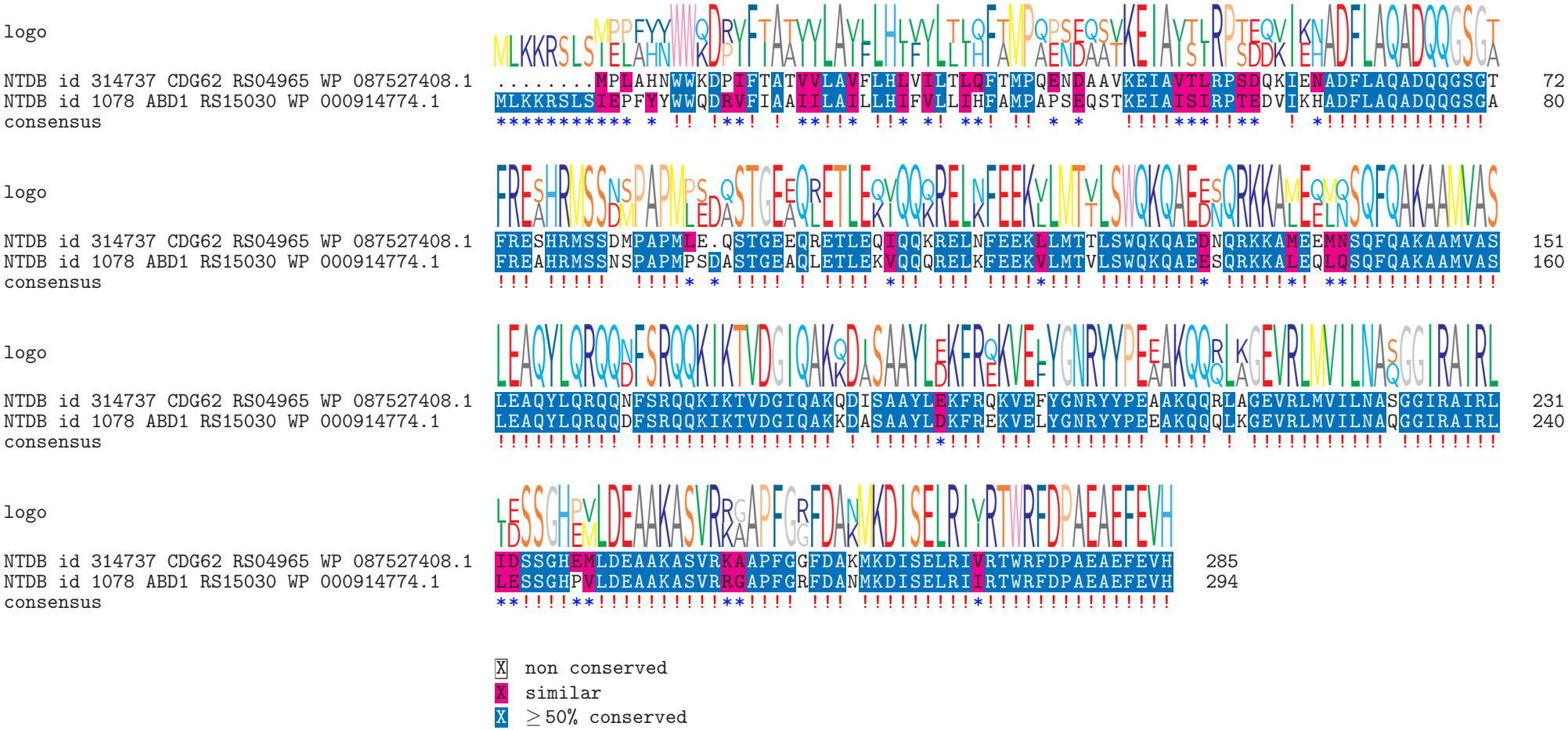
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NTDB id 314737 CDG62 RS04965 WP 087527408.1 ........MPLAHNWWKDPIFTATVVLAVFLHLVILTLQFTMPQENDAAVKEIAVTLRPSDQKIENADFLAQADQQGSGT 72
NTDB id 1078 ABD1 RS15030 WP 000914774.1 MLKKRSLSIEPFYYWWQDRVFIAAIILAILLHIFVLLIHFAMPAPSEQSTKEIAISIRPTEDVIKHADFLAQADQQGSGA 80
consensus ********** * !! !**! ! **!!* !!* *! **! !! * * !!!!***!!** ! *!!!!!!!!!!!!!
logo
NTDB id 314737 CDG62 RS04965 WP 087527408.1 FRESHRMSSDMPAPMLE.QSTGEEQRETLEQIQQKRELNFEEKLLMTTLSWQKQAEDNQRKKAMEEMNSQFQAKAAMVAS 151
NTDB id 1078 ABD1 RS15030 WP 000914774.1 FREAHRMSSNSPAPMPSDASTGEAQLETLEKVQQQRELKFEEKVLMTVLSWQKQAEESQRKKALEQLQSQFQAKAAMVAS 160
consensus !!! !!!!! !!!!* * !!!! ! !!!! *!! !!! !!!!*!!! !!!!!!!!* !!!!!*! **!!!!!!!!!!!!
logo
NTDB id 314737 CDG62 RS04965 WP 087527408.1 LEAQYLQRQQNFSRQQKIKTVDGIQAKQDISAAYLEKFRQKVEFYGNRYYPEEAKQQRLAGEVRLMVILNASGGIRAIRL 231
NTDB id 1078 ABD1 RS15030 WP 000914774.1 LEAQYLQRQQDFSRQQKIKTVDGIQAKKDASAAYLDKFREKVELYGNRYYPEEAKQQQLKGEVRLMVILNAQGGIRAIRL 240
consensus !!!!!!!!!! !!!!!!!!!!!!!!!! ! !!!!!*!!! !!! !!!!!!!! !!!! ! !!!!!!!!!!! !!!!!!!!
logo
NTDB id 314737 CDG62 RS04965 WP 087527408.1 IDSSGHEMLDEAAKASVRKAAPFGGFDAKMKDISELRIVRTWRFDPAEAEFEVH 285
NTDB id 1078 ABD1 RS15030 WP 000914774.1 LESSGHPVLDEAAKASVRRGAPFGRFDANMKDISELRIIRTWRFDPAEAEFEVH 294
consensus **!!!!**!!!!!!!!!!!**!!!! !!! !!!!!!!!!*!!!!!!!!!!!!!!!!
X non conserved
X similar
X ≥ 50% conserved