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logo
NTDB id 156089 RN86 RS05260 WP 048777855.1  MKKYIFMRILRSLVSIFLVTALTYTIIYTMVPRKLIFKQDPNYNKIATTKDKKANYENTIYERMGYVDYYDTKELQEKASKEDSS  85
NTDB id 476 HSISS4 RS06995 WP 013990225.1   MKKYILLRTLRSLLSIFVVTALTYTIIYTMVPRRLIFRQDPNYNKIAKTADSKANYENTIYERMGYIDYYDTKELQEKASSLDKS  85
NTDB id 325 STU RS16125 WP 002946410.1      MKKYILLRTLRSLLSIFVVTALTYTIIYTMVPRRLIFRQDPNYNKIAKTADSKANYENTIYERLGYIDYYDTKELQEKASSLDKS  85
NTDB id 293 STER RS06925 WP 011681416.1     MKKYILLRTLRSLLSIFVVTALTYTIIYTMVPRRLIFRQDPNYNKIAKTADSKANYENTIYERLGYIDYYDTKELQEKASSLDKS  85
consensus

logo
NTDB id 156089 RN86 RS05260 WP 048777855.1  VTVEGTDANKKIYEAYIKKLGNGWKLHQFEESKQFYAVREVPVLERVLGFYGNLIQFDHTGFVKDETNPKLKRYIRIENDPAIGW  170
NTDB id 476 HSISS4 RS06995 WP 013990225.1   VTTKPTKENKAIYQKYVDSLGHGWRLHQFKQSKQFYATRDVPVLERVFGFYTHLFEFDNTGWVKDKTNPNLKRYIRIENDPAIGW  170
NTDB id 325 STU RS16125 WP 002946410.1      VTTKPTKENKAIYQKYVDSLGRGWRLHQFKQSKQFYATRDVPVLERVFGFYTHLFEFDNTGWVKDKTNPNLKRYIRIENDPAIGW  170
NTDB id 293 STER RS06925 WP 011681416.1     VTTKPTKENKAVYQKYVDSLGRGWRLHQFKQSKQFYATRDVPVLERVFGFYTHLFEFDNTGWVKDKTNPNLKRYIRIENDPAIGW  170
consensus

logo
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NTDB id 325 STU RS16125 WP 002946410.1      SVVGSGTKHKYLLYFNSQFPFVHQNFVKMNLGTSYPTYAEQGVLDVITQGQGQTKSSVVNFPTGKKISSVDIYSRTYKSPSQADA  255
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consensus

logo
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NTDB id 293 STER RS06925 WP 011681416.1     RDRSYYG.NDPYTATKSRYQYPSMVTSSAIAGLIGLVLSYALAIPLGSYMALFKNTLFDSVSTGVLTFLLSLPTIALVYIIRLIG  339
consensus

logo
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NTDB id 476 HSISS4 RS06995 WP 013990225.1   SEIGLPDSFPILGAGDWRSYVLPSVILGLLSTPGLAIWIRRYMIDLQSQDFVRFARAKGLSEQEISNKHIFKNAMVSLVSGIPAS  424
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consensus

logo
NTDB id 156089 RN86 RS05260 WP 048777855.1  VIGVIQGATLTETVFAFPGMGKMLIDSVKASNNSMVVGLVFIFTGLSVFALLLGDILMTMLDPRIKLTTKGGK  498
NTDB id 476 HSISS4 RS06995 WP 013990225.1   IVSVITGATLTETIFAYPGMGKMLIDSVRASNNAMVVGLVFIFTALTIFSLLVGDILMTMIDPRIKLTSKGGK  497
NTDB id 325 STU RS16125 WP 002946410.1      IVGVITGATLTETIFAYPGMGKMLIDSVRASNNAMVVGLVFIFTALSIFSLLVGDILMTIIDPRIKLTSKGGK  497
NTDB id 293 STER RS06925 WP 011681416.1     IVGVITGATLTETIFAYPGMGKMLIDSVRASNNAMVVGLVFIFTALSIFSLLVGDILMTIIDPRIKLTSKGGK  497
consensus
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X non conserved
X similar
X $\geq$ 50% conserved