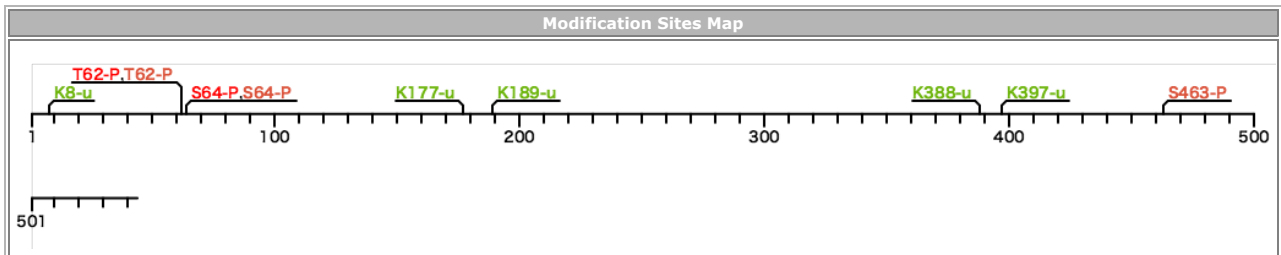


ID	Accession	GeneName	Chr.No.		Description
AGM1_HUMAN	O95394	PGM3	6q14.1	83870869..83903655	Phosphoacetylglucosamine mutase



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
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Protein Sequence

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MDLGAITKYS ALHAKPNGLI LQYGTAGFRT KAEHLDHVMF RMGLLAVLRS KQTKSTIGVM VTASHNPEED NGVKLVDPGLG EMLAPSWEEH ATCLANAEEQ DMQRVLIDIS EK
EAVNLQQD AFVVIGRDTR PSSEKLSQSV IDGVTVLGGQ FHDYGLLTPP QLHYMVYCRN TGGRYKATI EGYQKLSKA FVELTKQASC SGDEYRSLKV DCANGIGALK LREM
EHYFSQ GLSVQLFNDG SKGKLNHLCG ADFVKSHQKP PQGMEIKSNE RCCSFDGDAD RIVYYYHDAD GHFHLIDGDK IATLISSFLK ELLVEIGESL NIGVVQTAYA NGSSTR
YLEE VMKVPVYCTK TGVKHLHKA QEFDIGVYFE ANGHGTALFS TAVEMKIKQS AEQLEDKCRK AAKMLNIIID LFNQAAGDAI SDMLVIEAIL ALKGLTVQQW DALYTDLPN
R QLKVQVADRR VISITDAERQ AVTPPGLQEA INDLVKKYYKL SRAFVRPSGT EDVVRVYAEA DSQESADHLA HEVSLAVFQL AGGIGERPQP G

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