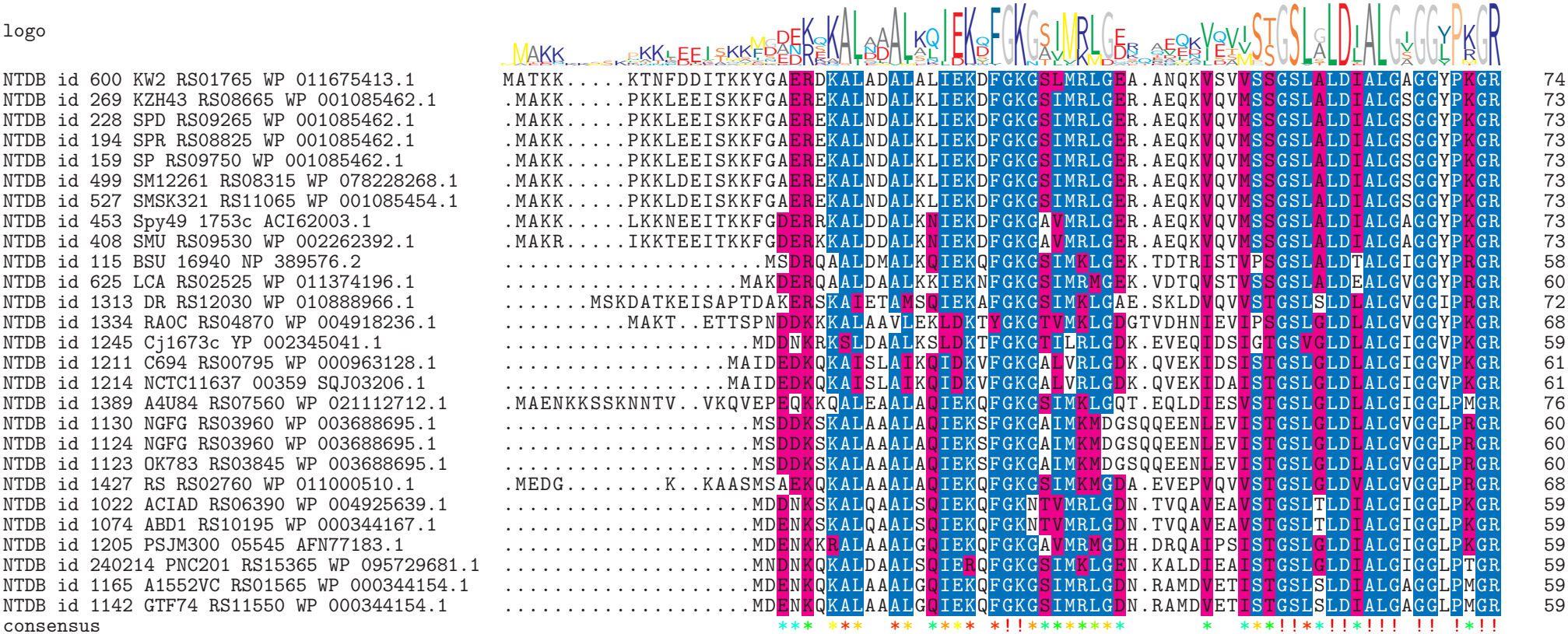
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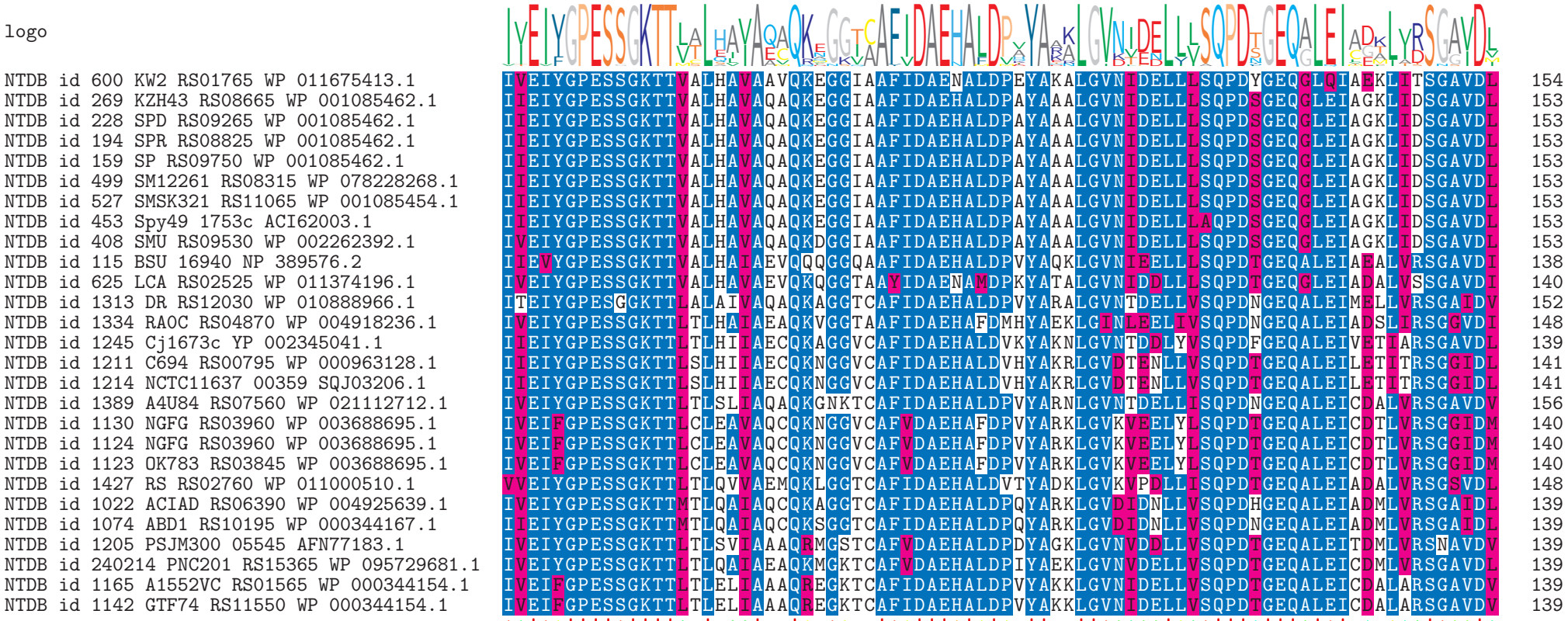
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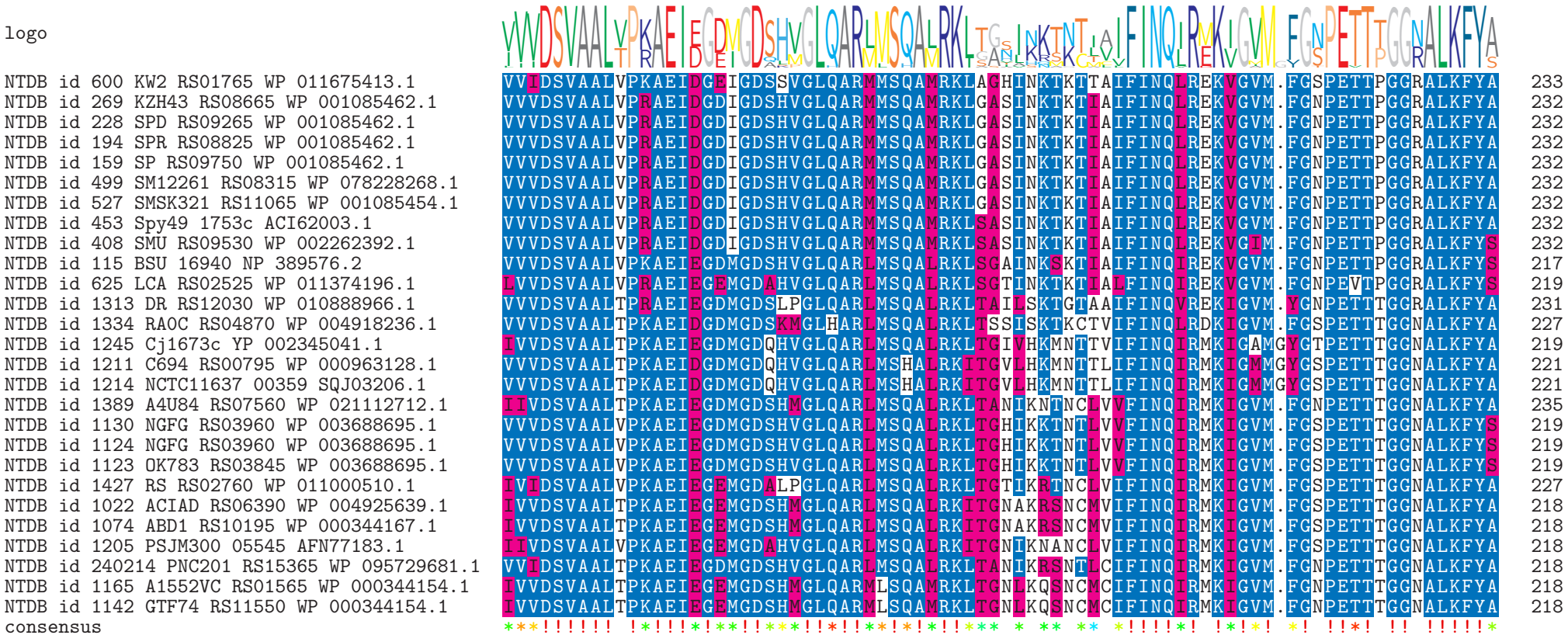
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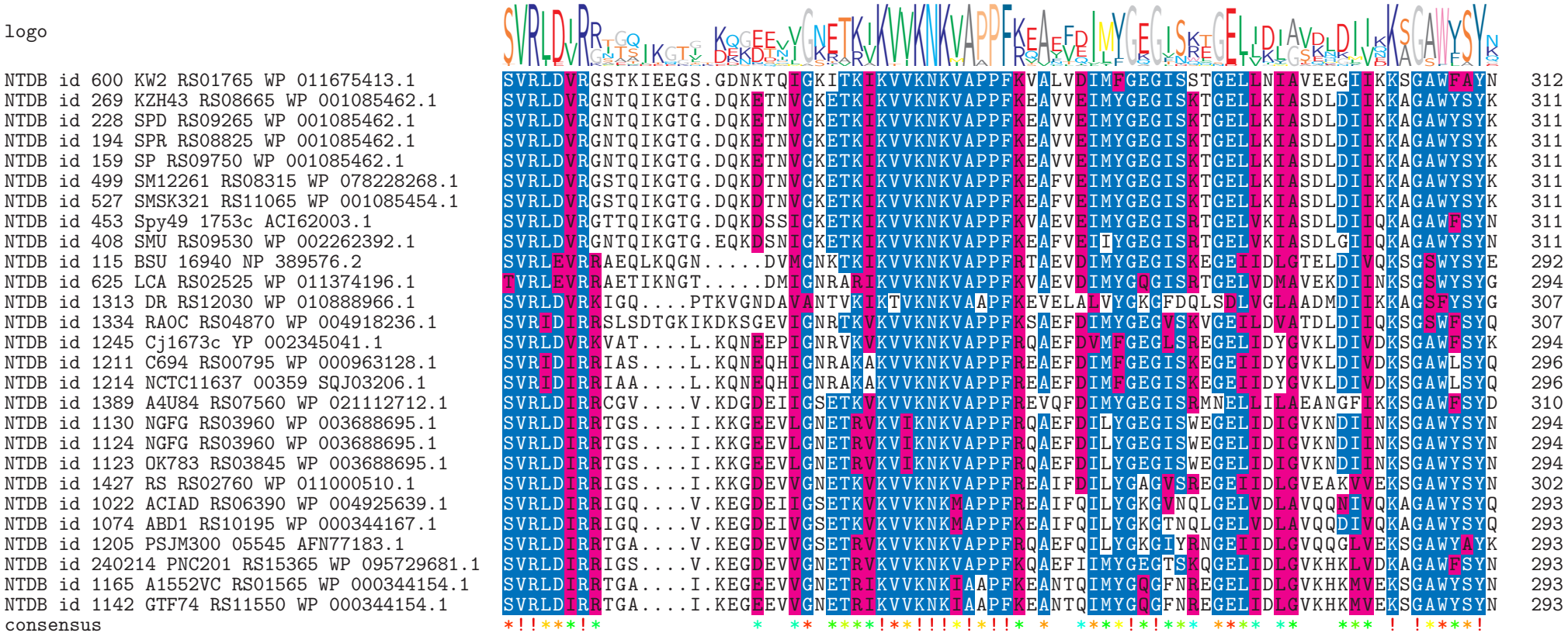
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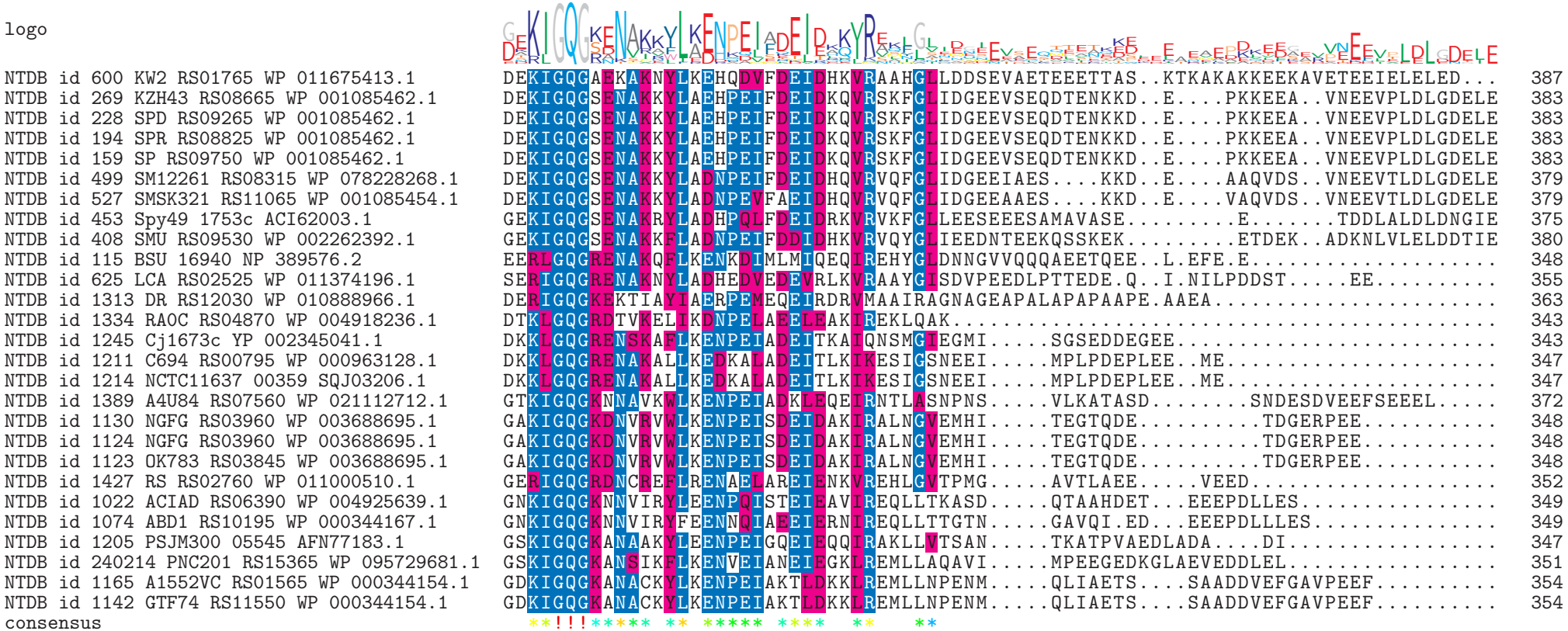
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NTDB id 408 SMU RS09530 WP 002262392.1 GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.........ETDEK..ADKNLVLELDDTIE 380
NTDB id 115 BSU 16940 NP 389576.2 EERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQAEETQEE..L.EFE.E..... 348
NTDB id 625 LCA RS02525 WP 011374196.1 SERIGQGRENAKNYLADHEDVEDEVRLKVRAAYGISDVPEEDLPTTEDE.Q..I.NILPDDST.....EE..... 355
NTDB id 1313 DR RS12030 WP 010888966.1 DERIGQGKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPE.AAEA..... 363
NTDB id 1334 RA0C RS04870 WP 004918236.1 DTKLGQGRDTVKELIKDNPELAEELEAKIREKLQAK..... 343
NTDB id 1245 Cj1673c YP 002345041.1 DKKLGQGRENSKAFLKENPEIADEITKAIQNSMGIEGMI.....SGSEDDEGEE..... 343
NTDB id 1211 C694 RS00795 WP 000963128.1 DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI.....MPLPDEPLEE..ME..... 347
NTDB id 1214 NCTC11637 00359 SQJ03206.1 DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI.....MPLPDEPLEE..ME..... 347
NTDB id 1389 A4U84 RS07560 WP 021112712.1 GTKIGQGKNNAVKWLKENPEIADKLEQEIRNTLASNPNS.....VLKATASD........SNDESDVEEFSEEEL..... 372
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NTDB id 1427 RS RS02760 WP 011000510.1 GERIGQGRDNCREFLRENAELAREIENKVREHLGVTPMG.....AVTLAEE.....VEED..... 352
NTDB id 1022 ACIAD RS06390 WP 004925639.1 GNKIGQGKNNVIRYLEENPQISTEIEAVIREQLLTKASD.....QTAAHDET...EEEPDLLES..... 349
NTDB id 1074 ABD1 RS10195 WP 000344167.1 GNKIGQGKNNVIRYFEENNQIAFEIERNIREQLLTTGTN.....GAVQI.ED...EEEPDLLLES..... 349
NTDB id 1205 PSJM300 05545 AFN77183.1 GSKIGQGKANAAKYLEENPEIGQEIEQQIRAKLLVTSAN.....TKATPVAEDLADA....DI..... 347
NTDB id 240214 PNC201 RS15365 WP 095729681.1 GSKIGQGKANSIKFLKENVEIANEIEGKLREMLLAQAVI.....MPEEGEDKGLAEVEDDLEL..... 351
NTDB id 1165 A1552VC RS01565 WP 000344154.1 GDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPENM.....QLIAETS....SAADDVEFGAVPEEF..... 354
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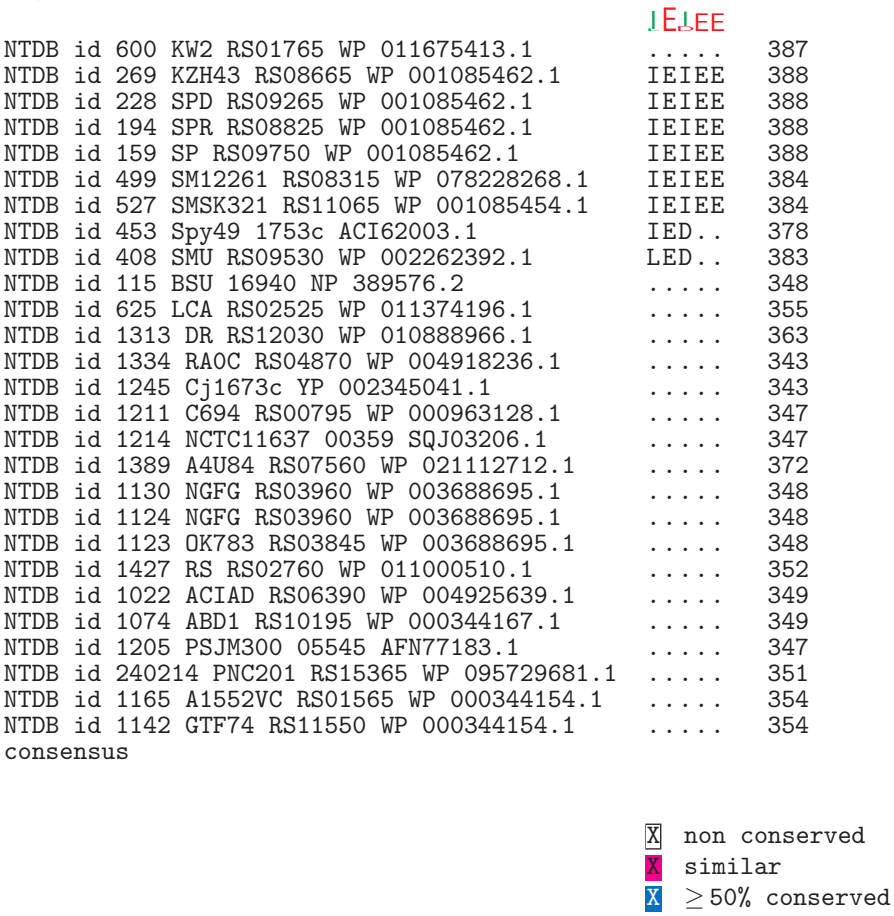
consensus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NTDB id 269 KZH43 RS08665 WP 001085462.1 IEIEE 388
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NTDB id 194 SPR RS08825 WP 001085462.1 IEIEE 388
NTDB id 159 SP RS09750 WP 001085462.1 IEIEE 388
NTDB id 499 SM12261 RS08315 WP 078228268.1 IEIEE 384
NTDB id 527 SMSK321 RS11065 WP 001085454.1 IEIEE 384
NTDB id 453 Spy49 1753c ACI62003.1 IED.. 378
NTDB id 408 SMU RS09530 WP 002262392.1 LED.. 383
NTDB id 115 BSU 16940 NP 389576.2 348
NTDB id 625 LCA RS02525 WP 011374196.1 355
NTDB id 1313 DR RS12030 WP 010888966.1 363
NTDB id 1334 RA0C RS04870 WP 004918236.1 343
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consensus
X non conserved
X similar
X ≥ 50% conserved