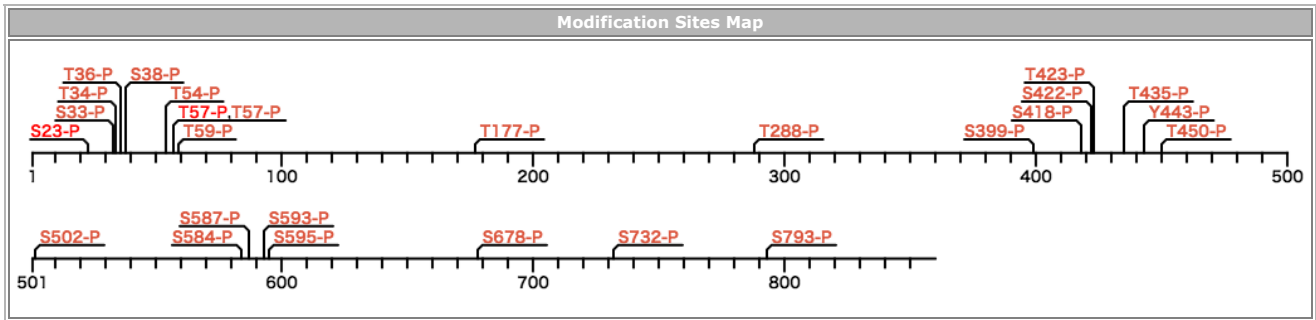


ID	Accession	GeneName	Chr.No.	Description
EF2_HUMAN	P13639	EEF2	19p13.3 3976054..3985467	Elongation factor 2



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
57	T	P	Lab	110218_pOVKATE_2.mgf[F017464]
57	T	P	Lab	110218_pOVKATE_3.mgf[F017465]
57	T	P	Lab	110218_pOVMANA_1.mgf[F017466]
57	T	P	Lab	110218_pOVMANA_2.mgf[F017467]
57	T	P	Lab	110218_pOVSAYO_1.mgf[F017469]
57	T	P	Lab	110218_pRMUGS_1.mgf[F017479]
57	T	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MVNFTVDQIR AIMDKKANIR NMSVIAHVDH GKSTLIDSLV CKAGIIASAR AGE^TRF^TIT^R KDEQERCITI KSTAISLFYE LSENDLNF^IK QSKDGAGFLI NLIDSPGHVD FSSE VTAALR VTDGALVVVD CVSGVCVQTE TVLRQAI^AER IKPVLMMNMK DRALLELQLE PEELYQ^TFQR IVENVNVII^S TYGEGESGPM GNIMIDPVLG TVGFGSGLHG WAFTLKQ FAE MYVAKFAAKG EGQLGPAERA KKVEDMMK^L WGD^RYFD^PAN GKFSK^SAT^SP EGK^LLP^RTFC QLILD^PIF^KV F^DAIMN^FK^E ETAKLIEKLD IKLDS^EDK^DK EGK^PLL^KAVM RRWL^PAGDAL LQMITI^HL^PS PVTAQ^KY^RCE LLYEG^PPD^DE AAMGI^KSC^DP KG^LPM^MY^IS^K MVPTSD^KGR^F YAFGR^VF^SGL V^ST^IGL^KVRIM GP^NY^TPG^KKE DL^LY^LK^PI^RQ^T ILM MGRY^VEP IEDV^PCGNIV GLV^GVD^QFLV KTGTIT^TFEH AHNMR^VM^KF^S V^SPV^RV^AVE AKNPAD^LP^KL VEGL^KR^LAK^S DPMV^QCI^EE SGEHII^AG^AG ELHLEI^CL^KD LEED^HAC IPI KKSD^PV^VSYR ET^VSEE^SSN^VL CL^SK^SPN^KHN RLYM^KAR^PFP DGLA^EID^ID^KG EVSAR^QEL^KQ RARY^LAE^KYE WDV^AE^AER^KI^W CFG^PD^GT^GPN ILTD^IT^KGV^Q YLNEI^KD^SVV AGF^QWAT^KEG ALCEEN^MRG^V RFDV^HD^VTLH ADAI^HR^GGG^GQ IIP^TARR^CLY A^SVL^TA^QP^RL MEPI^LY^LVEI^Q CPE^QV^VGGI^G VLN^RK^RGH^V FEES^QV^AG^TP MFV^VK^AY^LP^V NE SFG^TADL RSNT^GG^QAF^P QCV^FD^HW^QL PGD^PF^DN^SS^R PSQ^VVA^ETR^K RKGL^KEG^IPA LDN^FLD^KL

Backcolor of amino acid : Yellow -> site of modification, gray -> in front of processing