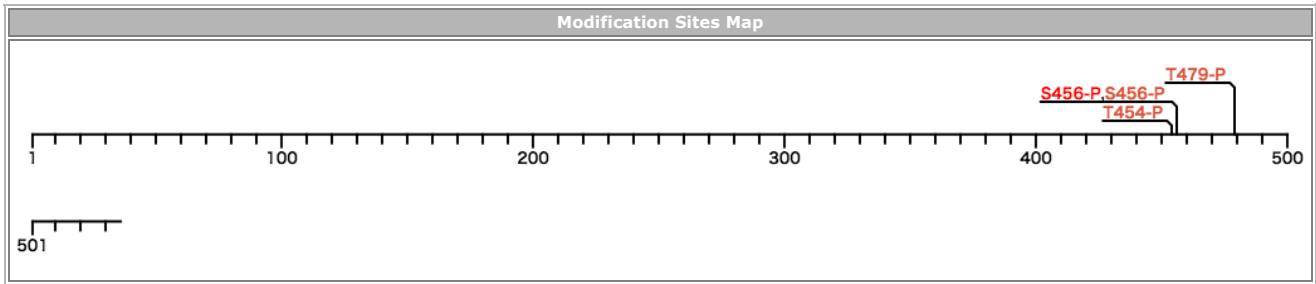


ID	Accession	GeneName	Chr.No.	Description
EHD1_HUMAN	Q9H4M9	EHD1	11q13.1 64619114..64655768	EH domain-containing protein 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
456	S	P	Lab	100627_akimura_pOVISe_2.mgf[F017440]
456	S	P	Lab	100627_akimura_pOVISe_3.mgf[F017443]
456	S	P	Paper	Sci Signal 2009, 2(84), ra46
456	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MFSWVSKDAR RKKEPELFQT VAEGLRQLYA QKLLPLEEHY RFHEFHSPAL EDADFDNKPM VLLVGQYSTG KTTFIRHLIE QDFPGMRIGP EPTTDSFIAV MHGPTGVVVP GNA LVVDPRR PFRKLNAGFN AFLNRFMCAQ LPNPVLDNIS IIDTPGILSG EKQIRISRGYD FAAVLEWFAE RVDRIILLFD AHKLDISDEF SEVIKALKNH EDKIRVVLNK ADQIETQQ LM RYVGALMWSL GKIINTPEVV RYVIGSFWSH PLLIPDNRKL FEAEEQDLFK DIQSLPRNAA LRKLNLIKR ARLAKVHAYI ISSLKKEMPN VFGKESKKKE LVNNLGEIYQ KI EREHQISP GDFPSLRKMQ ELLQTQDFSK FQALKPKLLD TVDDMLANDI ARLMVMVRQE ESLMPSQVVK GGAFDGTMNG PFGHGYGEGA GEGIDVVEWV VGKDKPTYDE IF YTLSPVNG KITGANAKKE MVKSKLPNTV LGKIWKLADV DKDGLLDDEE FALANHLIKV KLEGHELPAD LPHLVPPSK RRHE	

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