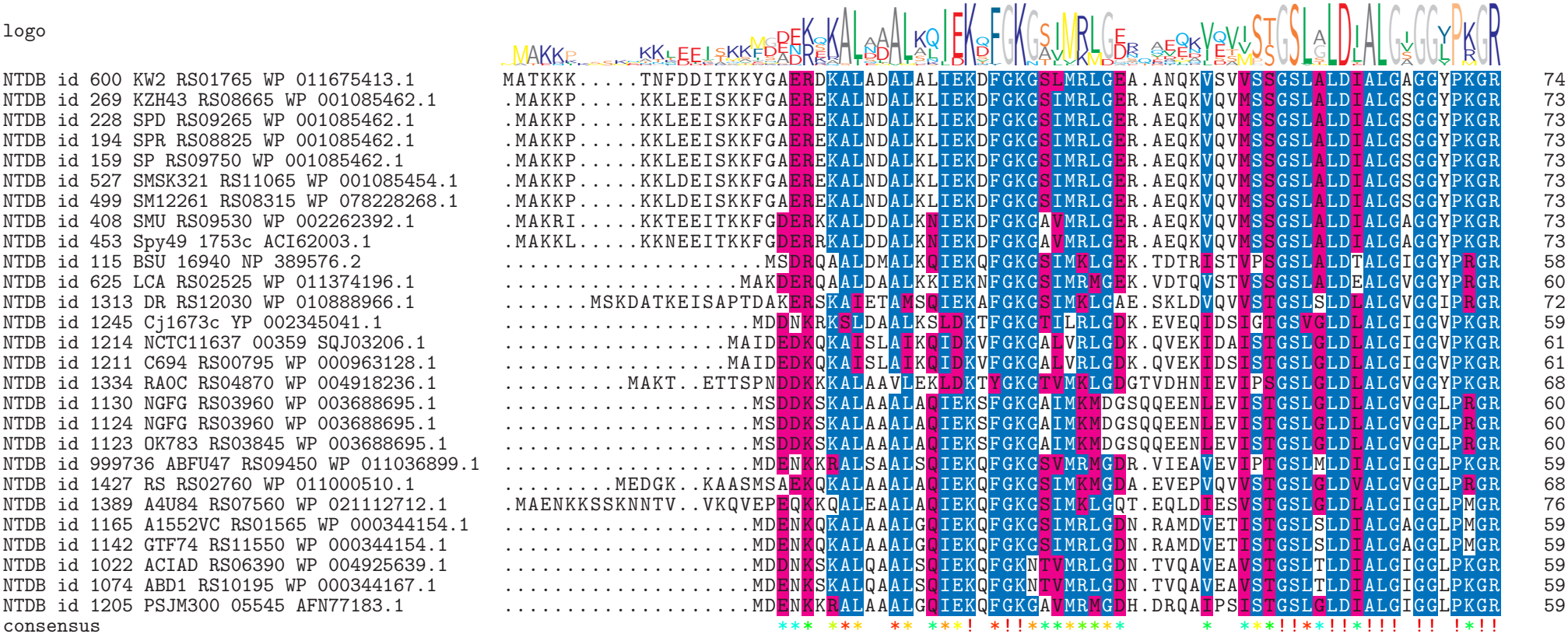
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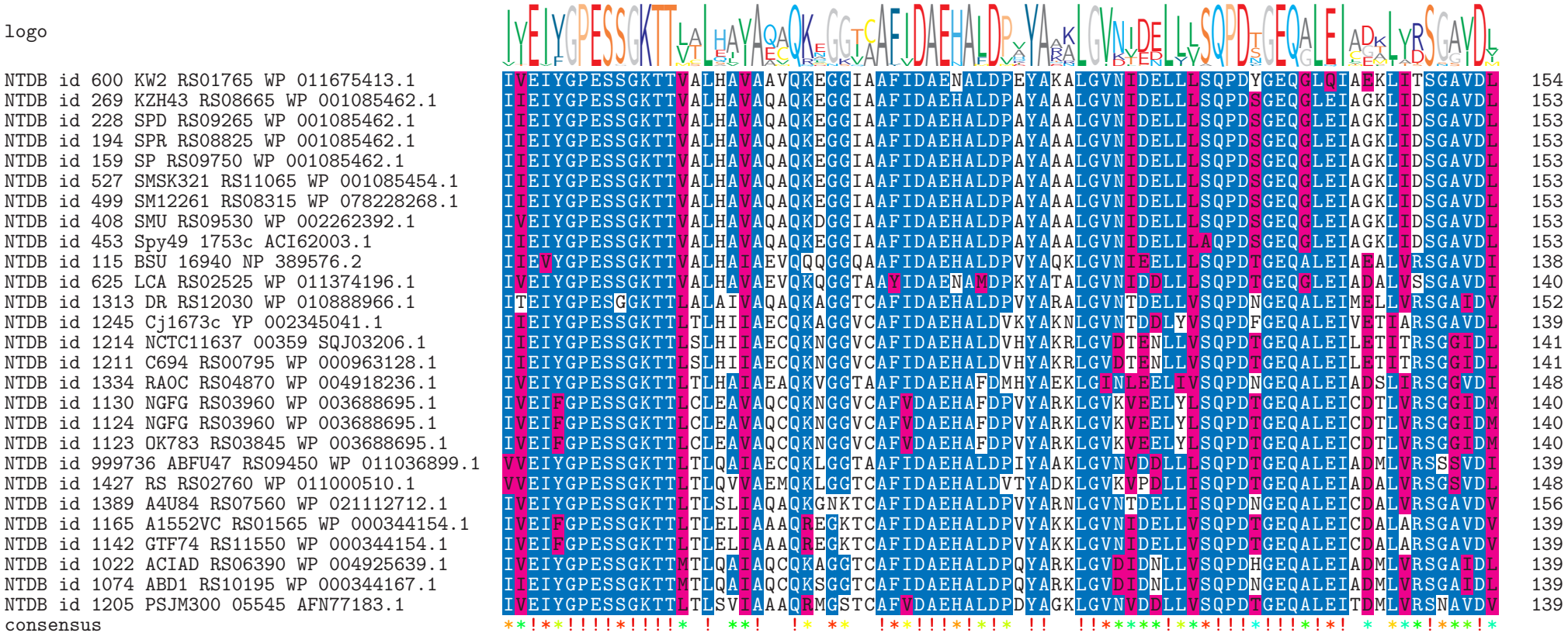
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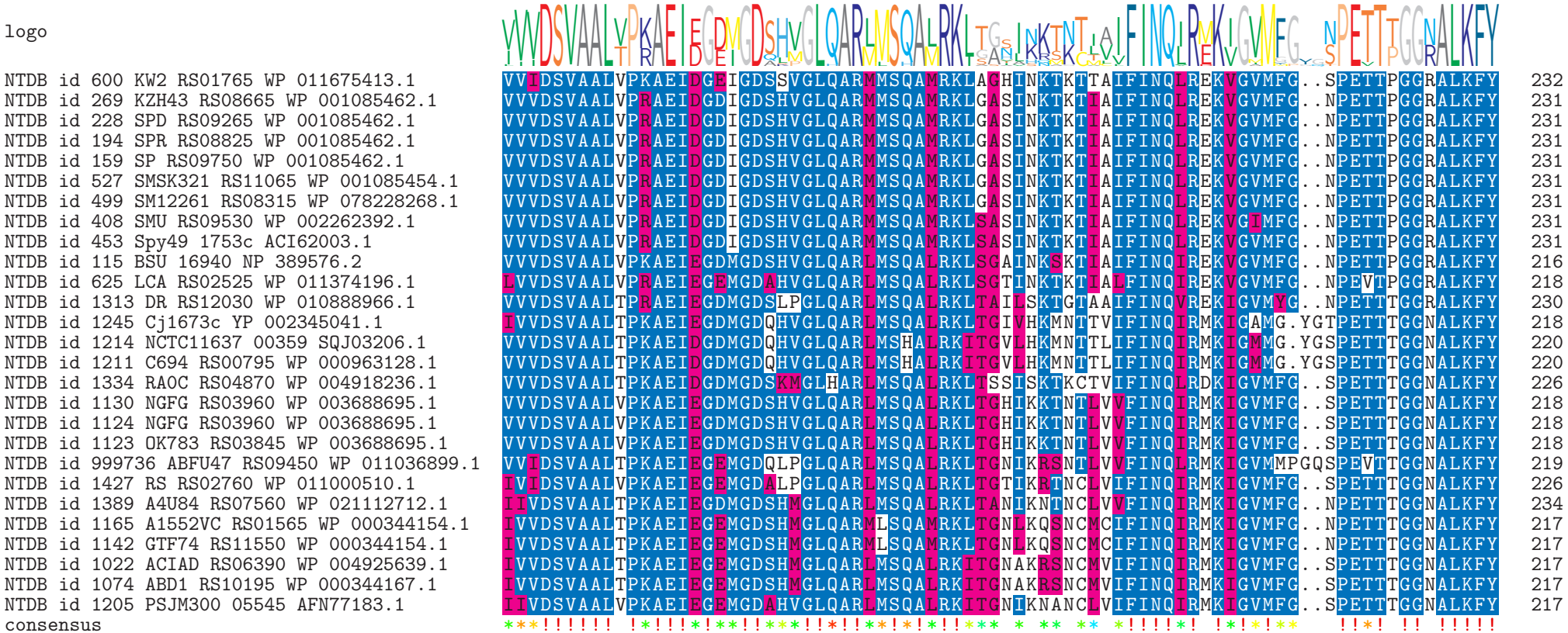
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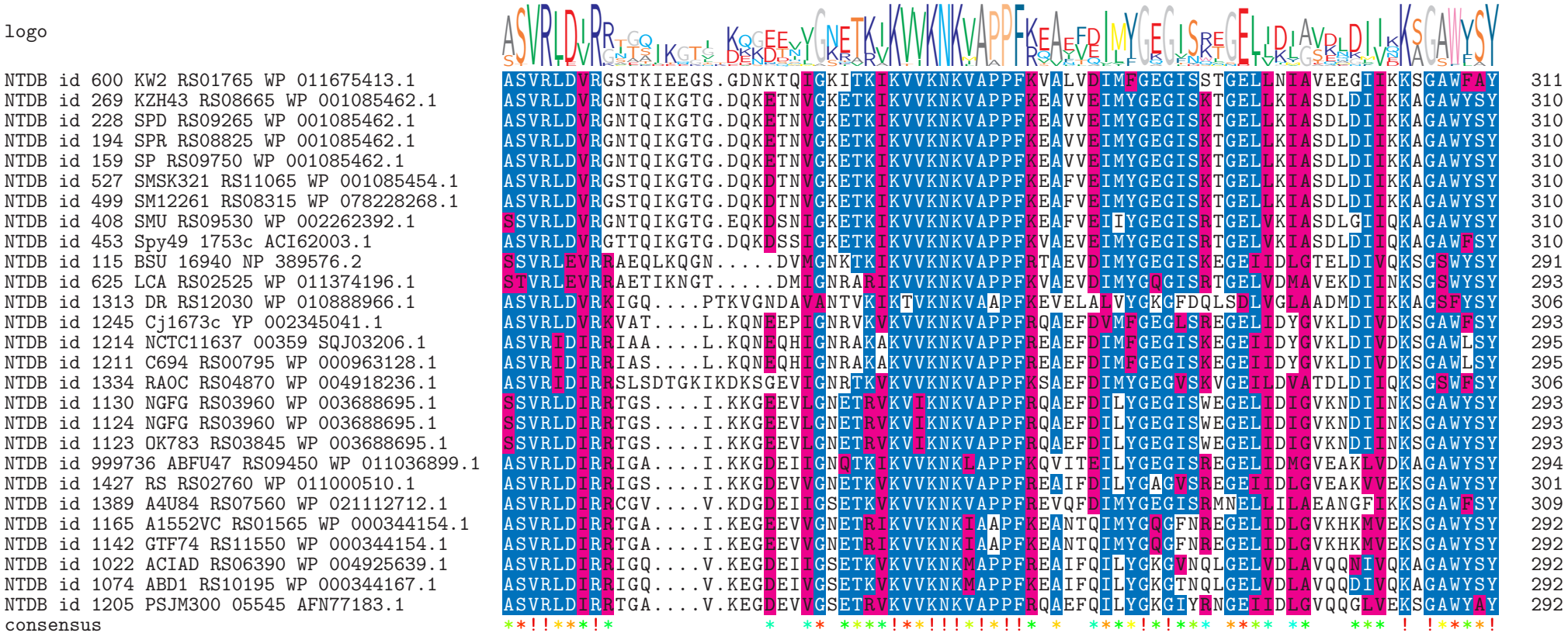
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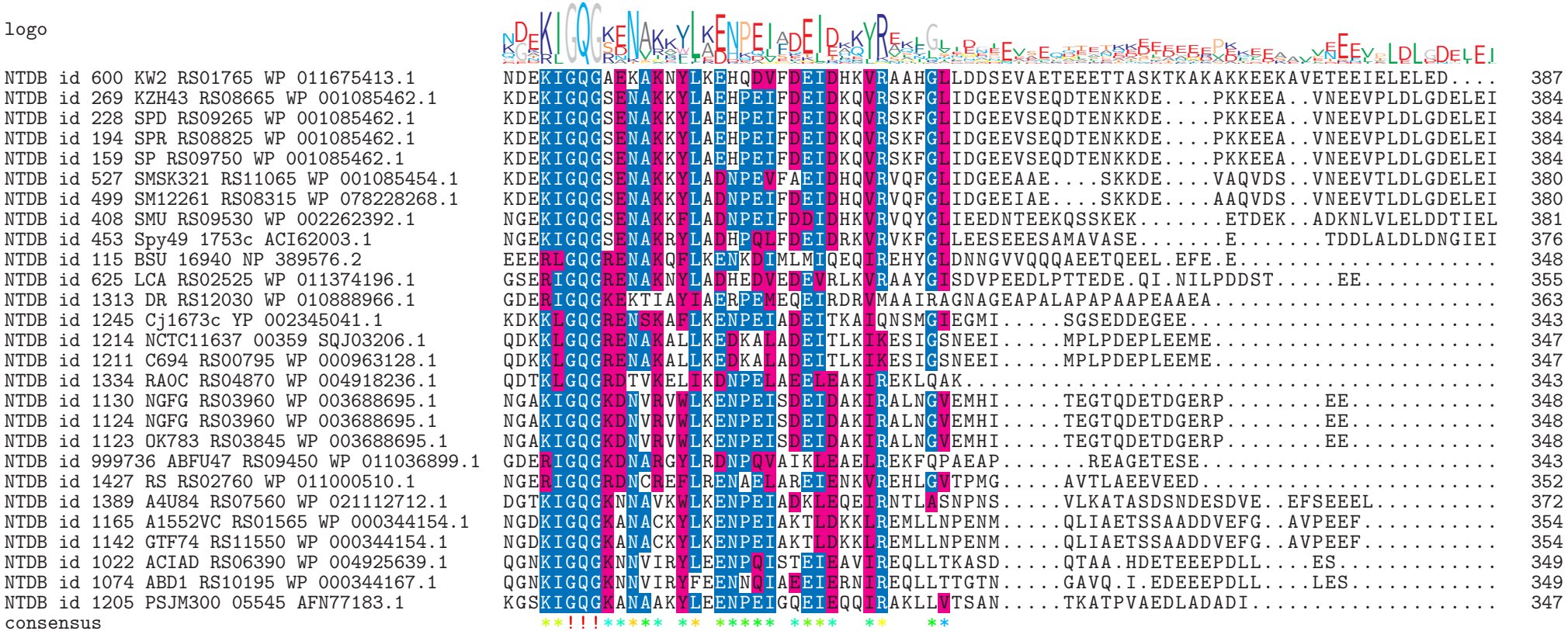
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NTDB id 453 Spy49 1753c ACI62003.1 NGEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEESAMAVASE.......E.......TDDLALDLDNGIEI 376
NTDB id 115 BSU 16940 NP 389576.2 EEERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQAEETQEEL.EFE.E..... 348
NTDB id 625 LCA RS02525 WP 011374196.1 GSERIGQGRENAKNYLADHEDVEDEVRLKVRAAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE..... 355
NTDB id 1313 DR RS12030 WP 010888966.1 GDERIGQGKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAEA..... 363
NTDB id 1245 Cj1673c YP 002345041.1 KDKKLGQGRENSKAFLKENPEIADEITKAIQNSMGIEGMI.....SGSEDDEGEE..... 343
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NTDB id 1334 RA0C RS04870 WP 004918236.1 QDTKLGQGRDTVKELIKDNPELAEELEAKIREKLQAK..... 343
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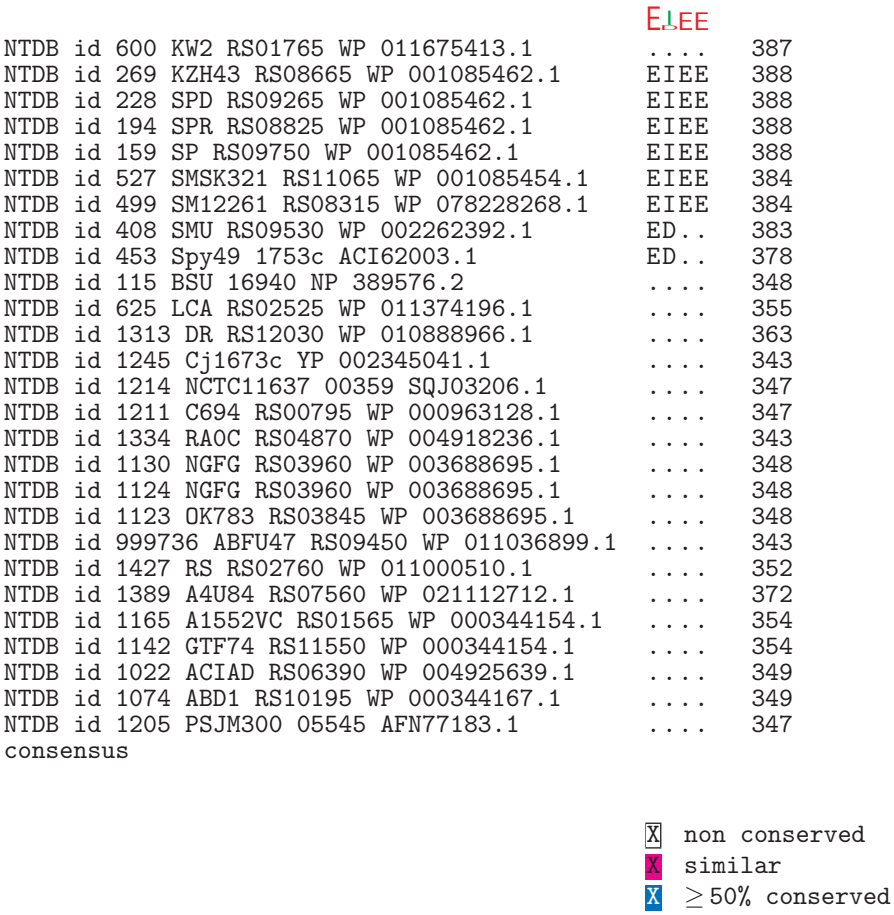
NTDB id 999736 ABFU47 RS09450 WP 011036899.1 GDERIGQGKDNARGYLRDNPQVAIKLEAELREKFQPAEAP.....REAGETESE..... 343
NTDB id 1427 RS RS02760 WP 011000510.1 NGERIGQGRDNCREFLRENAELAREIENKVREHLGVTPMG.....AVTLAEEVEED..... 352
NTDB id 1389 A4U84 RS07560 WP 021112712.1 DGTKIGQGKNNAVKWLKENPEIADKLEQEIRNTLASNPNS.....VLKATASDSNDESDVE..EFSEEEL..... 372
NTDB id 1165 A1552VC RS01565 WP 000344154.1 NGDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPENM.....QLIAETSSAADDVEFG..AVPEEF..... 354
NTDB id 1142 GTF74 RS11550 WP 000344154.1 NGDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPENM.....QLIAETSSAADDVEFG..AVPEEF..... 354
NTDB id 1022 ACIAD RS06390 WP 004925639.1 QGNKIGQGKNNVIRYLEENPQISTEIEAVIREQLLTKASD.....QTAA.HDETEEEPDLL....ES..... 349
NTDB id 1074 ABD1 RS10195 WP 000344167.1 QGNKIGQGKNNVIRYFEENNQIAEEIERNIREQLLTTGTN.....GAVQ.I.EDEEEPDLL....LES..... 349
NTDB id 1205 PSJM300 05545 AFN77183.1 KGSKIGQGKANAAKYLEENPEIGQEIEQQIRAKLLVTSAN.....TKATPVAEDLADADI..... 347
consensus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consensus
X non conserved
X similar
X ≥ 50% conserved