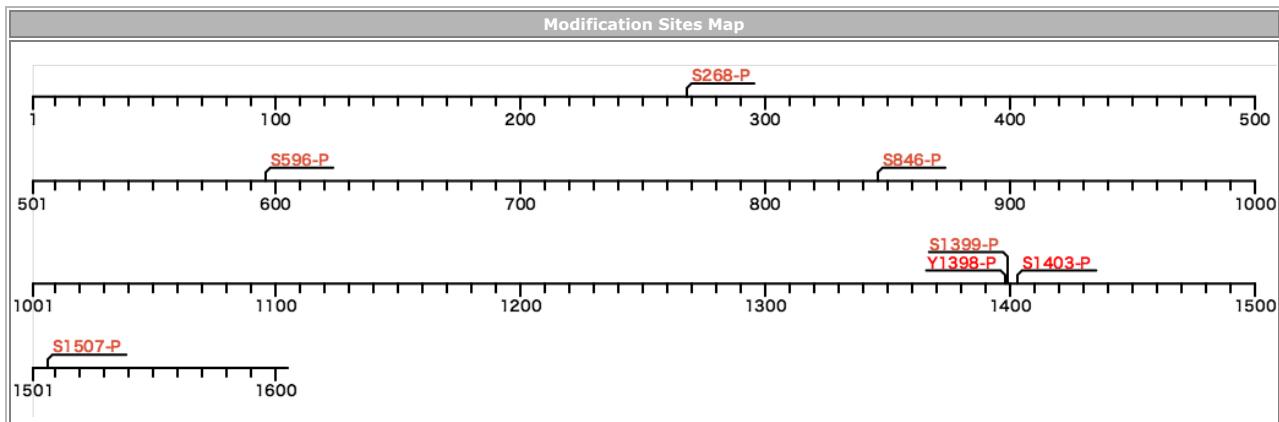


ID	Accession	GeneName	Chr.No.	Description
AHDC1_HUMAN	Q5TGY3	AHDC1	1p36.11	27860546..27930942 AT-hook DNA-binding motif-containing protein 1



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
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Protein Sequence	
MRVKPQGLVV TSSAVCSSPD YLREPKYYPG GPPTPRPLLP TRPPASPPDK AFSTHAFSEN PRPPPDRPS TRRPPVLAGK DDPLPRAAR PVSQARCPTV VGDGSSSRRC WDN GRVNLRP VVQLIDIMKD LTRLSQDLQH SGVHLDCGGI RLSRPPAPP GDLQYSFFSS PSLANSIRSP EERATPHAKS ERPSPHLYEP EPEPRDSPQP GQGHSPGATA AATGLP PEPE PDSTDYSELA DADILSELAS LTCPEAQLE AQALEPPSPE PEPQLLDPQP RLDPQALEP LGEAELPPL QPLADPLGLP GLALQALDTL PDSLESQLLD PQALDPLPKL LD VPGRRLEP QQPLGHCPA EPLRLDLCLP HGPPGPEGH KYALRRTDRP KILCRRRKAG RGRKADAGPE GRLLPLPMPT GLVAALAEPP PPPPPPPAL PGPGPVSVP E LPKESSQ TPV VSTRKGKCRG VRMVKRMAK IPVSLGRRNK TTGVSSLSS SLSVEGKELG LRVSAAEPTPL LKMKNNGRN VVVFPPEGMP IILKRKRGRP PKNLLLGPDK PKEPAVVAEE AATVAAATMA MPEVKRKKR KQKLASQPS YAADANDSKA EYSDVLAKLA FLNRQSQCAG RCSPPRCWTP SEPESVHQAP DTQSHFLH RVQGFRRRGK KAGGFGRRGG GHAAKSARCS FSDFFEIGK KKKVVAVAAA GVGGPLTEL GHPRKGRGE VDAVTGKPKR KRRSRKNGTL FPEQVPSGPG FGEAGAEWAG DKGGGWAPHH GHPGQQAGR N CGFQGEARA FASTGLESQA SGRCGYYSTG APSGQTELSQ ERQNLFTGYF RSLLSDDDSS DLLDFALSAS RPSRKASGT YAGPPTSALP AQRGLATFPS RGAKASPVA GSSGAGADPS FQPVL SARQT FPPGRAASYG LTPAASDCRA AETFPKLVPP PSAMARSPTT HPPANTYLPQ YGGYGAQGSV FAPTKPTGQ DCANSKDCSF AYGSGNSLPA SP SSAHSAGY APPPTGGPCL PPSKASFSSS SEGAPFSGA PTPLRCDRSA STVSPGGYMV PKGTTASATS AASAASSSSS SFQPSPENCQ QFAGASQWPF RQGYGLDW SEA FSQLYNP SFDCHVSEPN VILDISNYTP QKVQQTAVS ETFSESSSSD TQFNQPVGGG GFRRANSEAS SSEGQSSLSS LEKLMMDWNE ASSAPGYNWN QSVLFQSSSK PGR GRKKVD LFEASHLGFPS TSASAAASGY PSKRSTGPRQ PRGGRRGGAC SAKKERRGAA AKAKFIPKPKQ PVNPLFQDSP DLGLDYSGD SSMSPPLSQS RAFGVGERDP CDFI GPYSMN PSTPSDGTFG QGFHCDSPSL GAPELDGKHIF PPLAHPPTVF DAGLQKAYSP TC ^Y PTLGFKE ELRPPPTKLA ACEPLKHGLQ GASLGHAAA QAHLSCRDLQ LGQPHY DSPS CKGTAYWYPP GSAARSPPYE GKVGTLAAD FLGRTEAACL SAPHLASPPA TPkadKEPLE MARPPGPPRG PAAAAGYGC PLLSDLTSP VPRDSLLPLQ DTAYRPGF M PQAHPGLGGG PKSGFLGMA EPHPEDTFV TS	

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