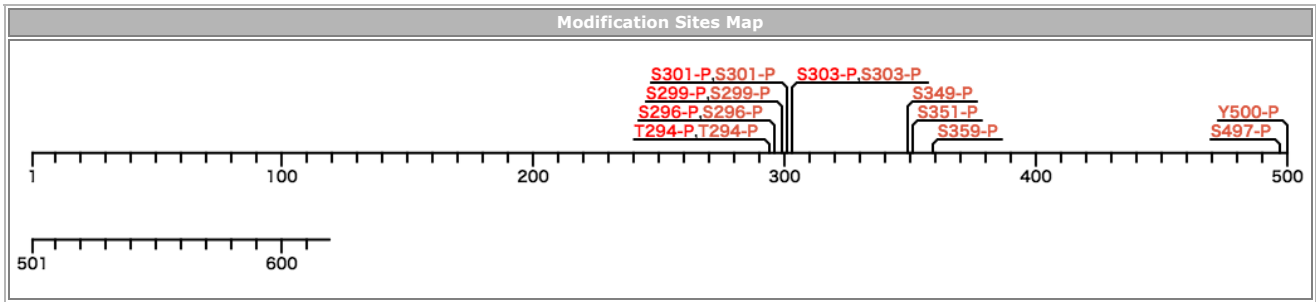


ID	Accession	GeneName	Chr.No.	Description
FNBP1_HUMAN	Q96RU3	FNBP1	9q34.11 132649466..132805473	Formin-binding protein 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
301	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
301	S	P	Lab	110218_pRMG1_3.mgf[F017474]
301	S	P	Lab	100627_akimura_pOVICE_1.mgf[F017437]
301	S	P	Lab	100628_akimura_pMCAS_3.mgf[F017456]
301	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MSWGTELWDQ FDNLEKHTQW GIDILEKYIK FVKERTEIEL SYAKQLRNLS KKYQPKNSK EEEYKYTSC KAFISNLNEM NDYAGQHEVI SENMASQIIV DLARYVQELK QE RKS NFHDG RKAQQHIETC WKQLESSKRR FERDCKEADR AQQYFEKMDA DINVTKADVE KARQQAQIRH QMAEDSKADY SSILQKFNHE QHEYHHTHIP NIFQKIQEME ER RIVRMGES MKTYAEVDRQ VIPIIGKCLD GIVKAAESID QKNDSQLVIE AYKSGFEPPG DIEFEDYTQP MKR T V S D N S L S N S R G E G K P D L K F G G K S K G K L W P F I K K N K L M S L L T S P H Q P P P P P A S A S P S A V P N G P Q S P K Q Q K E P L S H R F N E F M T S K P K I H C F R S L K R G L S L K L G A T P E D F S N L P P E Q R R K K L Q Q K V D E L N K E I Q K E M D Q R D A I T K M K D V Y L K N P Q M G D P A S L D H K L A E V S Q N I E K L R V E T Q K F E A W L A E V E G R L P A R S E Q A R R Q S G L Y D S Q N P P T V N N C A Q D R E S P D G S Y T E E Q S Q E S E M K V L A T D F D D E F D D E E P L P A I G T C K A L Y T F E G Q N E G T I S V V E G E T L Y V I E E D K G D G W T R I R R N E D E E G Y V P T S Y V E V C L D K N A K D S	

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