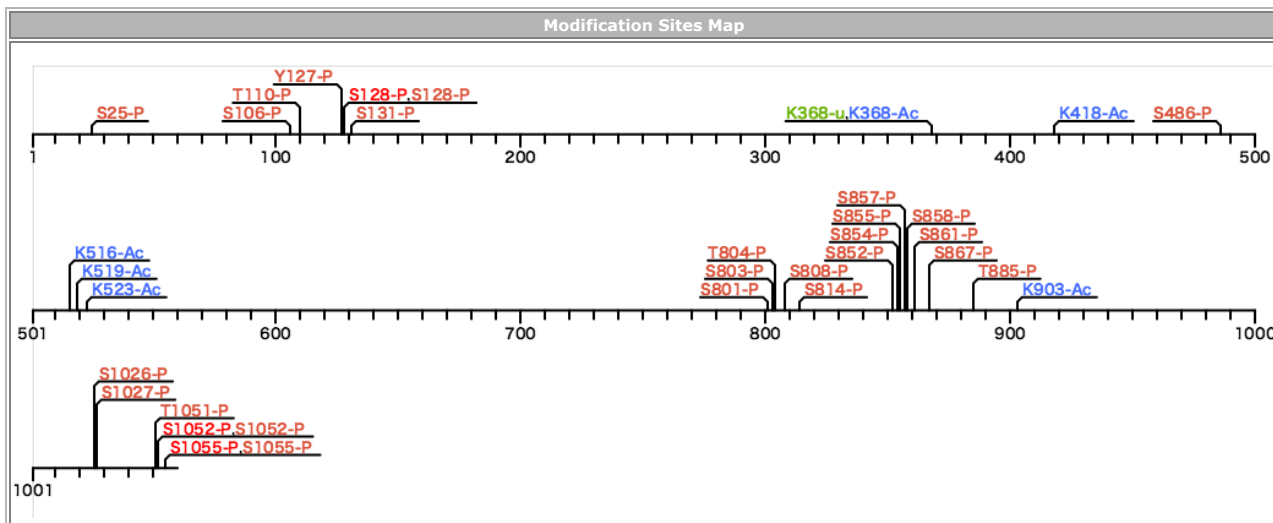


ID	Accession	GeneName	Chr.No.	Description
BRD1_HUMAN	O95696	BRD1	22q13.33 50166931..50221160	Bromodomain-containing protein 1



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
128	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
128	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
128	S	P	Lab	140320_Agarose_.mgf[F017423]
128	S	P	Lab	140320_Agarose_.mgf[F017423]
128	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
128	S	P	Lab	110218_pRMG2_1.mgf[F017475]
128	S	P	Lab	110218_pRMG2_3.mgf[F017477]
128	S	P	Lab	110218_pRMG2_4.mgf[F017478]
128	S	P	Lab	110711_titania_LNcaP_AI_4.mgf[F017445]
128	S	P	Lab	100628_akimura_pOVCAR3_3.mgf[F017459]
128	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
128	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
128	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
128	S	P	Paper	Sci Signal 2011, 4(164), rs3

Protein Sequence

MRRKGRCHRG SAARHPSSPC SVKHSPTRET LTYAQARMV EIEIEGLRHR ISIFDPLEII LEDDLTAQEM SECNSNKENS ERPPVCLRTK RHKNNRVKKK NEALPSAHGTPAS ASALPEP KVRIVEYSPPSAPRRPPVYVY KFIKSAEEL DNEVEYDME EDYAWLEIVN EKRGDCVPA VSQSMFEFLM DRFEKESHCE NQKQGEQQSL IDEDAVCCIC MDGEC QNSNV ILFCDMCNLA VHQECYGVPI IPEGQWLCRH CLQSRARPAD CVLCPNKGGA FKKTDDDRWG HVVCALWIPE VGFANTVFIE PIDGVRNIPP ARWKLTCYLC KQKGVG ACIQ CHKANCYAF HVTCAQKAGL YMKMEPVKEL TGGGTTFSVR KTAYCDVHTP PGCTRRPLNI YGDVEMKNGV CRKESVKTIV RSTSKVRKKA KKAKKALAEP CAVLPTVC AP YIPQRLNRI ANQVAIQKQK QFVERAHSYV LLKRLSRNGA PLLRRLQSSL QSQRSSQRE NDEEMKAAKE KLYWQRLRH DLERARLLIE LLRKREKLKR EQVKVEQVAM ELRLTPTVL LRSVLDQLQD KDPARIFAQP VSLKEVPDYL DHIKHPMDFA TMRKRLEAQQ YKNLHEFEED FDLIIDNCMK YNARDTVFYR AAVRLRDQGG VVLRQARREV DSI GLEEASG MHLPERPAAA PRRPFSWEDV DRLLDPANRA HLGLEELRE LLDMLDLTCA MKSSGSRSKR AKLLKKEIAL LRNKLSSQHS QPLPTGPGLG GFEEDGAALG PEAGE EVLPR LETLLQPRKR SRSTCGDSEV EEE SPGKRLD AGLTNGFGGA RSEQEPGGGL GRKATPRRRC AESSISSSN SPLCDSFNA PKCGRGKPAL VRRHTLEDRS ELISCIEN GN YAKAARIAAE VGQSSMWIST DAAASVLEPL KVVWAKCSGY PSYPALIIPD KMPRVPGHHN GVTIPAPPLD VLKIGEHMQT KSDEKLFVLV FFDNKRWSQW LPKSKMVLPLG IDETIDKLM MEGRNSIRK AVRIAFDRAM NHLSRVHGEP TSDLSDI

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