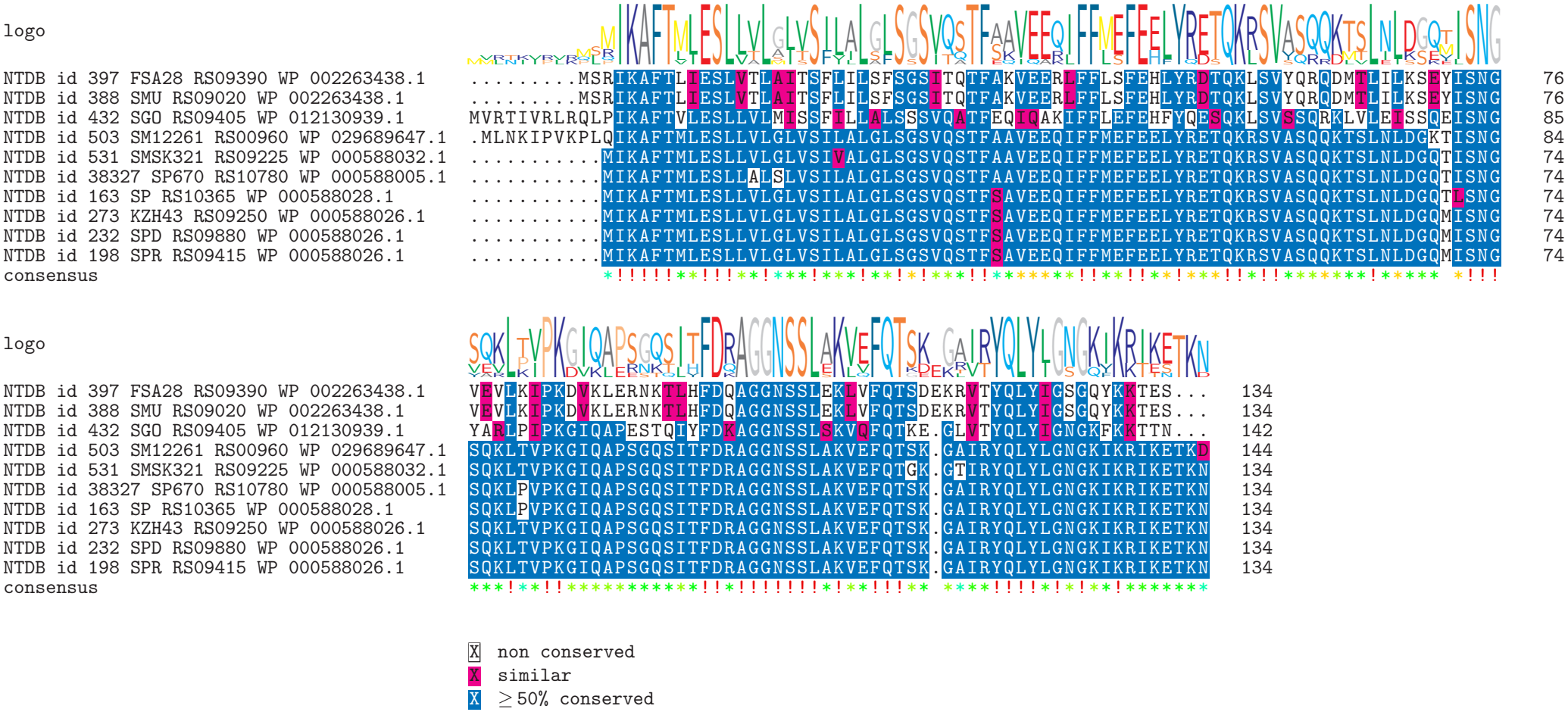
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NTDB id 397 FSA28 RS09390 WP 002263438.1 .....MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSEYISNG 76
NTDB id 388 SMU RS09020 WP 002263438.1 .....MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSEYISNG 76
NTDB id 432 SGO RS09405 WP 012130939.1 MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVLEISSQEISNG 85
NTDB id 503 SM12261 RS00960 WP 029689647.1 .MLNKIPVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGKTISNG 84
NTDB id 531 SMSK321 RS09225 WP 000588032.1 .....MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQTISNG 74
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NTDB id 198 SPR RS09415 WP 000588026.1 .....MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQMISNG 74
consensus *!!!!!**!!!**!***!***!**!*!***!!*******!!**!!**!***!!*!!*******!***** *!!!
logo
NTDB id 397 FSA28 RS09390 WP 002263438.1 VEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES... 134
NTDB id 388 SMU RS09020 WP 002263438.1 VEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES... 134
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NTDB id 503 SM12261 RS00960 WP 029689647.1 SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKD 144
NTDB id 531 SMSK321 RS09225 WP 000588032.1 SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTGK.GTIRYQLYLGNGKIKRIKETKN 134
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NTDB id 163 SP RS10365 WP 000588028.1 SQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN 134
NTDB id 273 KZH43 RS09250 WP 000588026.1 SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN 134
NTDB id 232 SPD RS09880 WP 000588026.1 SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN 134
NTDB id 198 SPR RS09415 WP 000588026.1 SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN 134
consensus ***!**!!**********!!*!!!!!!!*!**!!!** ****!!!!*!*!**!*******
X non conserved
X similar
X ≥ 50% conserved