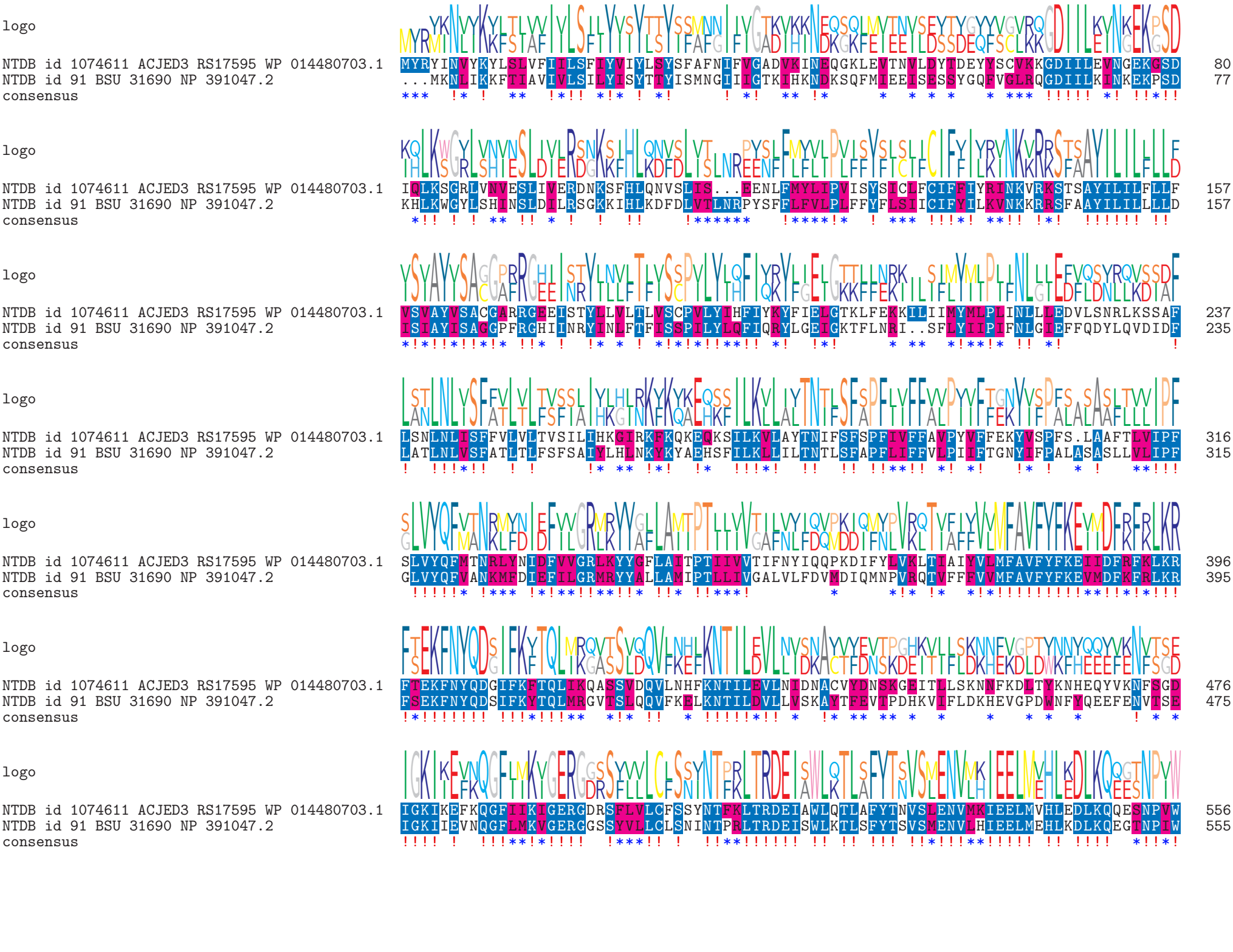
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NTDB id 91 BSU 31690 NP 391047.2 ...MKNLIKKFTIAVIVLSILYISYTTYISMNGIIIGTKIHKNDKSQFMIEEISESSYGQFVGLRQGDIILKINKEKPSD 77
consensus
logo
NTDB id 1074611 ACJED3 RS17595 WP 014480703.1 IQLKSGRLVNVESLIVERDNKSFHLQNVSLIS...EENLFMYLIPVISYSICLFCIFRIYRINKVRKSTSAYILILFLLF 157
NTDB id 91 BSU 31690 NP 391047.2 KHLKWGYLSHINSLDILRSGKKIHLKDFDLVTLNRPYSFFLFVLPLFFYFLSIICIFYILKVNKKRRSFAAYILILLLD 157
consensus
logo
NTDB id 1074611 ACJED3 RS17595 WP 014480703.1 VSVAYVSACGARRGEEISTYLLVLTLVSCPVLYIHFIYKYFIELGTKLFEKKILIIMYMLPLINLLLEDVLSNRLKSSAF 237
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consensus
logo
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NTDB id 91 BSU 31690 NP 391047.2 LATLNLVSFATLTLFSFSAIYLHLNKYKYAEHSFILKLLILTNTLSFAPFLIFFVLPIIFTGNYIFPALASASLLVLIPF 315
consensus
logo
NTDB id 1074611 ACJED3 RS17595 WP 014480703.1 SLVYQFMTNRLYNIDFVVGRLKYYGFLAITPTIIVVTIFNYIQQPKDIFYLVKLTIAIYVLMFAVFYFKEIIDFRFKLKR 396
NTDB id 91 BSU 31690 NP 391047.2 GLVYQFVANKMFDIEFILGRMRYYALLAMIPTLLIVGALVLFDVMDIQMNPVRQTVFFFVVMFAVFYFKEVMDFKFRLKR 395
consensus
logo
NTDB id 1074611 ACJED3 RS17595 WP 014480703.1 FTEKFNYQDGIFKFTQLIKQASSVDQVLNHFKNTILEVLNIDNACVYDNSKGEITLLSKNNFKDLTYKNHEQYVKNFSGD 476
NTDB id 91 BSU 31690 NP 391047.2 FSEKFNYQDSIFKYTQLMRGVTSLQQVFKELKNTILDVLLVSKAYTFEVTPDHKVIFLDKHEVGPDWNFYQEEFENVTSE 475
consensus
logo
NTDB id 1074611 ACJED3 RS17595 WP 014480703.1 IGKIKEFKQGFIIKIGERGDRSFLVLCFSSYNTFKLTRDEIAWLQTLAFYTNVSLENVMKIEELMVHLEDLKQQESNPVW 556
NTDB id 91 BSU 31690 NP 391047.2 IGKIIEVNQGFLMKVGERGGSSYVLLCLSNINTPRLTRDEISWLKTLSFYTSVSMENVLHIEELMEHLKDLKQEGTNPIW 555
consensus

logo

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NTDB id 1074611 ACJED3 RS17595 WP 014480703.1  LKKLMYTIEEKQRSDLARDLHDSVLQDLISLKRQCELFLGDFKKGDNPCREEVQDKLVQMNEQMSDVISMTRETCHELRP  636
NTDB id 91 BSU 31690 NP 391047.2               LKKLMFAIEEKQRSGLARDLHDSVLQDLISLKRQCELFLGDFKKDDNPCREEVQDKLVQMNEQMSDVISMTRETCHELRP  635
consensus                                      !!!!!* !!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

logo

```
NTDB id 1074611 ACJED3 RS17595 WP 014480703.1  QLLYDLGLVKALSKLVAQQQERVPFHIRLNTGRFTASLDLDSQLNLYRIIQEFLSNAVKHSQATDVLIMLISIQNKIVLH  716
NTDB id 91 BSU 31690 NP 391047.2               QLLYDLGLVKALSKLVAQQQERVPFHIRLNTGRFTASLDLDSQLNLYRIIQEFLSNAVKHSQATDVLIMLISIQNKIVLH  715
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

logo

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NTDB id 1074611 ACJED3 RS17595 WP 014480703.1  YEDDGVGFDQEKNTEHSMSMGLSGIKERVRALDGRLRIETSEGKGFKADIEIEL  770
NTDB id 91 BSU 31690 NP 391047.2               YEDDGVGFDQEKNTEHSMSMGLSGIKERVRALDGRLRIETSEGKGFKADIEIEL  769
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved