logo

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NTDB id 146 SP RS08570 WP 000614538.1      MIQIGKIFAGRYRIVKQIGRGGMADVYLAKDLILDGEEVAVKVLRTNYQTDPIAVARFQREARAMADLDHPHIVRITDIGEEDGQ  85
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NTDB id 384 SMU RS02325 WP 002263039.1     MIQIGKLFAGRYRILQSIGRGGMADVYLANDLILDNEEVAIKVLRTNYQTDQVAVTRFQREARAMAELNHPNIVAIRDIGEEDGQ  85
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NTDB id 467 HSISS4 RS06915 WP 021143821.1  MIQVGKLFAGRYRILKSIGRGGMADVYLANDLILDNEEVAIKVLRTNYQTDQVAVARFQREARAMAELSHPNIVAIRDIGEEDGQ  85
consensus                                  !!!*!!*!!!!!!!***!!!!!!!!!!!!!*!!!!!*!*!!*!!!!!!!!!!!!**!!*!!!!!!!!!!**!*!!*!!*!*!!!!!!!!!
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NTDB id 182 SPR RS07820 WP 000614552.1     QYLAMEYVAGLDLKRYIKEHYPLSNEEAVRIMGQILLAMRLAHTRGIVHRDLKPQNILLTPDGTAKVTDFGIAVAFAETSLTQTN  170
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NTDB id 26667 STER RS06845 WP 011681404.1  QFLVMEYVDGSDLKKFIQDHAPLSNQDVVRIMGEVLSAMTLAHQKGIIHRDLKPQNVLLTKDGRAKVTDFGIAVAFAETSLTQTN  170
NTDB id 467 HSISS4 RS06915 WP 021143821.1  QFLVMEYVDGSDLKKYIQDHAPLSNQDVVRIMGEVLSAMTLAHQKGIIHRDLKPQNVLLTKDGTAKVTDFGIAVAFAETSLTQTN  170
consensus                                  !*!*!!!!*!*!!!!*****!*!!!!****!!!***!*!!*!!!**!!*!!!!!!!!*!!!*!!*!!!!!!!!!!!!!!!!!!!!!
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NTDB id 26667 STER RS06845 WP 011681404.1  SMLGSVHYLSPEQARGSKATIQSDIYAMGIMLFEMLTGHIPFDGDSAVTIALQHFQKPLPSILAENRNVPQALENVVIRATAKKL  255
NTDB id 467 HSISS4 RS06915 WP 021143821.1  SMLGSVHYLSPEQARGSKATIQSDIYAMGIMLFEMLTGHIPYDGDSAVTIALQHFQKPLPSILAENRNVPQALENVVIRATAKKL  255
consensus                                  !!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!***!!!!!!!!*!!!!!!!!!!!!!!!*!!!!**!!!  *!!!!!!!!!*!*!!!!*!
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NTDB id 146 SP RS08570 WP 000614538.1      TNRYRSVSEMYVDLSSSLSYNRRNESKLIFDETSKADTKTLPKVSQSTLTSIPKVQAQTEH.KSIKNPSQAVTEETYQPQAPKKH  339
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NTDB id 182 SPR RS07820 WP 000614552.1     TNRYRSVSEMYVDLSSSLSYNRRNESKLIFDETSKADTKTLPKVSQSTLTSIPKVQAQTEH.KSIKNPSQAVTEETYQPQAPKKH  339
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NTDB id 26667 STER RS06845 WP 011681404.1  ENRYNSTLEMSRDLATSLHPSHSRDSKVVFDDMT..DTKTLPKVDSVSSVGPEK.KISPKPSEPTPAPSKQPREKASLS..KKKK  335
NTDB id 467 HSISS4 RS06915 WP 021143821.1  ENRYNSTLEMSRDLVTSLHPSHRRDAKVVFDDMT..DTKTLPKVDPVPSASLEKKAVAAEPSEPTPAPSKQPRKKTTPAKKKKK  338
consensus                                  **!!*!**!!**!!*!!*!!*********!**!******!!!*!!!*********************  ***********************!!*
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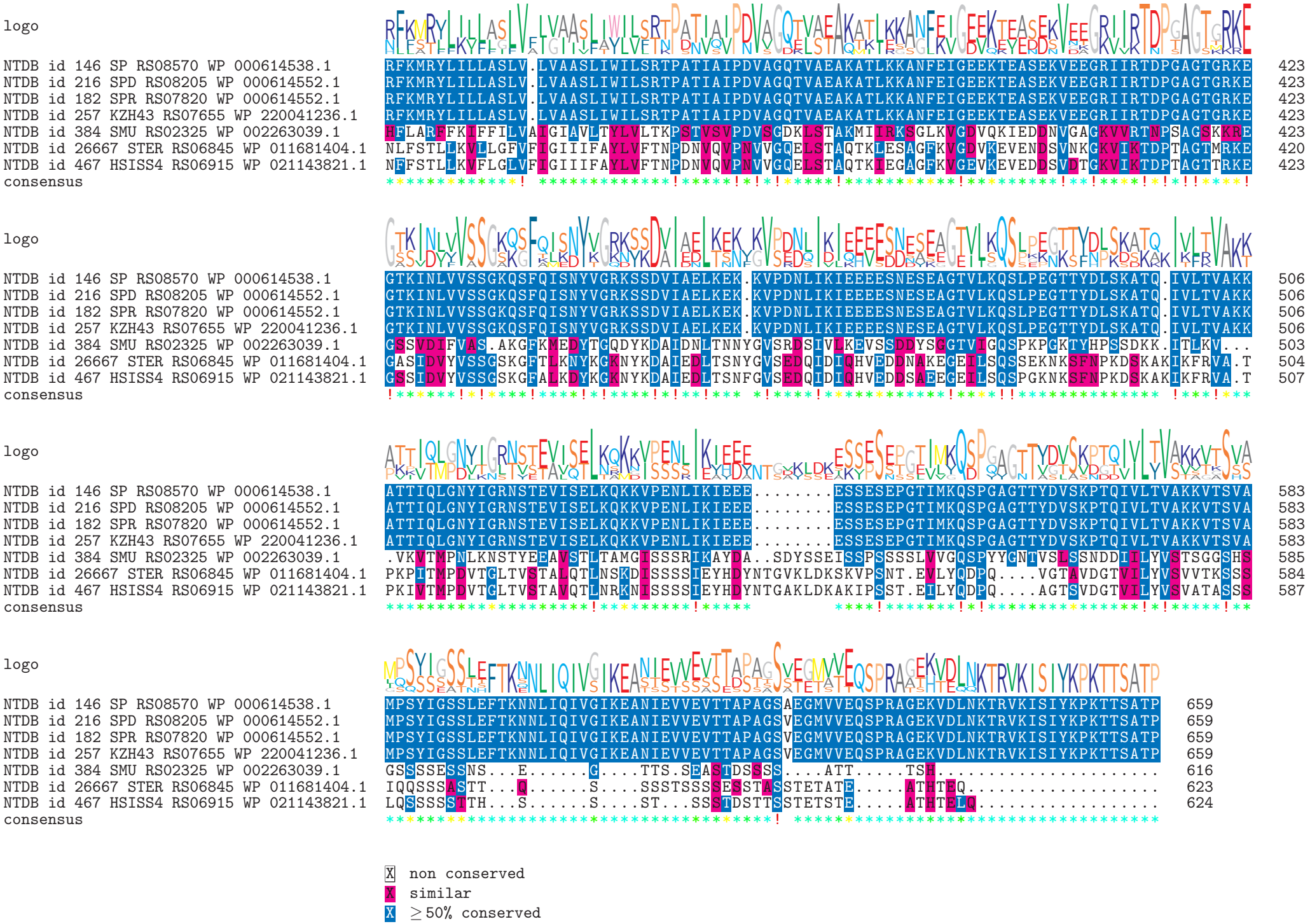
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NTDB id 216 SPD RS08205 WP 000614552.1 RFKMRYLILLASLV.LVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTDPGAGTGRKE 423
NTDB id 182 SPR RS07820 WP 000614552.1 RFKMRYLILLASLV.LVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTDPGAGTGRKE 423
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consensus
logo
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consensus
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NTDB id 384 SMU RS02325 WP 002263039.1 GSSSSESSNS...E......G....TTS.SEASTDSSSS....ATT.....TSH..... 616
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consensus
X non conserved
X similar
X ≥ 50% conserved