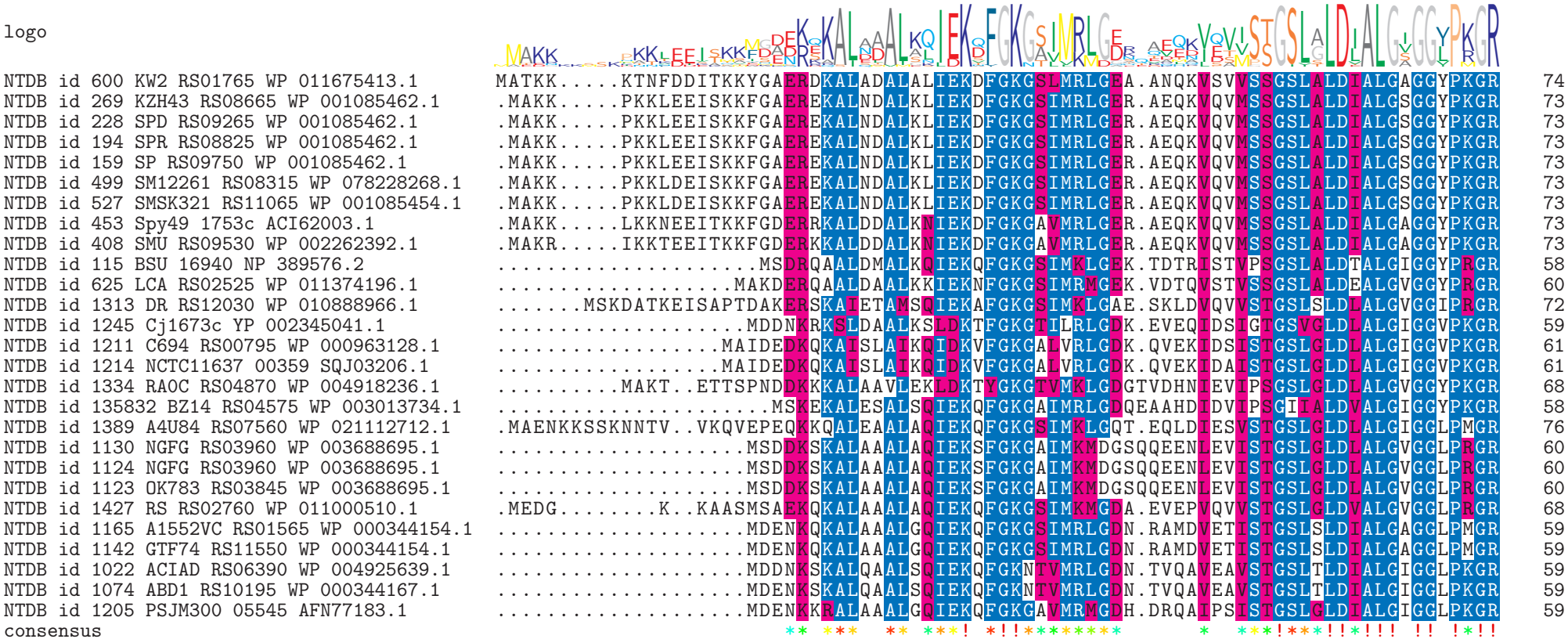
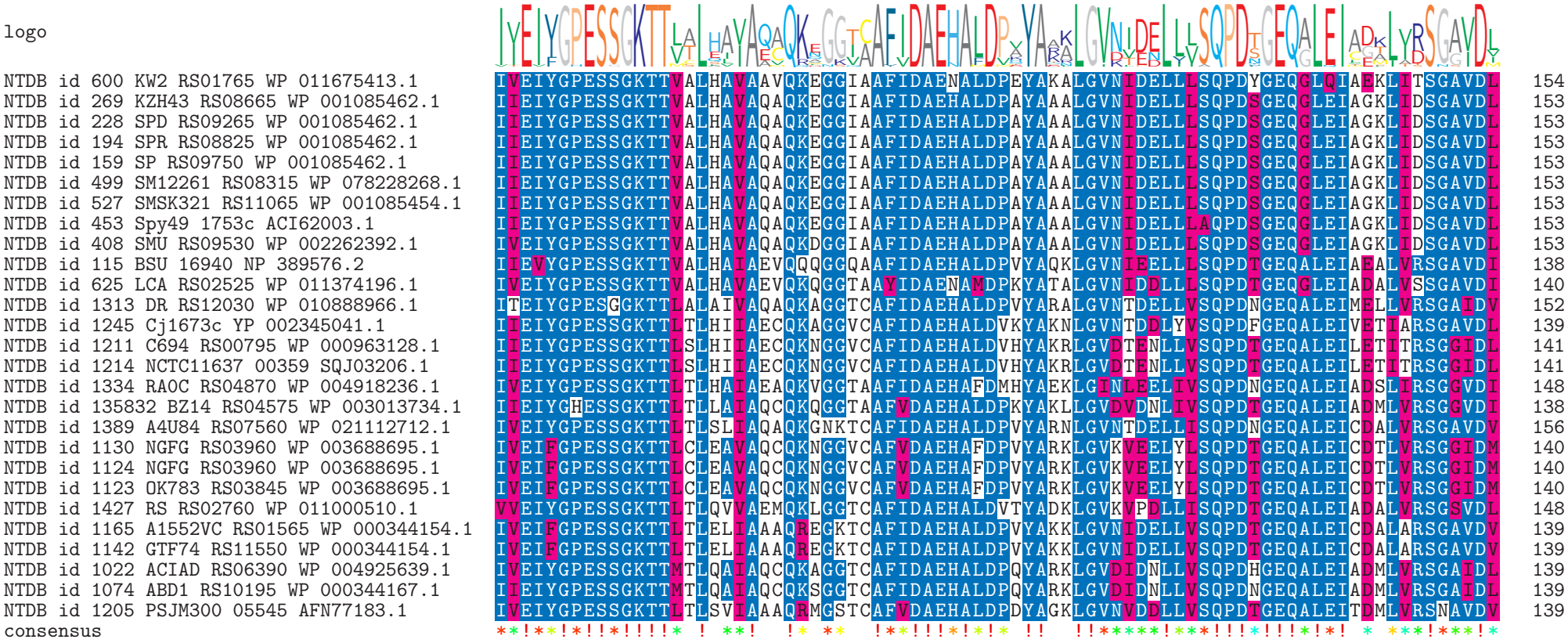
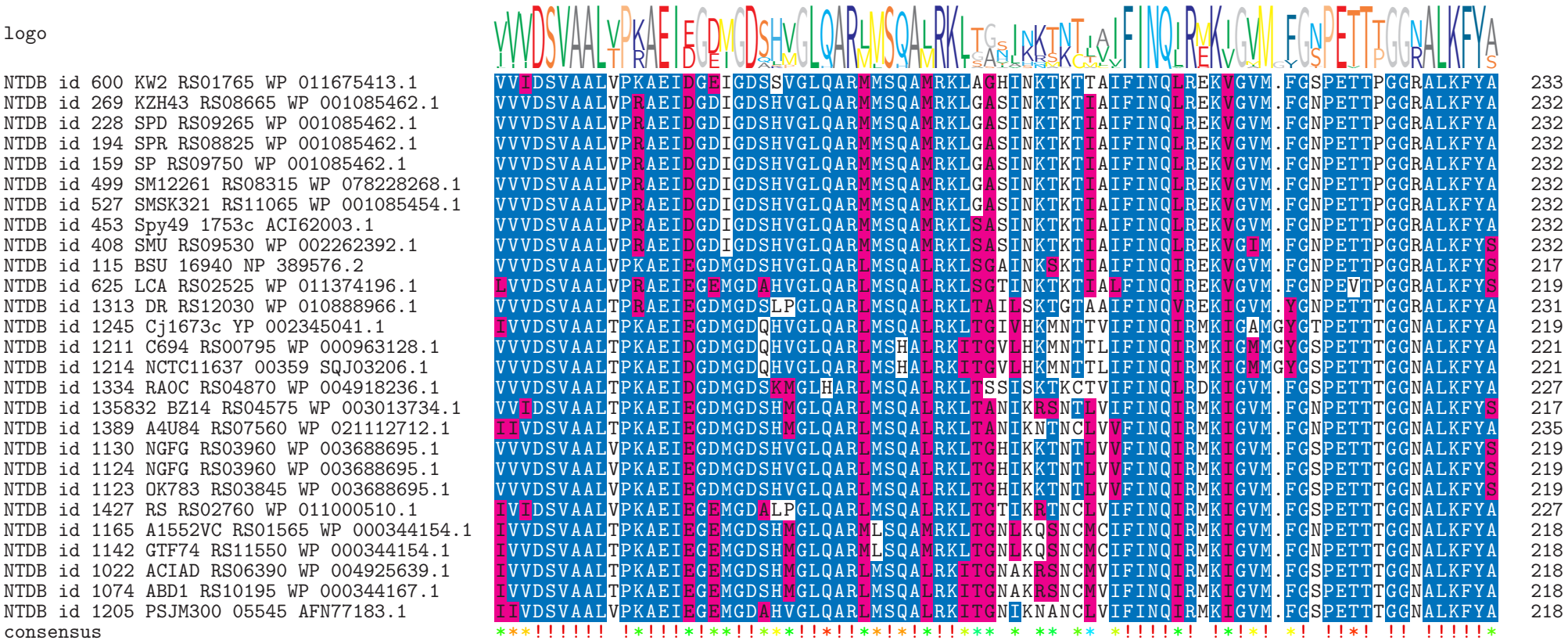
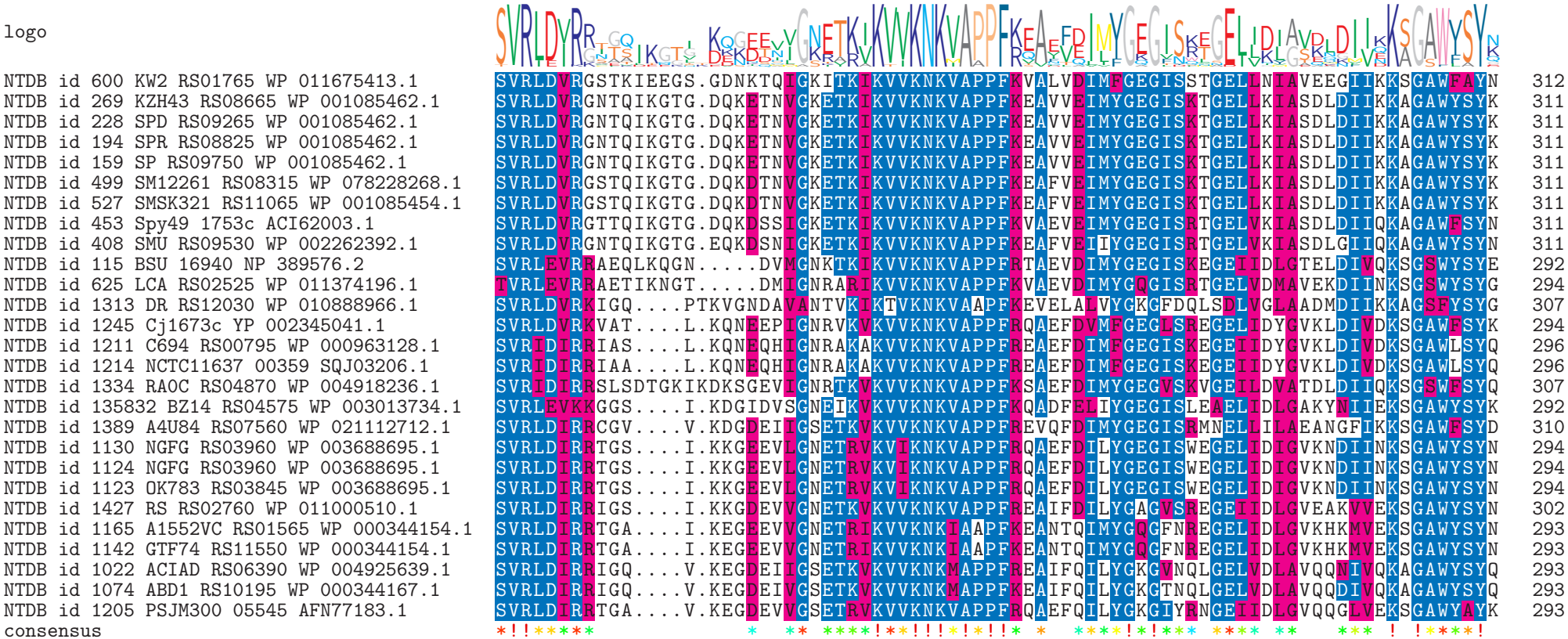
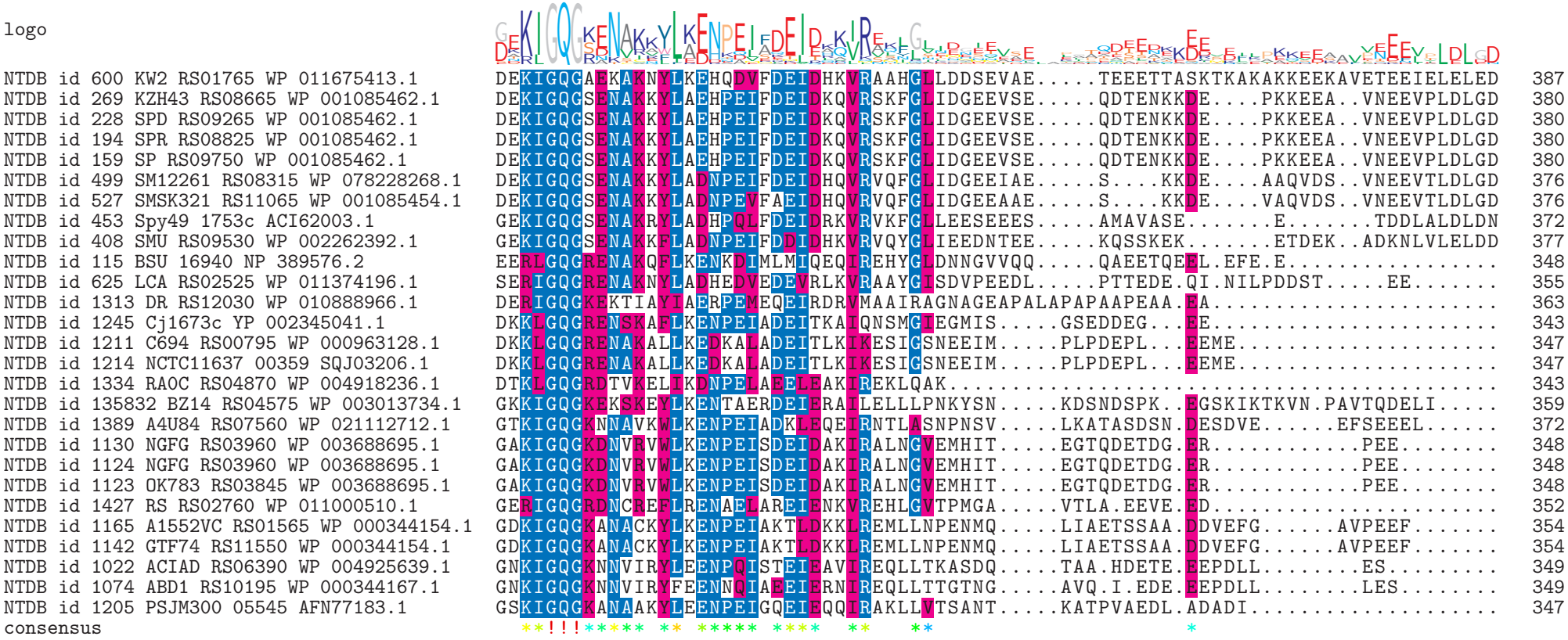
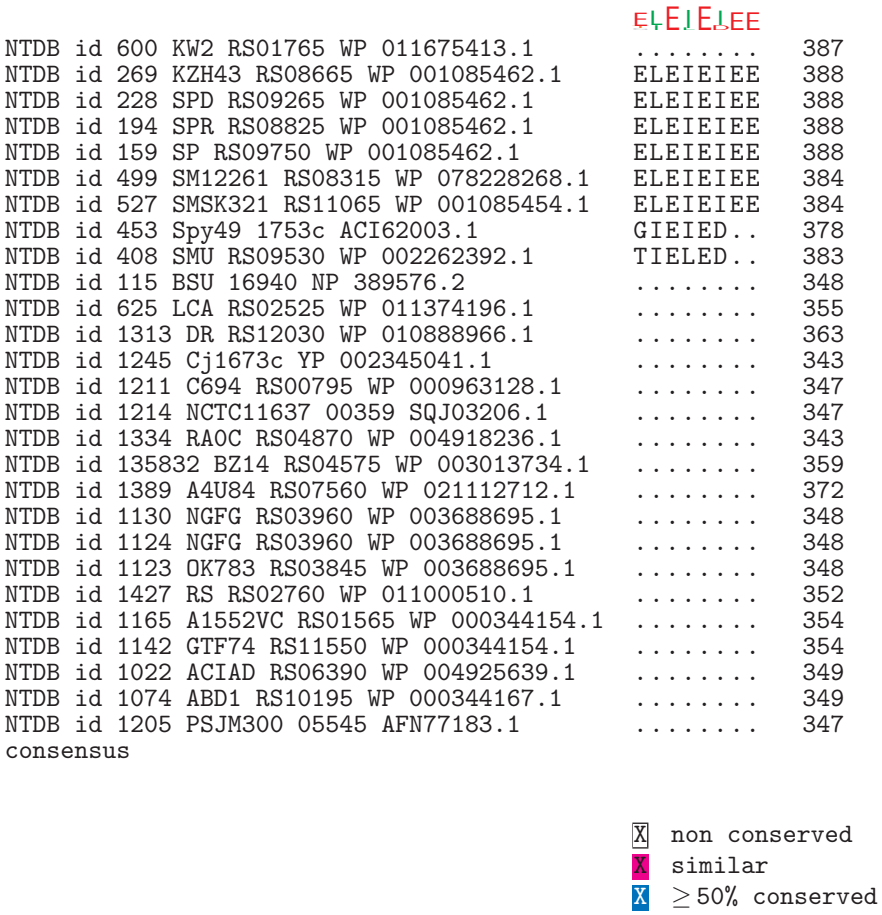

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NTDB id 499 SM12261 RS08315 WP 078228268.1 DEKIGQGSENAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAE.....S....KKDE....AAQVDS..VNEEVTLDLGD 376
NTDB id 527 SMSK321 RS11065 WP 001085454.1 DEKIGQGSENAKKYLADNPEVFAEIDHQVRVQFGLIDGEEAAE.....S....KKDE....VAQVDS..VNEEVTLDLGD 376
NTDB id 453 Spy49 1753c ACI62003.1 GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEES.....AMAVASE.......E.......TDDLALDLDN 372
NTDB id 408 SMU RS09530 WP 002262392.1 GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEE.....KQSSKEK.......ETDEK..ADKNLVLELDD 377
NTDB id 115 BSU 16940 NP 389576.2 EERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQ.....QAEETQEEL.EFE.E................ 348
NTDB id 625 LCA RS02525 WP 011374196.1 SERIGQGRENAKNYLADHEDVEDEVRLKVRAAYGISDVPEEDL.....PTTEDE.QI.NILPDDST.....EE....... 355
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NTDB id 1334 RA0C RS04870 WP 004918236.1 DTKLGQGRDTVKELIKDNPELAEELEAKIREKLQAK.. 343
NTDB id 135832 BZ14 RS04575 WP 003013734.1 GKKIGQGKEKSKEYLKENTAERDEIERAILELLLPNKYSN.....KDSNDSPK..EGSKIKTKVN.PAVTQDELI..... 359
NTDB id 1389 A4U84 RS07560 WP 021112712.1 GTKIGQGKNNAVKWLKENPEIADKLEQEIRNTLASNPNSV.....LKATASDSN.DESDVE......EFSEEEL..... 372
NTDB id 1130 NGFG RS03960 WP 003688695.1 GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEMHIT.....EGTQDETDG.ER.............PEE........ 348
NTDB id 1124 NGFG RS03960 WP 003688695.1 GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEMHIT.....EGTQDETDG.ER.............PEE........ 348
NTDB id 1123 OK783 RS03845 WP 003688695.1 GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEMHIT.....EGTQDETDG.ER.............PEE........ 348
NTDB id 1427 RS RS02760 WP 011000510.1 GERIGQGRDNCREFLRENAELAREIENKVREHLGVTPMGA.....VTLA.EEVE.ED..................... 352
NTDB id 1165 A1552VC RS01565 WP 000344154.1 GDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPENMQ.....LIAETSSAA.DDVEFG......AVPEEF....... 354
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NTDB id 1074 ABD1 RS10195 WP 000344167.1 GNKIGQGKNNVIRYFEENNQIAEEIERNIREQLLTTGTNG.....AVQ.I.EDE.EEPDLL.........LES......... 349
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