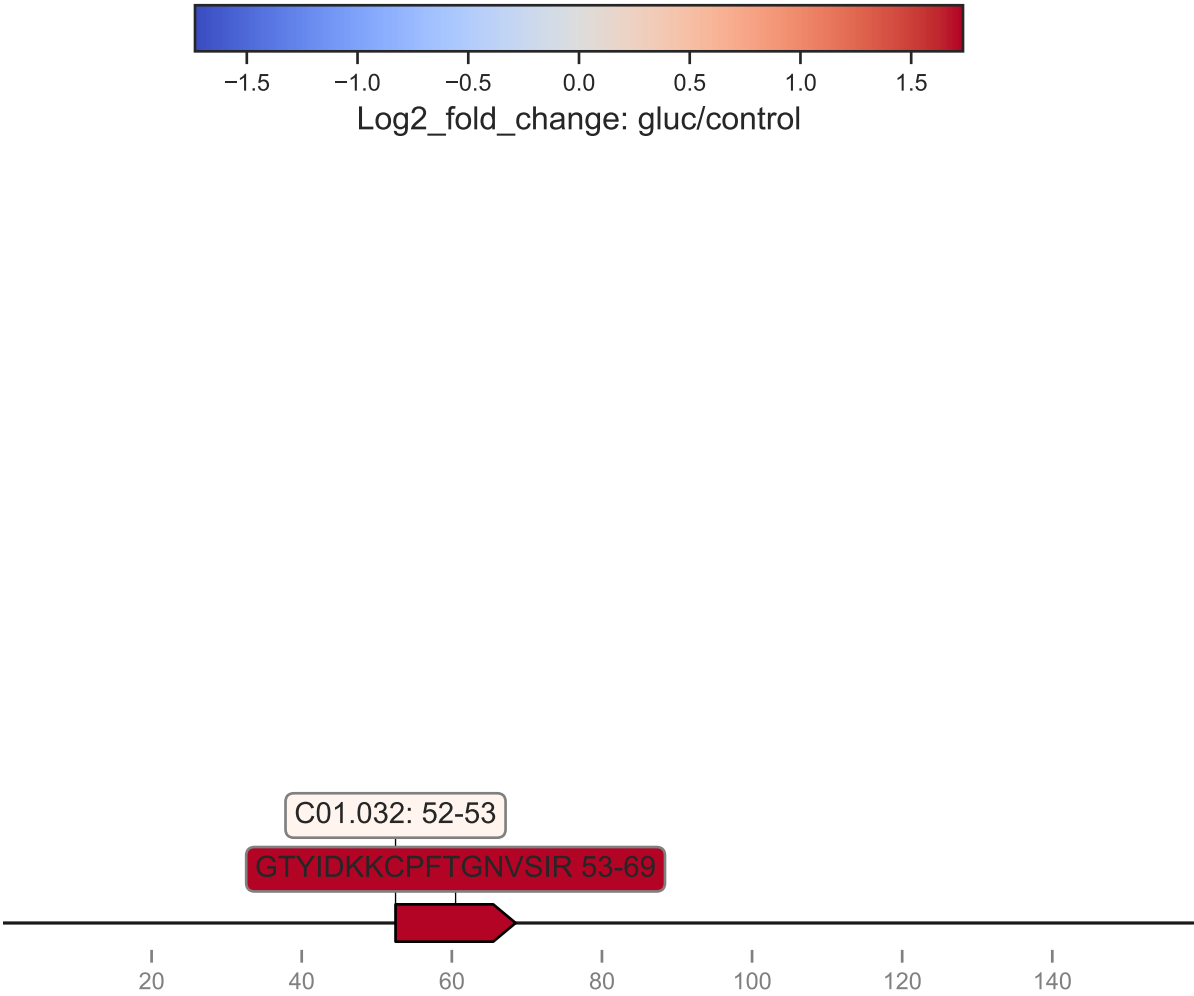
P09382-LGALS1-Galectin-1



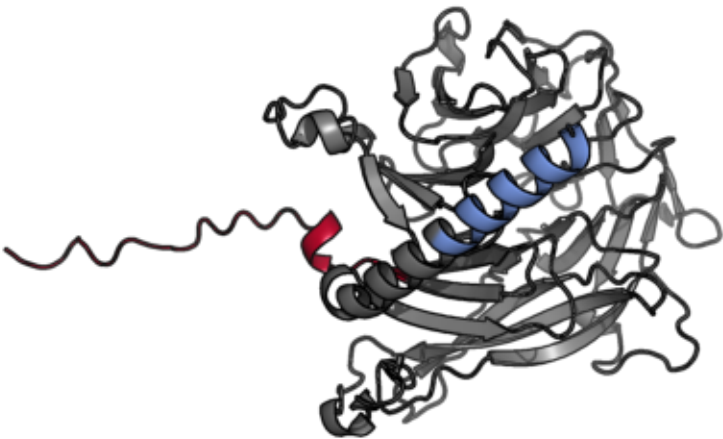
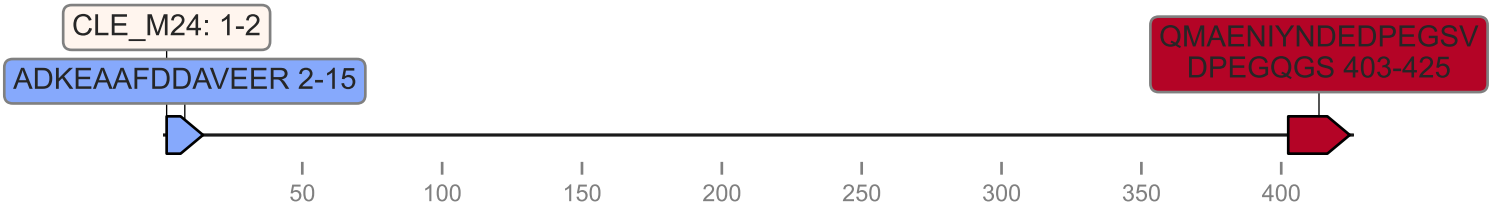
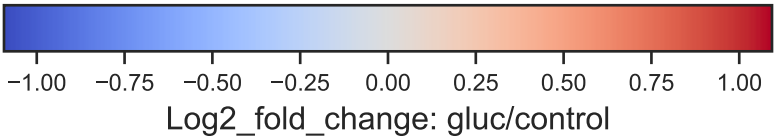
P08238-HSP90AB1-Heat shock protein HSP 90-beta



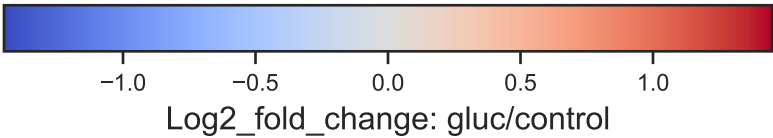
P62280-RPS11-40S ribosomal protein S11



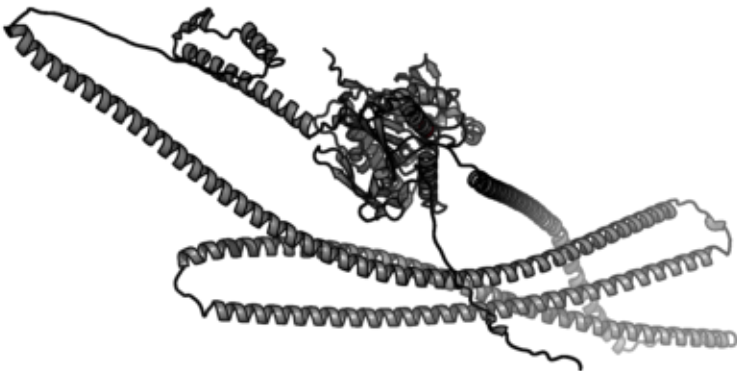
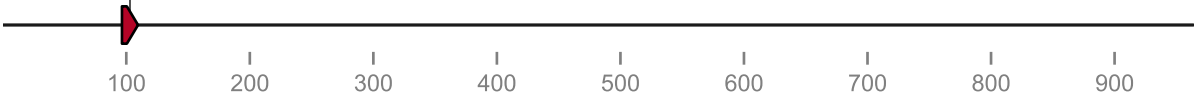
Q09028-RBBP4-Histone-binding protein RBBP4



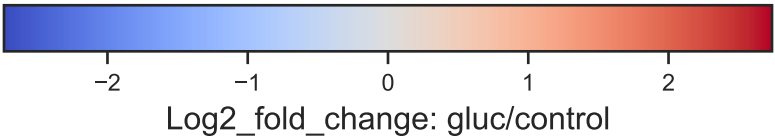
P33176-KIF5B-Kinesin-1 heavy chain {ECO:0000305}



GKLHDPEGMGIIPR 97-110



P47755-CAPZA2-F-actin-capping protein subunit alpha-2



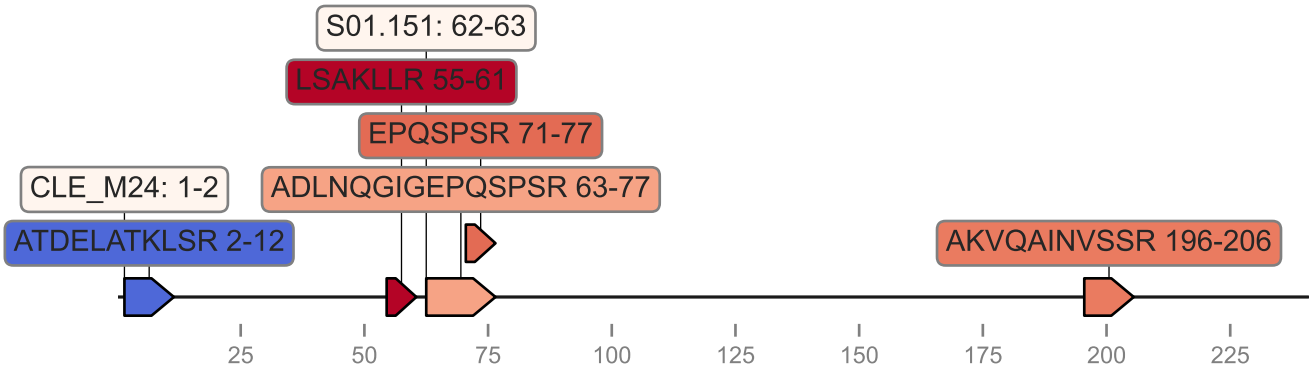
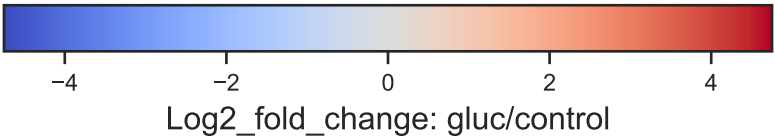
ADLEEQLSDEEKVR 2-15

CLE_M24: 1-2

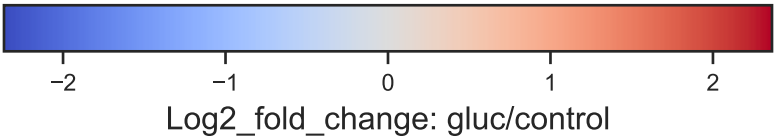
HGDLGNGKFLDPKNR 79-93



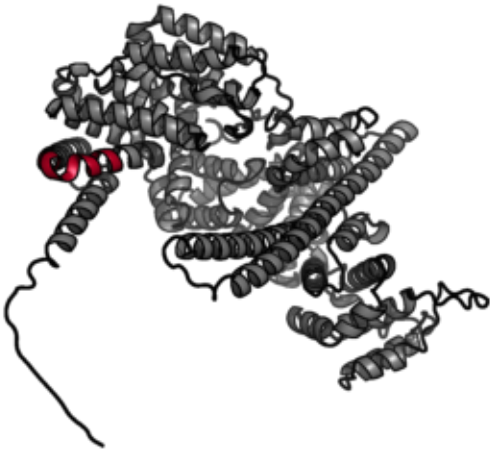
Q96C19-EFHD2-EF-hand domain-containing protein D2



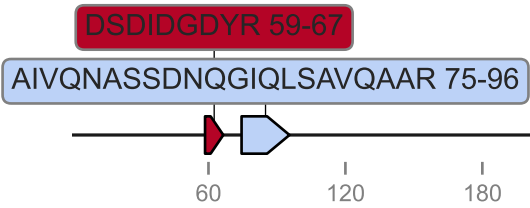
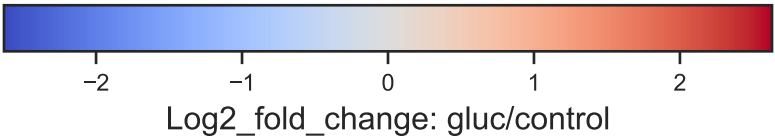
Q09161-NCBP1-Nuclear cap-binding protein subunit 1



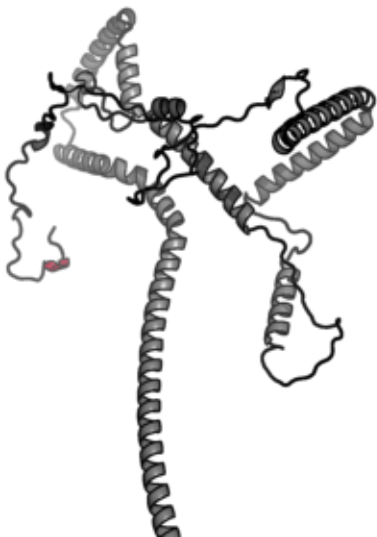
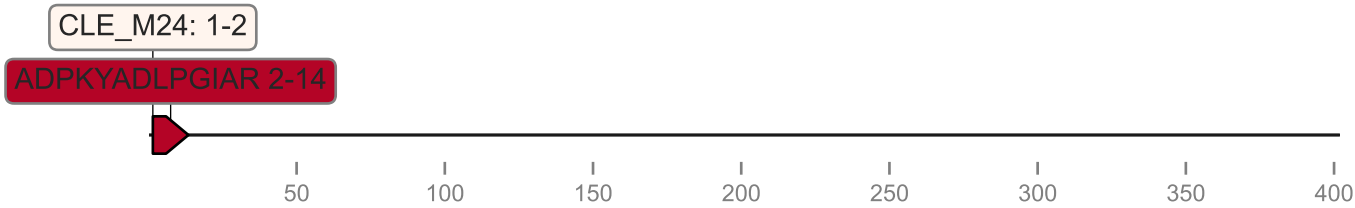
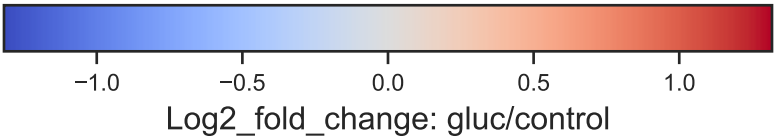
ADLPNYKSKILR 59-70



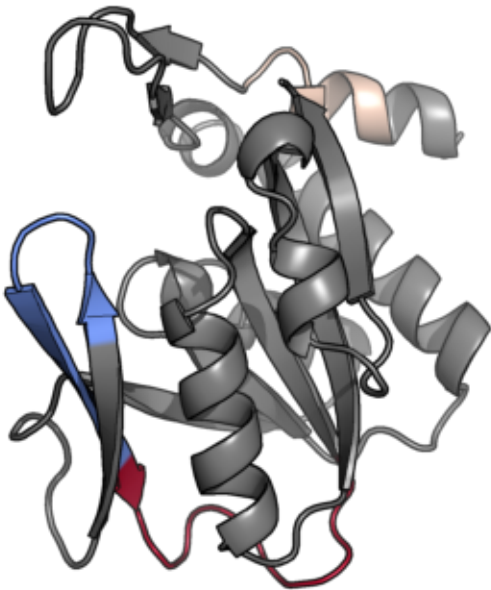
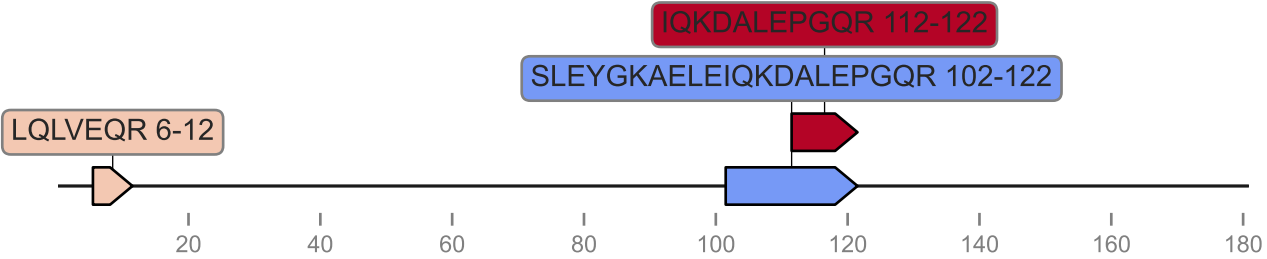
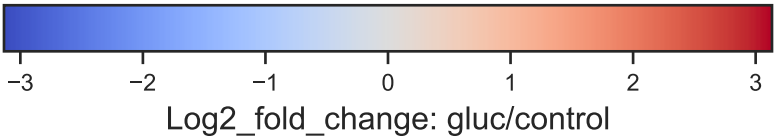
O00629-KPNA4-Importin subunit alpha-3



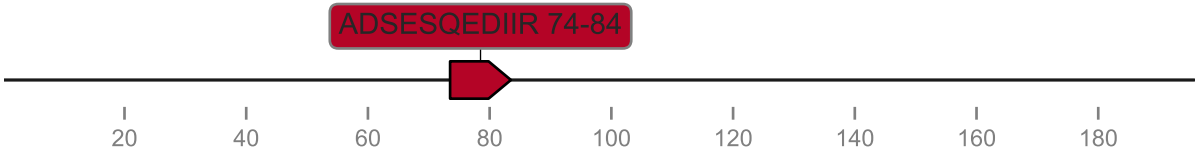
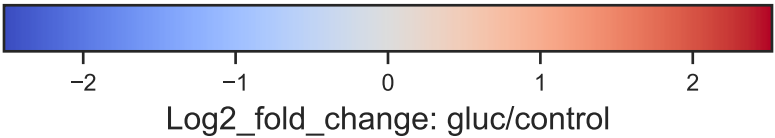
Q13561-DCTN2-Dynactin subunit 2



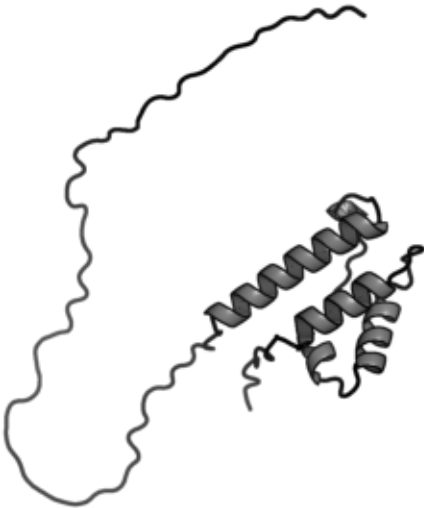
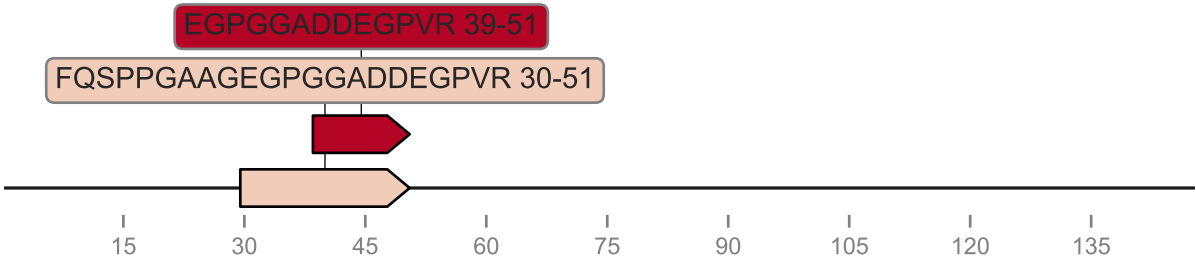
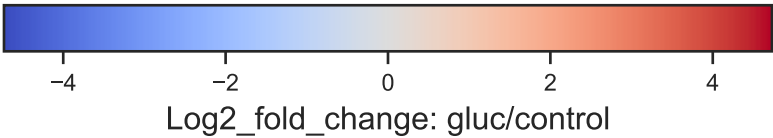
P07741-APRT-Adenine phosphoribosyltransferase {ECO:0000305}



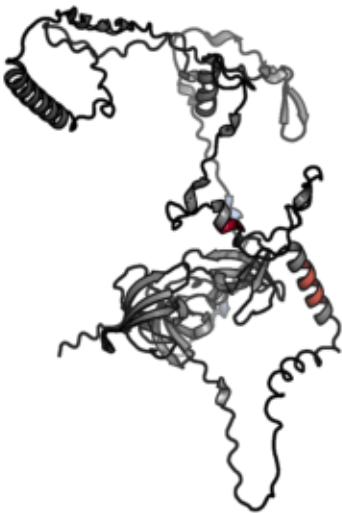
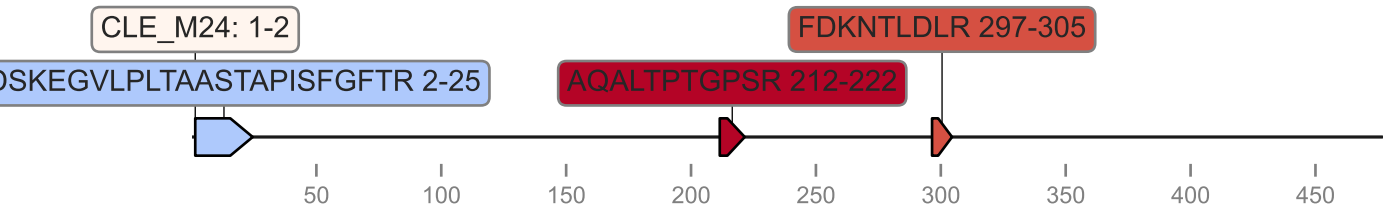
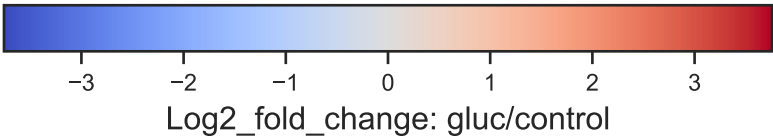
P55957-BID-BH3-interacting domain death agonist



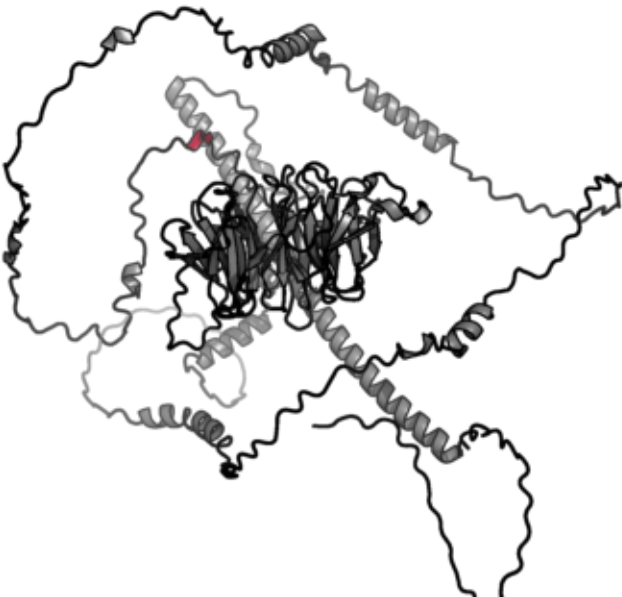
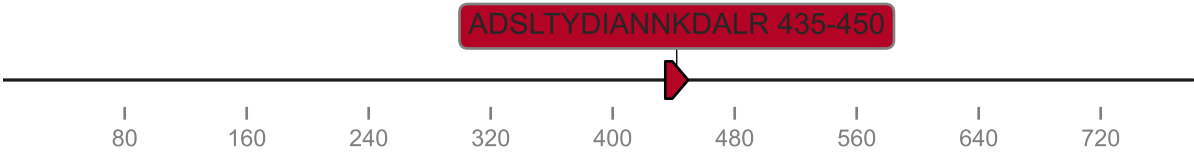
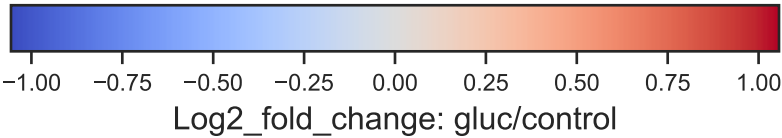
Q96C90-PPP1R14B-Protein phosphatase 1 regulatory subunit 14B



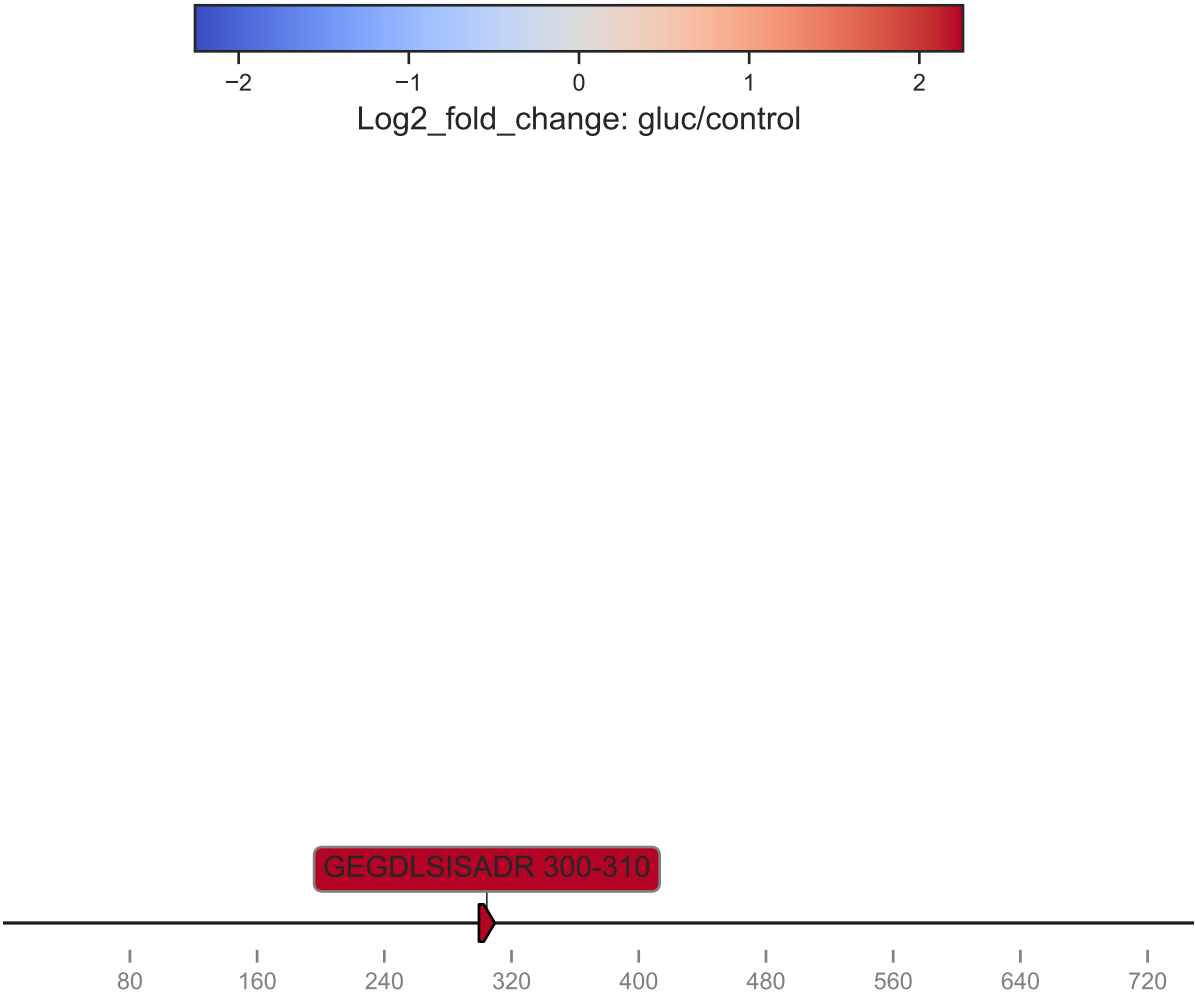
Q92917-GPKOW-G-patch domain and KOW motifs-containing protein



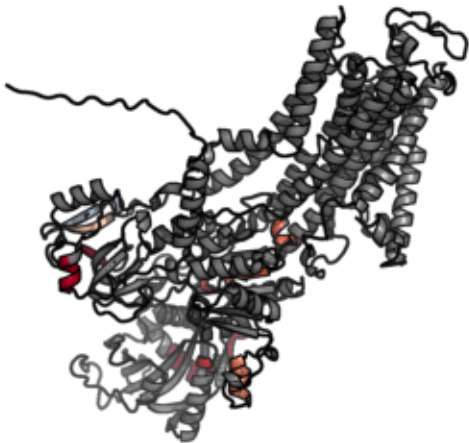
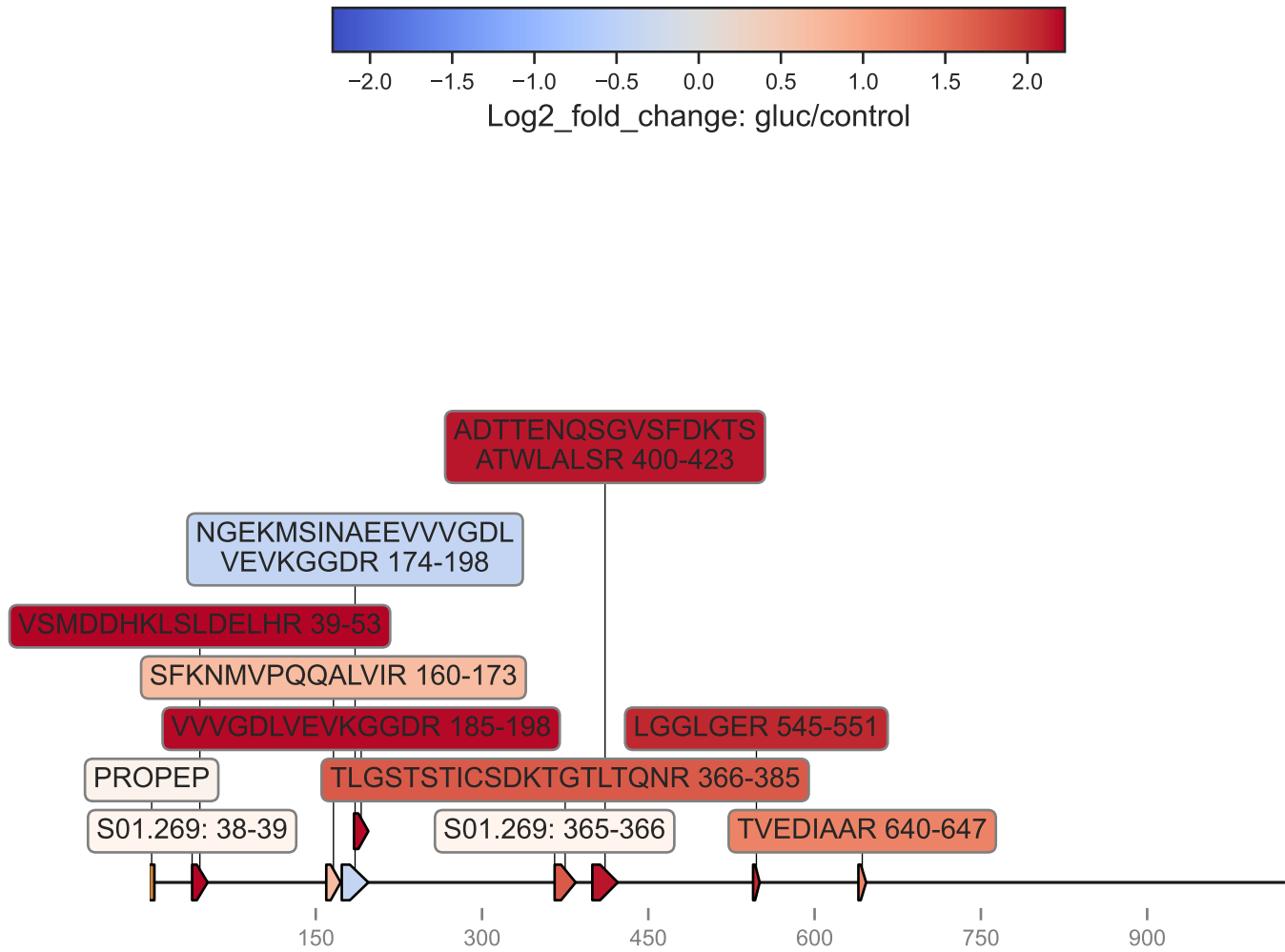
O43815-STRN-Striatin



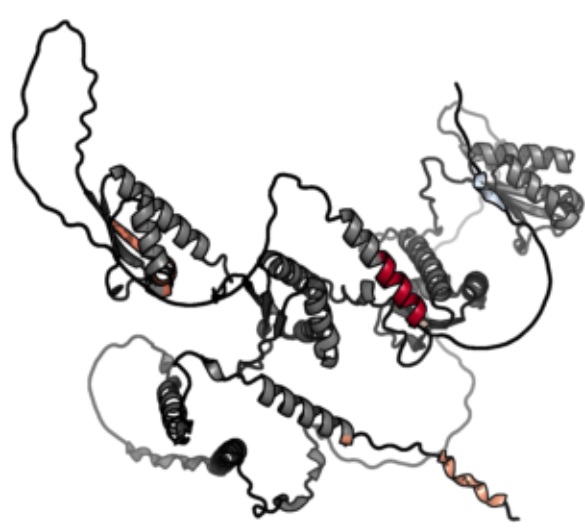
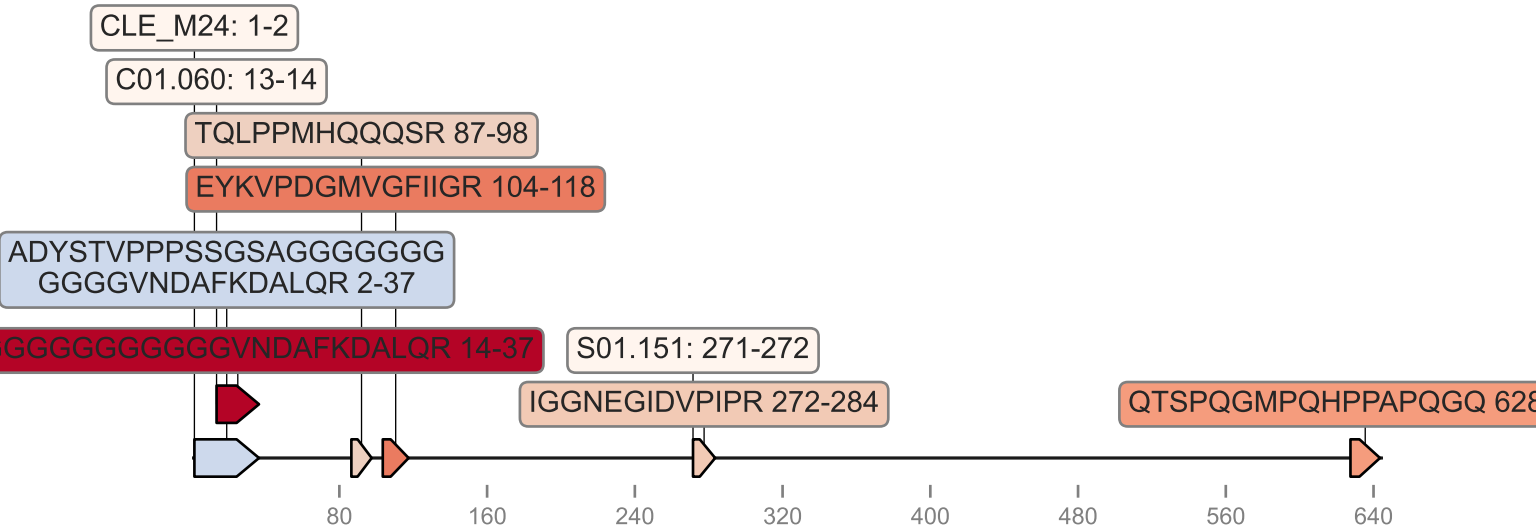
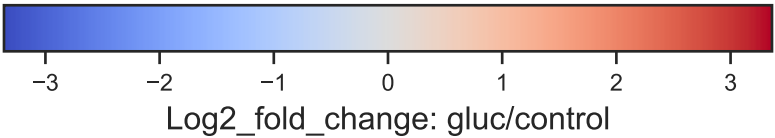
P49589-CARS1-Cysteine--tRNA ligase, cytoplasmic



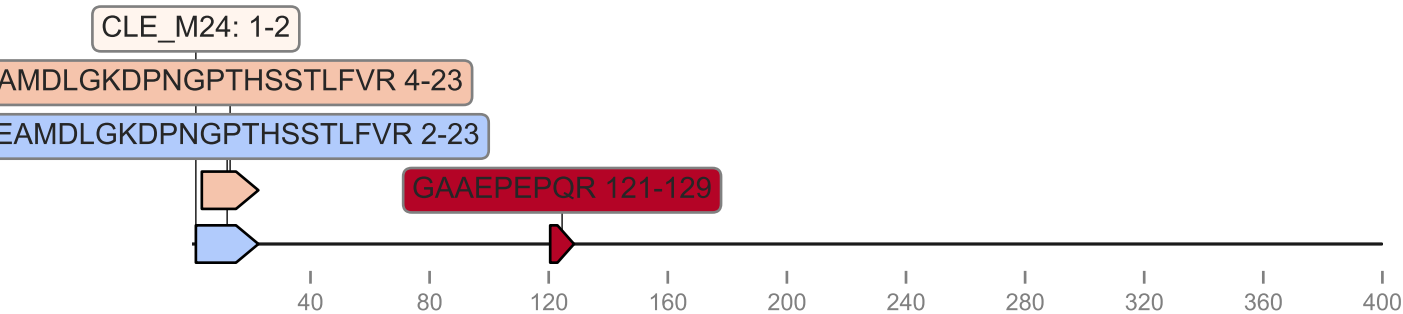
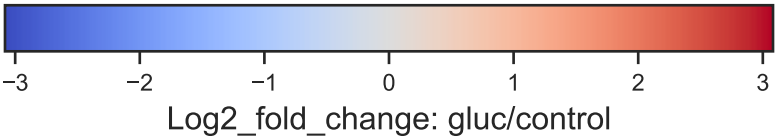
P05023-ATP1A1-Sodium/potassium-transporting ATPase subunit alpha-1



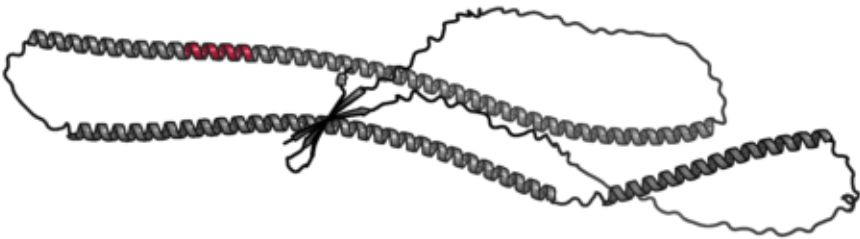
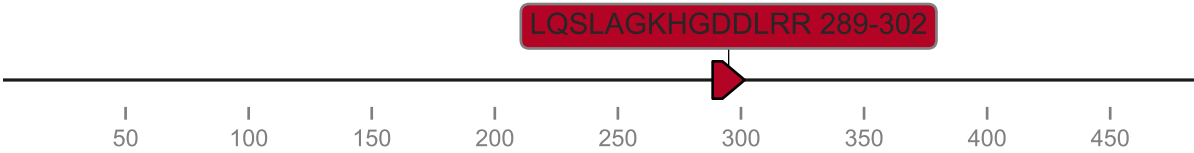
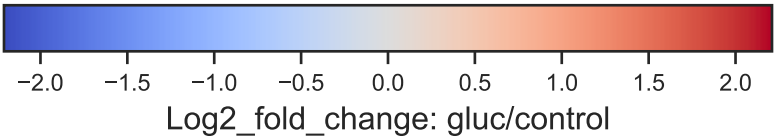
Q96AE4-FUBP1-Far upstream element-binding protein 1



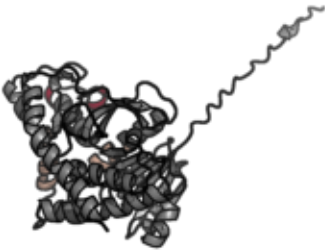
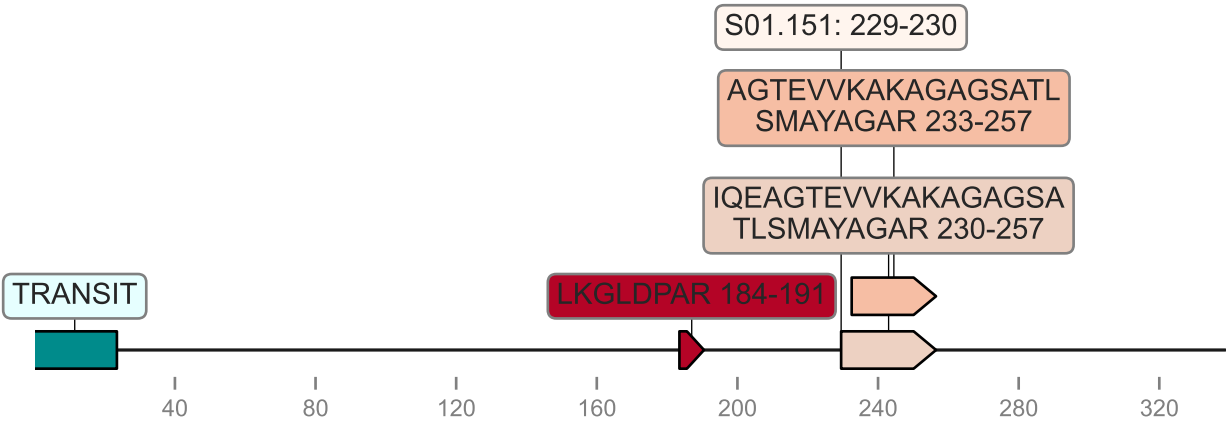
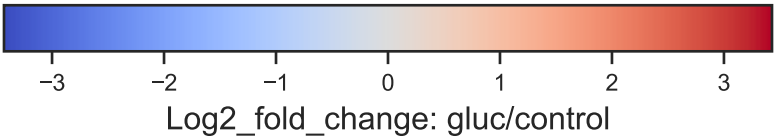
Q9NYB0-TERF2IP-Telomeric repeat-binding factor 2-interacting protein 1



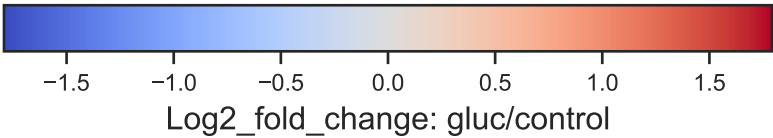
P05787-KRT8-Keratin, type II cytoskeletal 8



P40926-MDH2-Malate dehydrogenase, mitochondrial



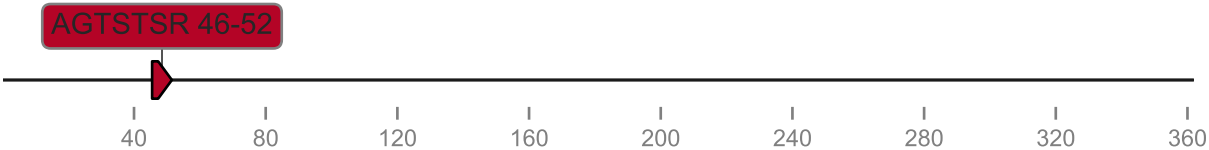
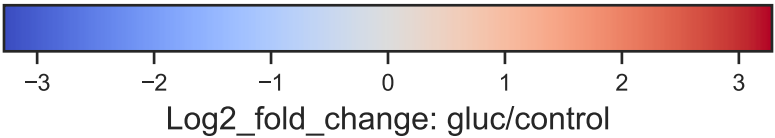
O43290-SART1-U4/U6.U5 tri-snRNP-associated protein 1



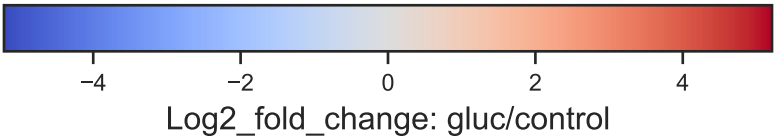
AGTKEEPVTADVNPALR 144-162



O96008-TOMM40-Mitochondrial import receptor subunit TOM40 homolog

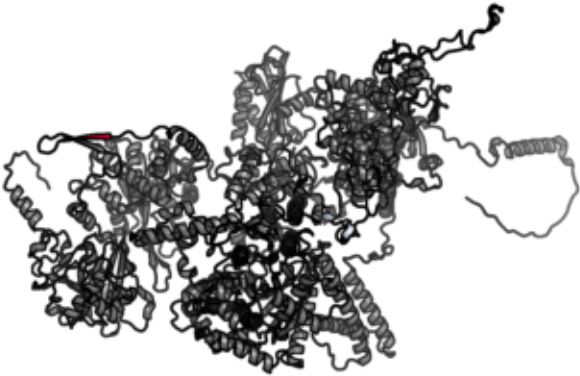


Q6P2Q9-PRPF8-Pre-mRNA-processing-splicing factor 8

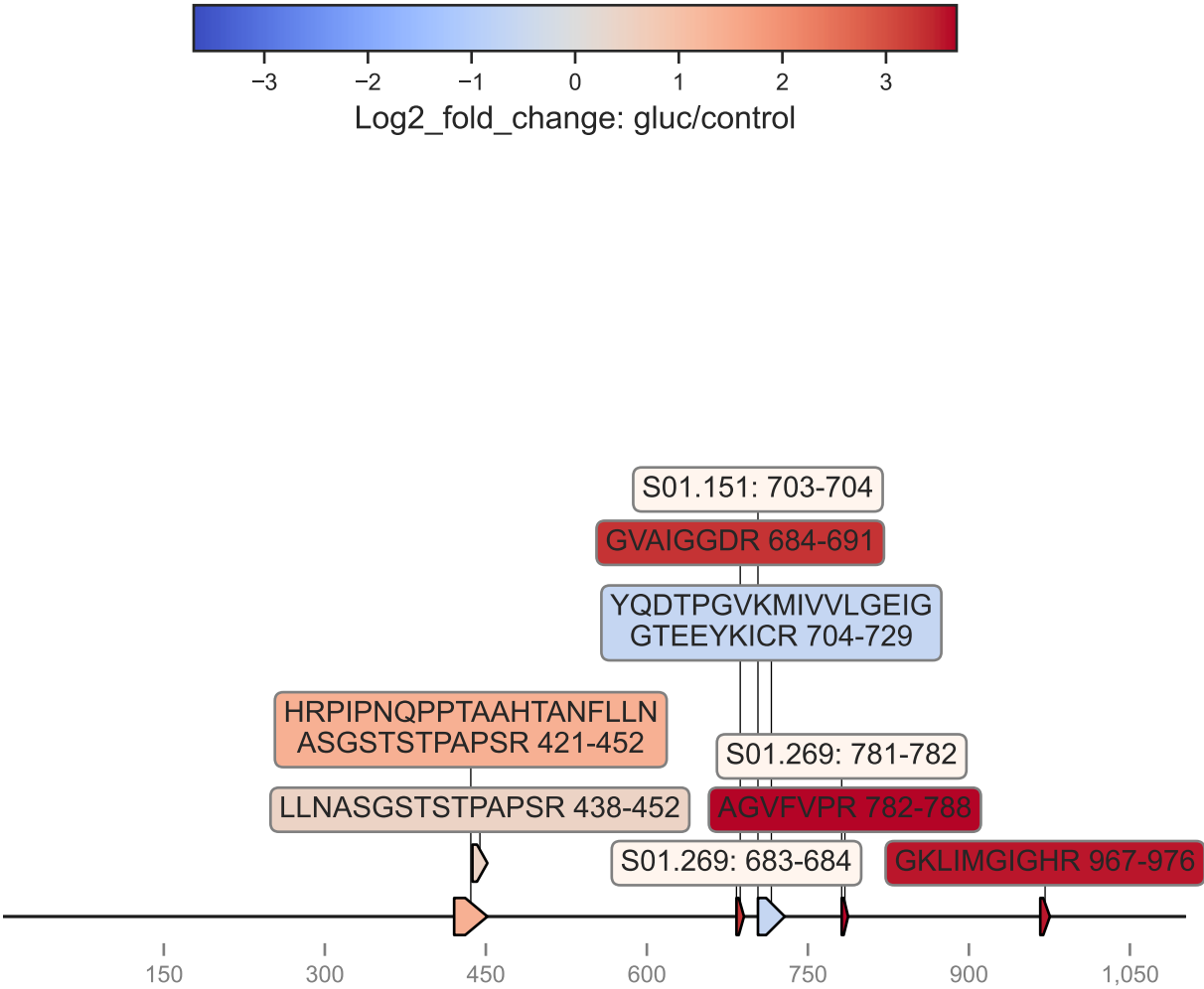


DINLQDEDWNEFNDINKIIR 285-305

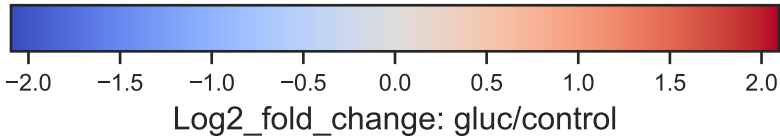
QSQLTATQTR 2036-2045



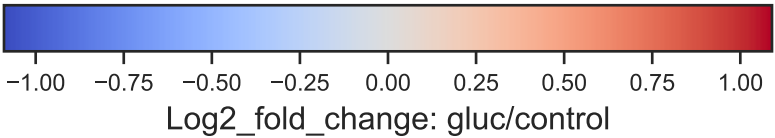
P53396-ACLY-ATP-citrate synthase



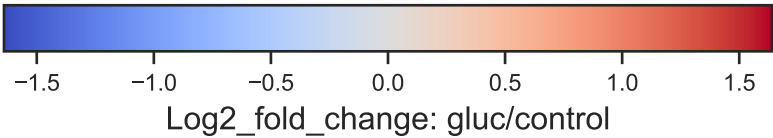
Q16643-DBN1-Drebrin



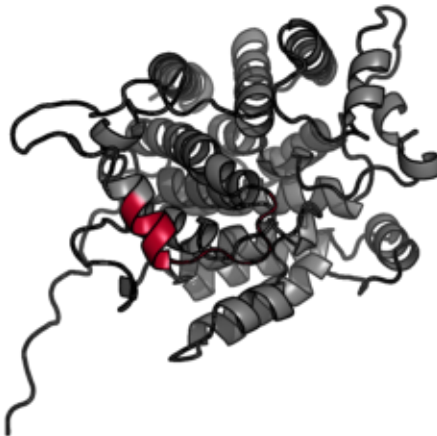
Q7Z7E8-UBE2Q1-Ubiquitin-conjugating enzyme E2 Q1



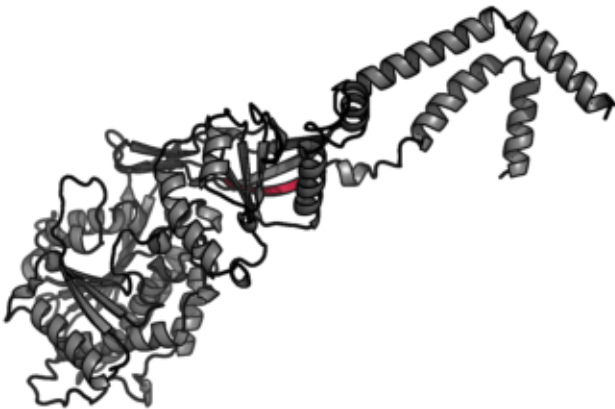
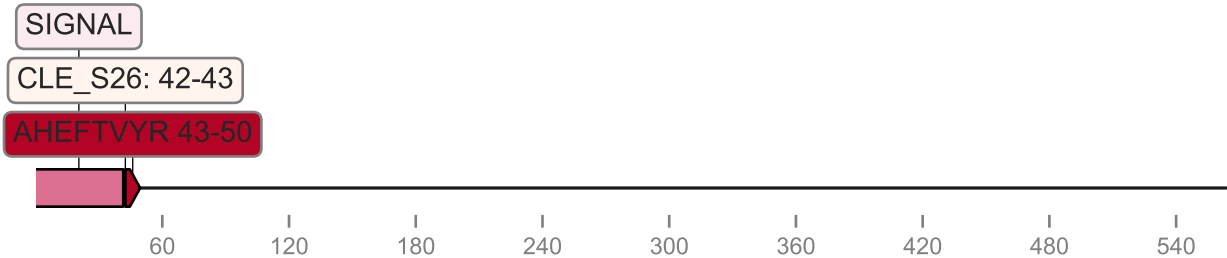
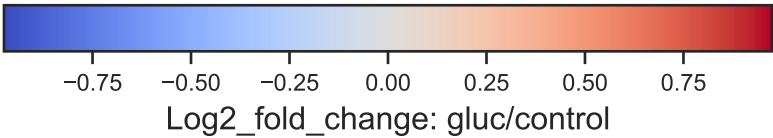
Q9NX46-ADPRS-ADP-ribosylhydrolase ARH3 {ECO:0000305}



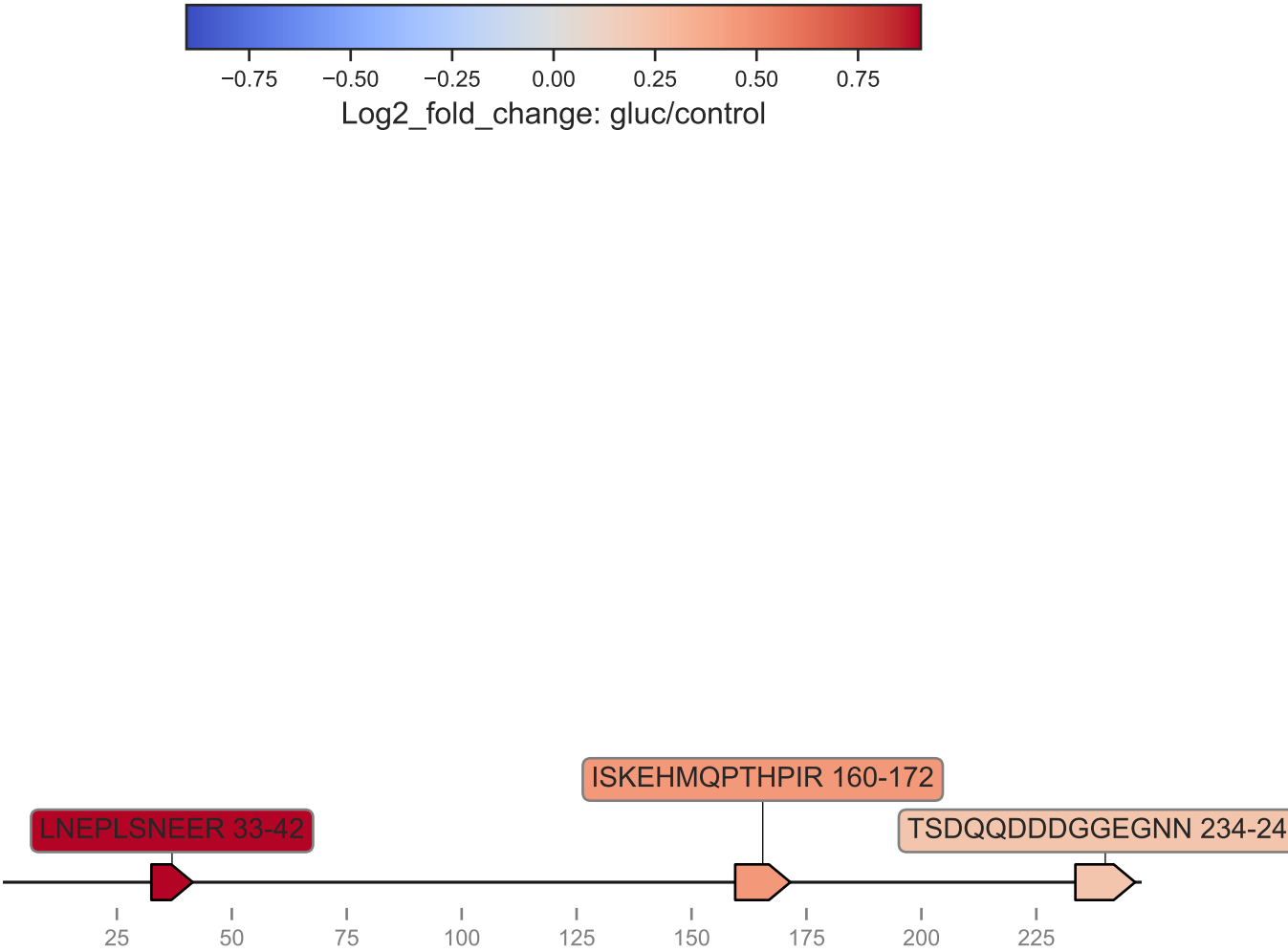
AHDTVDLTSVLR 42-53



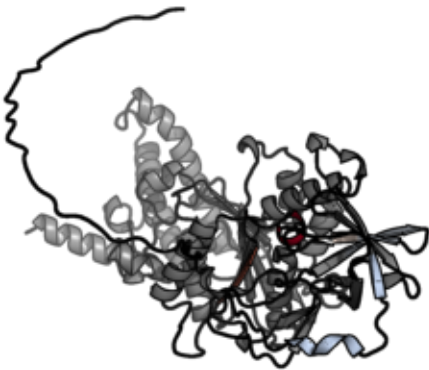
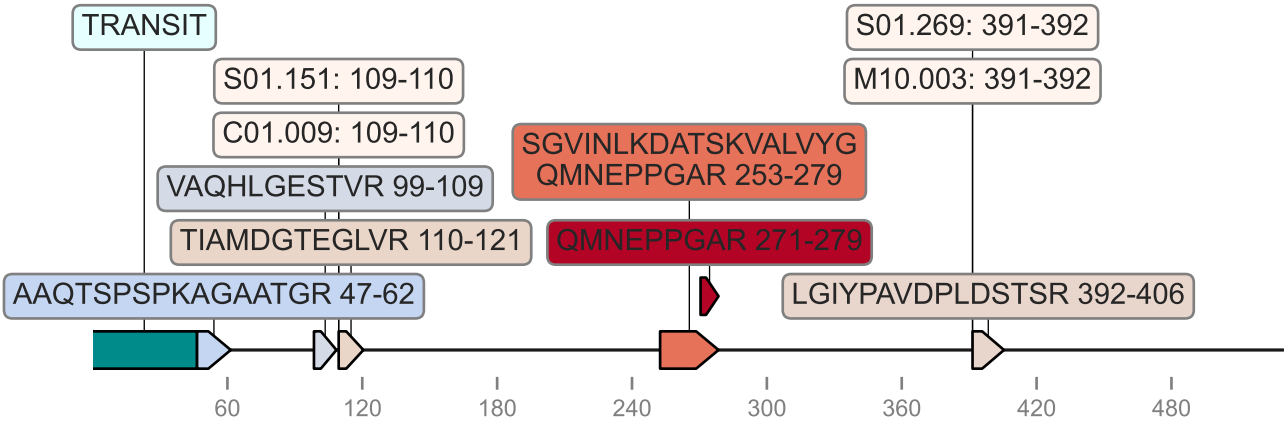
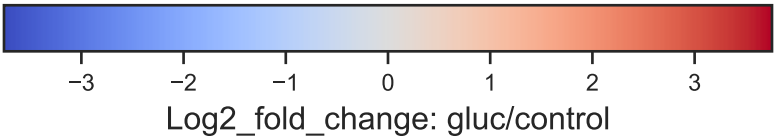
Q969V3-NCLN-BOS complex subunit NCLN {ECO:0000305}



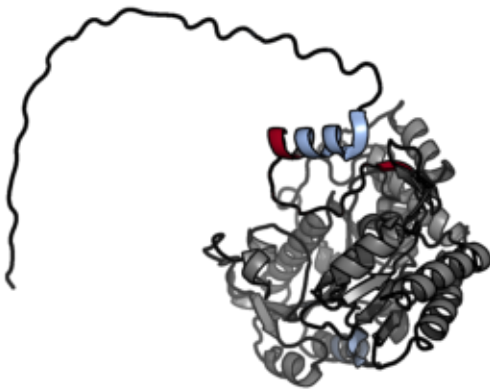
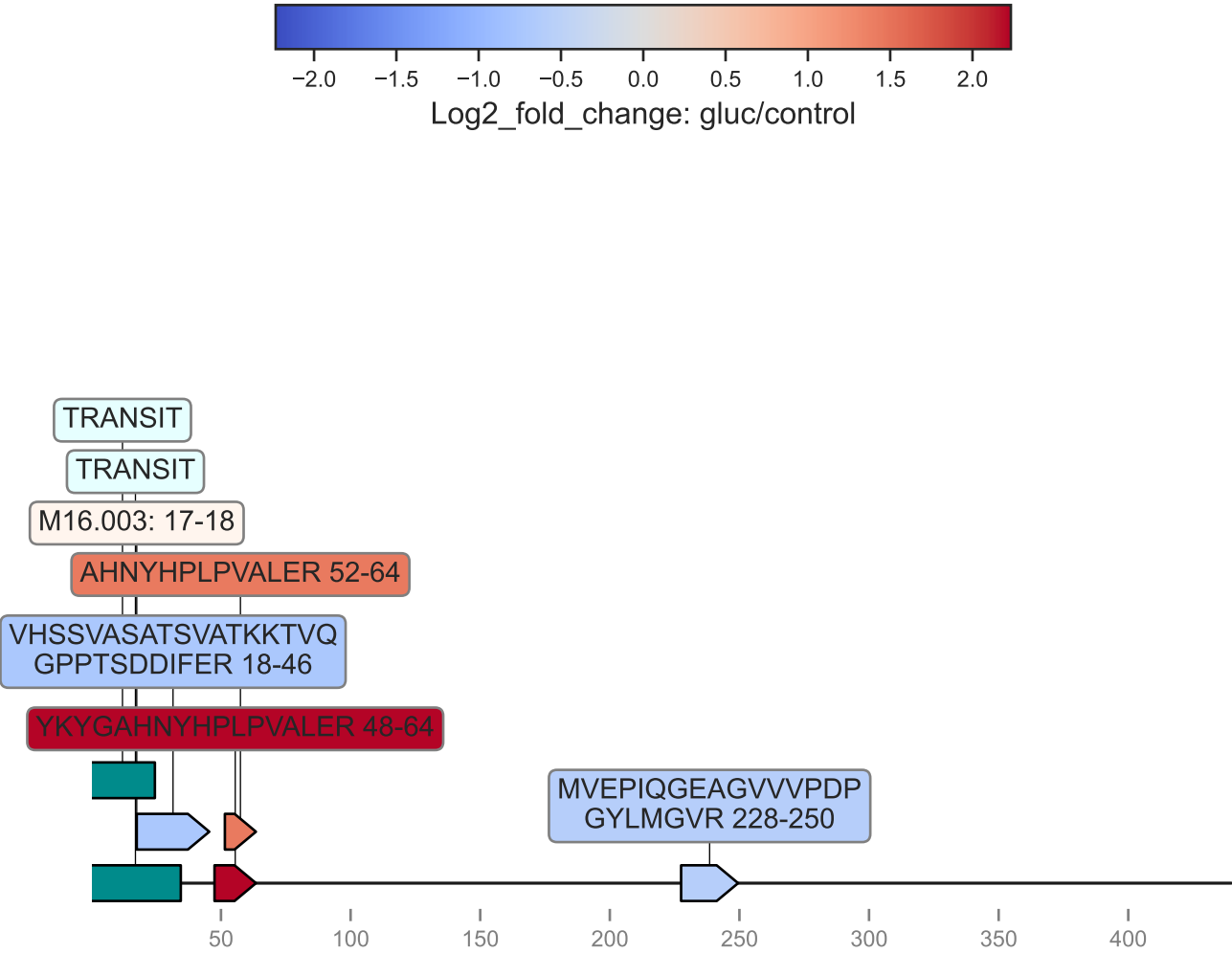
P61981-YWHAG-14-3-3 protein gamma



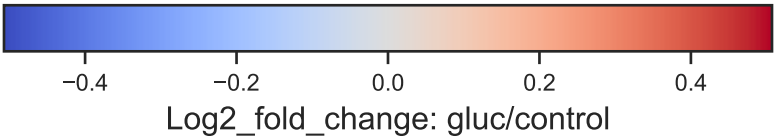
P06576-ATP5F1B-ATP synthase subunit beta, mitochondrial {ECO:0000305}



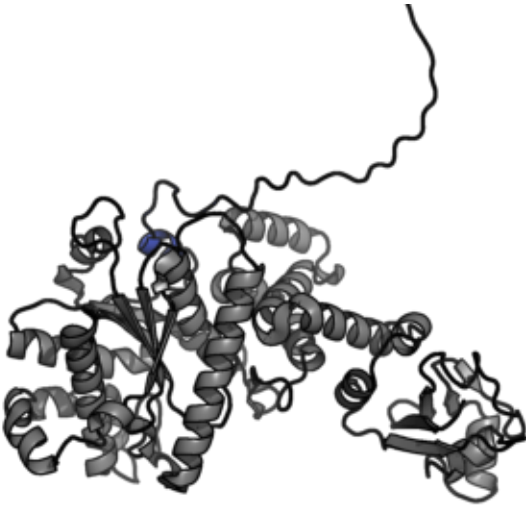
P04181-OAT-Ornithine aminotransferase, mitochondrial



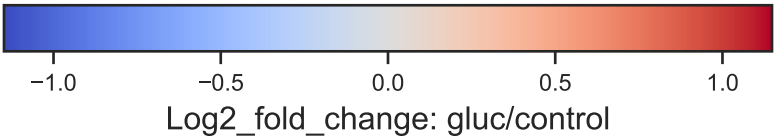
Q9Y2Z4-YARS2-Tyrosine--tRNA ligase, mitochondrial



TRANSIT
AHSGAQGLLAAQKAR 31-45

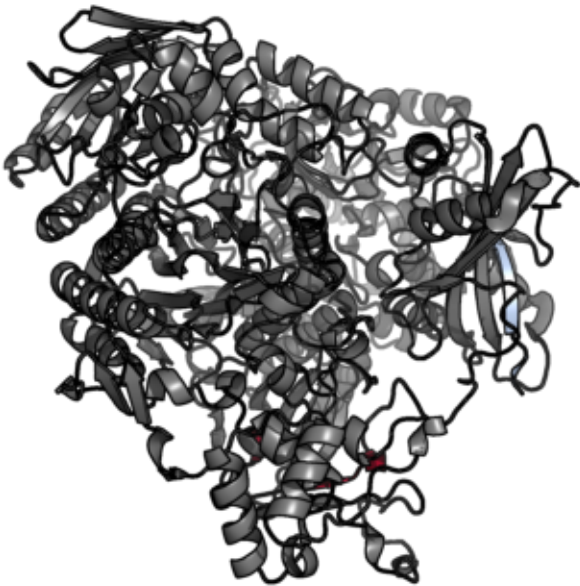


O15067-PFAS-Phosphoribosylformylglycinamidine synthase



CLE_M24: 1-2
SPVLH FYVR 2-10

S01.269: 528-529
LSDPAGAIYTSR 529-541



P63220-RPS21-40S ribosomal protein S21

