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logo
                                                
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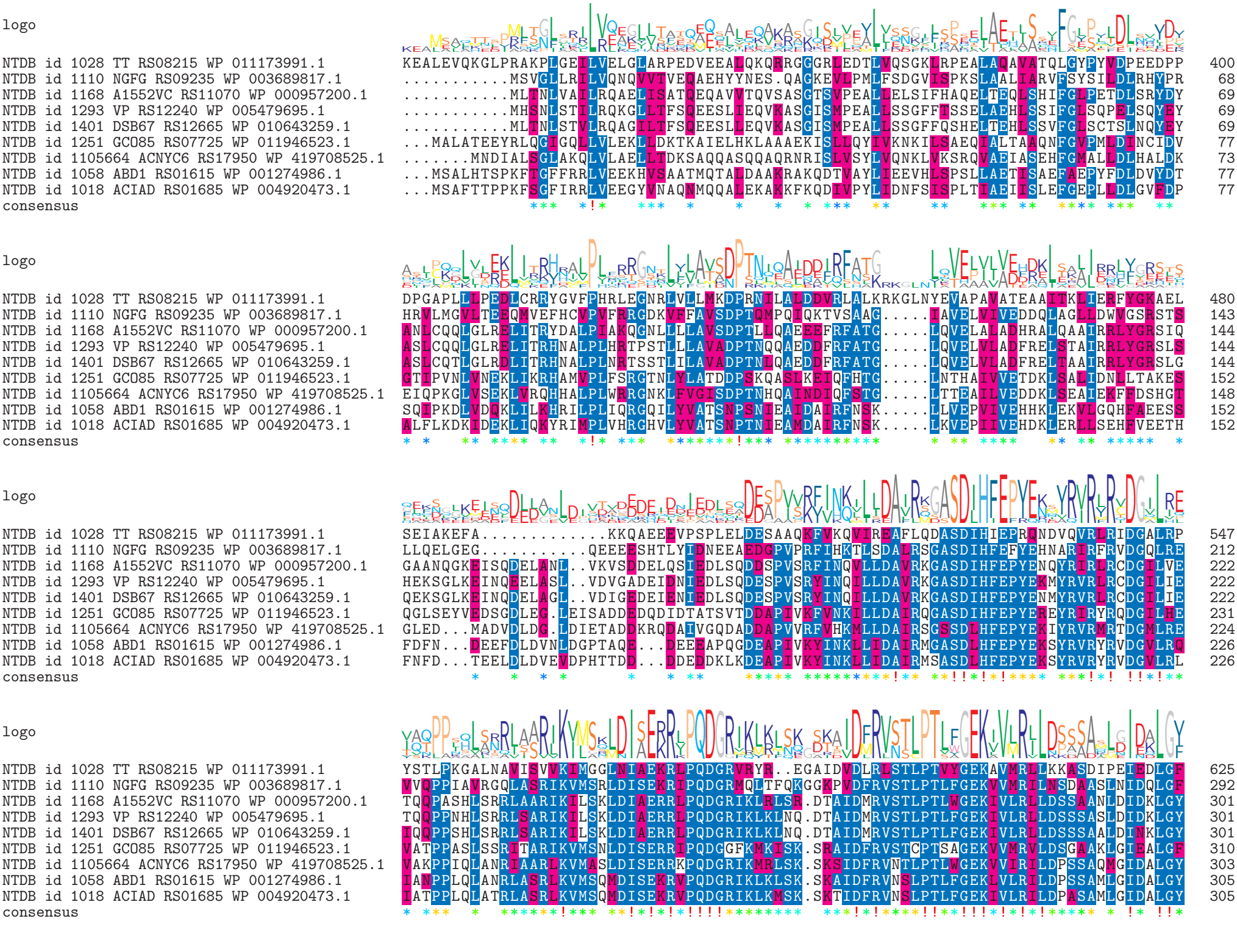
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NTDB id 1293 VP RS12240 WP 005479695.1 .....MHSNLSTILRQKGLLTFSQEESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQYEY 69
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NTDB id 1251 GCO85 RS07725 WP 011946523.1 ...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQNFGVPMLDINCIDV 77
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consensus
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consensus
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consensus
logo
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consensus

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consensus                                        * ** ****** *!*!**!*!!!!!!!!*******!***** **!**!*!!!*!*** !*!!*********!**!* !!**
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NTDB id 1293 VP RS12240 WP 005479695.1          FLRQDPDVVMVGEIRDLDTAEIAIKASQTGHLVLSTLHTNSAAETIIRLSNMGVESFNLASSLSLIIAQRLARKLCPYCK  461
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NTDB id 1251 GCO85 RS07725 WP 011946523.1       FLRQDPDIIMVGEIRDLETAEIAIKAAQTGHLVLSTLHTNSAAETLNRLVNMGIPTFNIASSVTLIIAQRLARKLCNQCK  470
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NTDB id 1058 ABD1 RS01615 WP 001274986.1        FLRQDPDIIMVGEIRDLETAEIAIKAAQTGHMVMSTLHTNSAPETLTRLRNMGVPSFNIATSVNLVIAQRLARRLCSQCK  465
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consensus                                       !!!!!!!**!!!!!!!**!!*!!**!**!!!*!**!!!!!*!******!* *!!* *!!*********!!!*!**!  !*
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NTDB id 1168 A1552VC RS11070 WP 000957200.1     IAVR.PSALLQSQFAFQPNEI....LYEANAAGCNECTG.GYSGRVGIYEVMAFNTELAEAIMQRASIHQIERLAKANGM  535
NTDB id 1293 VP RS12240 WP 005479695.1          QPQE.HTVQL.QHLGIQTTDN....IFRANPDGCNECTH.GYSGRTGIYEVMRFDESLSEALIKGASVHELEKLAIANGM  534
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NTDB id 1018 ACIAD RS01685 WP 004920473.1       RP.IQVPERSLLEMGFTPEDLAQPEFQIFEPVGCHDCRE.GYKGRVGIYEVMKITPEISKIIMEDGNALEIAATAETLGF  543
consensus                                        *          *  *** * **       ** **!! *!   !!*!!*****!***** **  ***  **  ***  **   !*
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NTDB id 1168 A1552VC RS11070 WP 000957200.1     QTLQESGLEKLREGITSFAELQRVLYF  562
NTDB id 1293 VP RS12240 WP 005479695.1          STLQMSGIEKLKQGITSFSELQRVLYF  561
NTDB id 1401 DSB67 RS12665 WP 010643259.1       QTLQMSGIEKLKQGITSFRELQRVLYF  561
NTDB id 1251 GCO85 RS07725 WP 011946523.1       LTIFQSGIEKVKEGITTIEEVNRVTVD  574
NTDB id 1105664 ACNYC6 RS17950 WP 419708525.1   NDLRTSGLLKAMQGITSLEEINRVSKD  566
NTDB id 1058 ABD1 RS01615 WP 001274986.1        NNLRRSGLKKVMQGVTSLQEVNRVTSE  570
NTDB id 1018 ACIAD RS01685 WP 004920473.1       NNLRRSGLKKVMQGVTSLQEINRVTSE  570
consensus                                       **** *!* !***!*!** !***** *
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X non conserved
X similar
X ≥ 50% conserved