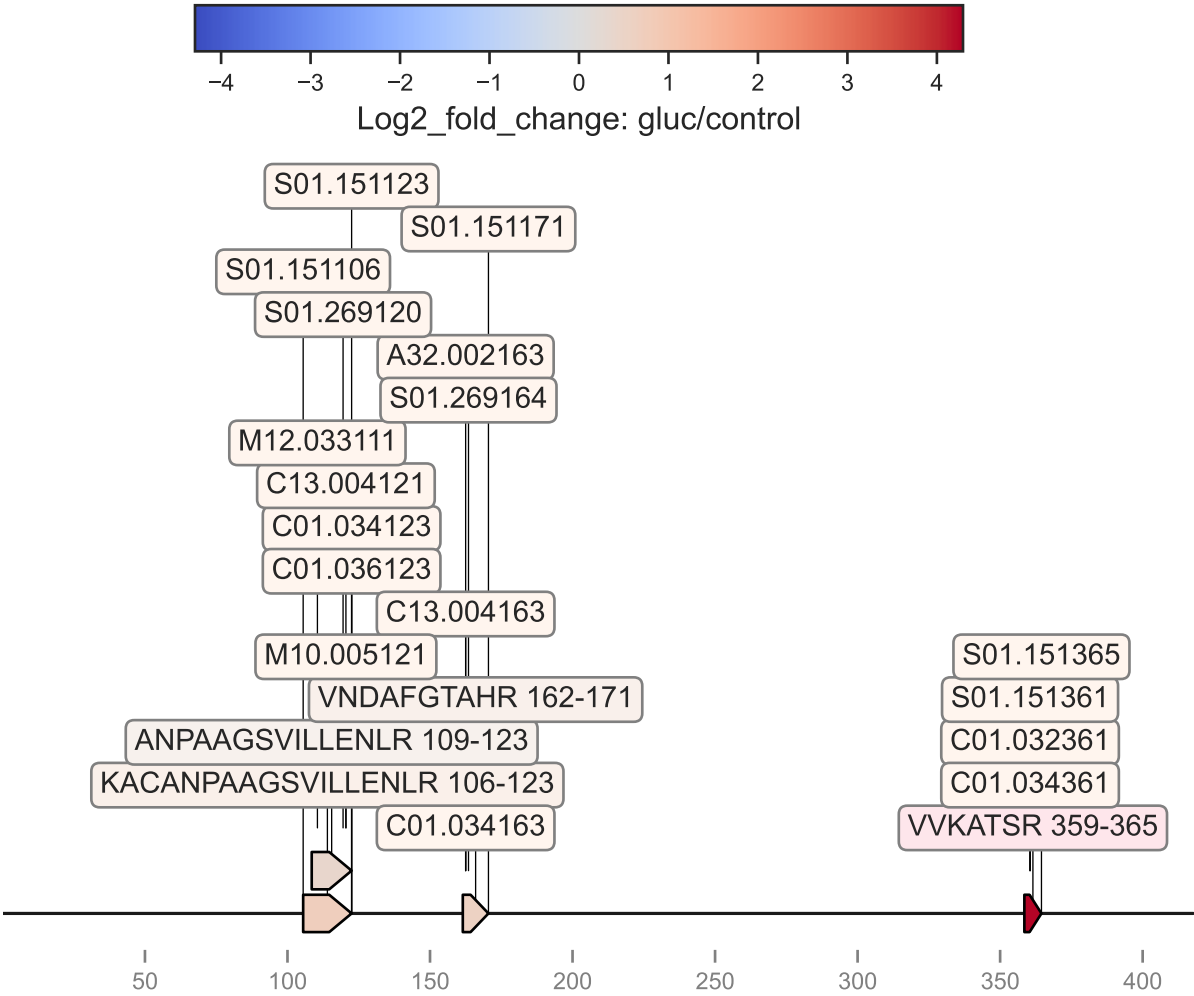
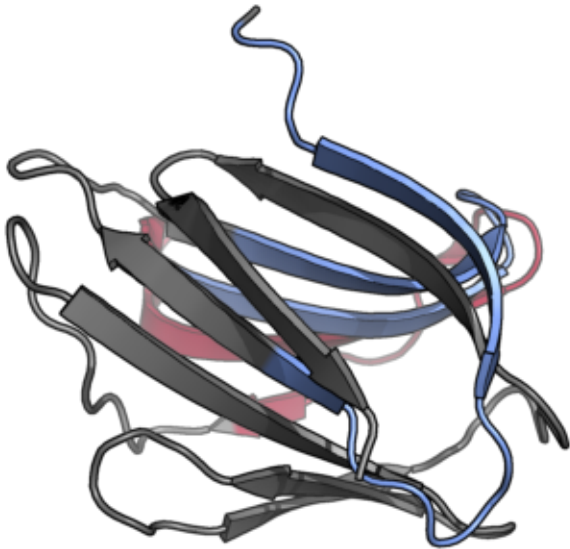
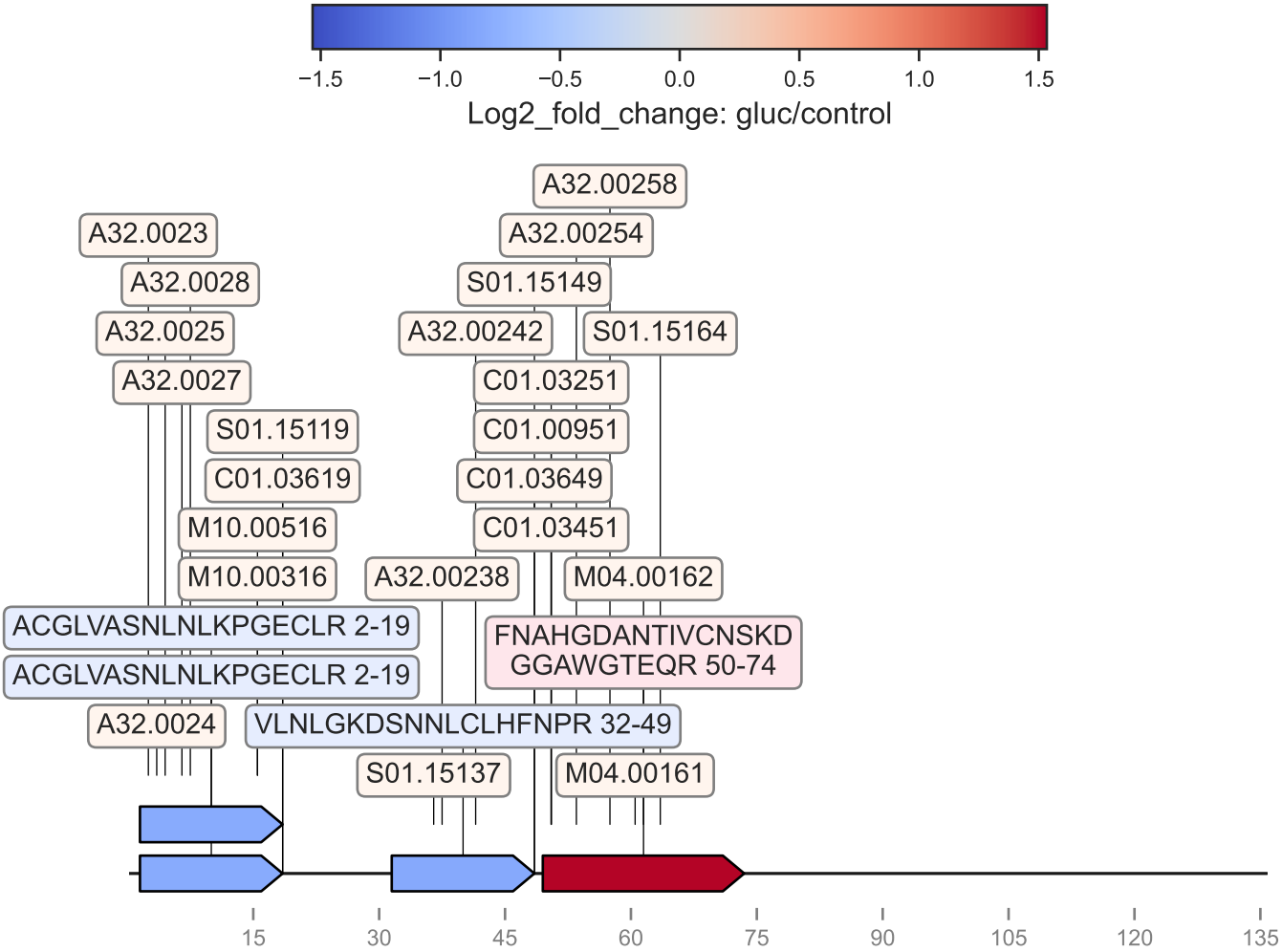


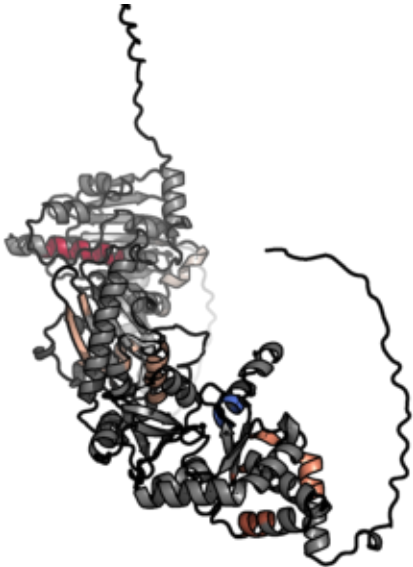
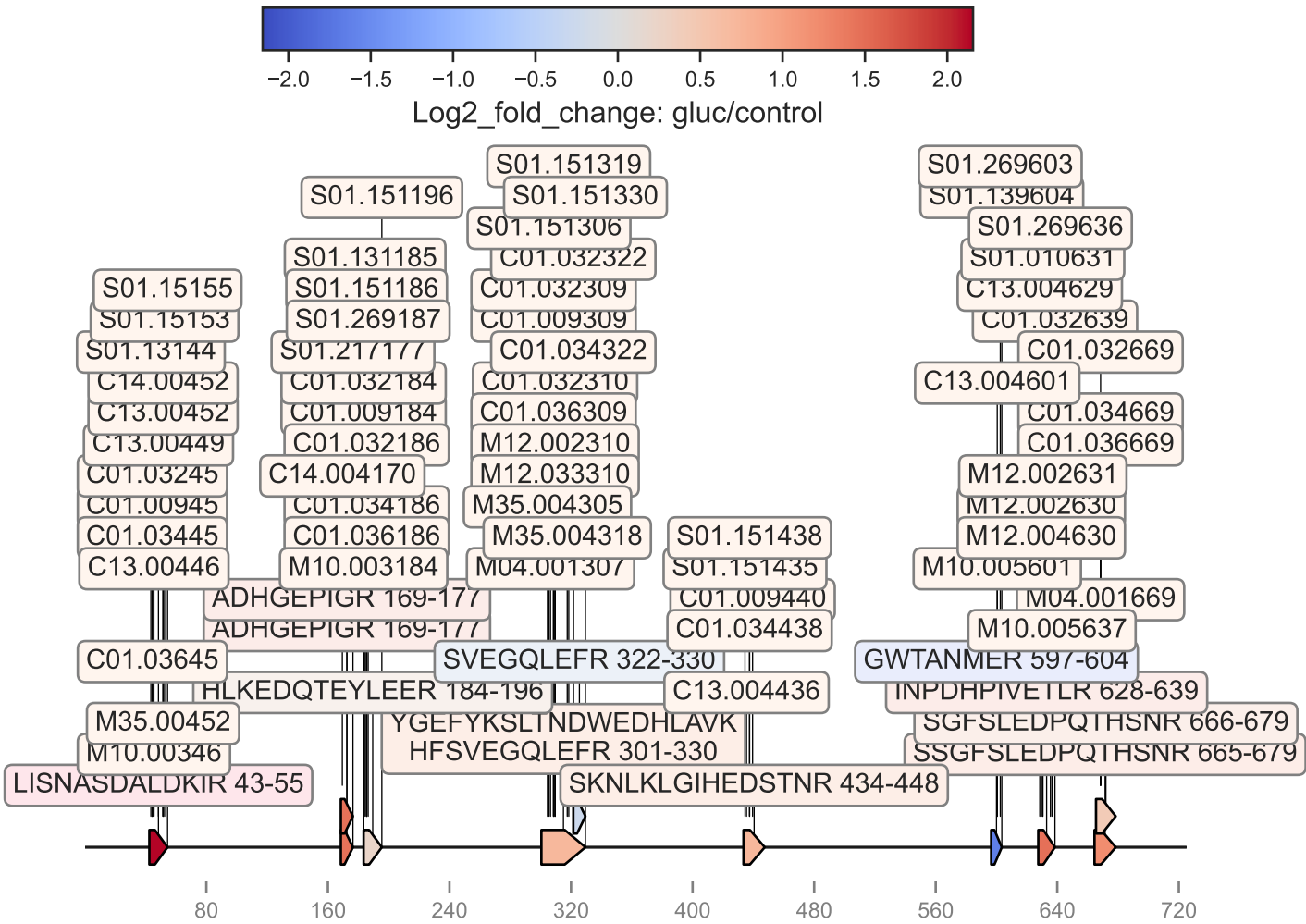
P00558-PGK1-Phosphoglycerate kinase 1

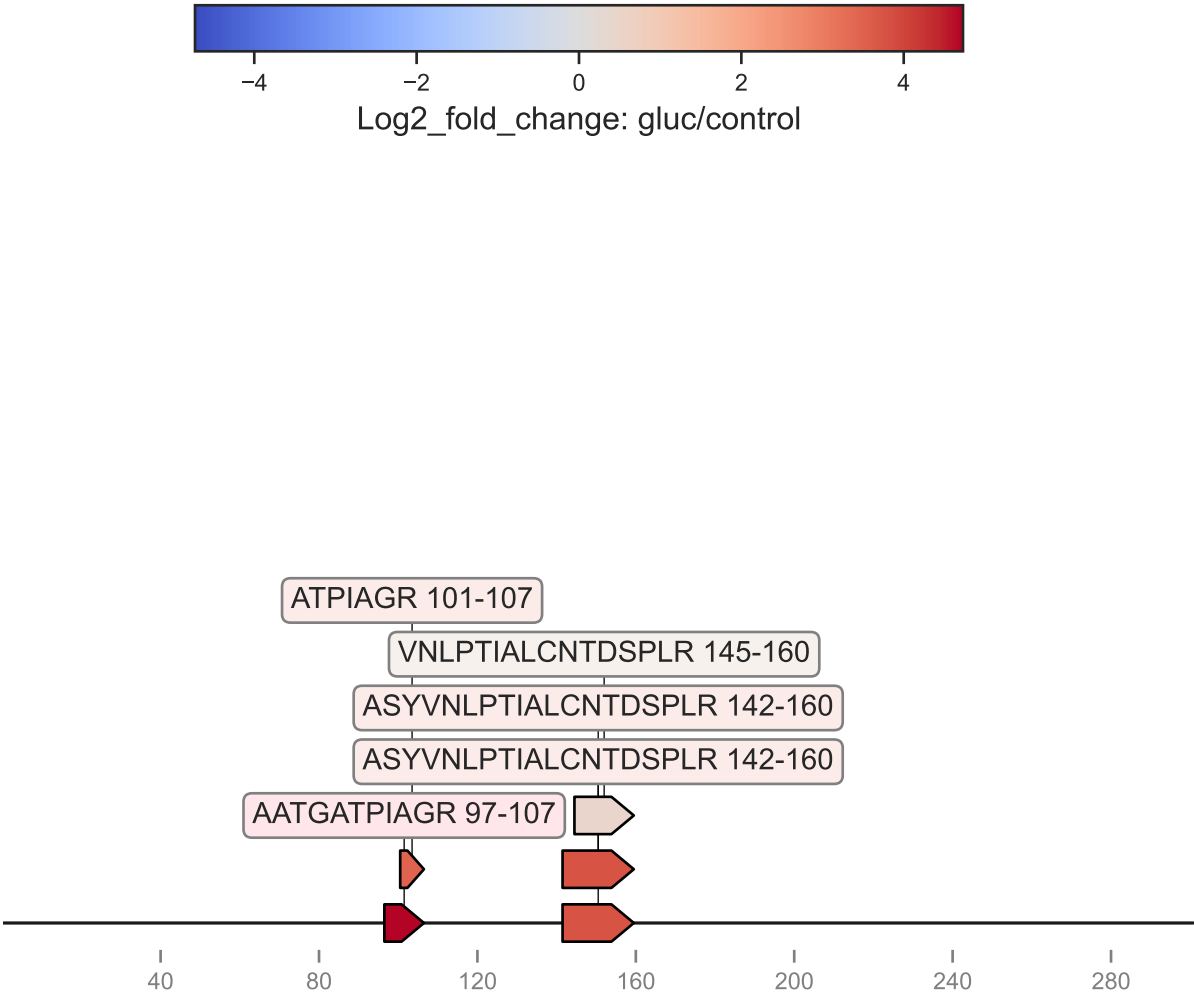


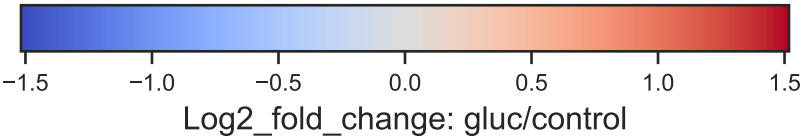
P09382-LGALS1-Galectin-1



P08238-HSP90AB1-Heat shock protein HSP 90-beta

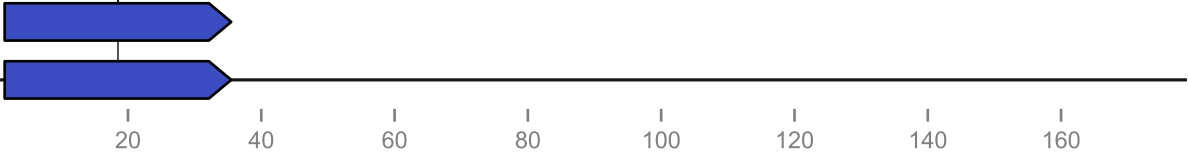




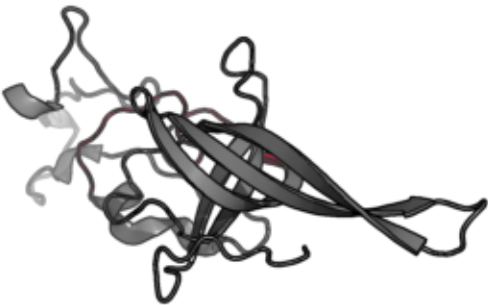
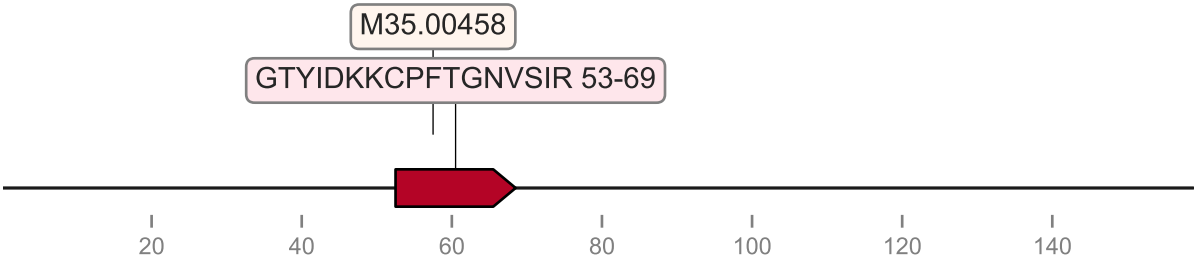
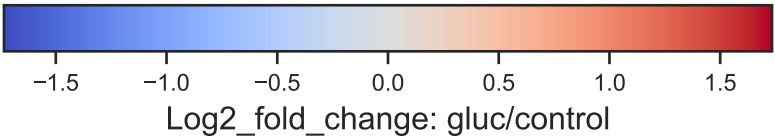


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SNTTEKLTQVMQNPR 2-36

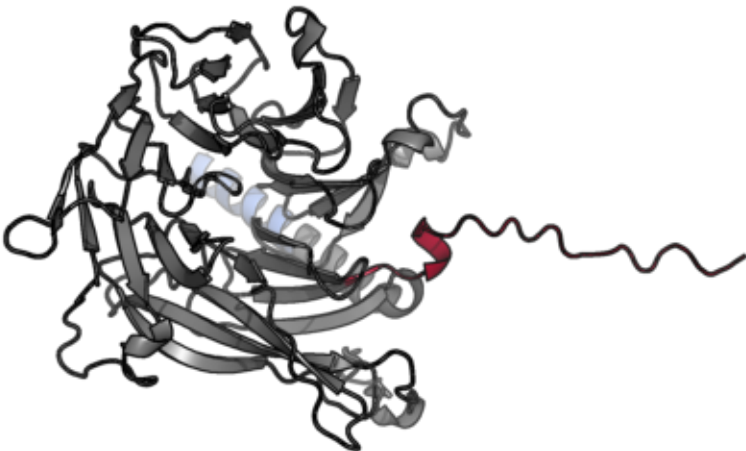
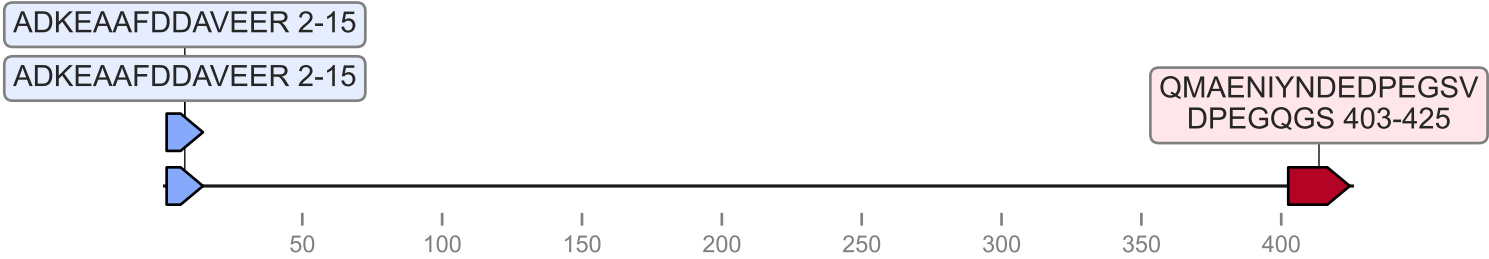
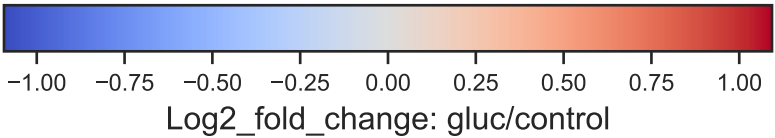
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SNTTEKLTQVMQNPR 2-36

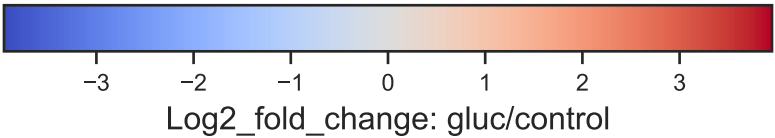


P62280-RPS11-40S ribosomal protein S11



Q09028-RBBP4-Histone-binding protein RBBP4

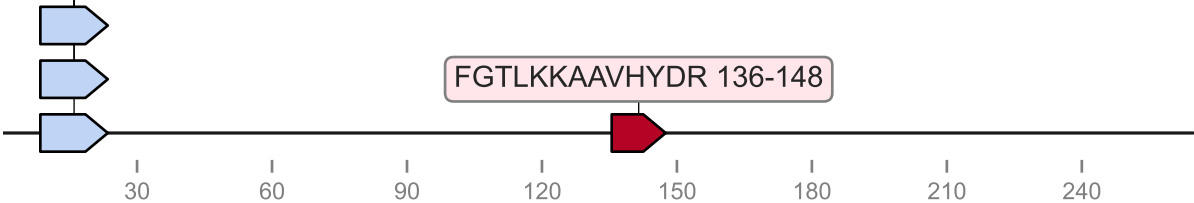




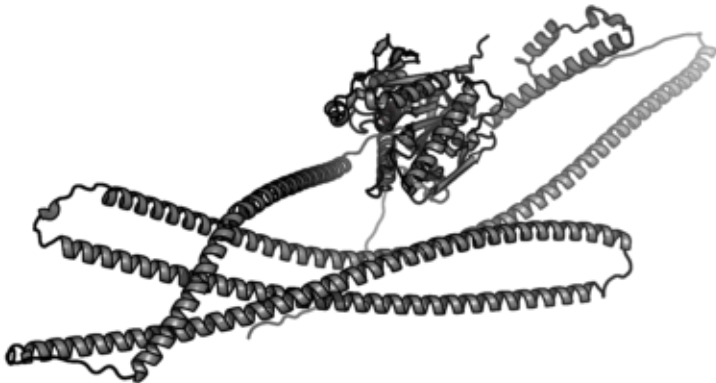
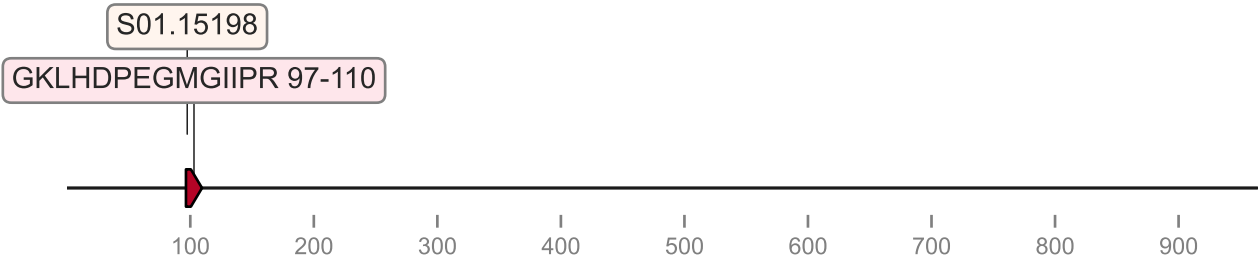
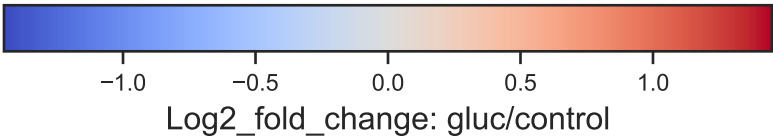
ADKMDMSLDDIIKLNR 9-24

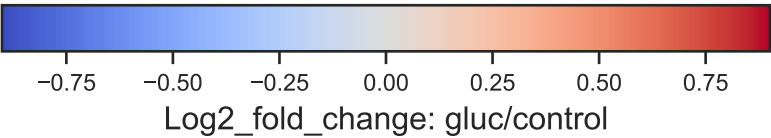
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ADKMDMSLDDIIKLNR 9-24



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



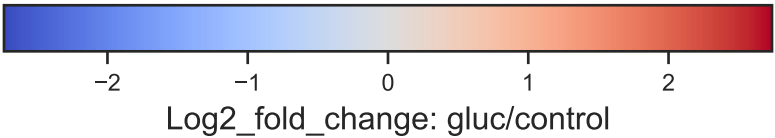


LDSPPKLSGVQQPSEGVGGGR 6-28



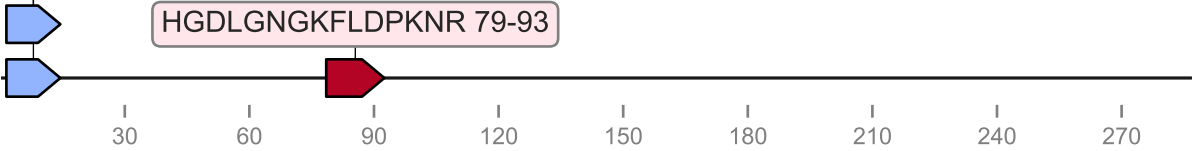
80 160 240 320 400 480 560 640 720

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

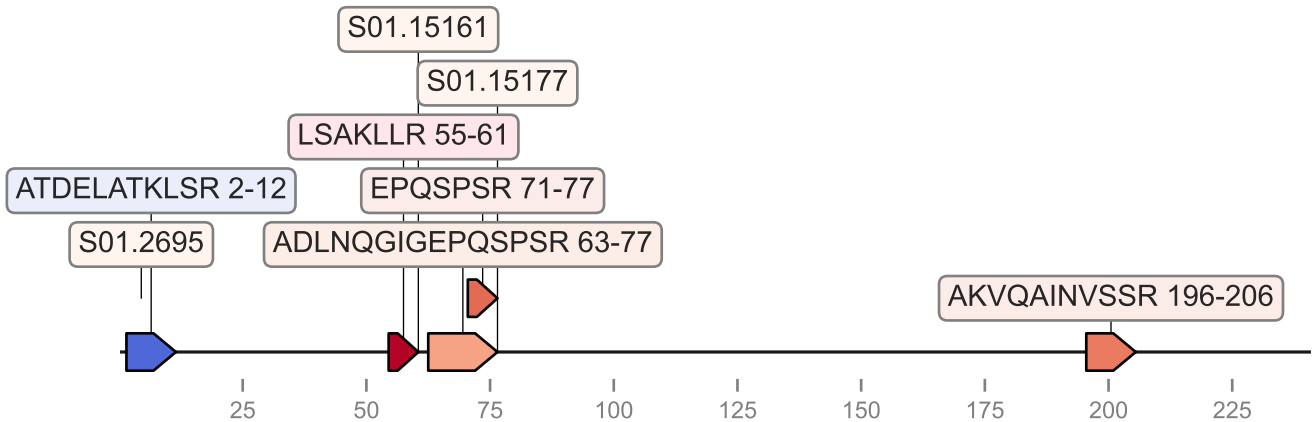
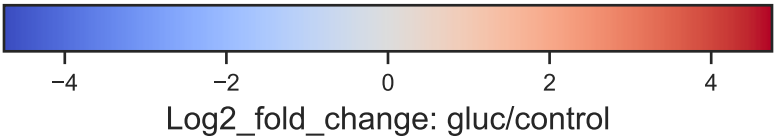


ADLEEQLSDEEKVR 2-15
ADLEEQLSDEEKVR 2-15

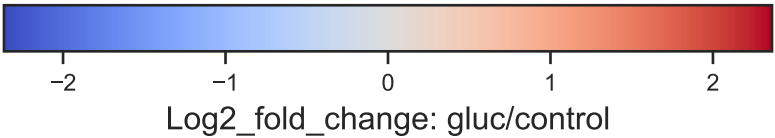
HGDLGNGKFLDPKNR 79-93



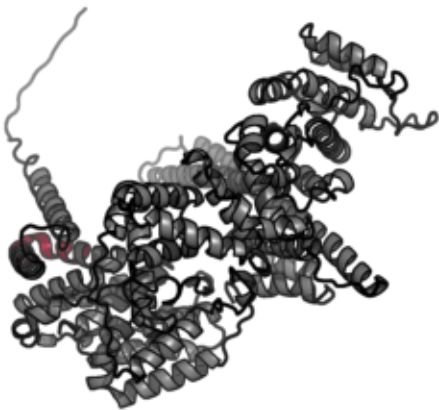
Q96C19-EFHD2-EF-hand domain-containing protein D2

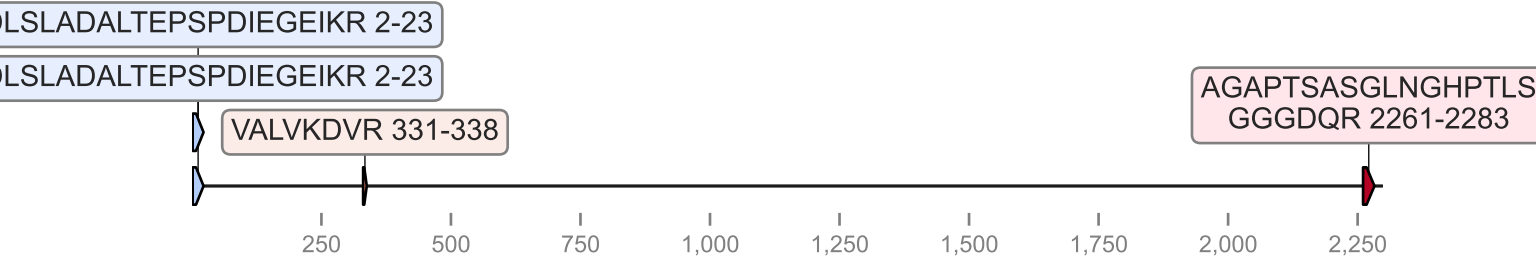
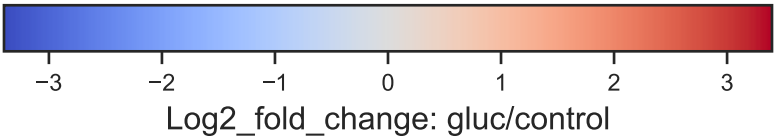


Q09161-NCBP1-Nuclear cap-binding protein subunit 1

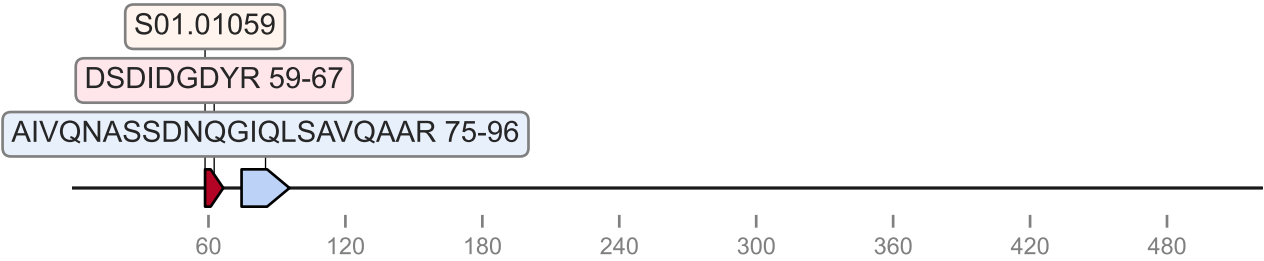
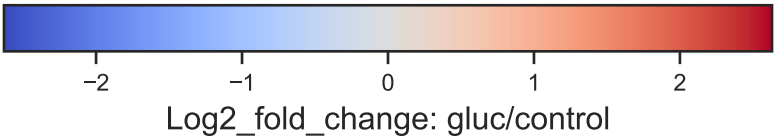


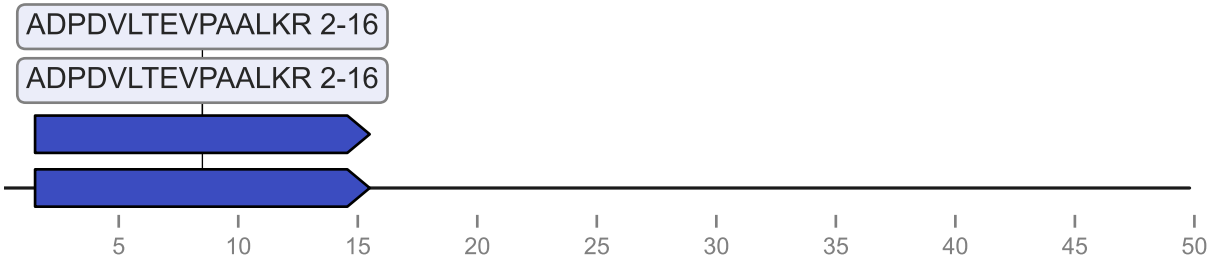
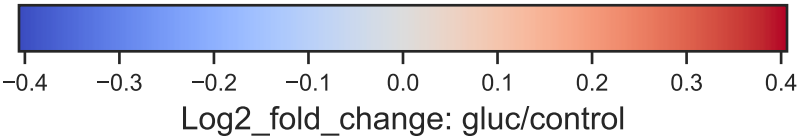
ADLPNYKSKILR 59-70

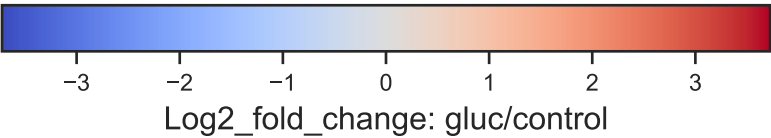




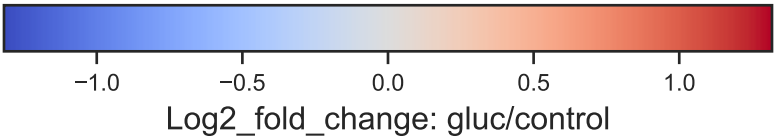
O00629-KPNA4-Importin subunit alpha-3





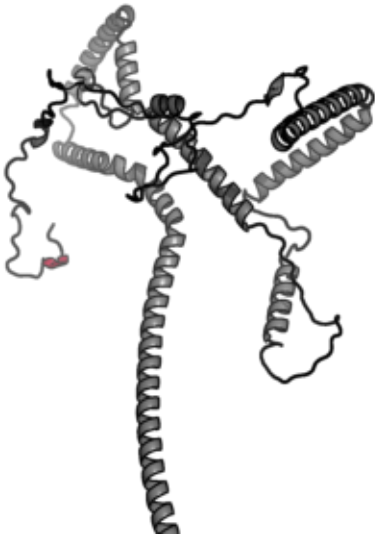


Q13561-DCTN2-Dynactin subunit 2

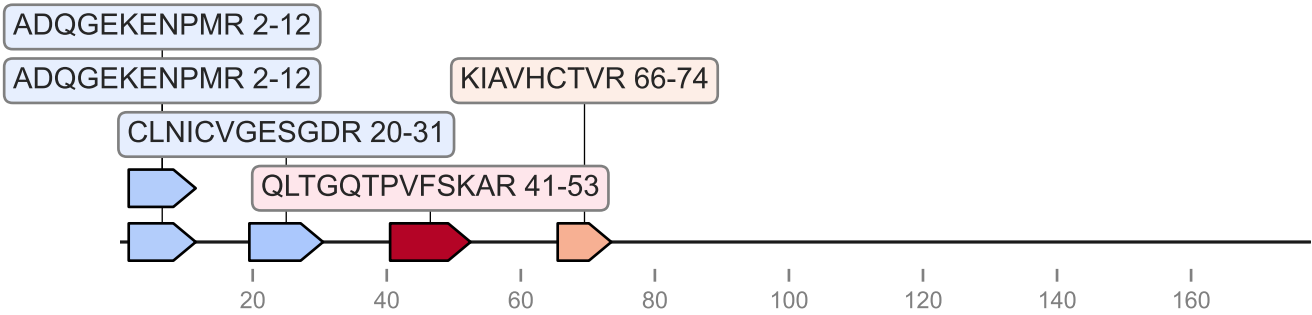
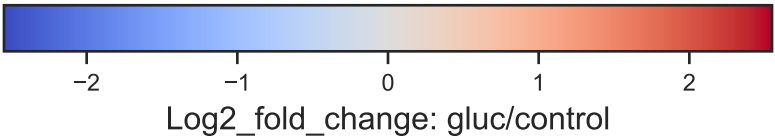


ADPKYADLPGIAR 2-14

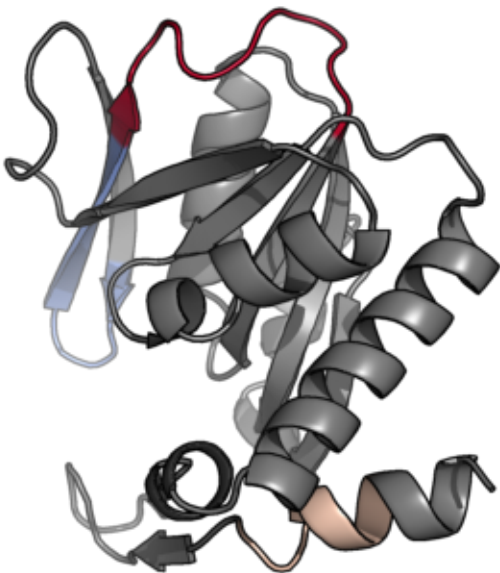
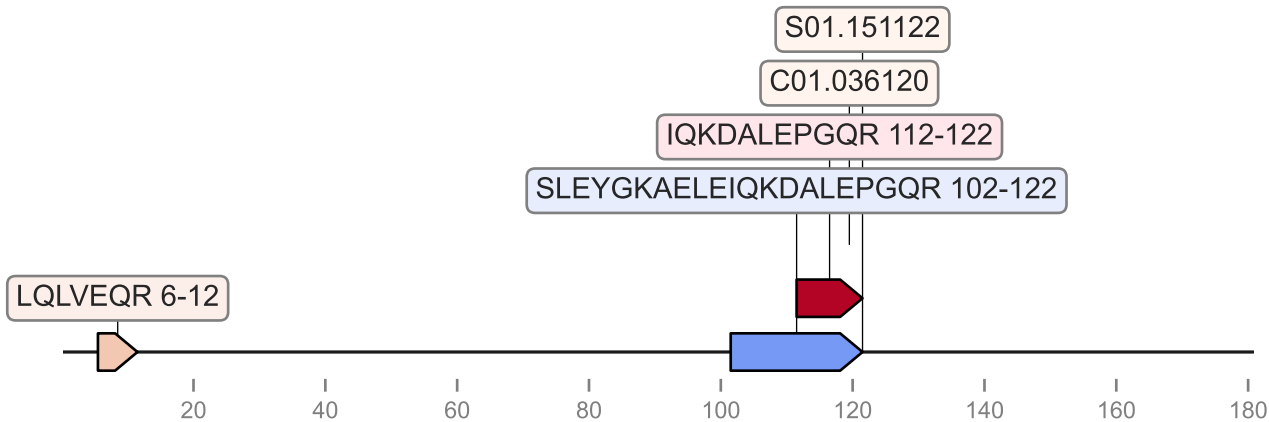
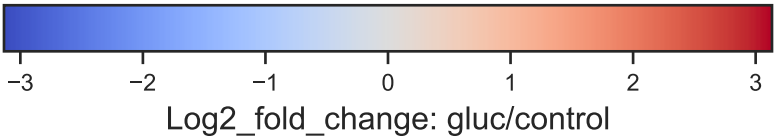
ADPKYADLPGIAR 2-14



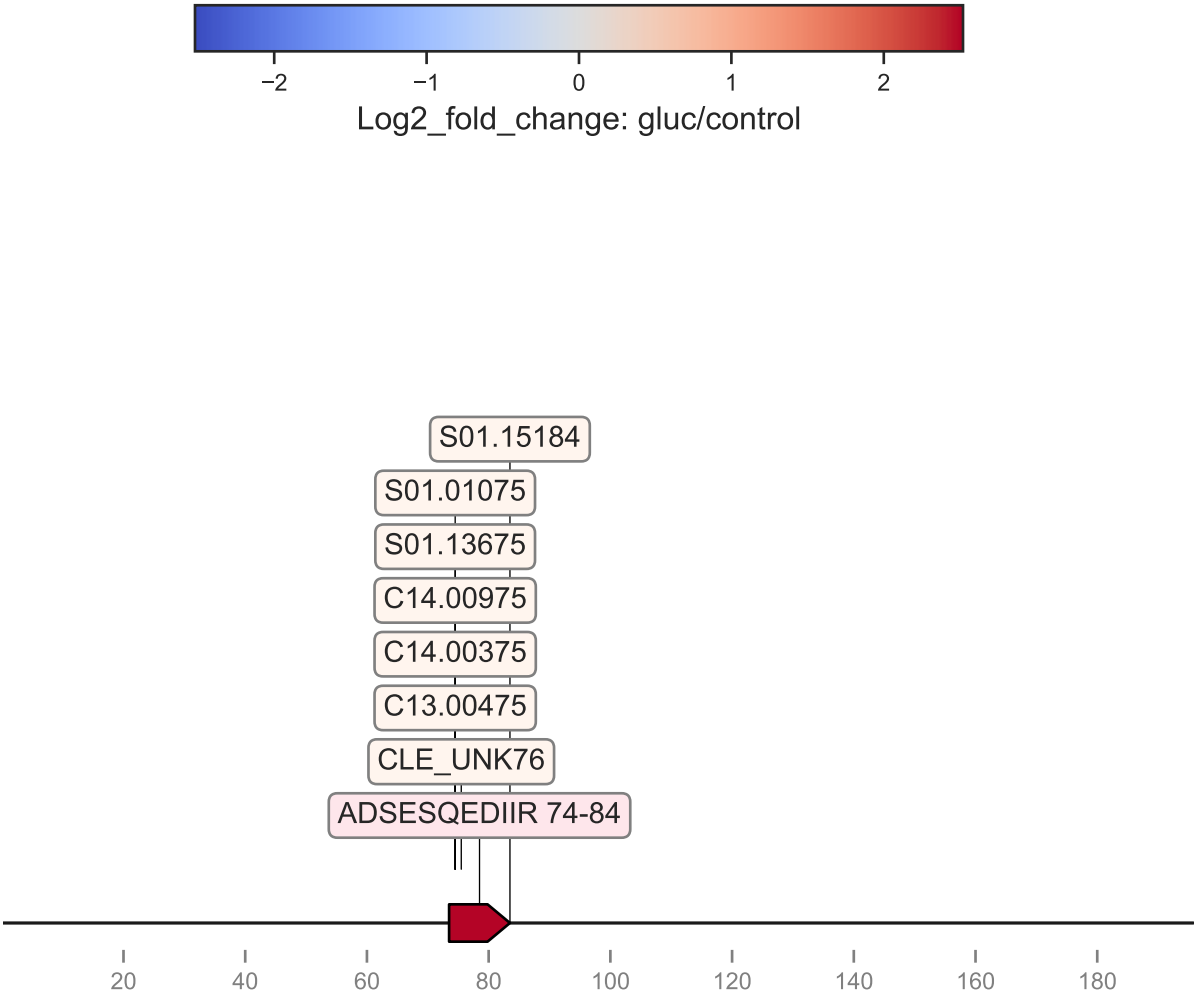
P62913-2-RPL11-Isoform 2 of 60S ribosomal protein L11



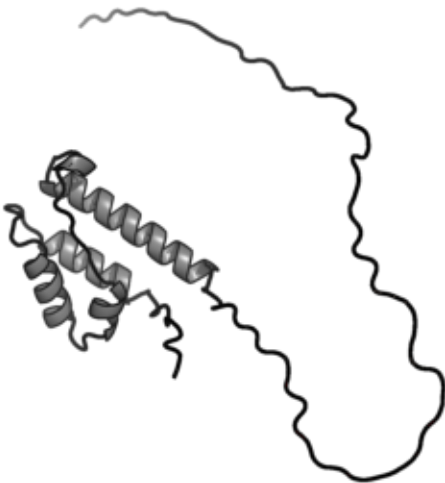
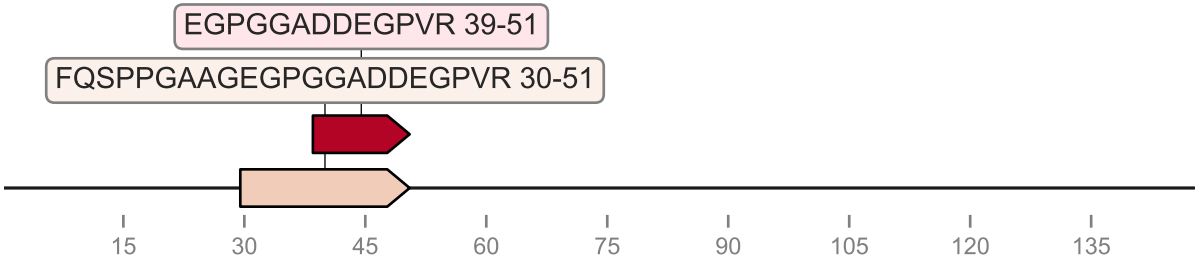
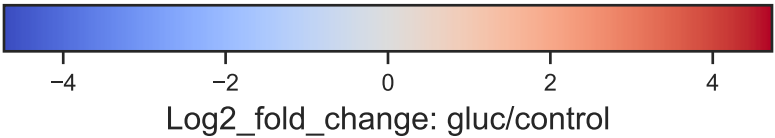
P07741-APRT-Adenine phosphoribosyltransferase {ECO:0000305}



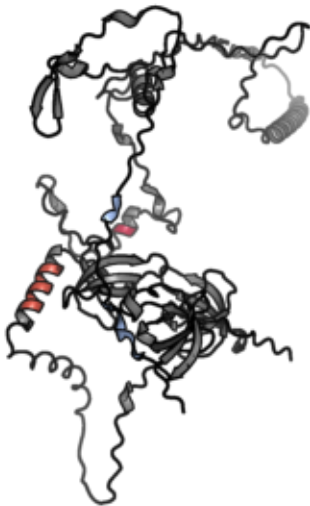
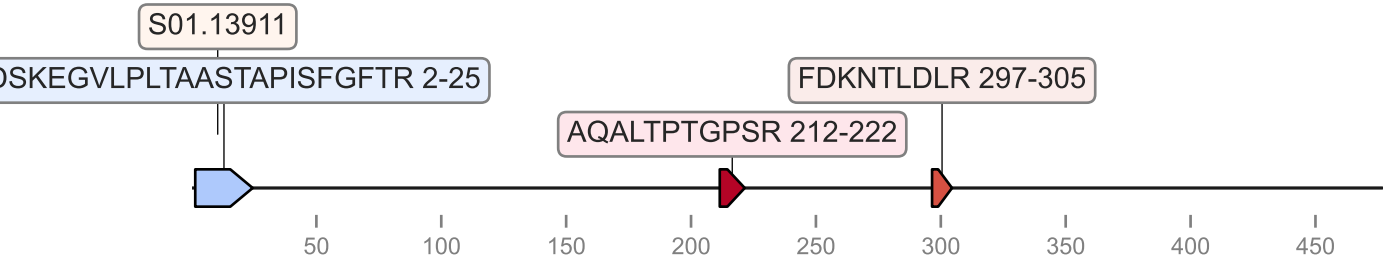
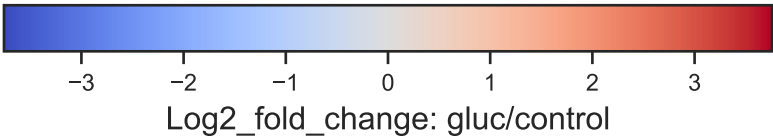
P55957-BID-BH3-interacting domain death agonist



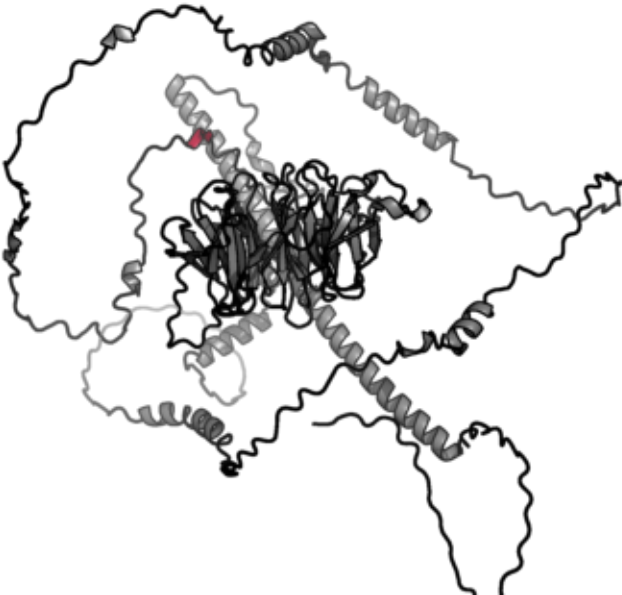
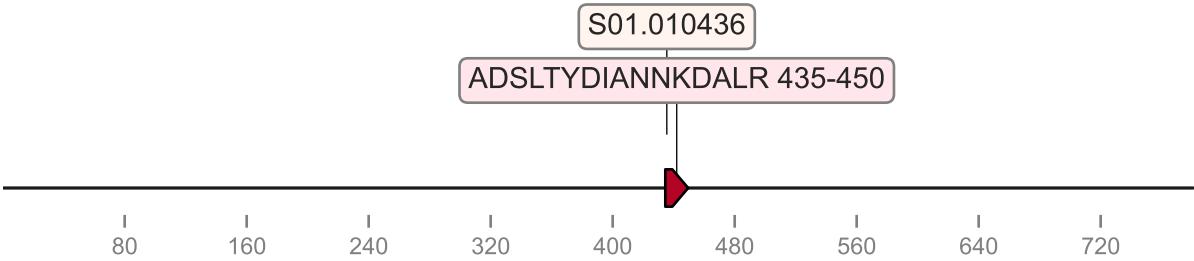
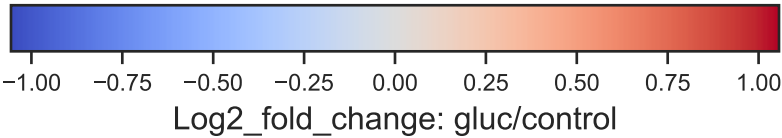
Q96C90-PPP1R14B-Protein phosphatase 1 regulatory subunit 14B



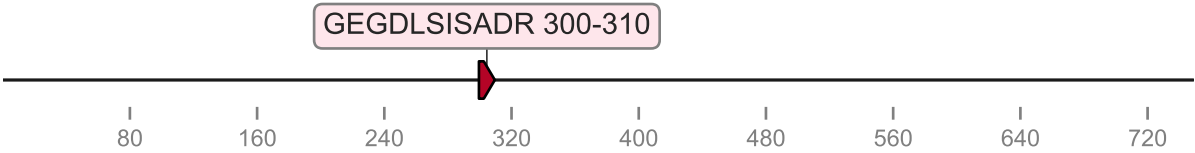
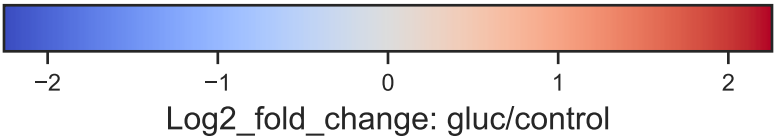
Q92917-GPKOW-G-patch domain and KOW motifs-containing protein



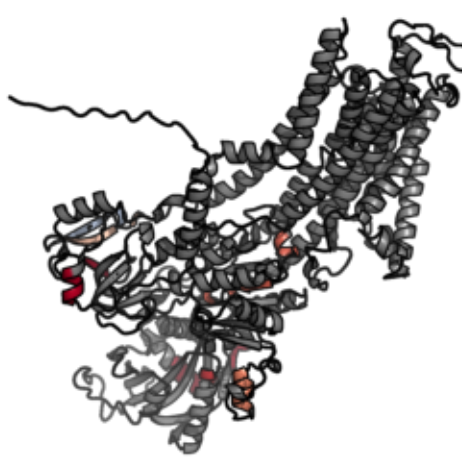
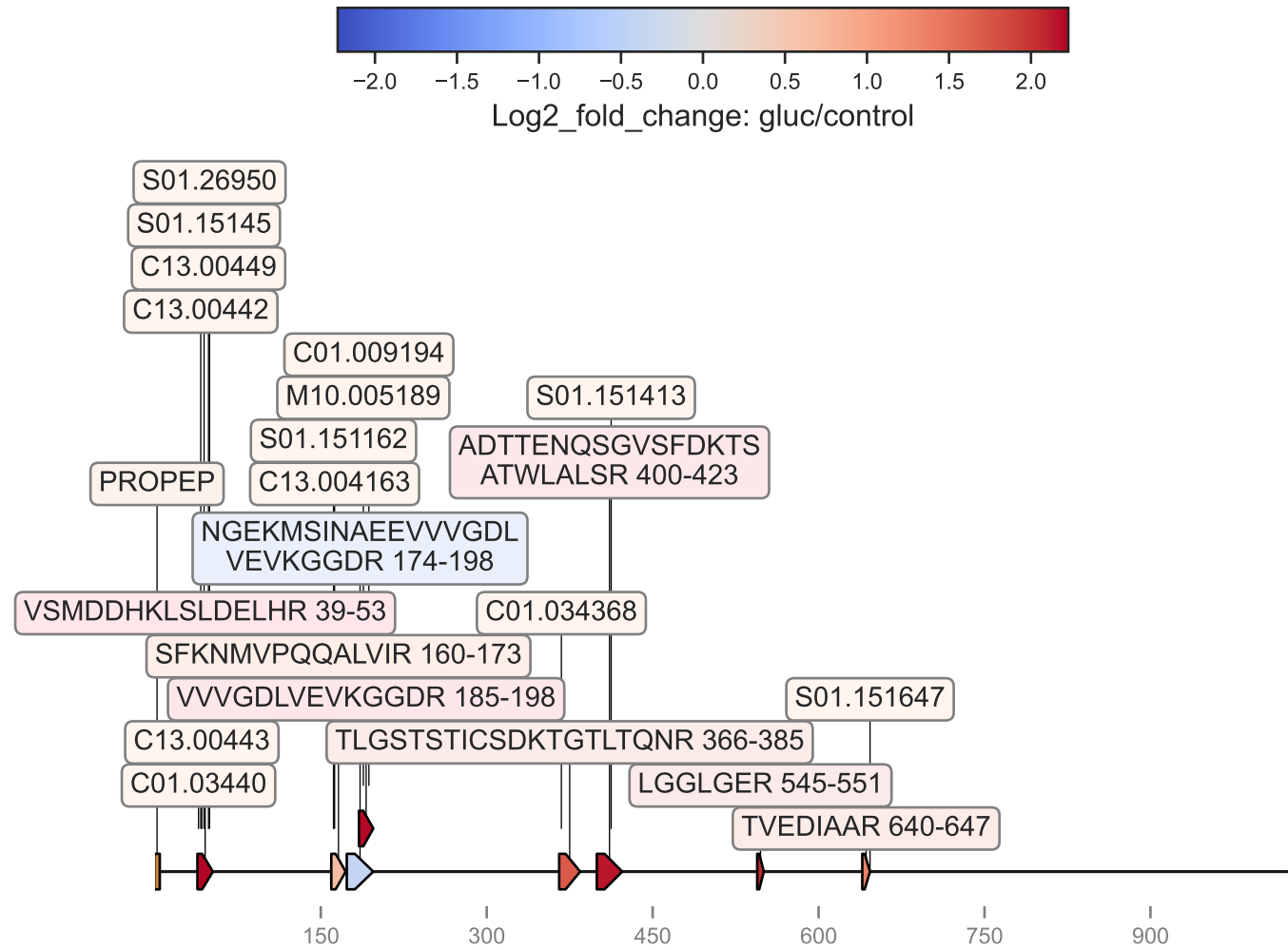
O43815-STRN-Striatin

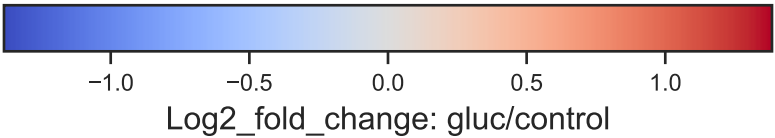


P49589-CARS1-Cysteine--tRNA ligase, cytoplasmic



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

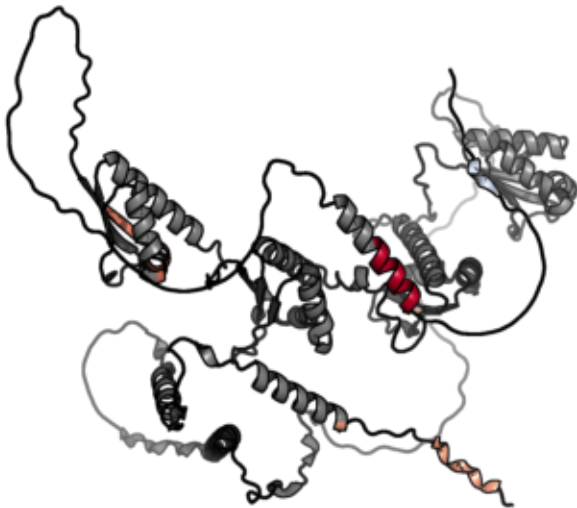
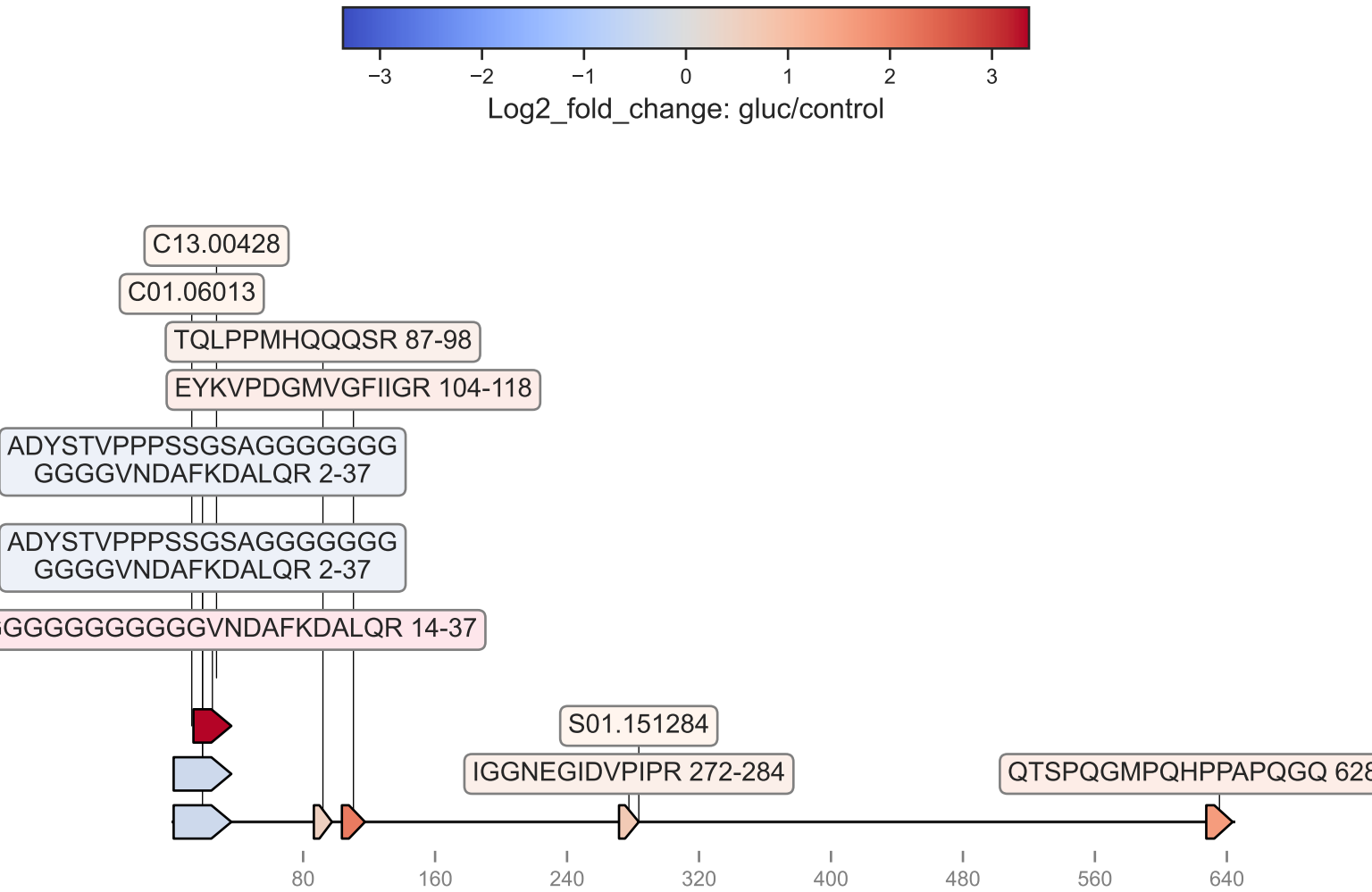




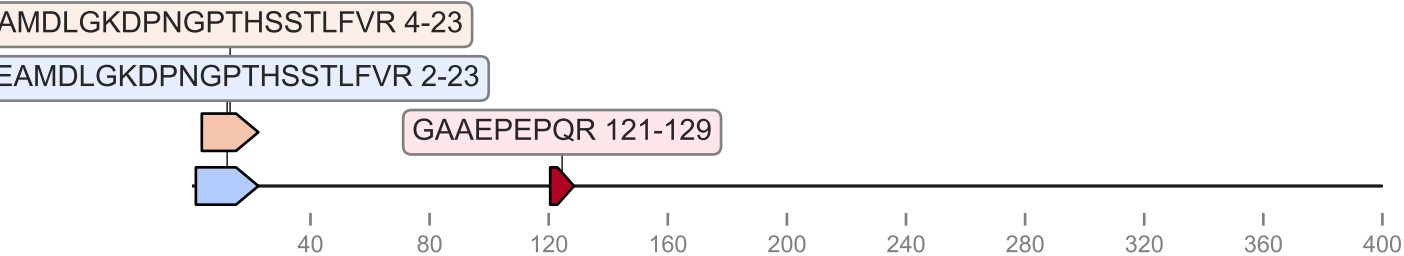
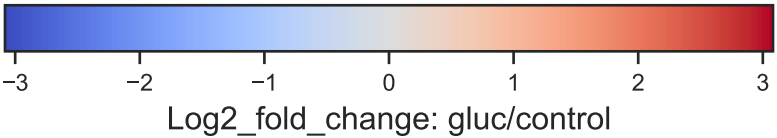
ADV VVGKDKGGEQR 2-15



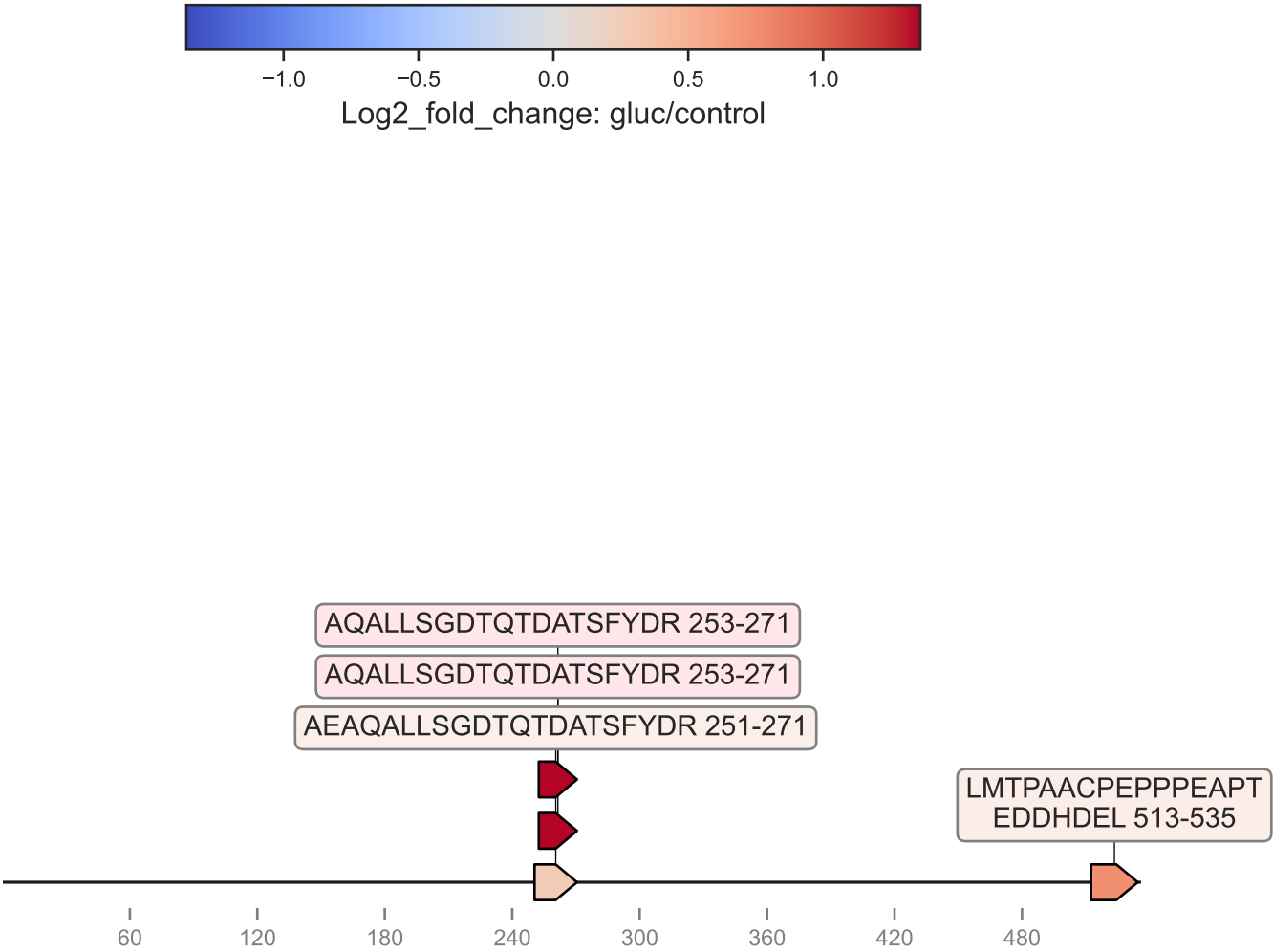
Q96AE4-FUBP1-Far upstream element-binding protein 1

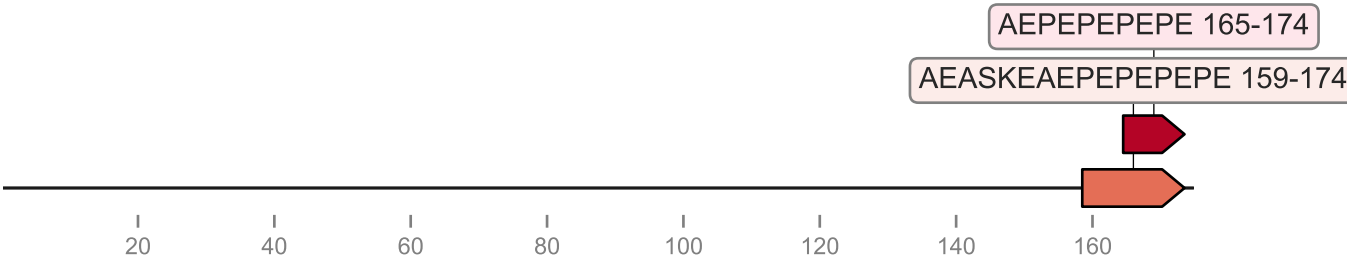
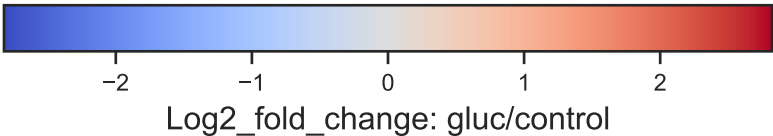


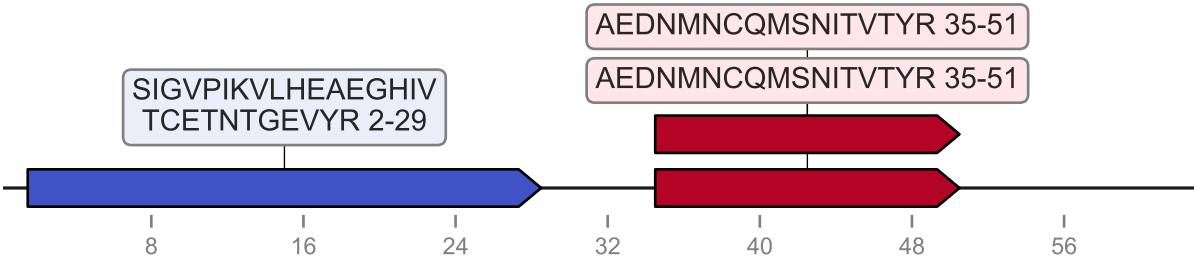
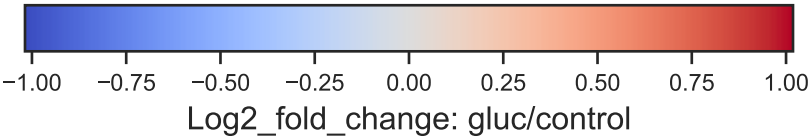
Q9NYB0-TERF2IP-Telomeric repeat-binding factor 2-interacting protein 1

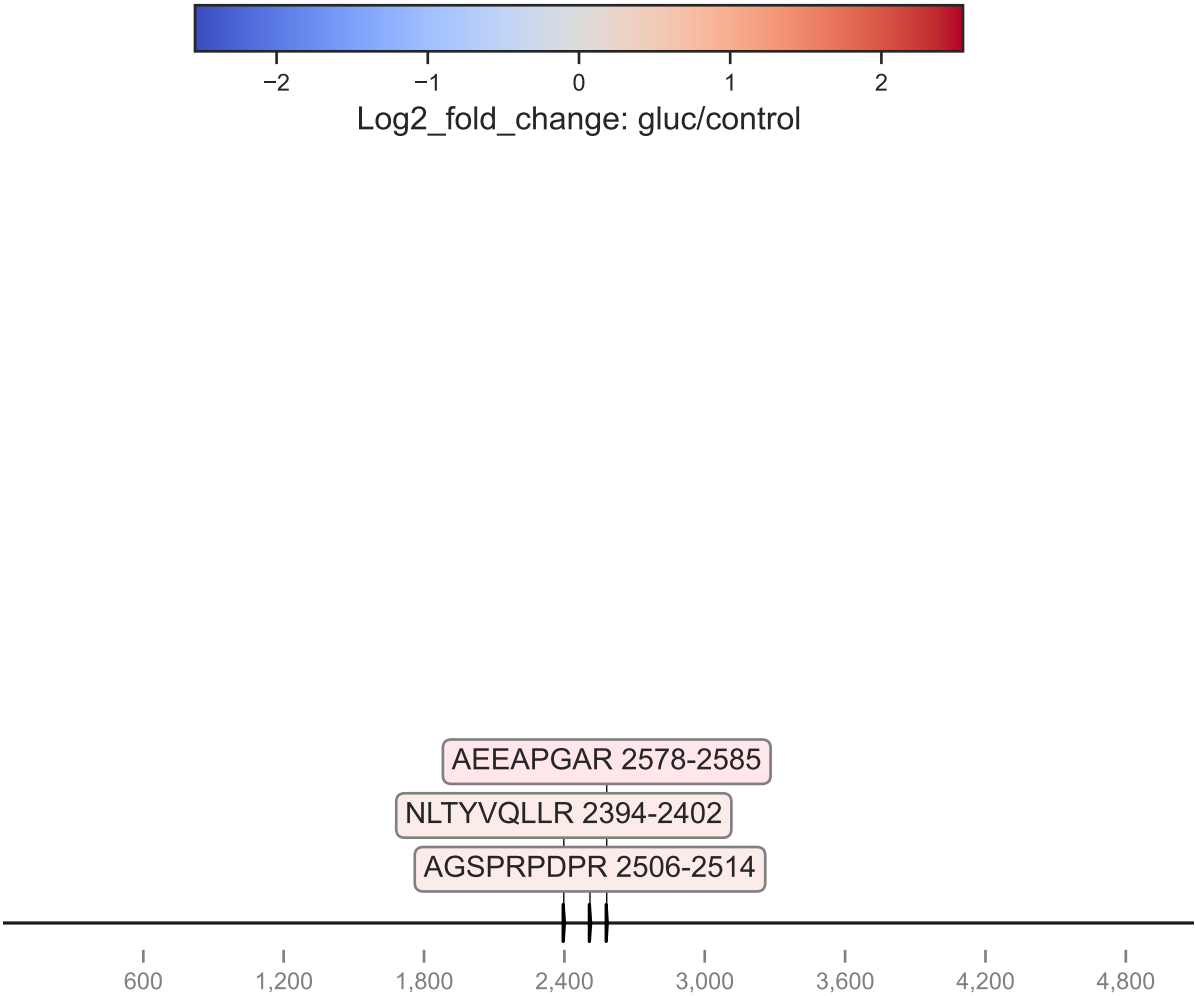


K7ELL7-PRKCSH-Glucosidase 2 subunit beta {ECO:0000256|ARBA:ARBA00022387}

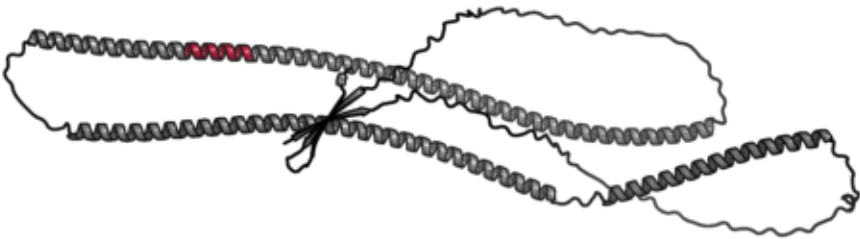
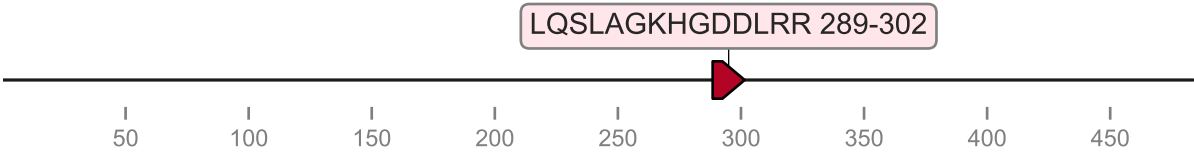
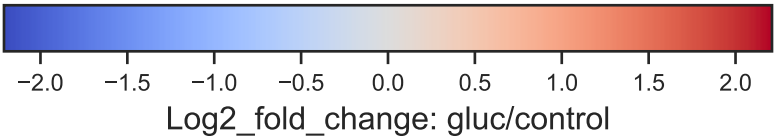




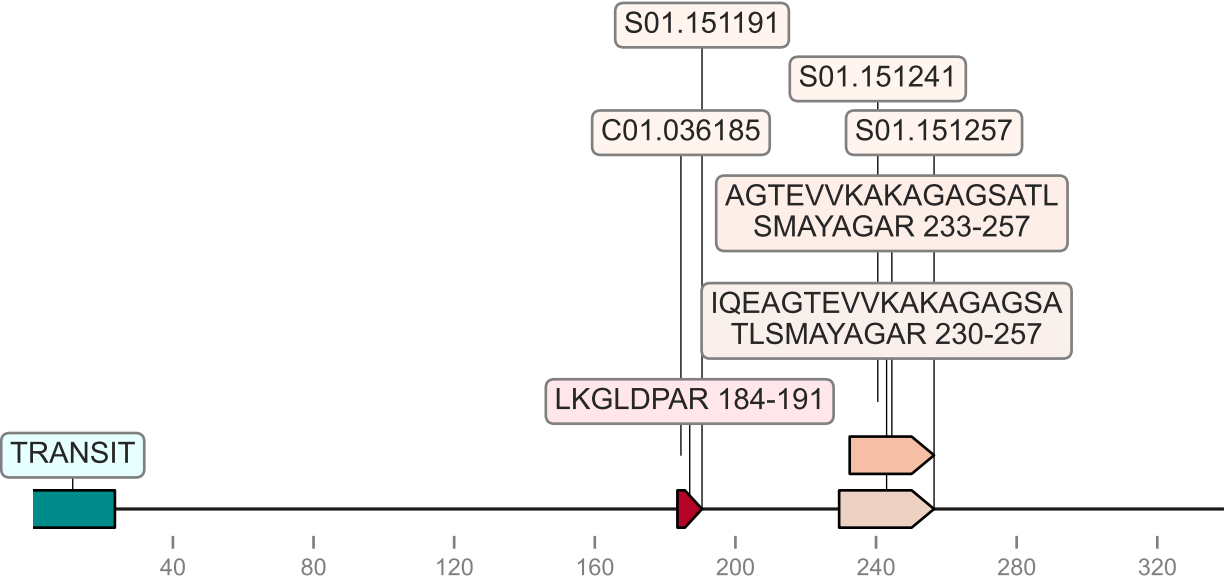
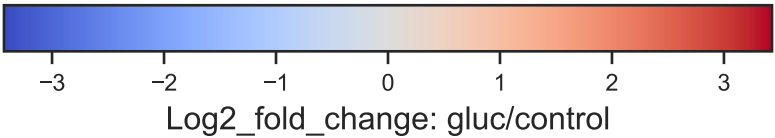




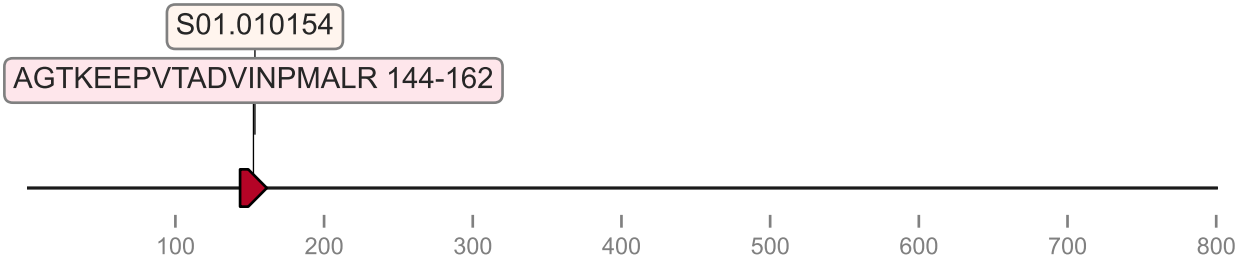
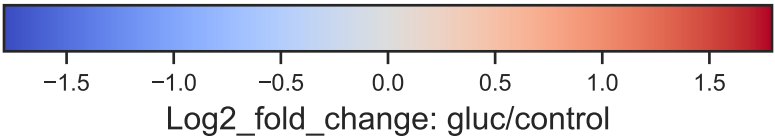
P05787-KRT8-Keratin, type II cytoskeletal 8



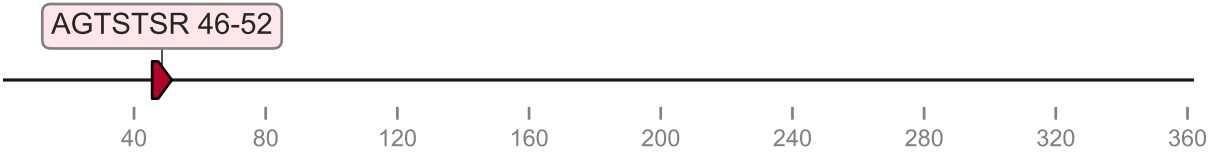
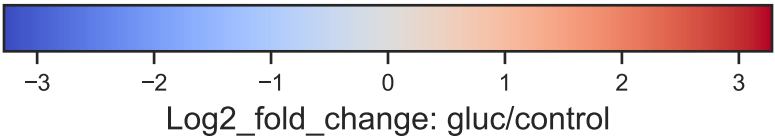
P40926-MDH2-Malate dehydrogenase, mitochondrial



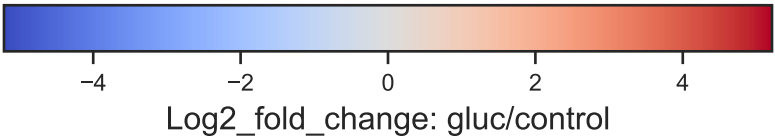
O43290-SART1-U4/U6.U5 tri-snRNP-associated protein 1



O96008-TOMM40-Mitochondrial import receptor subunit TOM40 homolog

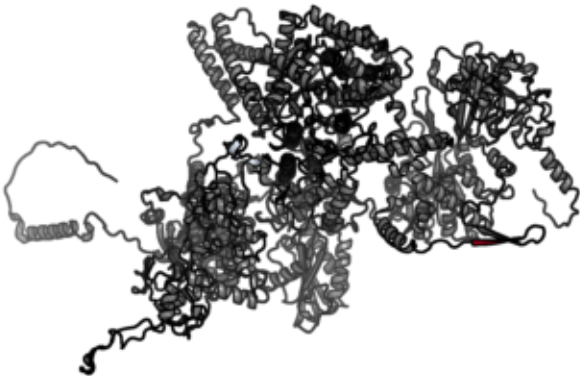


Q6P2Q9-PRPF8-Pre-mRNA-processing-splicing factor 8

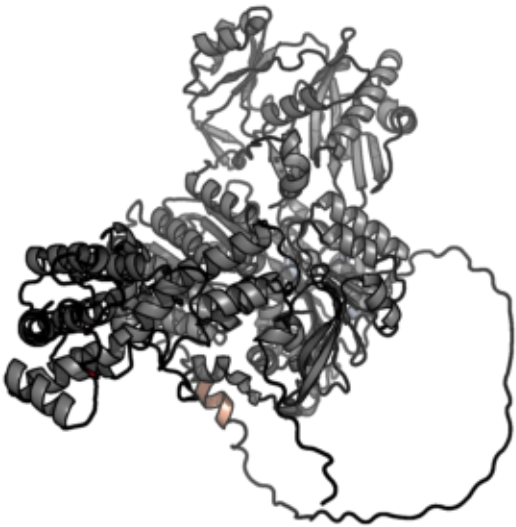
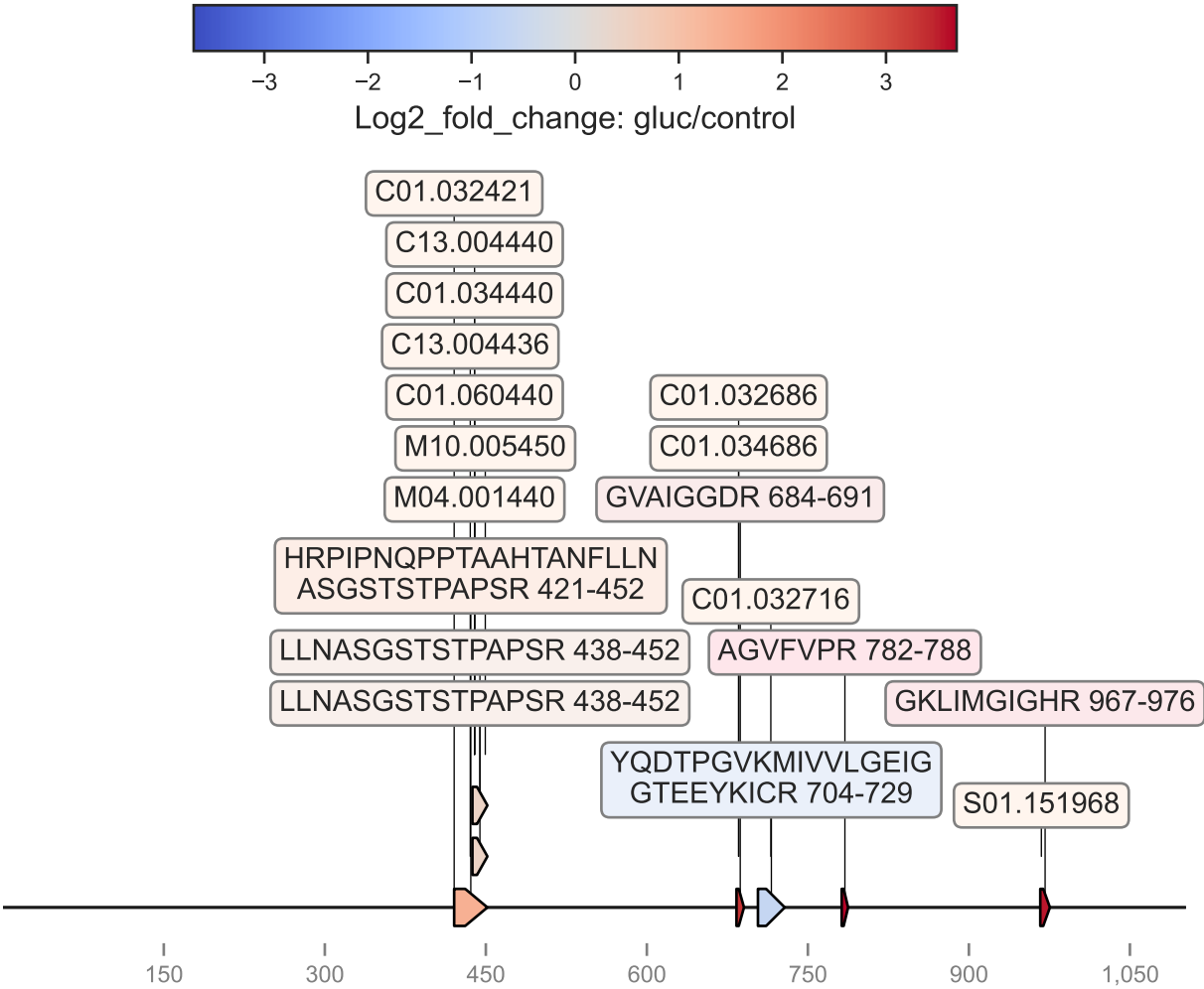


DINLQDEDWNEFNDINKIIR 285-305

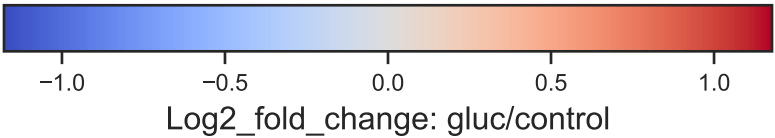
QSQLTATQTR 2036-2045



P53396-ACLY-ATP-citrate synthase

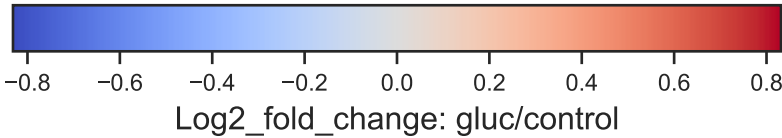


F8WAS3-NDUFA5-NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 {EC

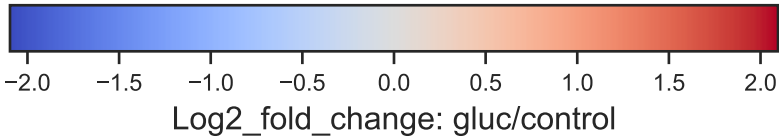


AGVLKKTGLVGLAVCNTPHER 2-23

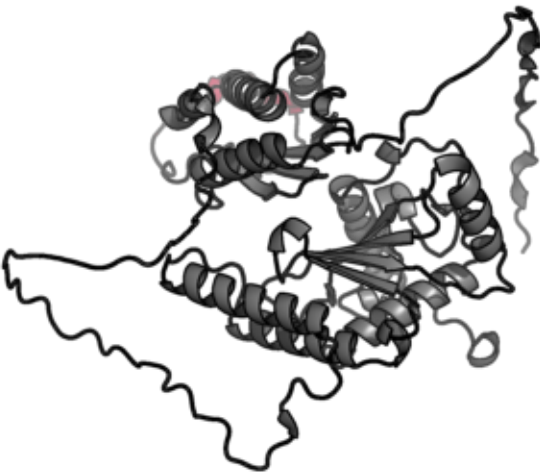
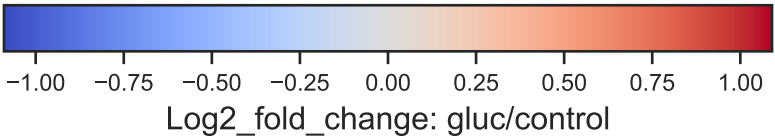




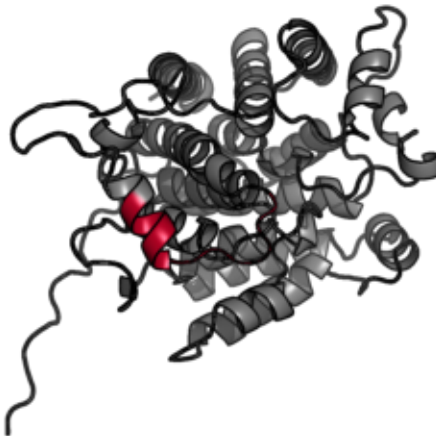
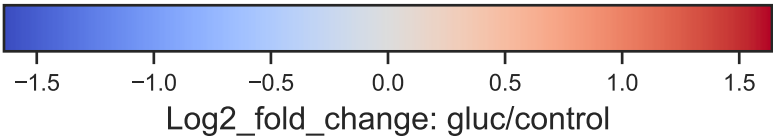
Q16643-DBN1-Drebrin



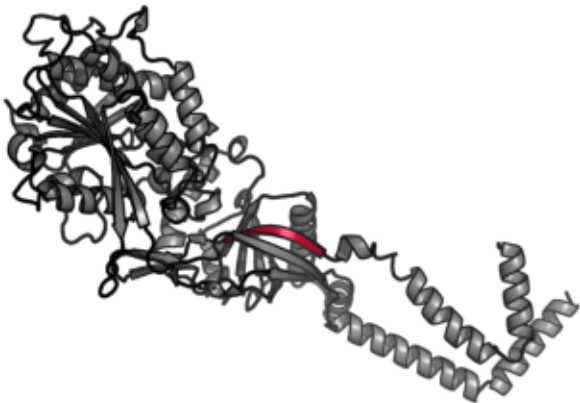
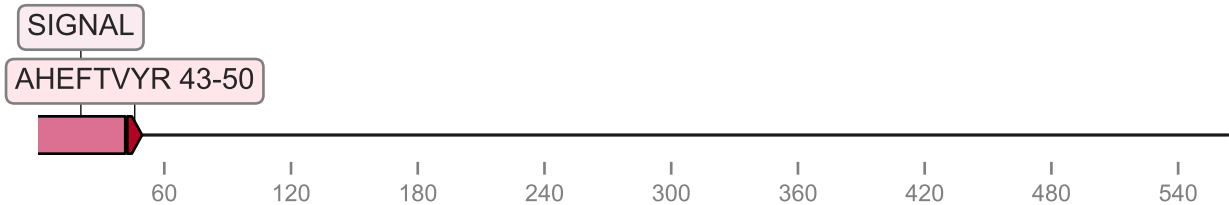
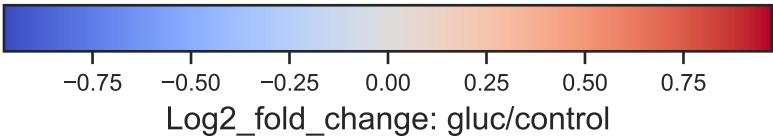
Q7Z7E8-UBE2Q1-Ubiquitin-conjugating enzyme E2 Q1



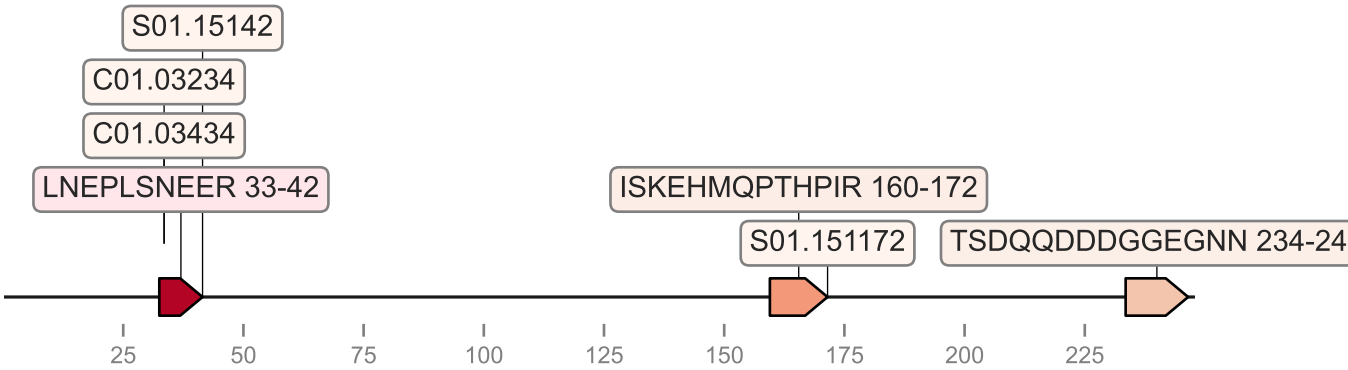
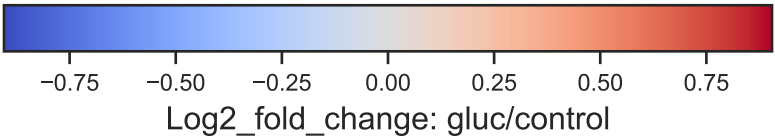
Q9NX46-ADPRS-ADP-ribosylhydrolase ARH3 {ECO:0000305}

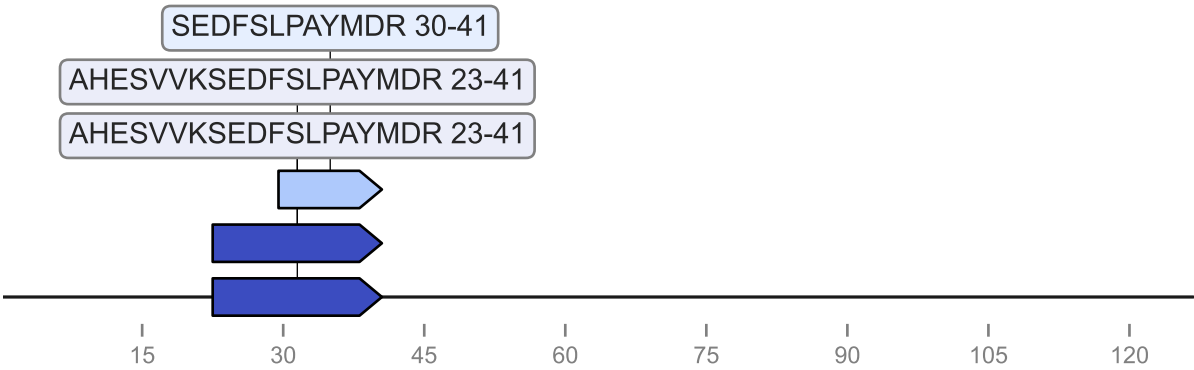
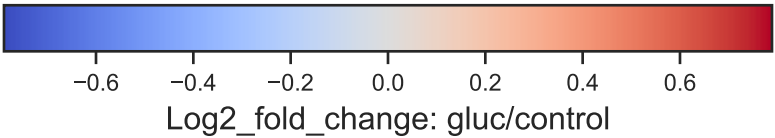


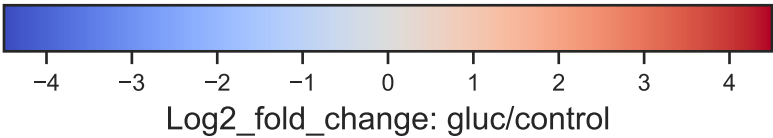
Q969V3-NCLN-BOS complex subunit NCLN {ECO:0000305}

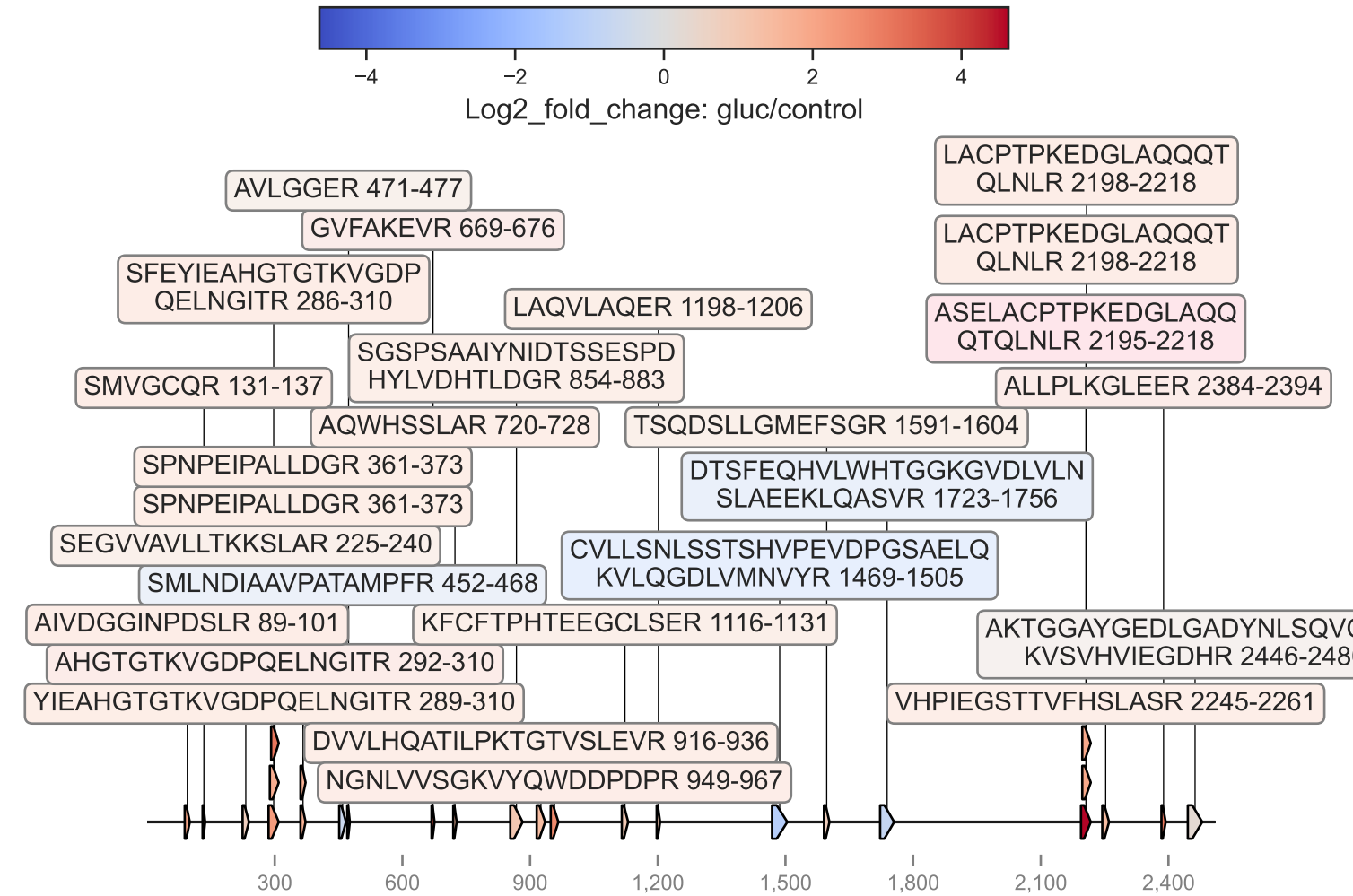


P61981-YWHAG-14-3-3 protein gamma

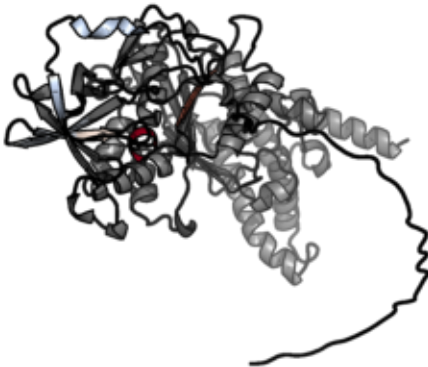
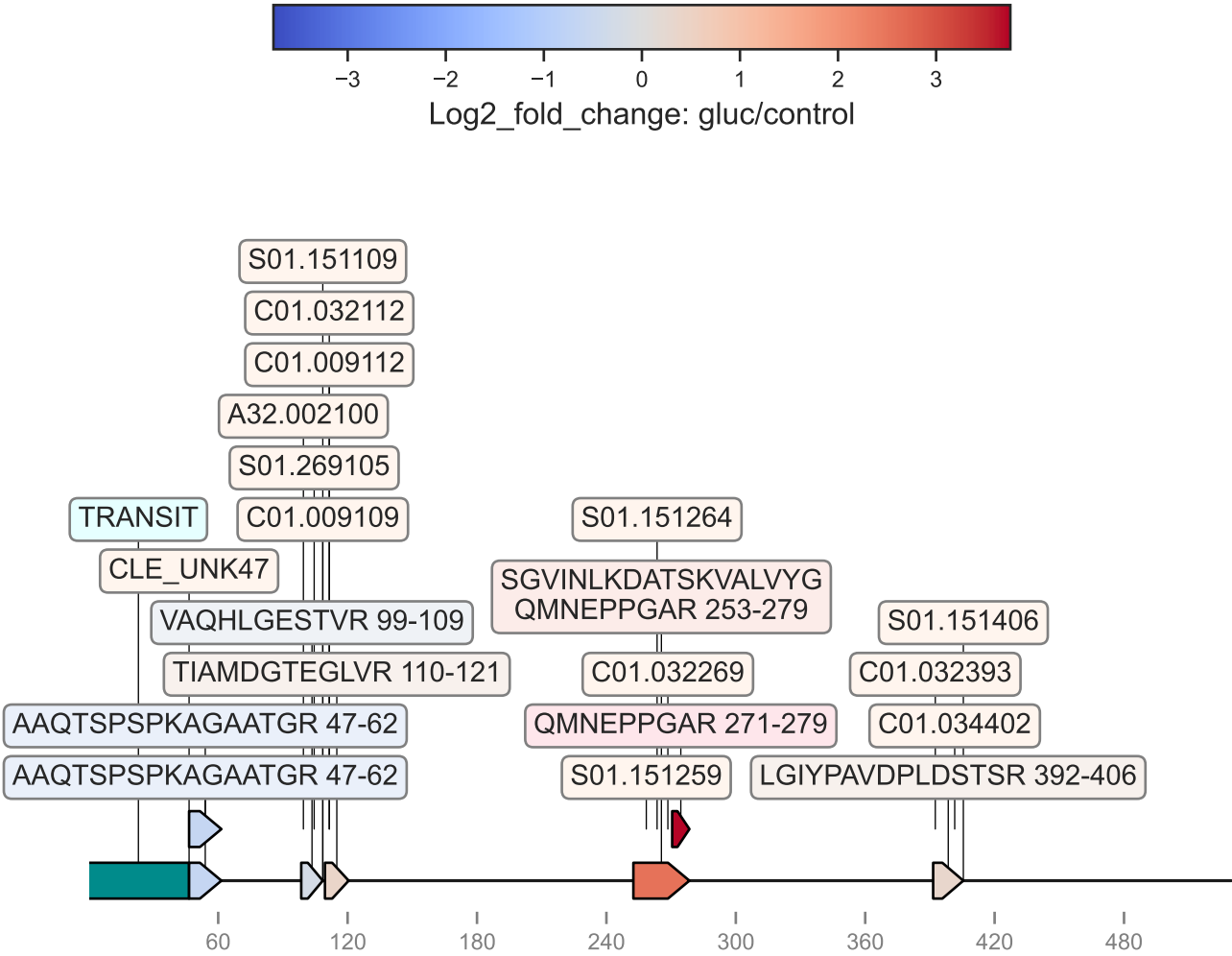




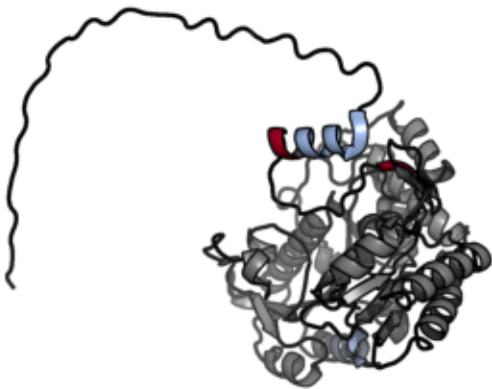
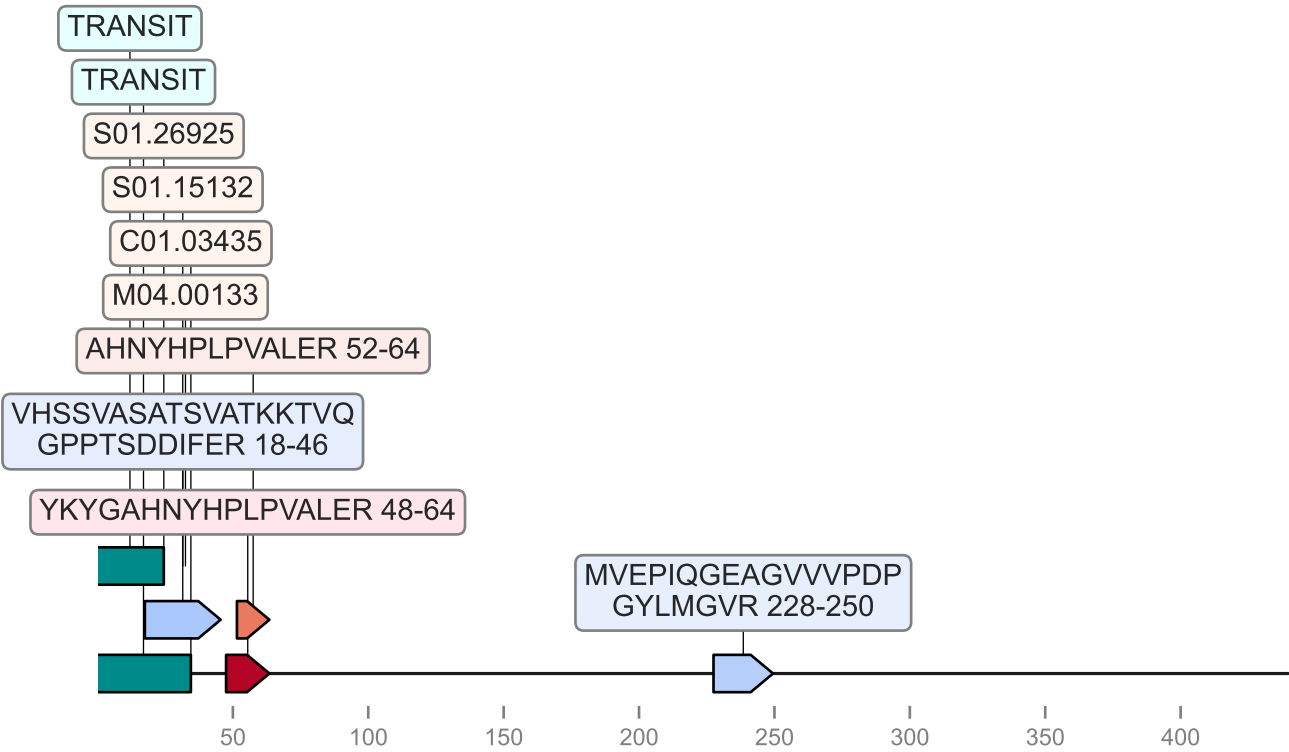
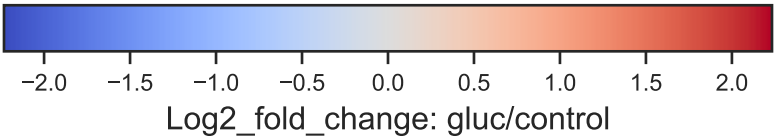




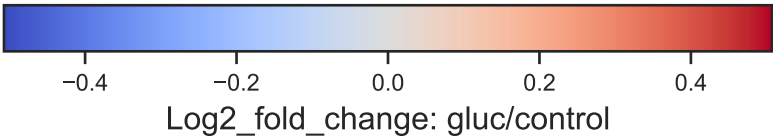
P06576-ATP5F1B-ATP synthase subunit beta, mitochondrial {ECO:0000305}



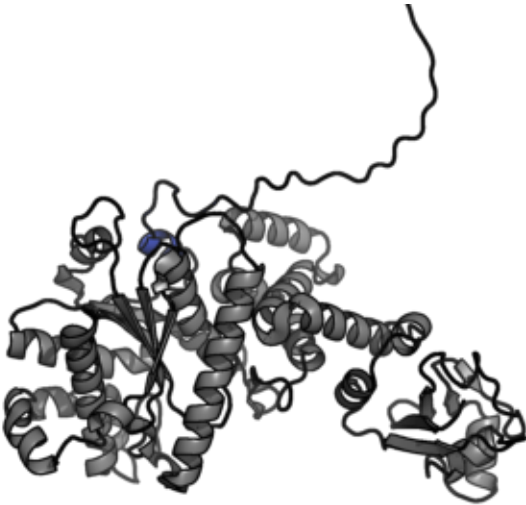
P04181-OAT-Ornithine aminotransferase, mitochondrial

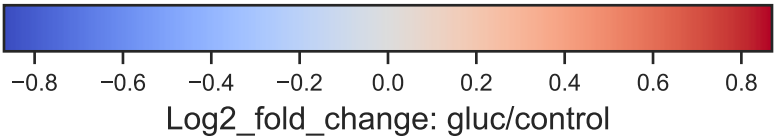


Q9Y2Z4-YARS2-Tyrosine--tRNA ligase, mitochondrial



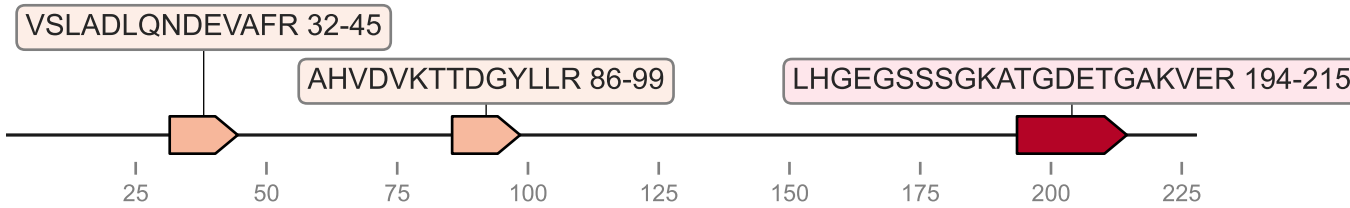
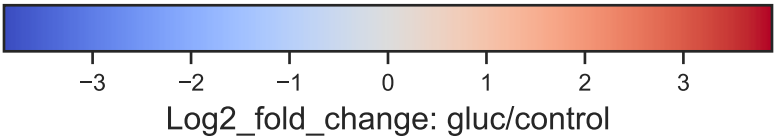
TRANSIT
AHSGAQGLLAAQKAR 31-45

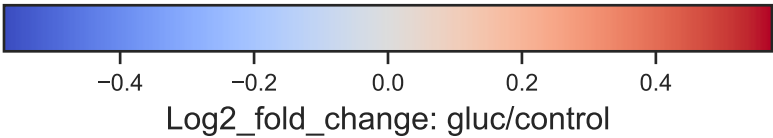




AHSPVAVQVPGMQNNIA
DPEELFTKLER 2-29



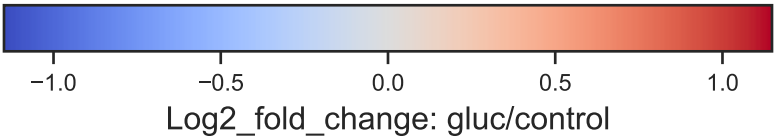




AHYKAADSKR 2-11

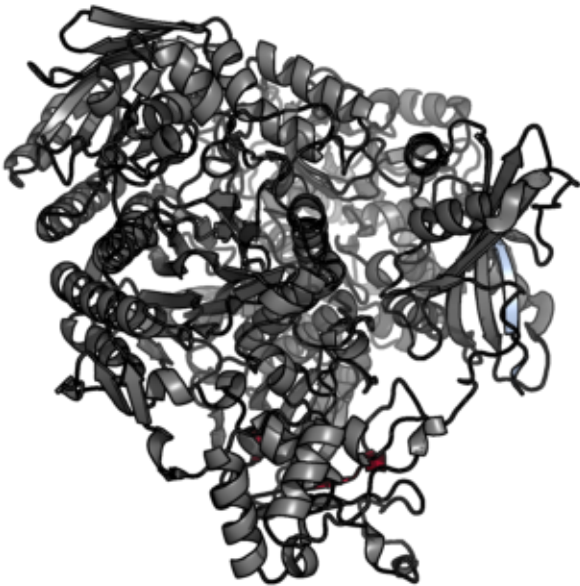


O15067-PFAS-Phosphoribosylformylglycinamide synthase

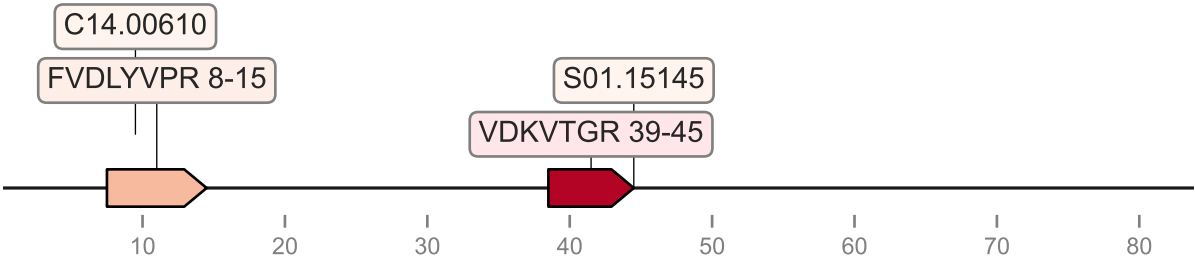
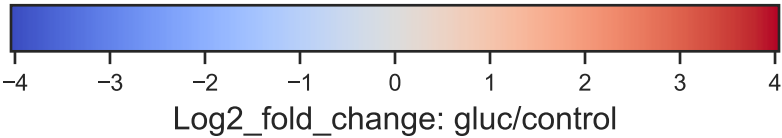


SPVLHFYVR 2-10

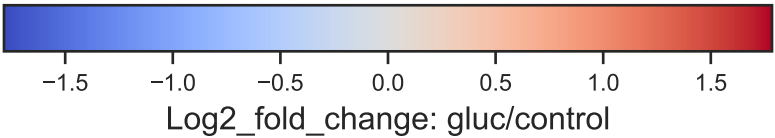
LSDPAGAIITSR 529-541



P63220-RPS21-40S ribosomal protein S21



Q01105-2-SET-Isoform 2 of Protein SET



TSEKEQQEAIEHIDEVQNEIDR 23-44

