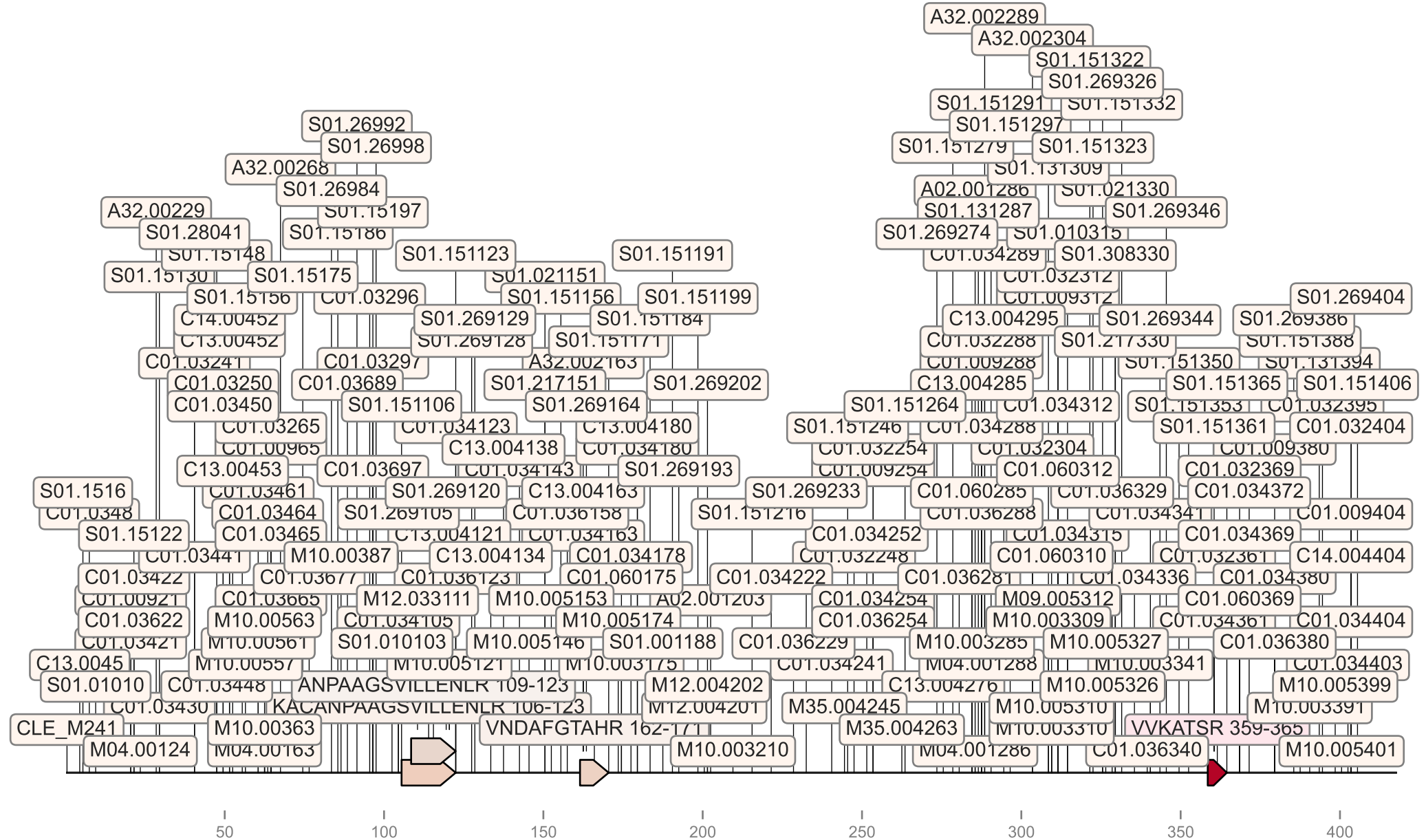
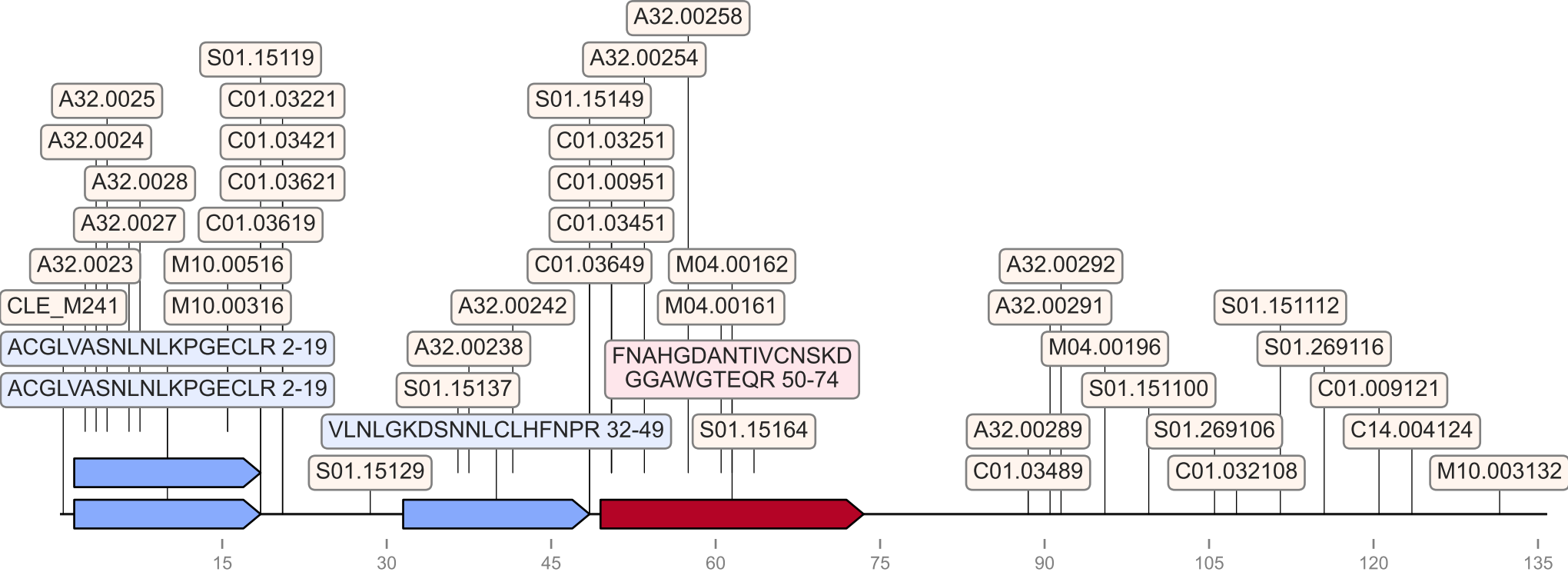


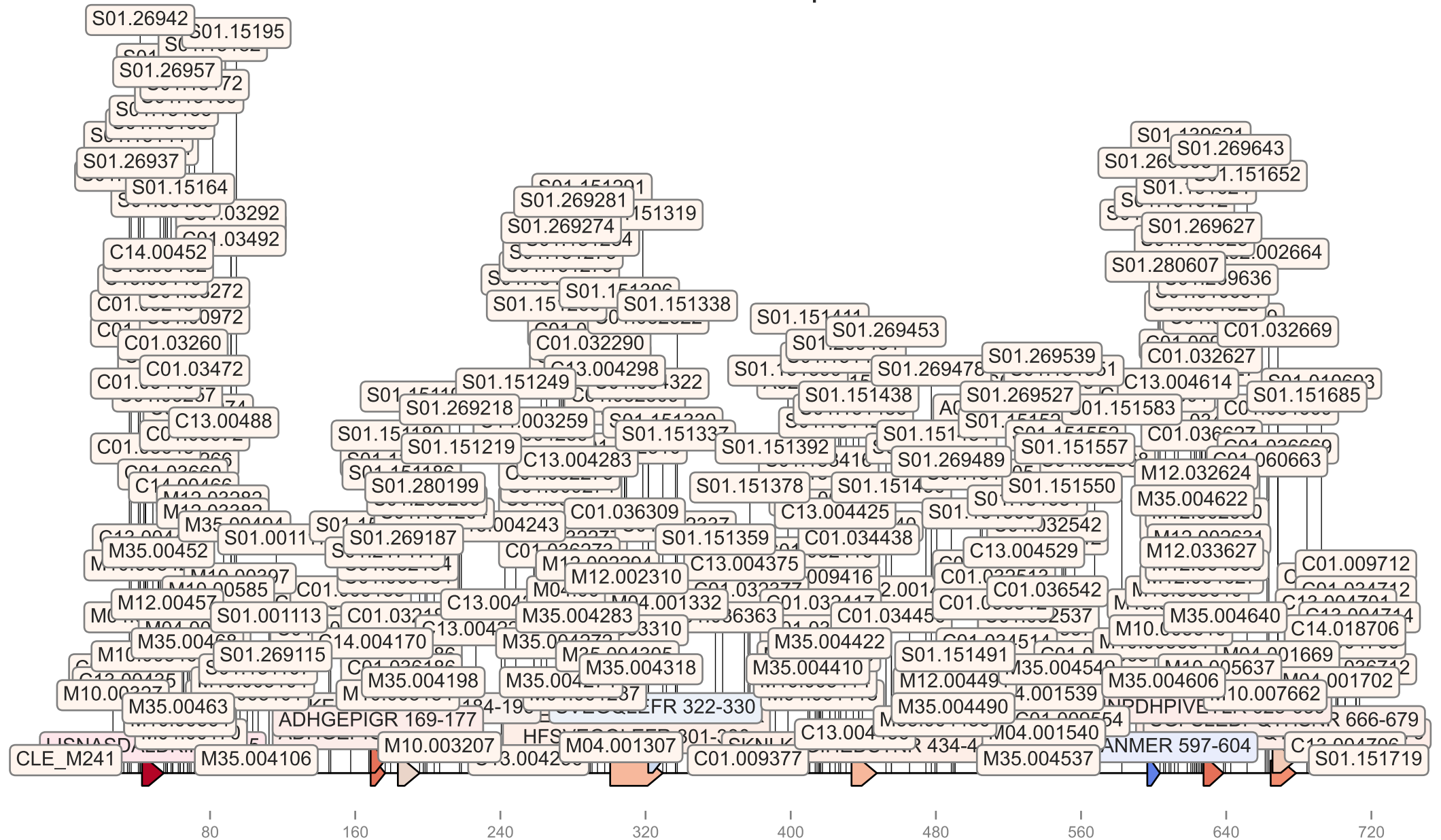
## P00558-PGK1-Phosphoglycerate kinase 1



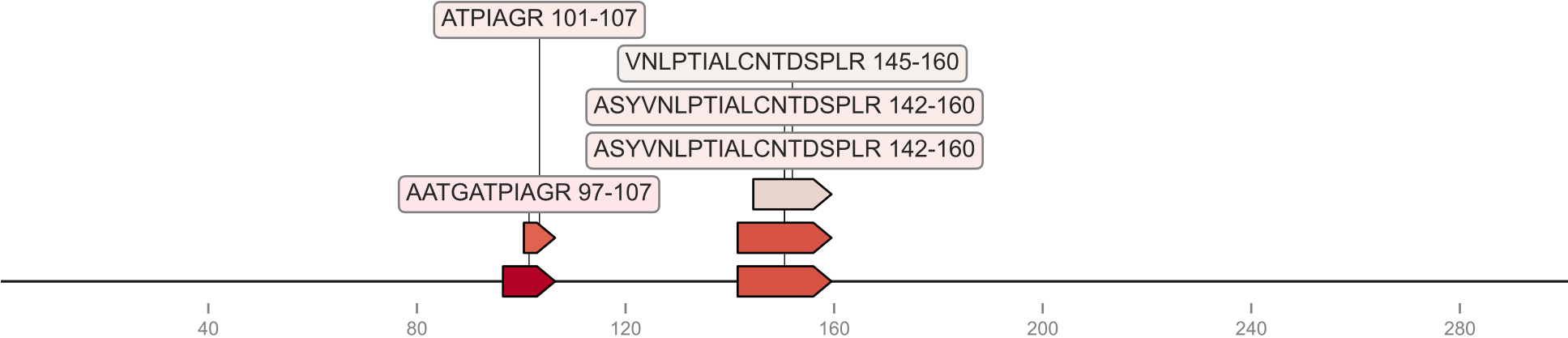
P09382-LGALS1-Galectin-1



# P08238-HSP90AB1-Heat shock protein HSP 90-beta



A0A0C4DG17-RPSA-40S ribosomal protein SA {ECO:0000256|HAMAP-Rule:MF\_03016}



4-Nucleosome assembly protein 1-like 4 {ECO:0000313|Ensembl:ENSP00000405912.1, ECO:0000313|Ensembl:ENS

ADHSFSDGVPSDSVEAAKNA  
SNTTEKLTQVMQNPR 2-36

ADHSFSDGVPSDSVEAAKNA  
SNTTEKLTQVMQNPR 2-36



20

40

60

80

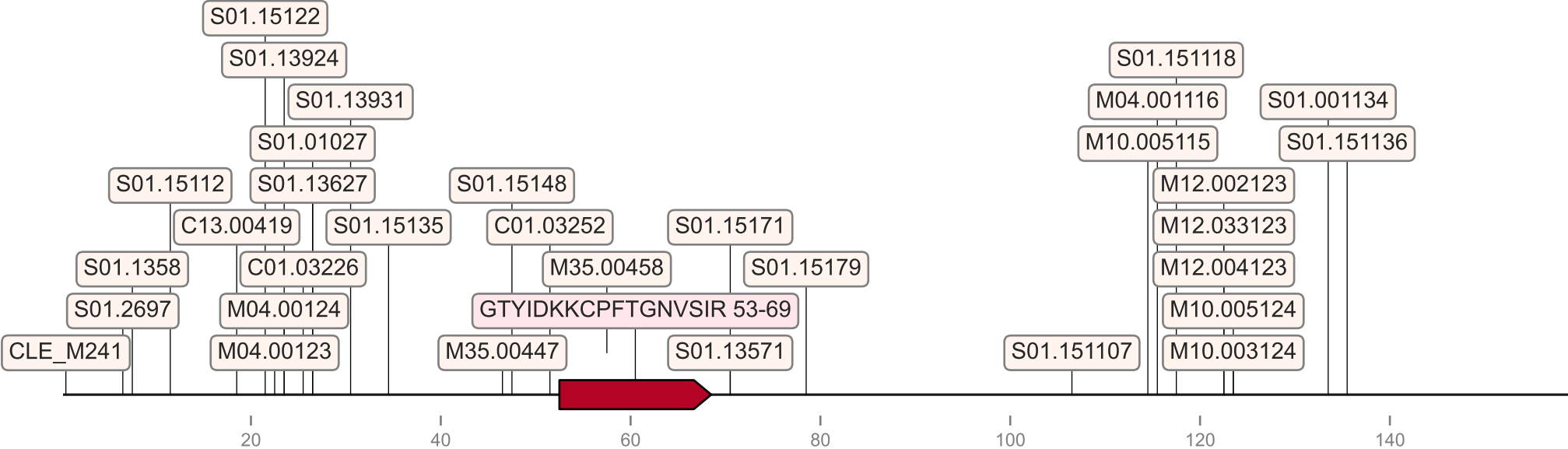
100

120

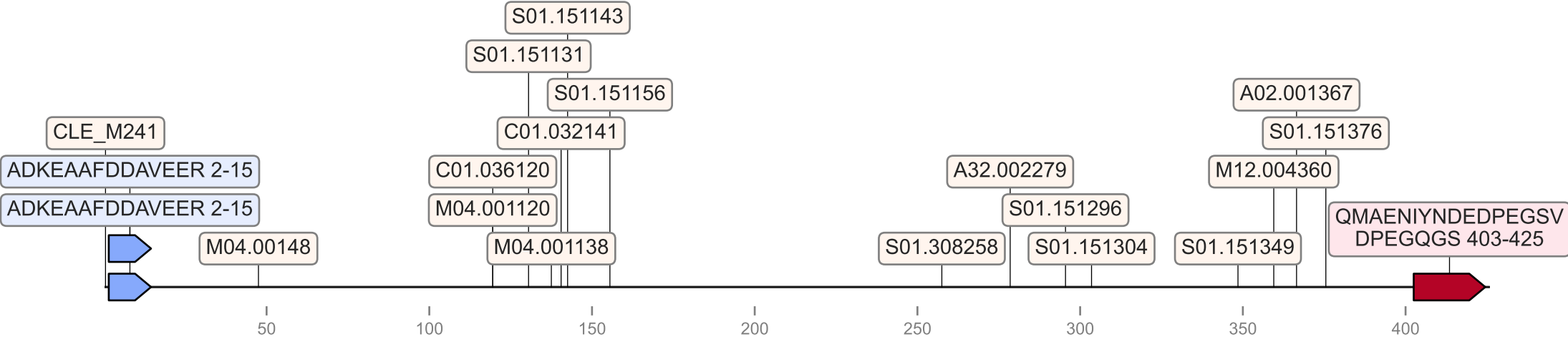
140

160

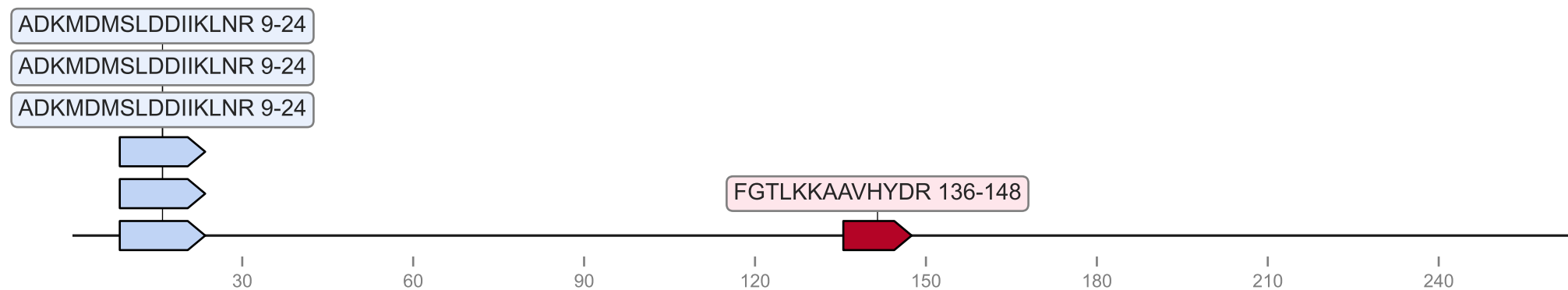
P62280-RPS11-40S ribosomal protein S11



Q09028-RBBP4-Histone-binding protein RBBP4

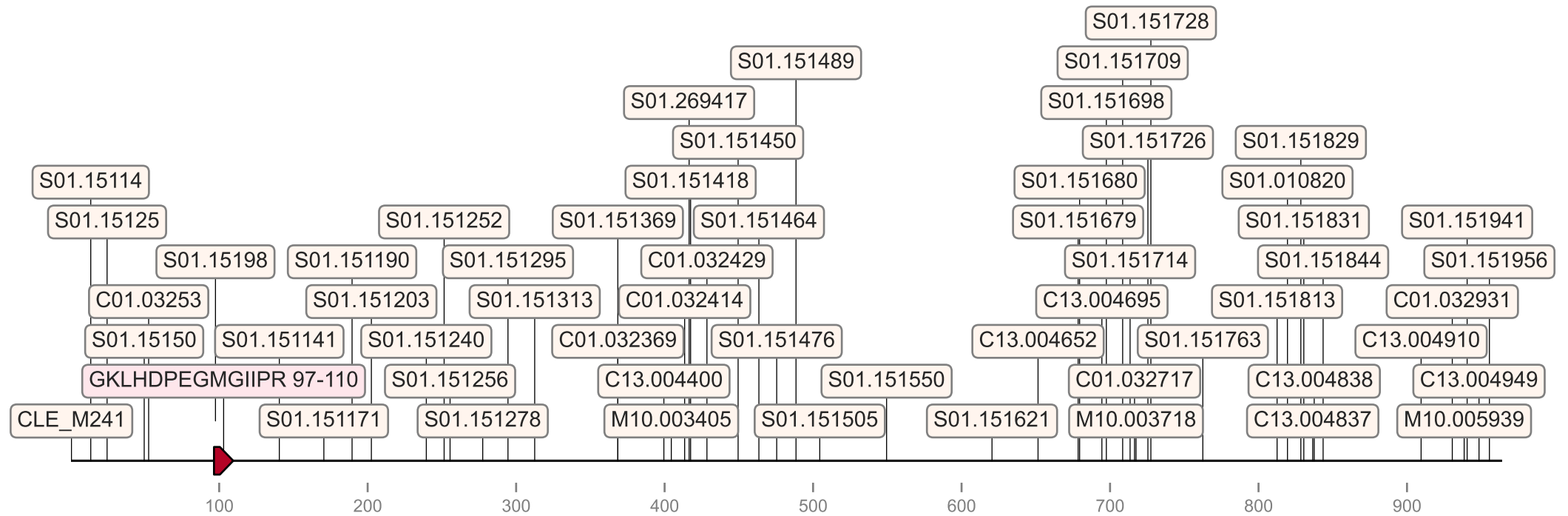


# E9PB61-ALYREF-Aly/REF export factor {ECO:0000313|Ensembl:ENSP00000421592.2}

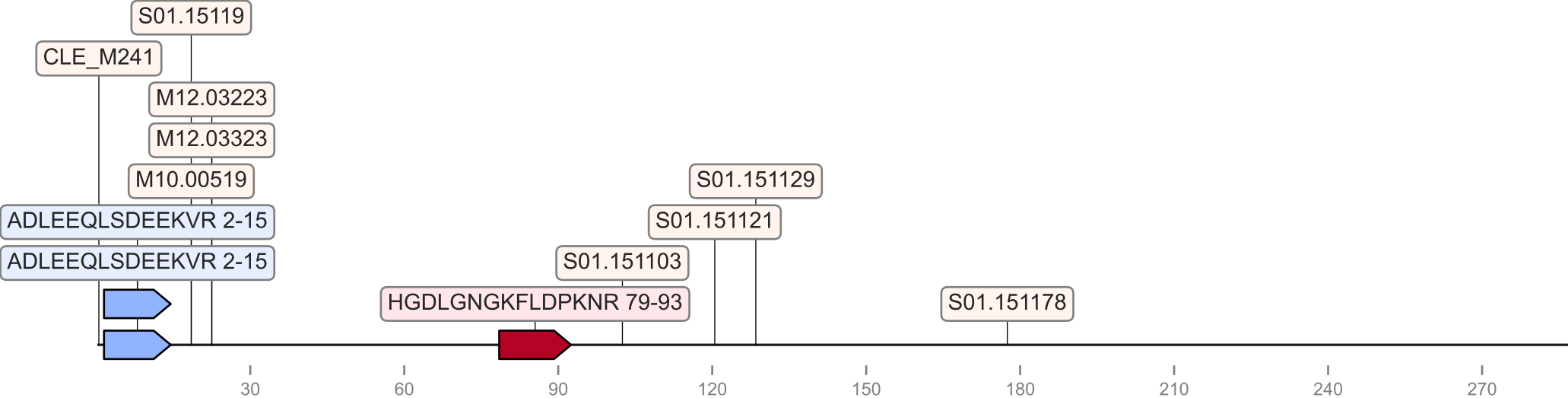




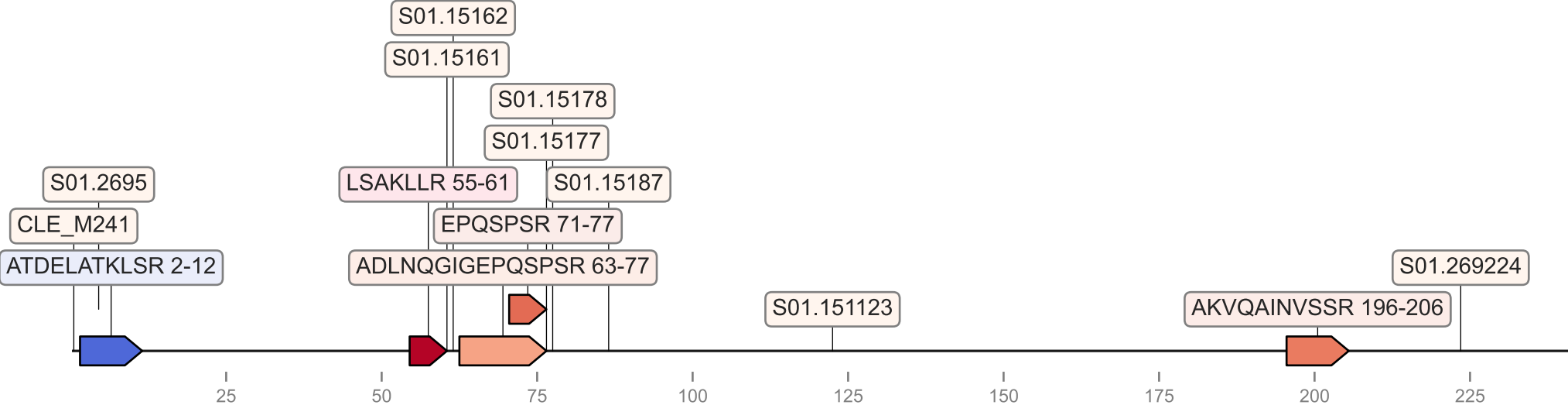
P33176-KIF5B-Kinesin-1 heavy chain {ECO:0000305}



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



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

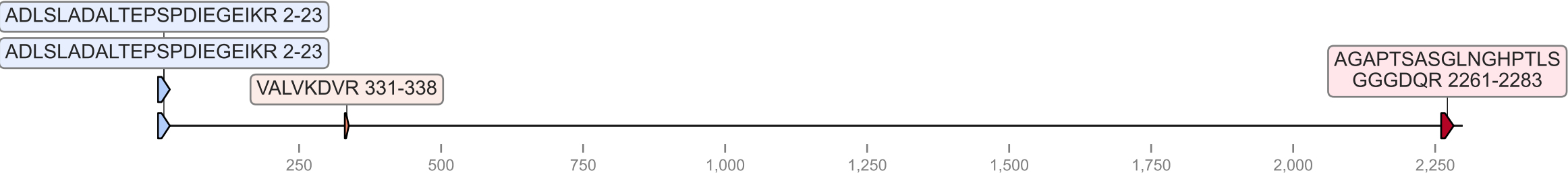


# Q09161-NCBP1-Nuclear cap-binding protein subunit 1

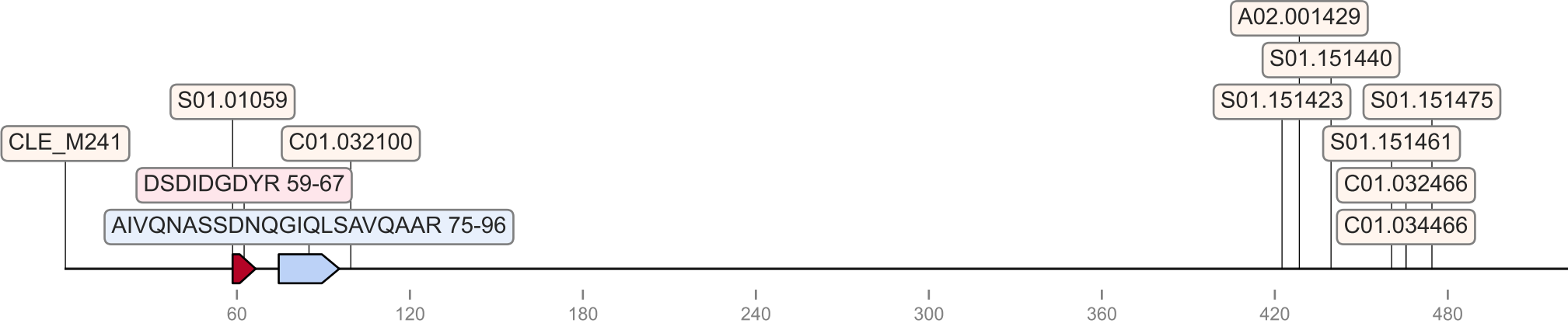
ADLPNYKSKILR 59-70



E7EVA0-MAP4-Microtubule-associated protein {ECO:0000256|RuleBase:RU000686}



O00629-KPNA4-Importin subunit alpha-3



C9J329-GTF2E1-General transcription factor IIE subunit 1 {ECO:0000313|Ensembl:ENSP00000417342.1}

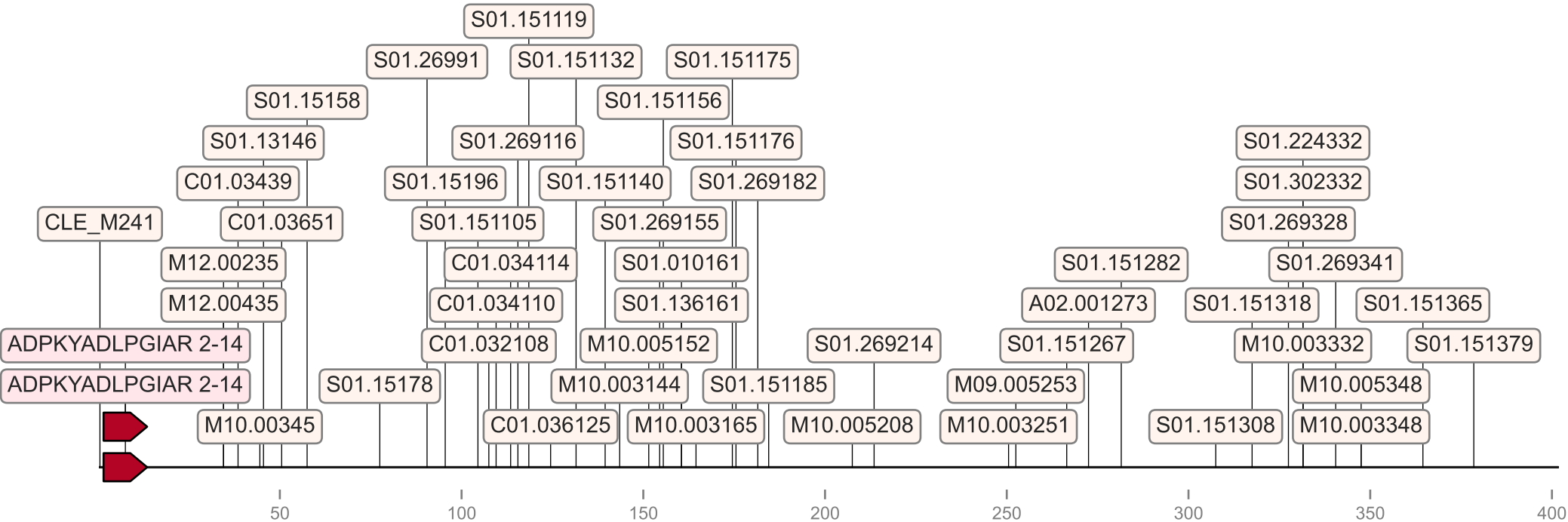


A0A087X2B5-BSG-Basigin (Ok blood group) {ECO:0000313|Ensembl:ENSP00000484849.2}

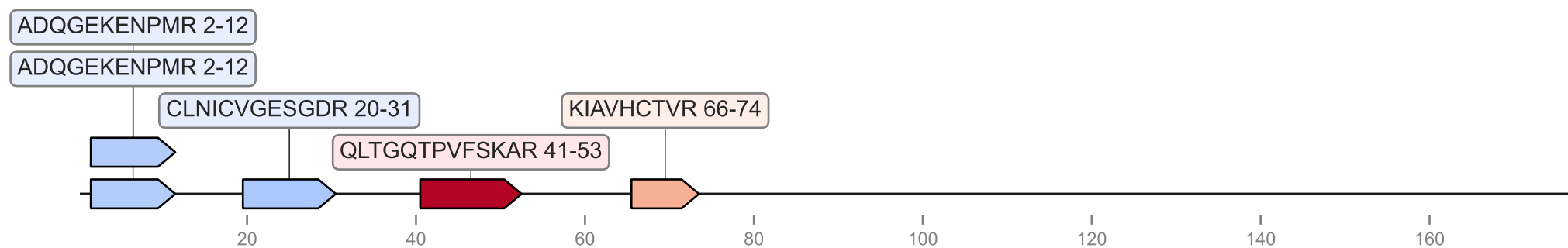




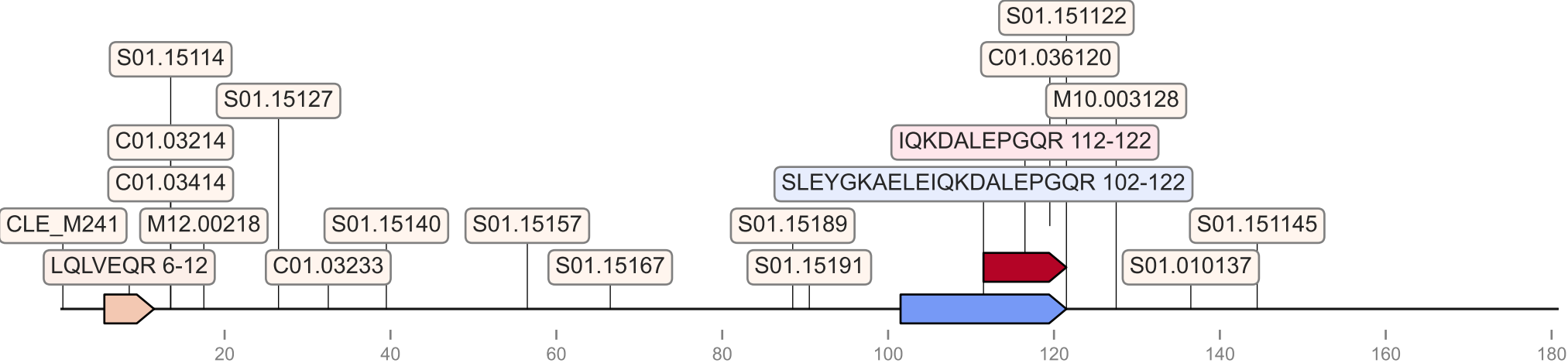
Q13561-DCTN2-Dynactin subunit 2



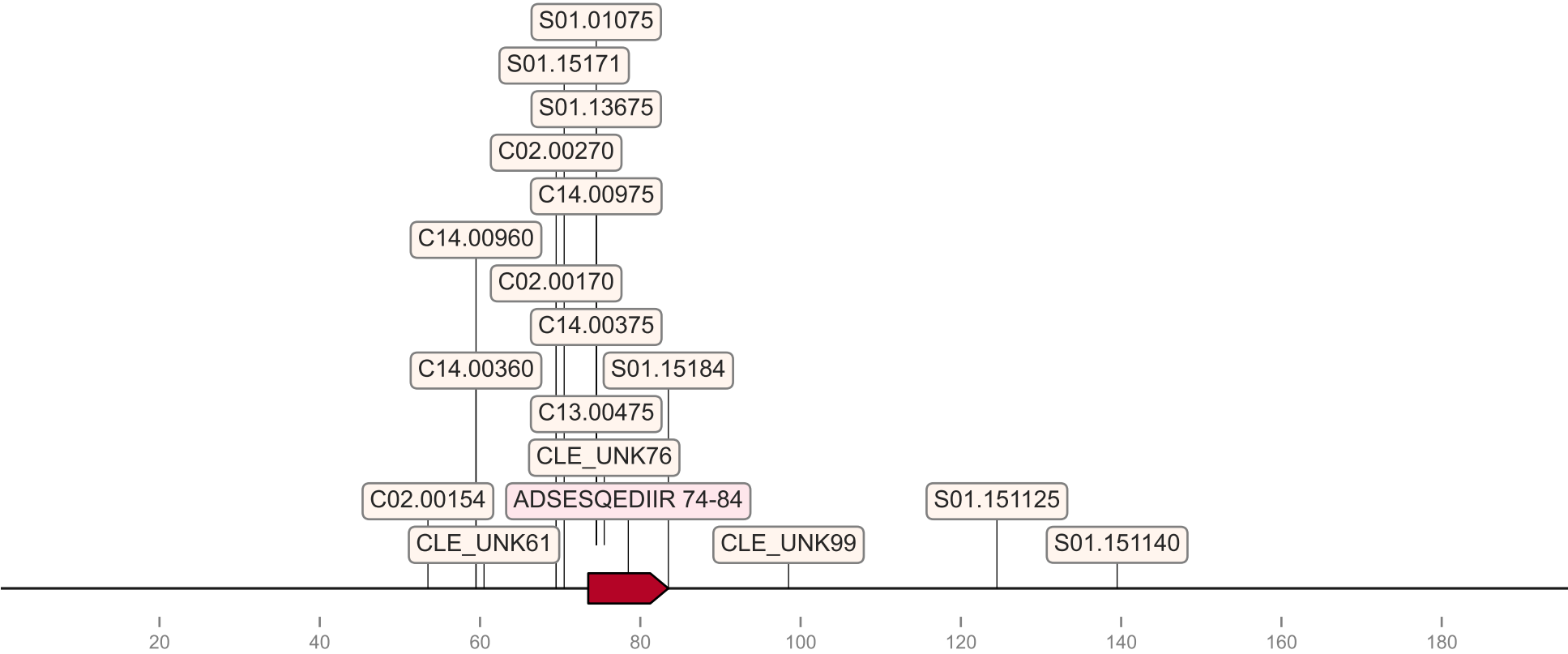
## P62913-2-RPL11-Isoform 2 of 60S ribosomal protein L11



P07741-APRT-Adenine phosphoribosyltransferase {ECO:0000305}



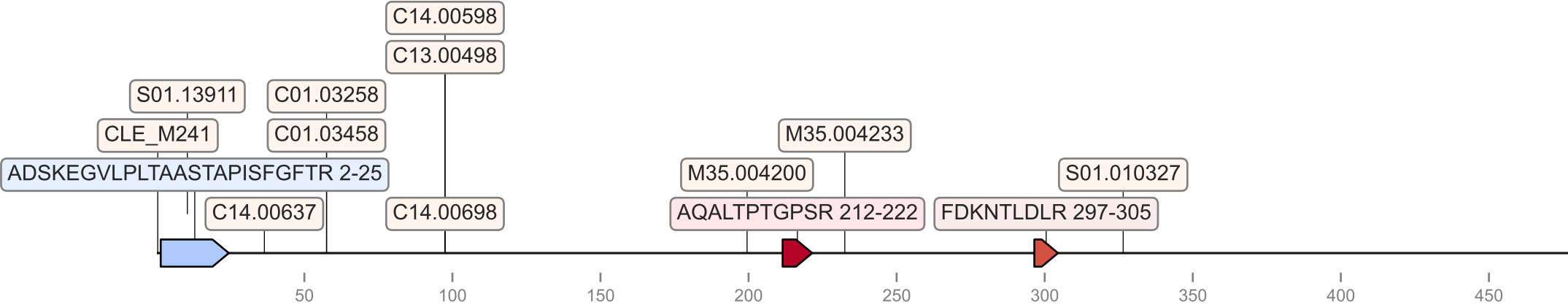
P55957-BID-BH3-interacting domain death agonist



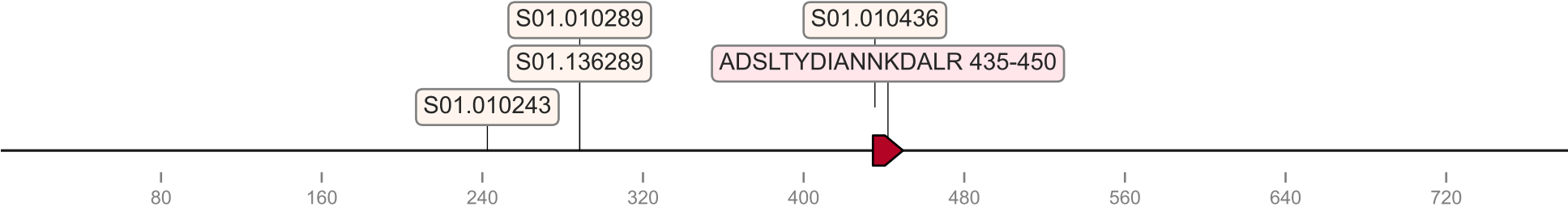
Q96C90-PPP1R14B-Protein phosphatase 1 regulatory subunit 14B



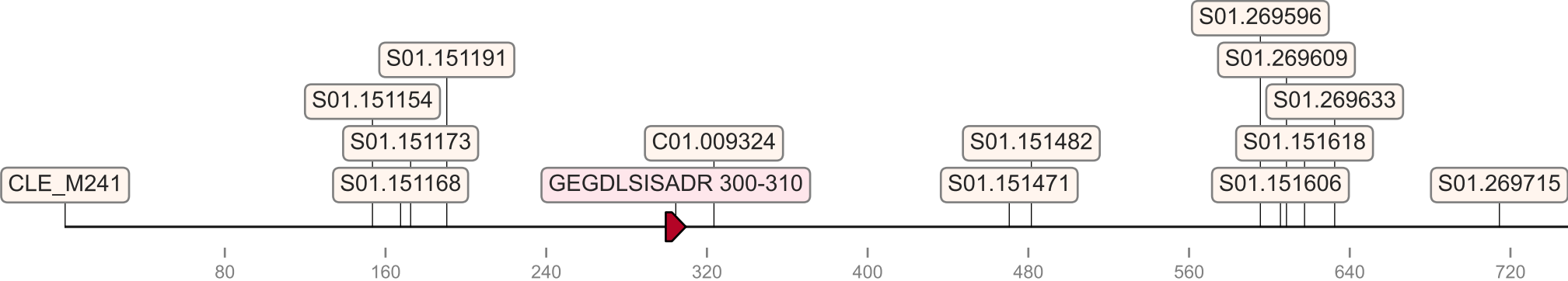
Q92917-GPKOW-G-patch domain and KOW motifs-containing protein



O43815-STRN-Striatin

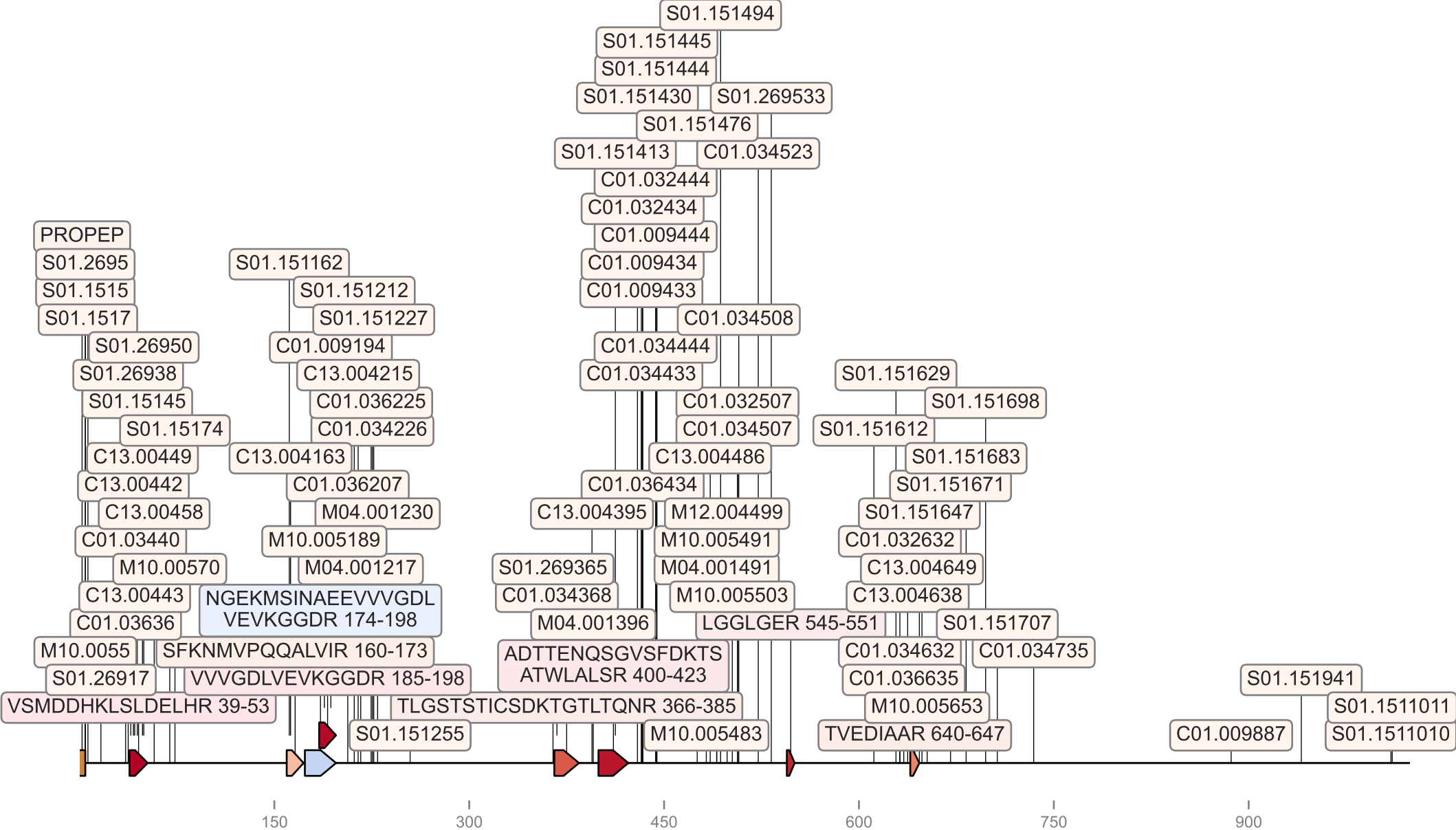


P49589-CARS1-Cysteine--tRNA ligase, cytoplasmic

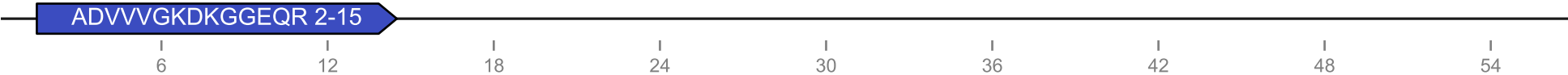




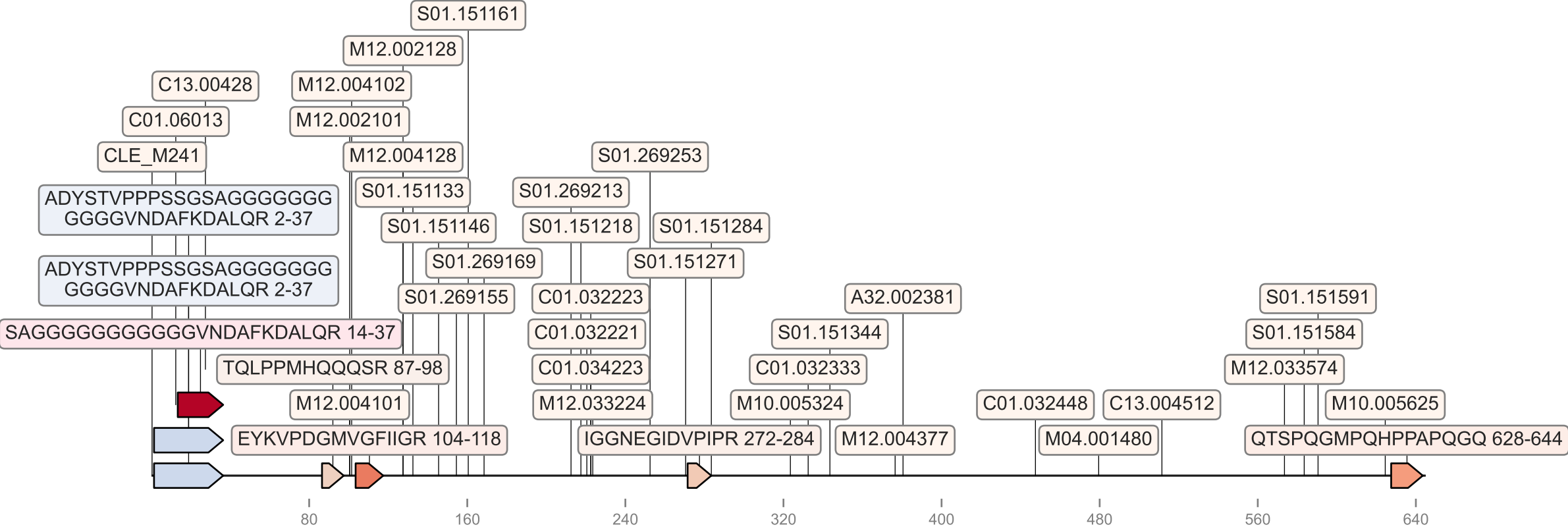
P05023-ATP1A1-Sodium/potassium-transporting ATPase subunit alpha-1



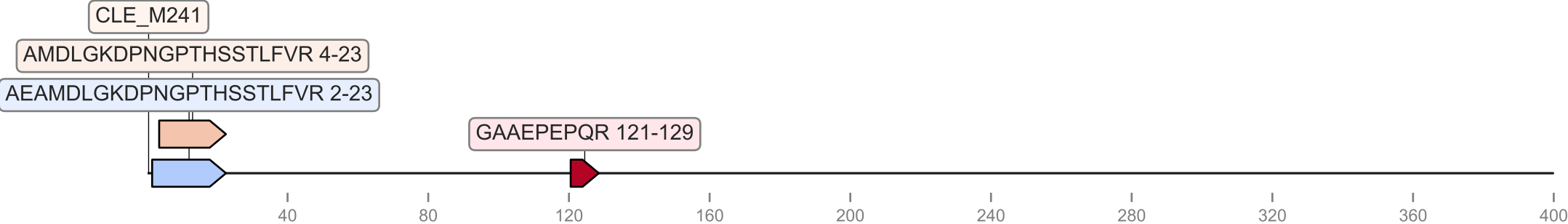
E5RGS9-CHRA1-Chromatin accessibility complex subunit 1 {ECO:0000313|Ensembl:ENSP00000428697.1}



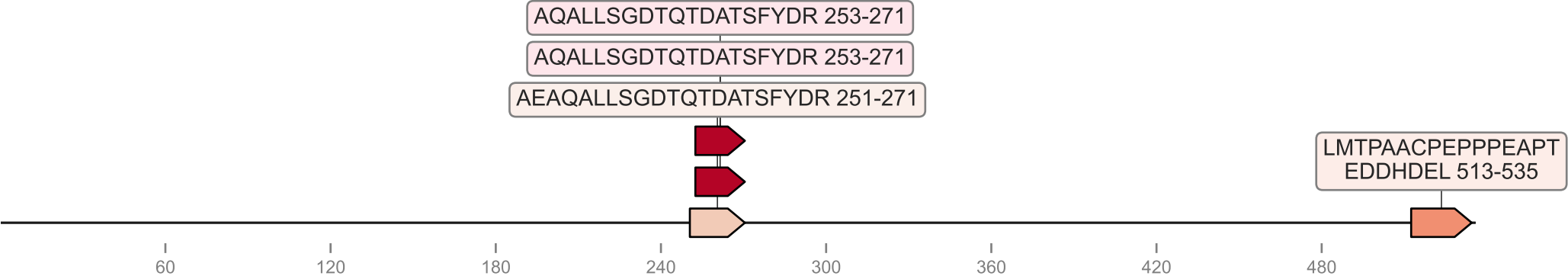
Q96AE4-FUBP1-Far upstream element-binding protein 1



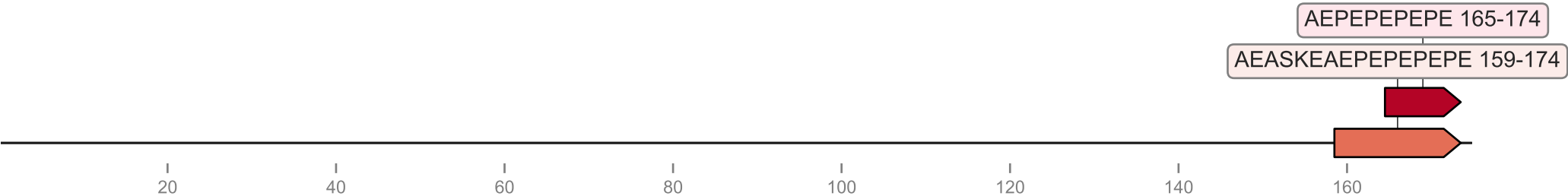
Q9NYB0-TERF2IP-Telomeric repeat-binding factor 2-interacting protein 1



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



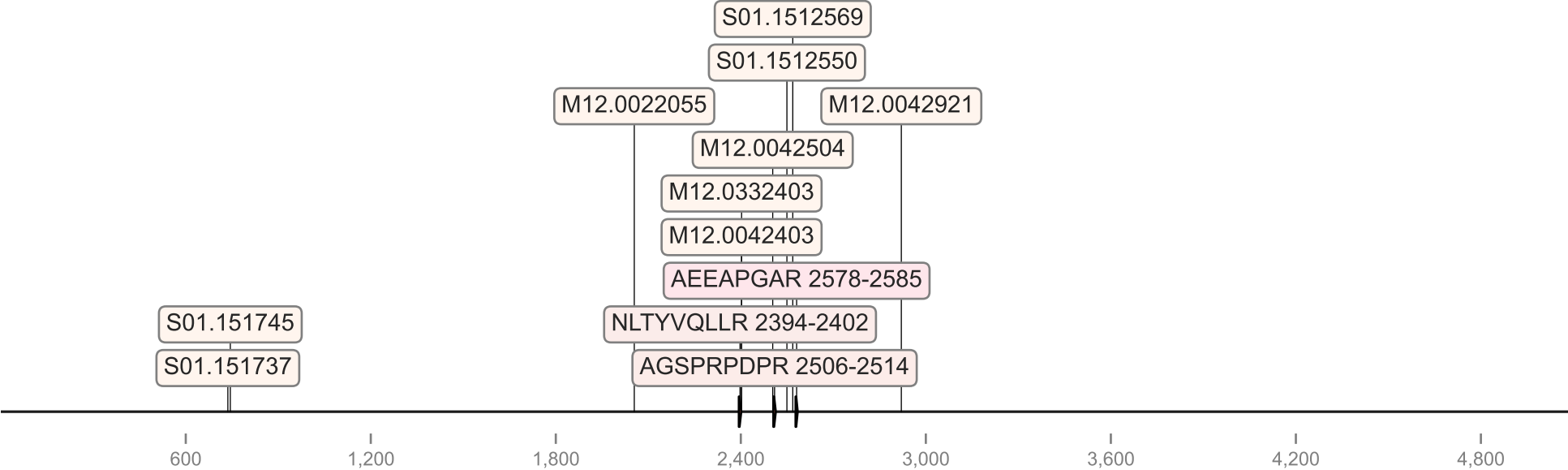
H3BUP2-NOL3-Nucleolar protein 3 {ECO:0000313|Ensembl:ENSP00000457732.1}



3-SNRPD3-Small nuclear ribonucleoprotein Sm D3 {ECO:0000256|ARBA:ARBA00020160, ECO:0000256|RuleBase:R

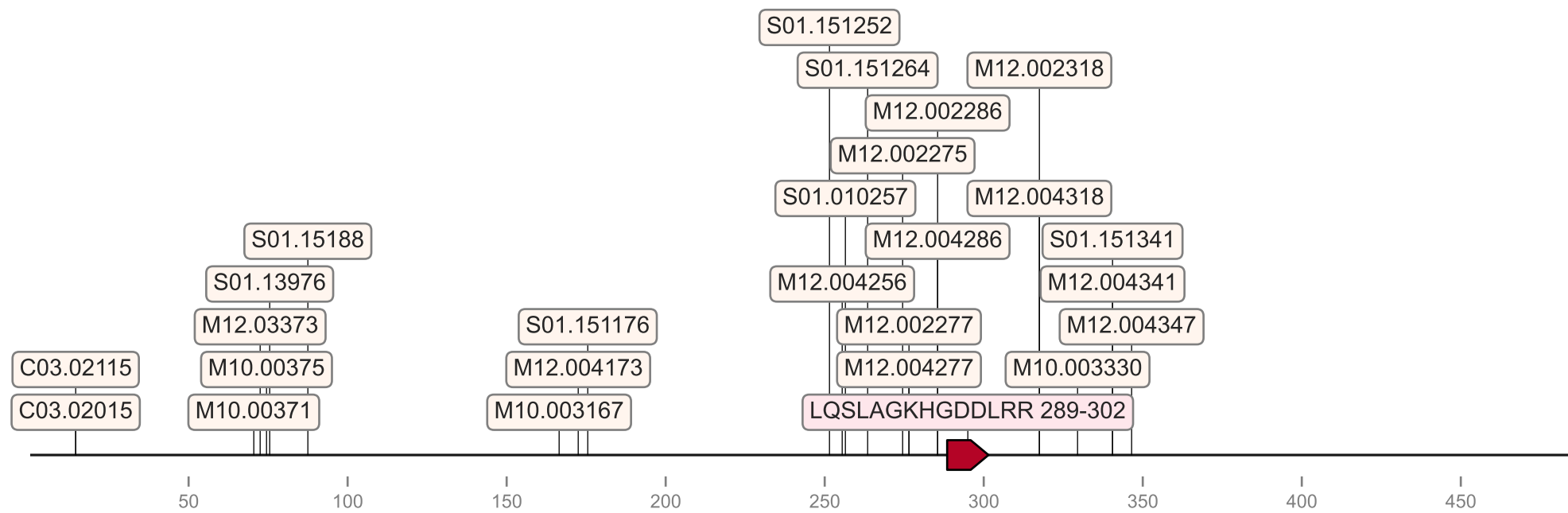


P58107-EPPK1-Epiplakin {ECO:0000303|PubMed:11278896}

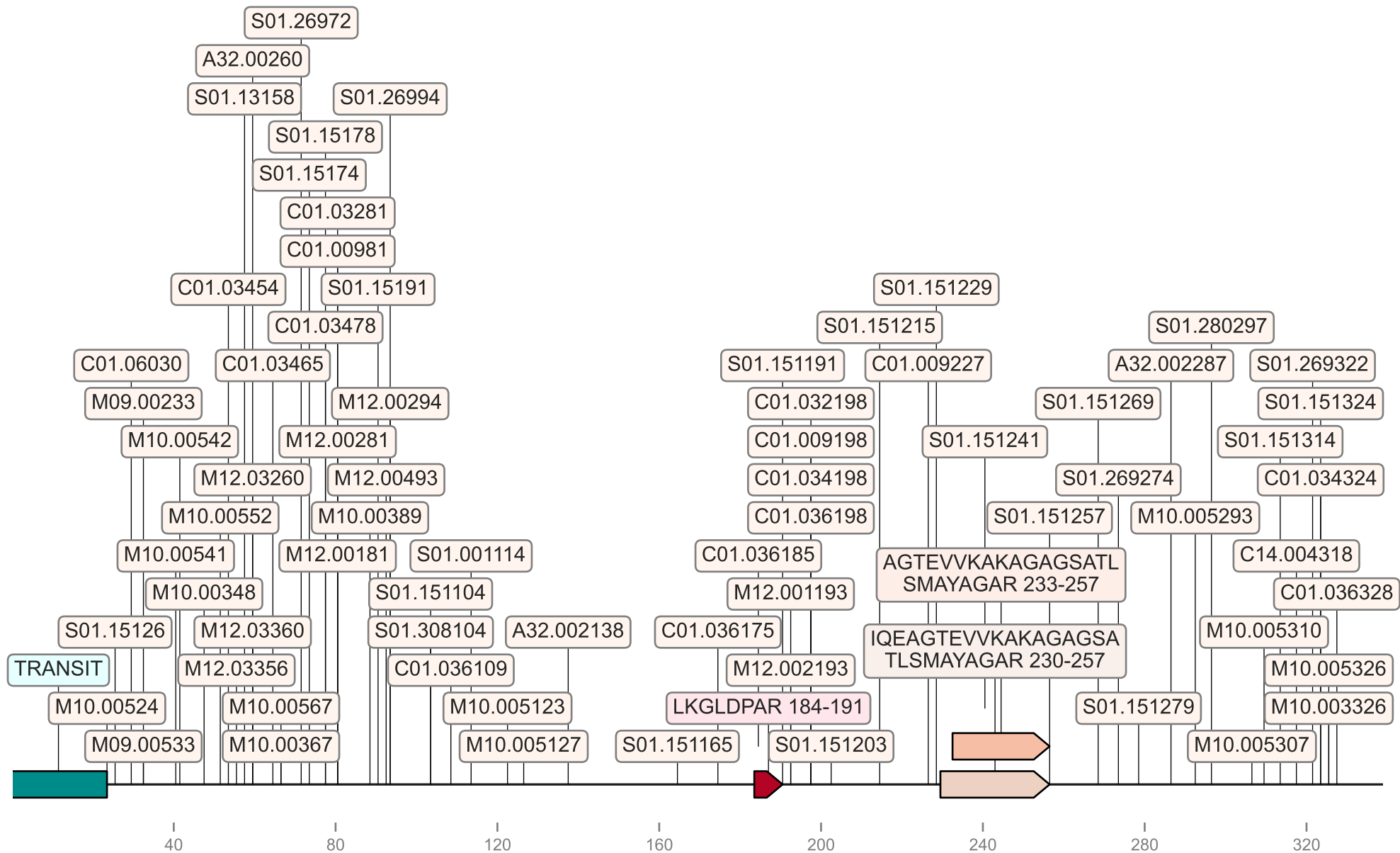




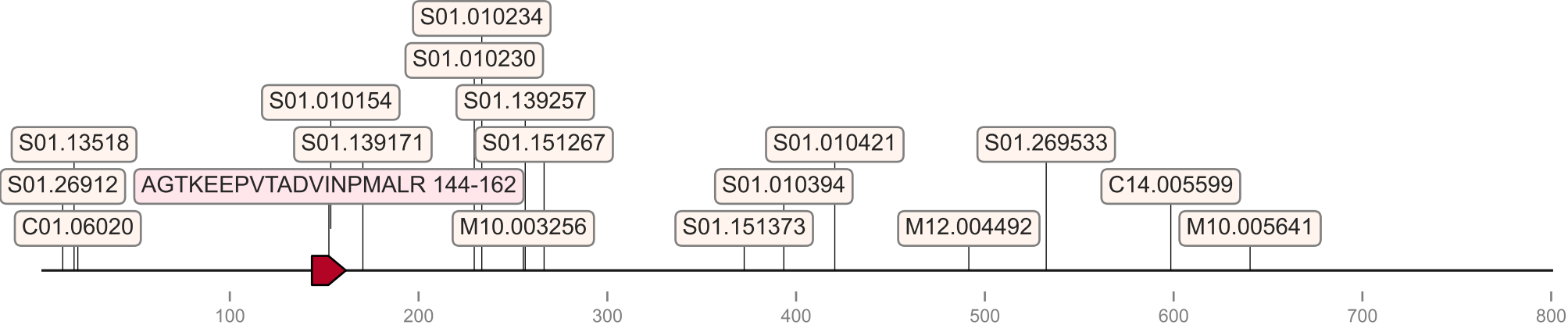
P05787-KRT8-Keratin, type II cytoskeletal 8



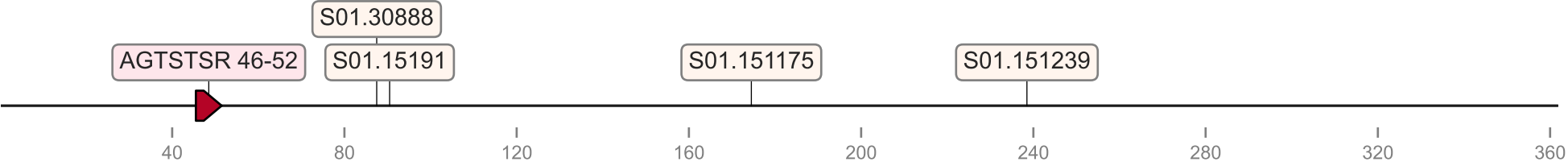
P40926-MDH2-Malate dehydrogenase, mitochondrial



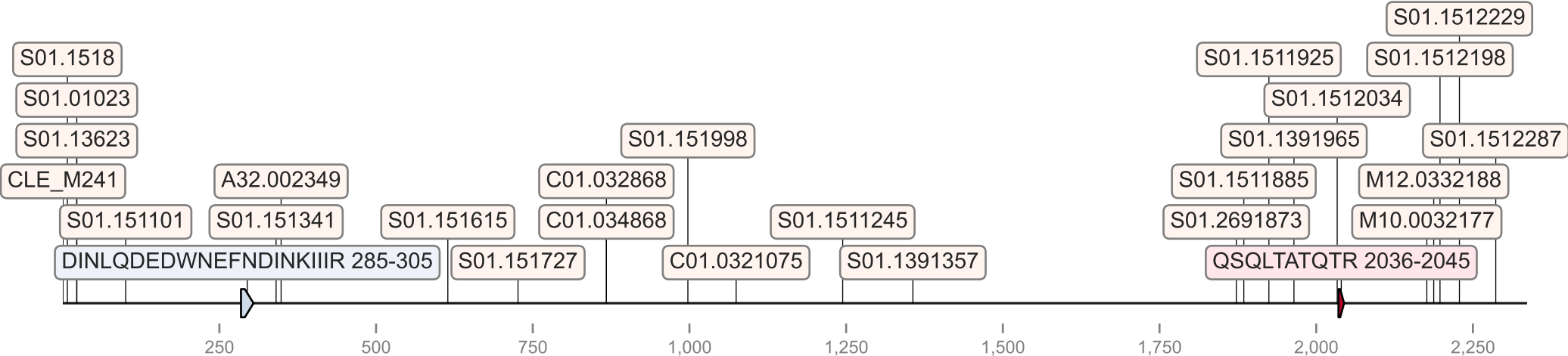
O43290-SART1-U4/U6.U5 tri-snRNP-associated protein 1



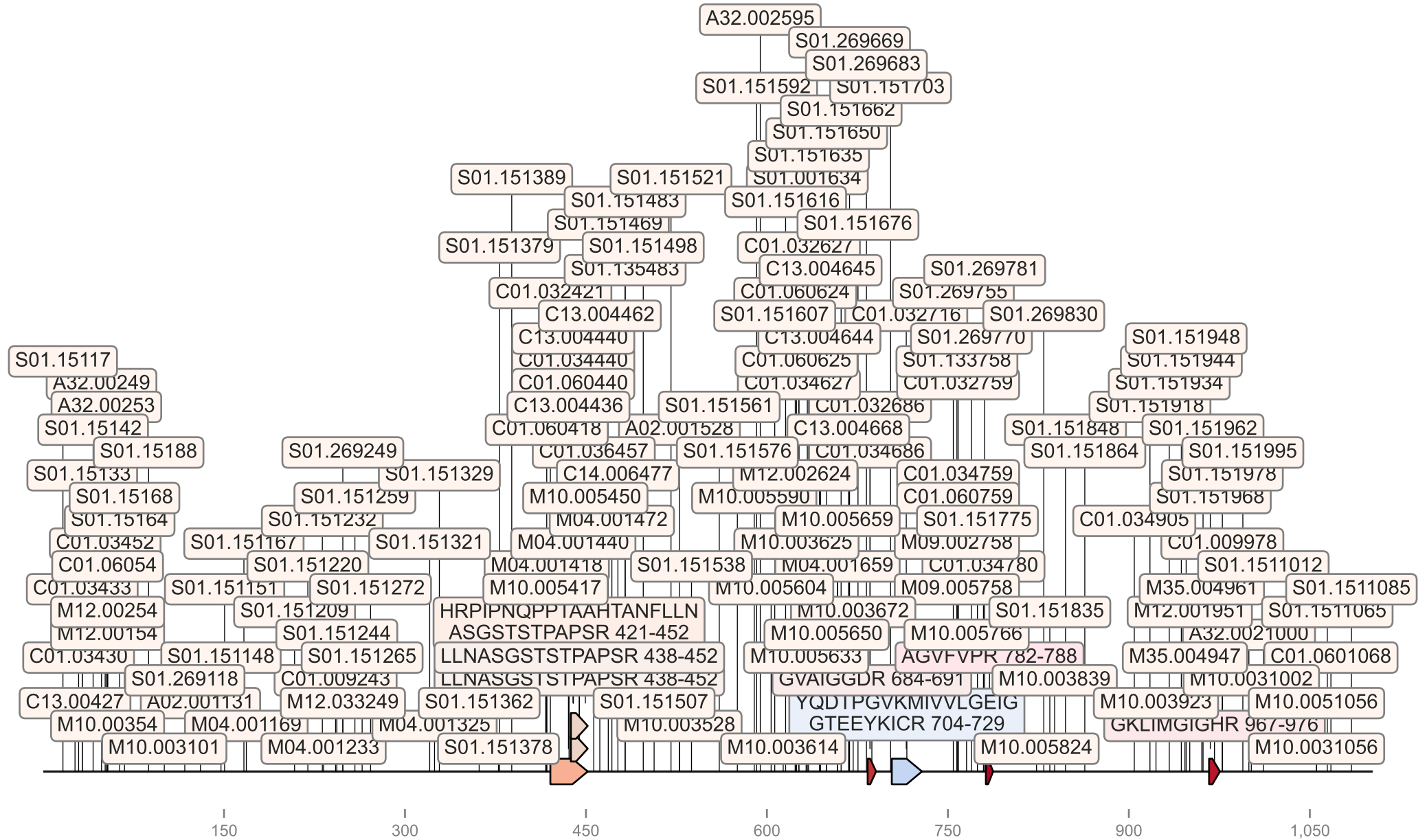
O96008-TOMM40-Mitochondrial import receptor subunit TOM40 homolog



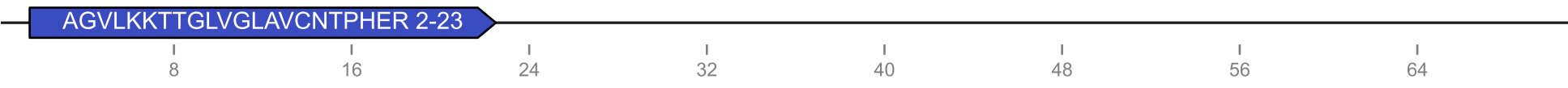
Q6P2Q9-PRPF8-Pre-mRNA-processing-splicing factor 8



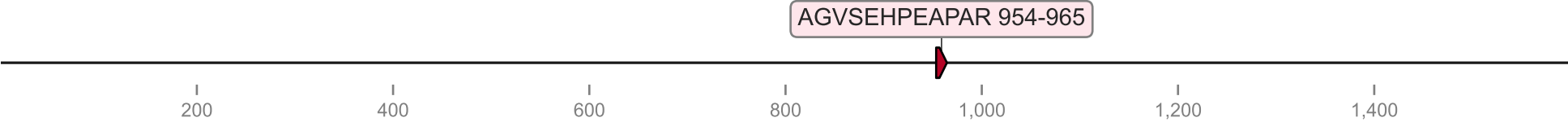
## P53396-ACLY-ATP-citrate synthase



WAS3-NDUFA5-NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 {ECO:0000256|ARBA:ARBA00010

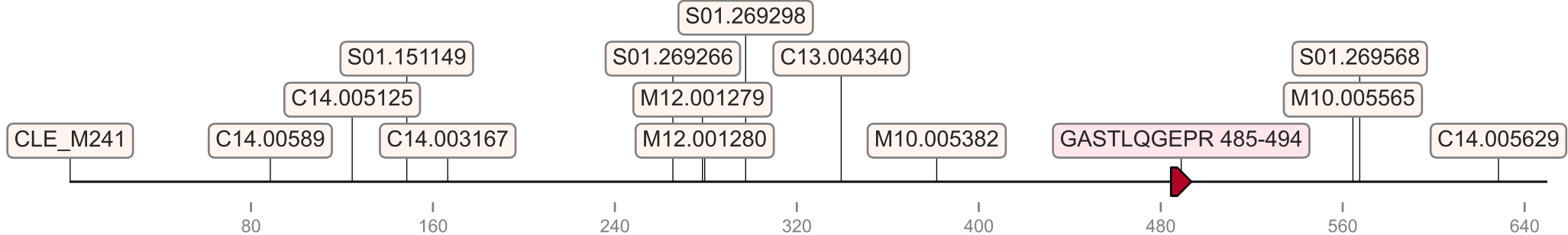


A0A2R8YF49-ARHGEF5-Rho guanine nucleotide exchange factor 5 {ECO:0000313|Ensembl:ENSP00000494816.1





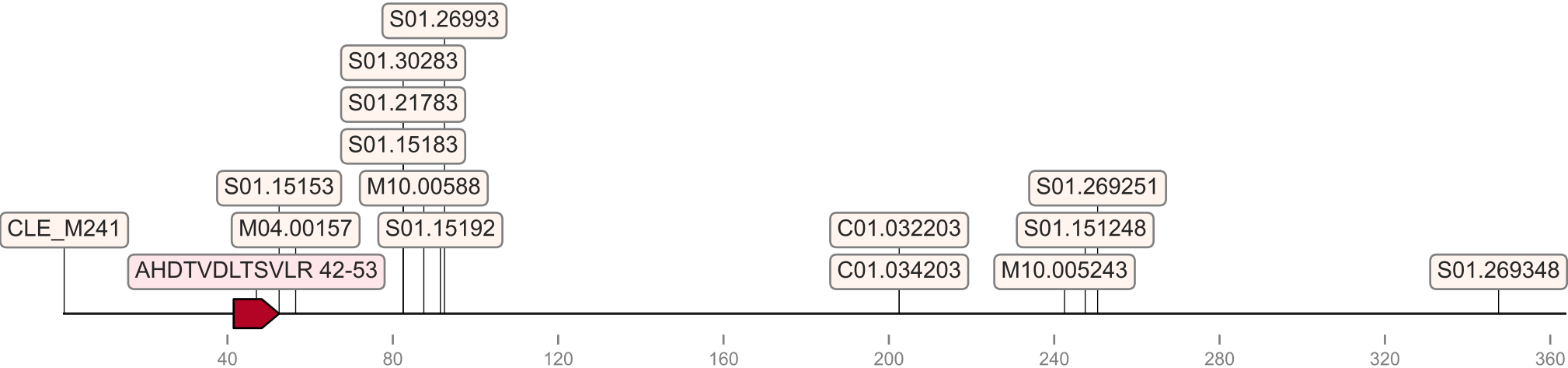
Q16643-DBN1-Drebrin



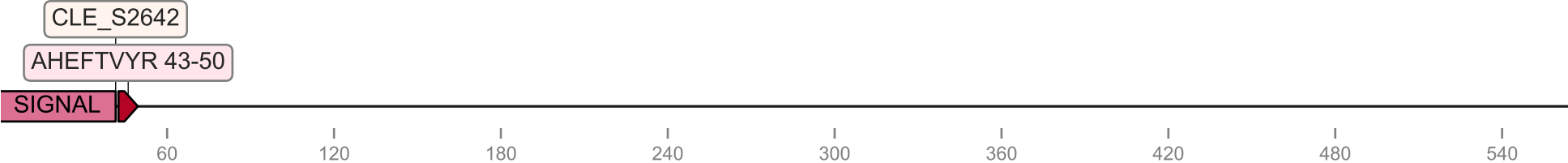
Q7Z7E8-UBE2Q1-Ubiquitin-conjugating enzyme E2 Q1



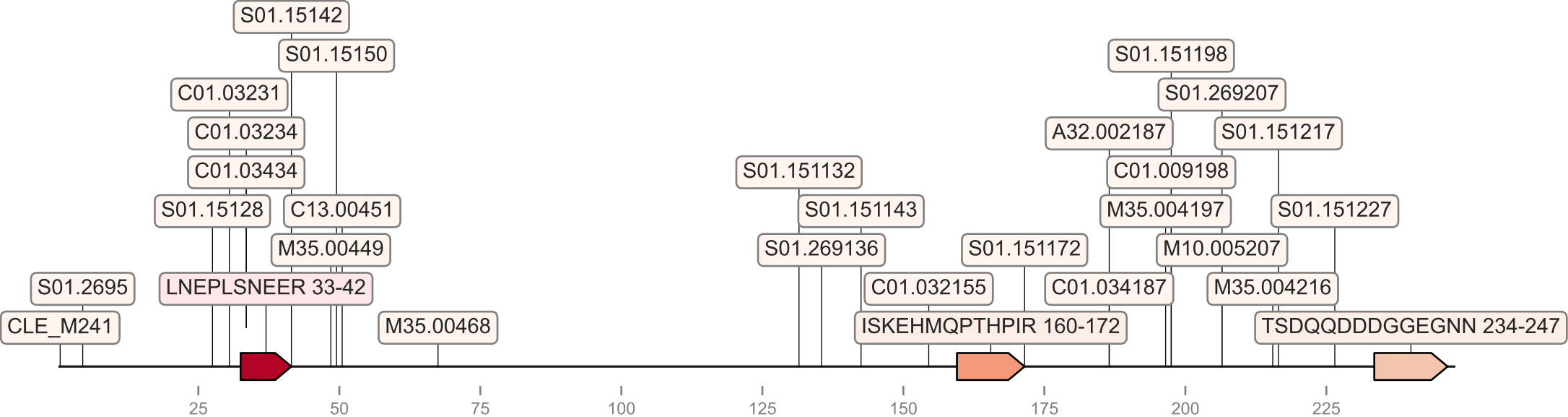
Q9NX46-ADPRS-ADP-ribosylhydrolase ARH3 {ECO:0000305}



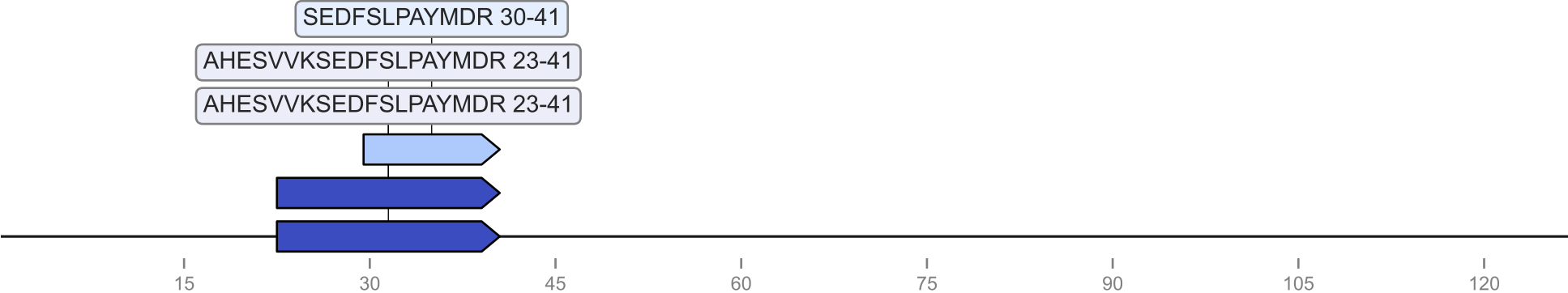
Q969V3-NCLN-BOS complex subunit NCLN {ECO:0000305}



P61981-YWHAG-14-3-3 protein gamma



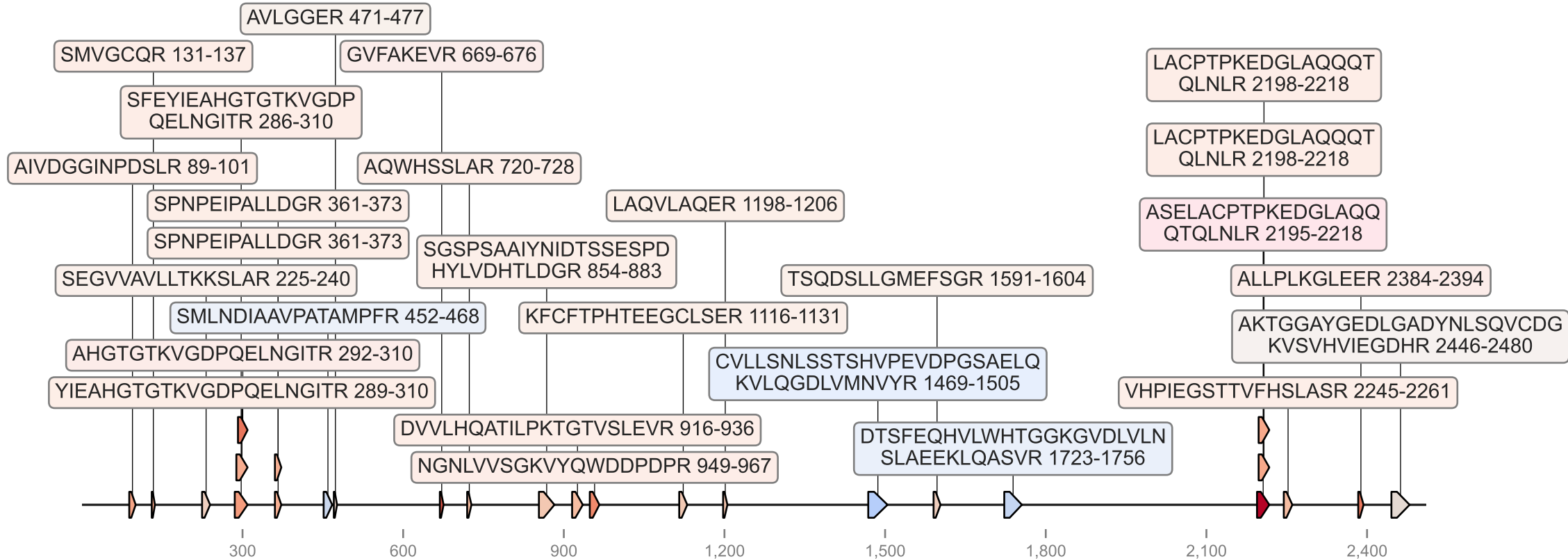
H3BN72-COX4I1-Cytochrome c oxidase subunit 4 {ECO:0000256|RuleBase:RU367145}



C9J9C1-PPP2R1A-Protein phosphatase 2 scaffold subunit Aalpha {ECO:0000313|Ensembl:ENSP00000391905.3}

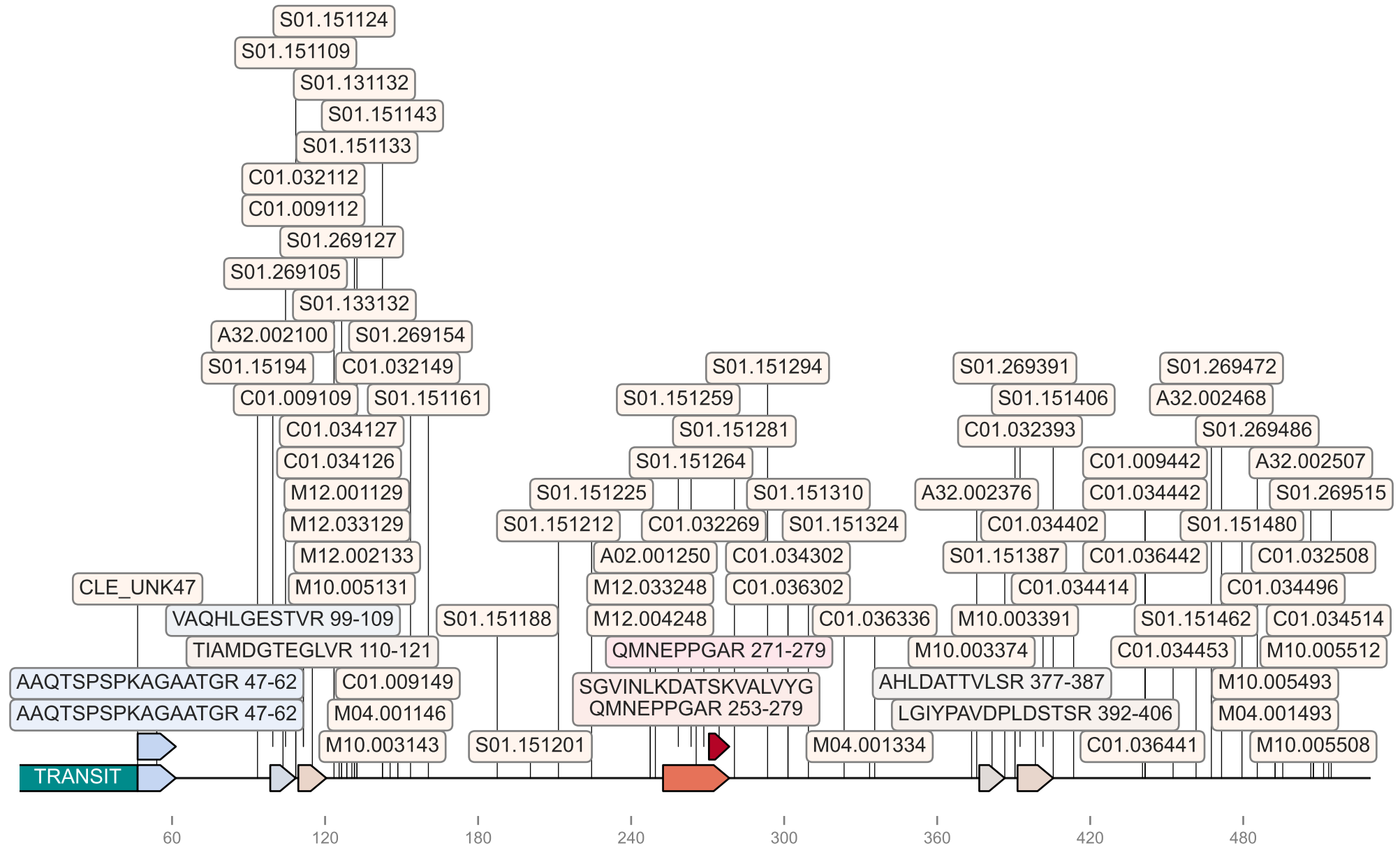


A0A0U1RQF0-FASN-Fatty acid synthase {ECO:0000256|ARBA:ARBA00018769}

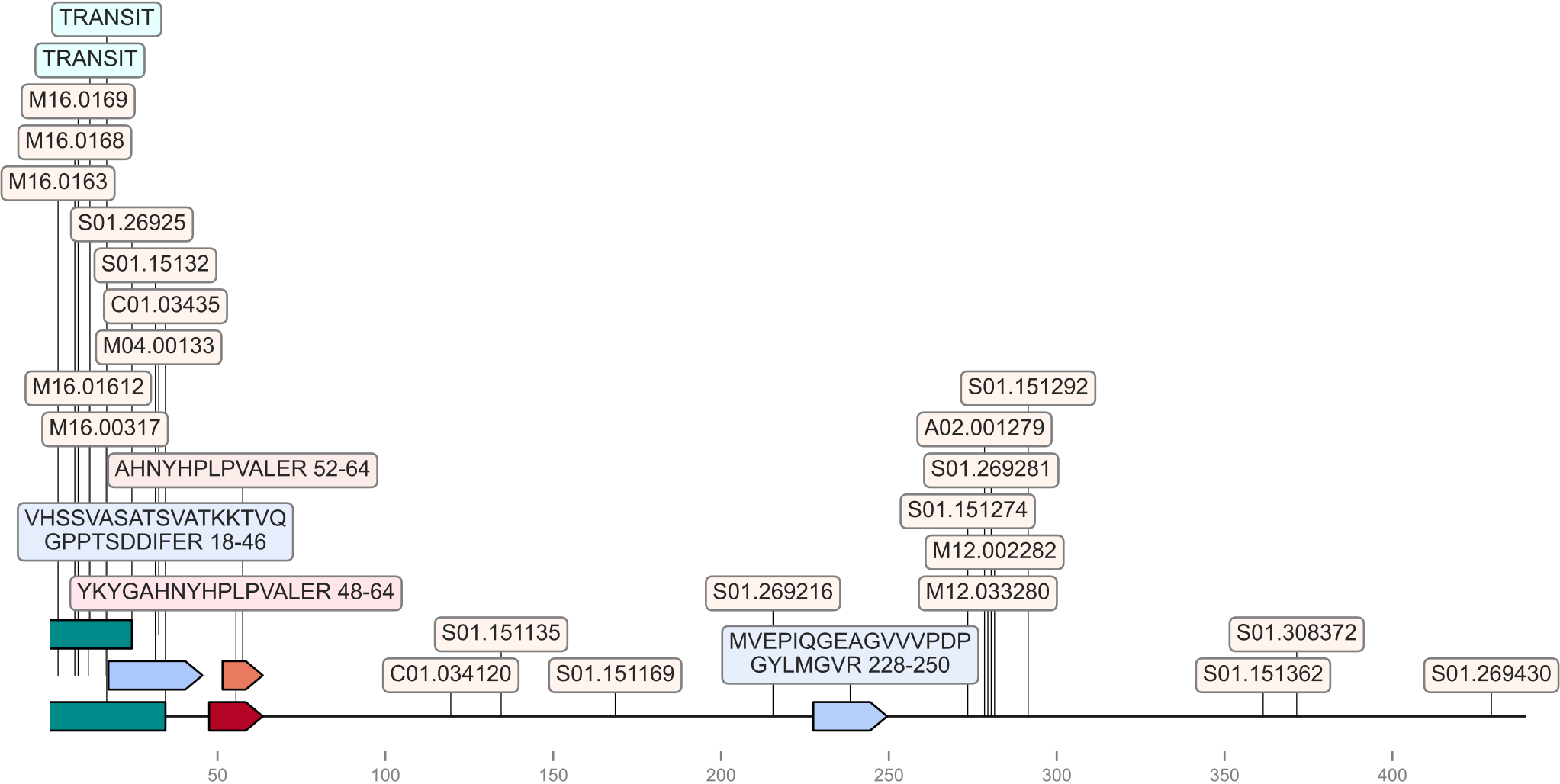




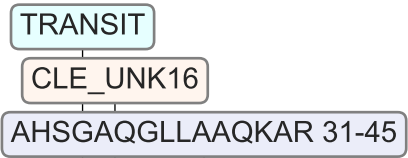
P06576-ATP5F1B-ATP synthase subunit beta, mitochondrial {ECO:0000305}



P04181-OAT-Ornithine aminotransferase, mitochondrial



Q9Y2Z4-YARS2-Tyrosine--tRNA ligase, mitochondrial



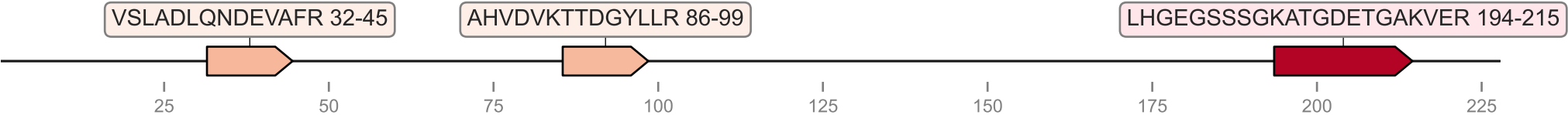
B4E0Y9-STK26-Serine/threonine kinase 26 {ECO:0000313|Ensembl:ENSP00000418753.1}

AHSPVAVQVPGMQNNIA  
DPEELFTKLER 2-29

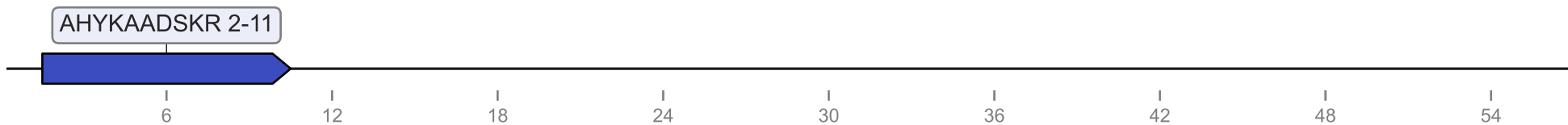


50 100 150 200 250 300 350 400

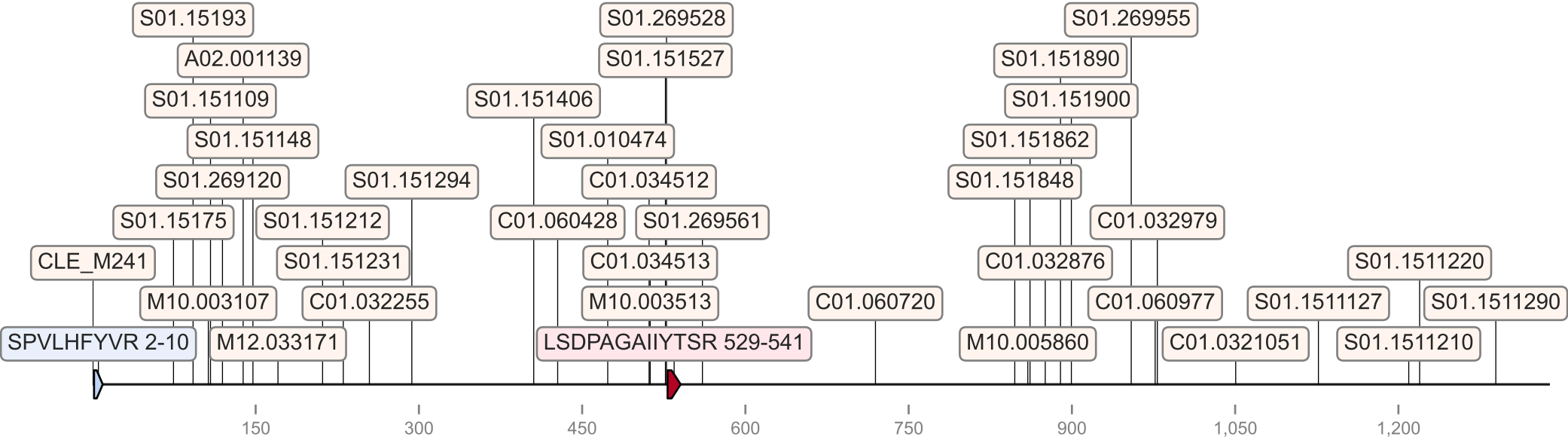
D6RAT0-RPS3A-40S ribosomal protein S3a {ECO:0000256|HAMAP-Rule:MF\_03122}



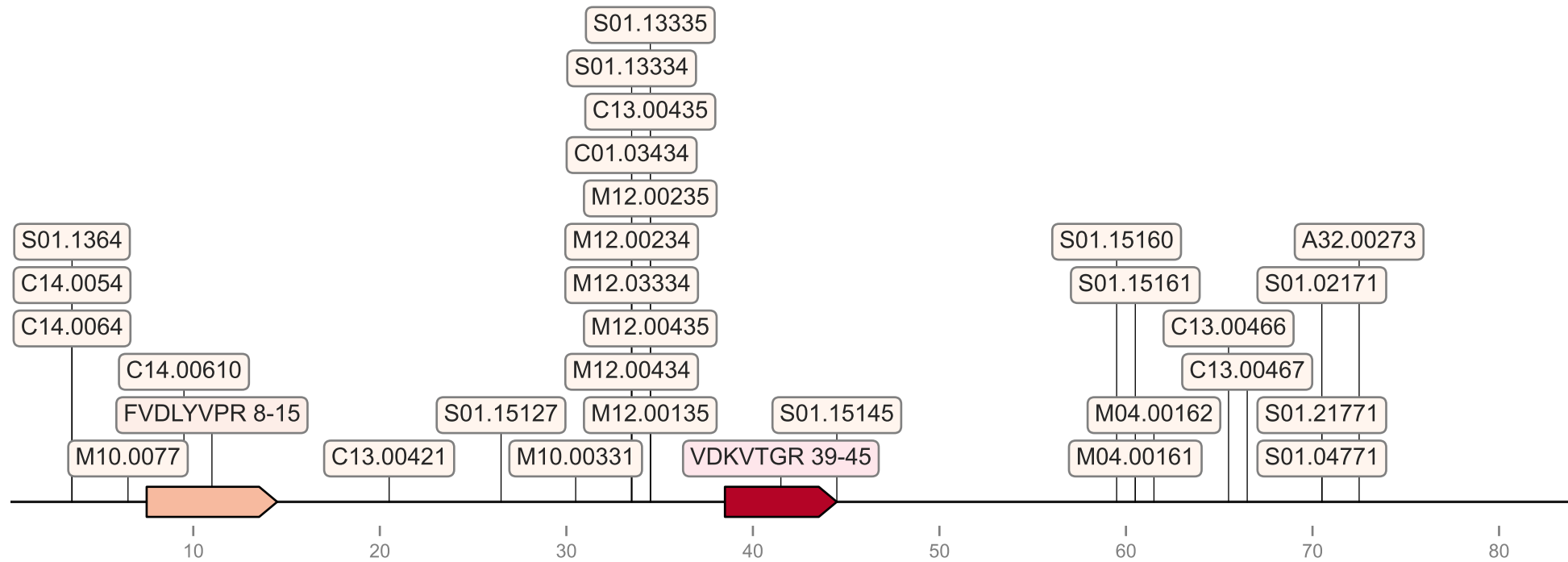
A0A0D9SEI7-MYCBP-MYC binding protein {ECO:0000313|Ensembl:ENSP00000485717.1}



O15067-PFAS-Phosphoribosylformylglycinamide synthase



# P63220-RPS21-40S ribosomal protein S21





Q01105-2-SET-Isoform 2 of Protein SET

