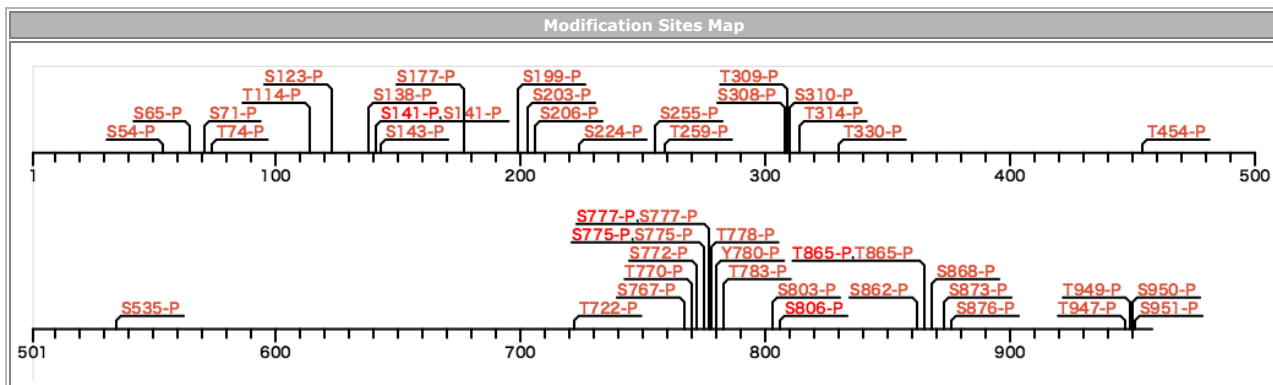


| ID | Accession | GeneName | Chr.No. | Description |
|-------------|-----------|----------|--------------------------|---------------------------------------|
| CAF1A_HUMAN | Q13111 | CHAF1A | 19p13.3 4402659..4445015 | Chromatin assembly factor 1 subunit A |



Click a modification site to display information in detail.

| Site no | Amino acid | Type | Division | Detail |
|---------|------------|------|----------|-------------------------------------|
| 141 | S | P | Lab | 130327_HEK_CE_pphos.mgf[F015007] |
| 141 | S | P | Lab | 130415_HEK_CE_tphos.mgf[F015009] |
| 141 | S | P | Paper | Cell Rep 2014, 8(5), 1583-1594 |
| 141 | S | P | Paper | J Proteome Res 2013, 12(1), 260-271 |
| 141 | S | P | Paper | Sci Signal 2010, 3(104), ra3 |
| 141 | S | P | Paper | Sci Signal 2011, 4(164), rs3 |

| Protein Sequence | |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--|
| <p>MLEEECGAP GARGAATAMD CKDRPAFPVK KLIQARLPFK RLNLVPGKGA DDMSDDQGTS VQKSPDLEA SLDTLENNCH VGSDIDFRPK LVNGKGPLDN FLRNRIETSI GQ STVIIDLT ED SNEQPDSL VDHNKLNS EA SPSREAINQ REDTGDQQL LKAIQNDKLA FPGETLSDIP CKTEEEGVGC GGAGRRGDSQ EC SPRSCP EL TSGPRMCPRK EQD SWSEAGG ILFKGKPMV VLQDILAVRP PQIKSLPATP QGKNMTP ESE VLESFPEEDS VLSHSSLSSP SSTSSPEGPP APPKQHSSTS PFPTSTPLRR ITKKFVKGS I EKNKLR L QRD QERLGKQLKL RAEREEKEKL KEEAKRAKEE AKKKKEEKEE LKEKERREKR EKDEKEKA EK QRLKEERRKE RQEALEAKLE EKRRKKEEKR LREEEKRIKA EKAEITRRFQ KPKTPQAPKT LAGSCGKFP FEIKEHMLVA PRRRTAFHPD LCSQLDQLL QQSGEFSFLK DLKGRQPLRS GPTHVSTRNA DIFNSDVVIV ERGKGDGVP E RRFKGRMKLL QF CENHRPAY WGTWNKKTAL IRARDPWAQD TKLLDYEVD S DEEWEEEPG ESLSHSEGGD DDDMGEDEDE DDGFFVPHGY LSEDEGVTEE CADPENHKVR QKLKAKEWDE FLAKGKRFRV LQPVKIGCVW AADRDCAGDD LKVLQQFAAC FLETLP AQEE QTPKASKRER RDEQILAQLL PLLHGNVNGS KVIIREFQEH CRRGLLSNHT GSPRSPSTTY LHT PTPSEDA AIPSKSRLKR LISENS VYEK RPDFRMCWYV HPQVLQSFQQ EHLVPCQWS YVTSVPSAPK EDSGSVPSTG PSQGTPI SLK RKSAGSMCIT QFMKKRRHDG QIGA EDMDGF QADTEEEEEEGDCMIVDP DAAEVQAPCG AASGAGGGVG VDTGKATLTS SPLGA</p> | |

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