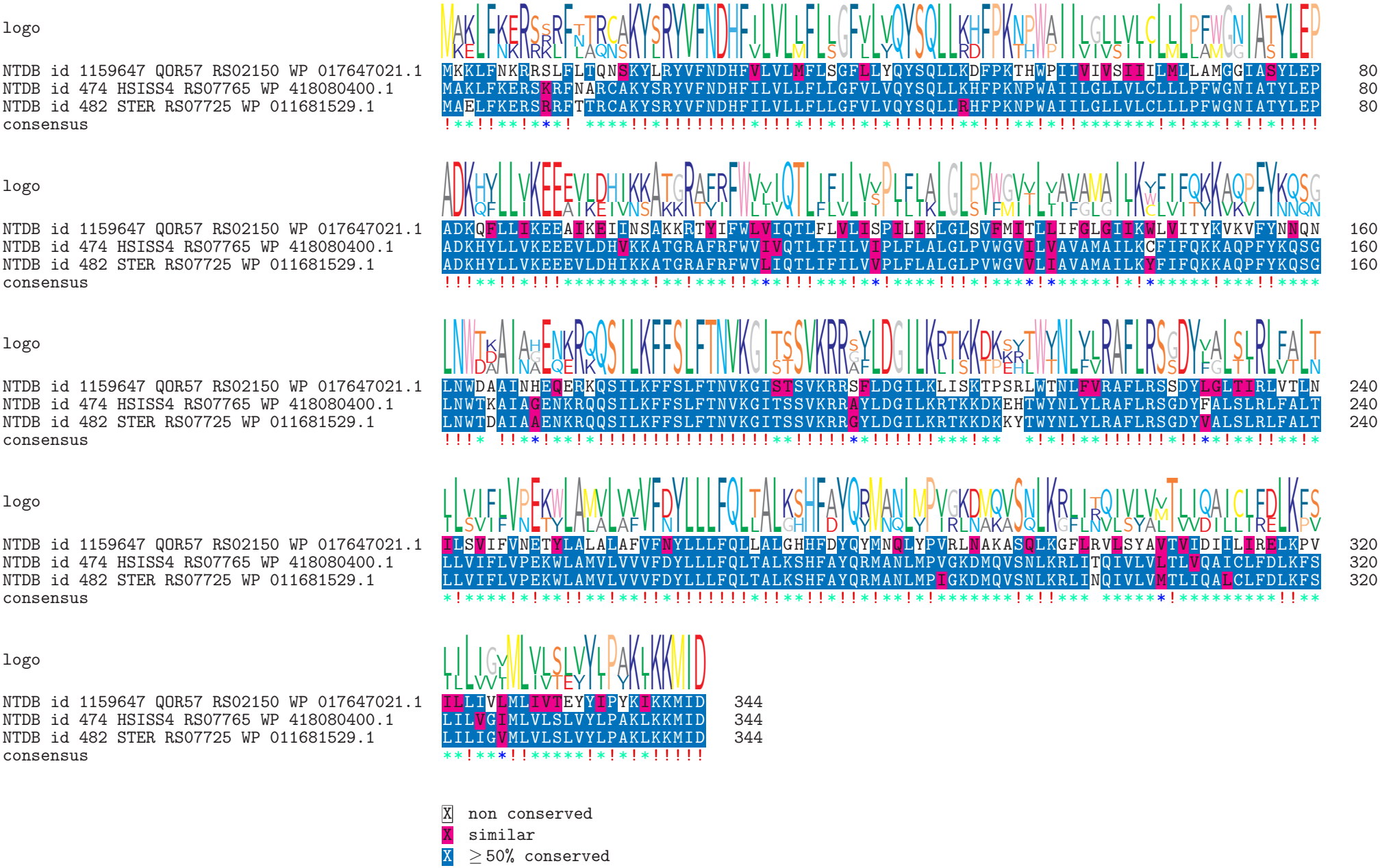
logo
NTDB id 1159647 QOR57 RS02150 WP 017647021.1 MKKLFNKRRSLFLTQNSKYLRYVFNDHFVLVLMFLSGFLLYQYSQLLKDFPKTHWPIIVISIILMLLAMGGIASYLEP 80
NTDB id 474 HSISS4 RS07765 WP 418080400.1 MAKLFKERSKRFNARCAKYSRYVFNDHFILVLLFLLGFVLVQYSQLLKHFPKNPWAIILGLLVLCLLLPFWGNIATYLEP 80
NTDB id 482 STER RS07725 WP 011681529.1 MAELFKERSRRFTTRCAKYSRYVFNDHFILVLLFLLGFVLVQYSQLLRHFPKNPWAIILGLLVLCLLLPFWGNIATYLEP 80
consensus
logo
NTDB id 1159647 QOR57 RS02150 WP 017647021.1 ADKQFLLIKEEAIKEIINSAKKRTYIFWLVIQTLFLVLISPILIKLGLSVFMITLLIFGLGIIKWLVITYKVKVFYNNQN 160
NTDB id 474 HSISS4 RS07765 WP 418080400.1 ADKHYLLVKEEEVLDHVKKATGRAFRFWVIVQTLIFILVIPLFLALGLPVWGVILVAVAMAILKCFIFQKKAQPFYKQSG 160
NTDB id 482 STER RS07725 WP 011681529.1 ADKHYLLVKEEEVLDHIKKATGRAFRFWVLIQTLIFILVVPLFLALGLPVWGVVLIAVAMAILKYFIFQKKAQPFYKQSG 160
consensus
logo
NTDB id 1159647 QOR57 RS02150 WP 017647021.1 LNWDAAINHEQERKQSILKFFSLFTNVKGISTSVKRRSFLDGILKLISKTPSRLWTNLFVRAFLRSSDYLGLTIRLVTLN 240
NTDB id 474 HSISS4 RS07765 WP 418080400.1 LNWTKAIAGENKRQQSILKFFSLFTNVKGITSSVKRRAYLDGILKRTKKDKEHTWYNLYLRAFLRSGDYFALSLRLFALT 240
NTDB id 482 STER RS07725 WP 011681529.1 LNWTDAIAAENKRQQSILKFFSLFTNVKGITSSVKRRGYLDGILKRTKKDKKYTWYNLYLRAFLRSGDYVALSLRLFALT 240
consensus
logo
NTDB id 1159647 QOR57 RS02150 WP 017647021.1 ILSVIFVNETYLALALAFVFNYLLLFQLLALGHHFDYQYMNQLYPVRLNAKASQLKGFLRVLSYAVTVIDILIRELKPV 320
NTDB id 474 HSISS4 RS07765 WP 418080400.1 LLVIFLVPEKWLAMVLVVVFDYLLLFQLTALKSHFAYQRMANLMPVGKDMQVSNLKRLITQIVLVLTLVQAICLFDLKFS 320
NTDB id 482 STER RS07725 WP 011681529.1 LLVIFLVPEKWLAMVLVVVFDYLLLFQLTALKSHFAYQRMANLMPIGKDMQVSNLKRLINQIVLVMTLIQALCLFDLKFS 320
consensus
logo
NTDB id 1159647 QOR57 RS02150 WP 017647021.1 ILLIVLMLIVTEYYIPYKIKKMID 344
NTDB id 474 HSISS4 RS07765 WP 418080400.1 LILVGIMLVLSLVYLPAKLKKMID 344
NTDB id 482 STER RS07725 WP 011681529.1 LILIGVMLVLSLVYLPAKLKKMID 344
consensus
X non conserved
X similar
X ≥ 50% conserved