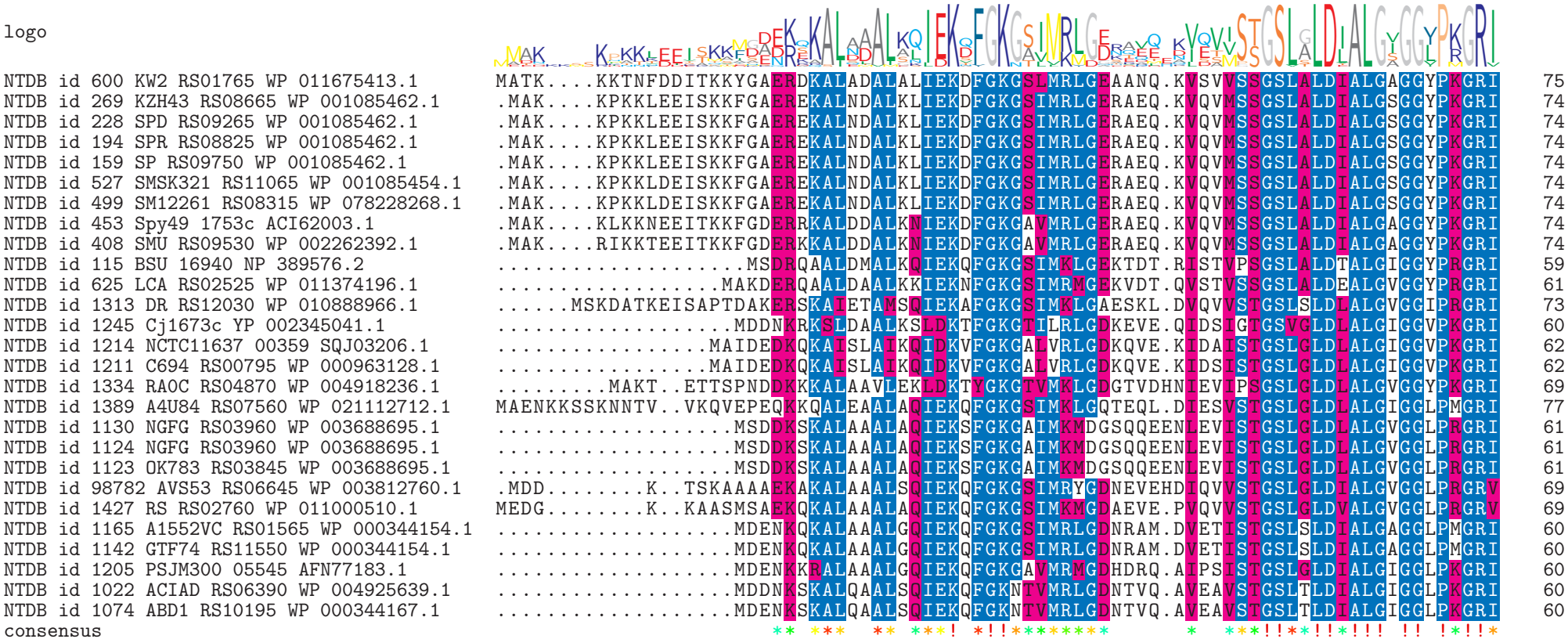
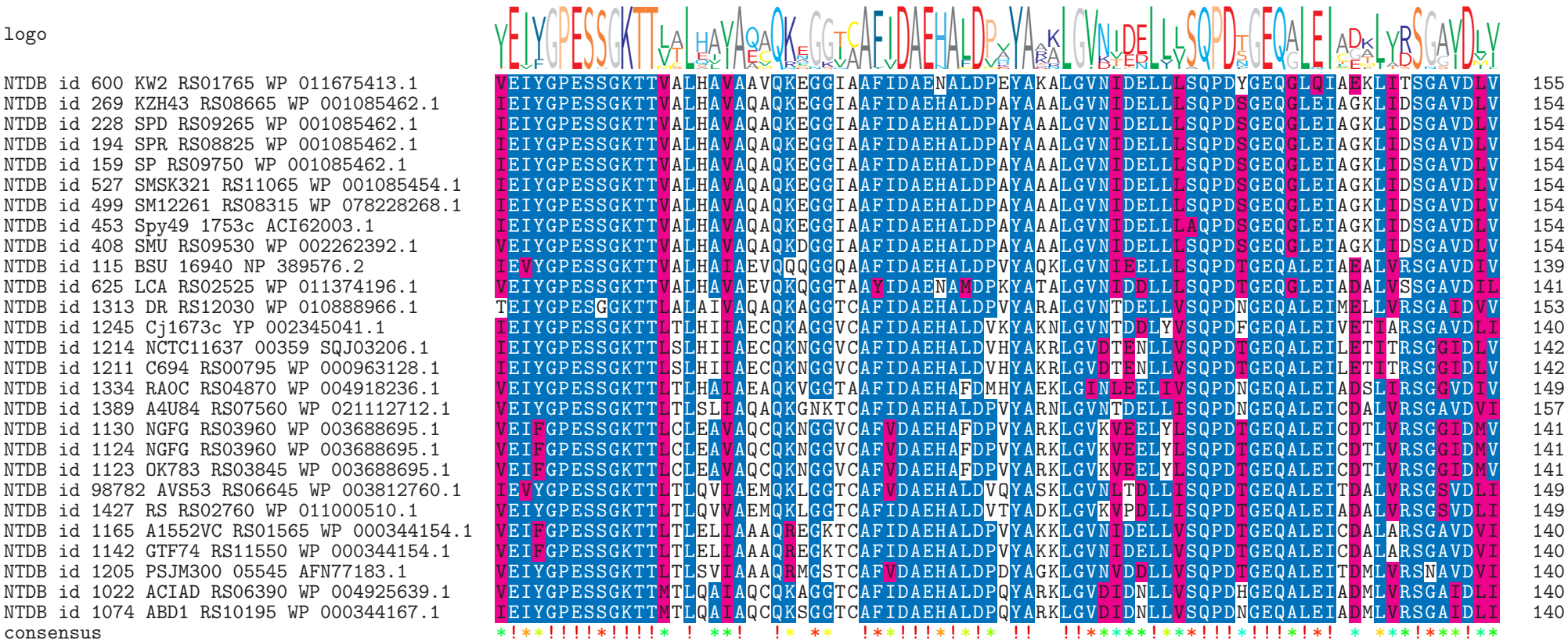
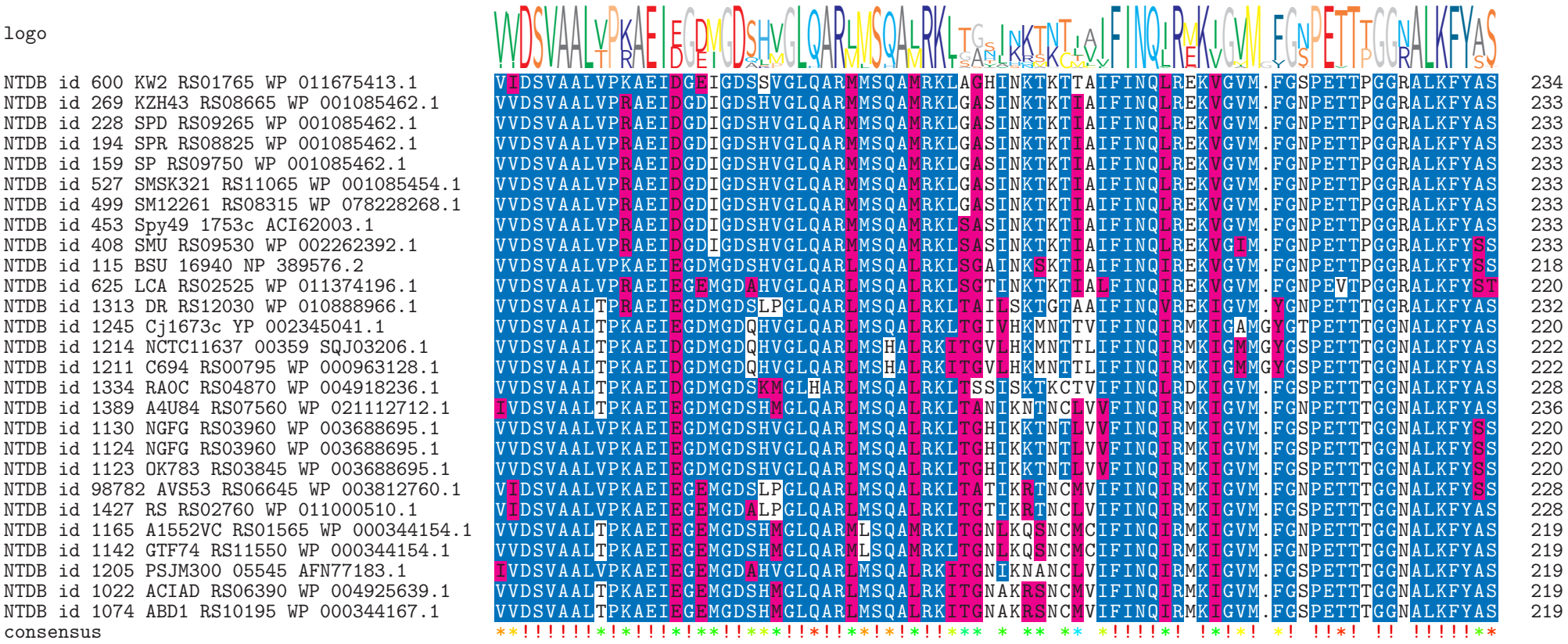
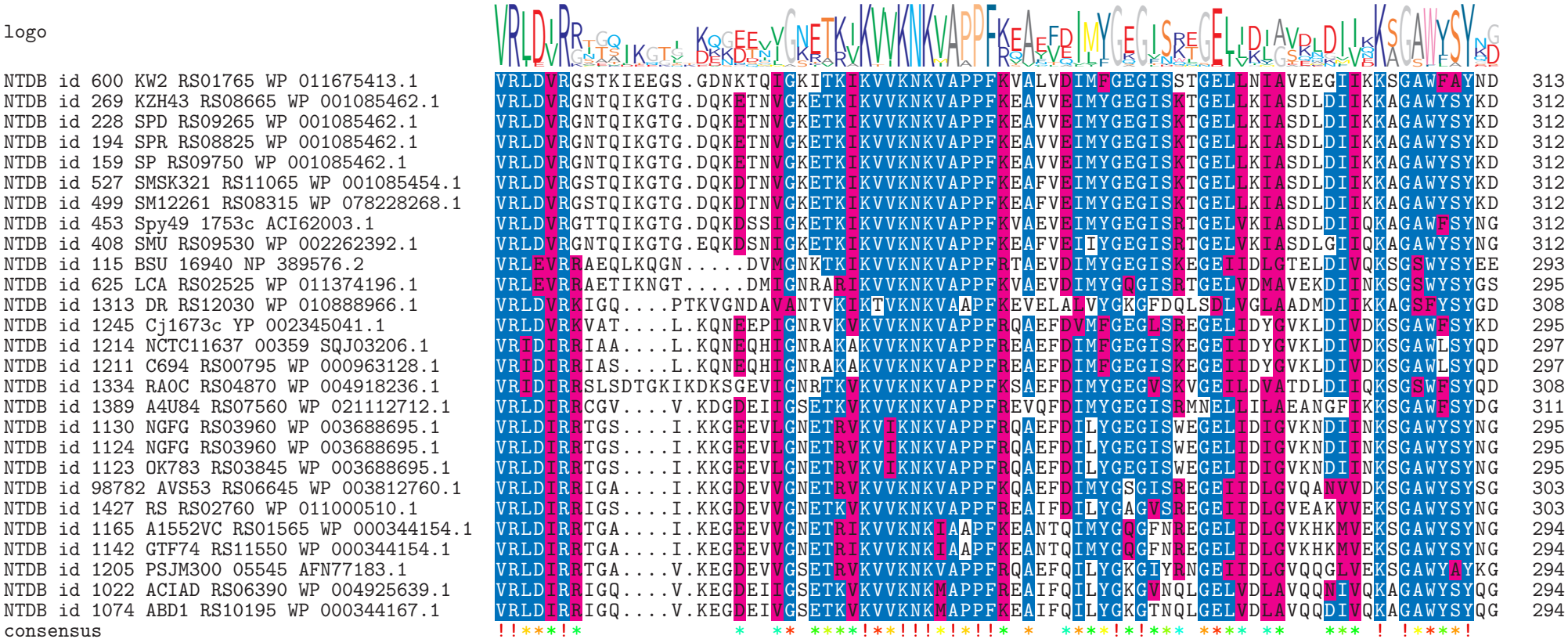
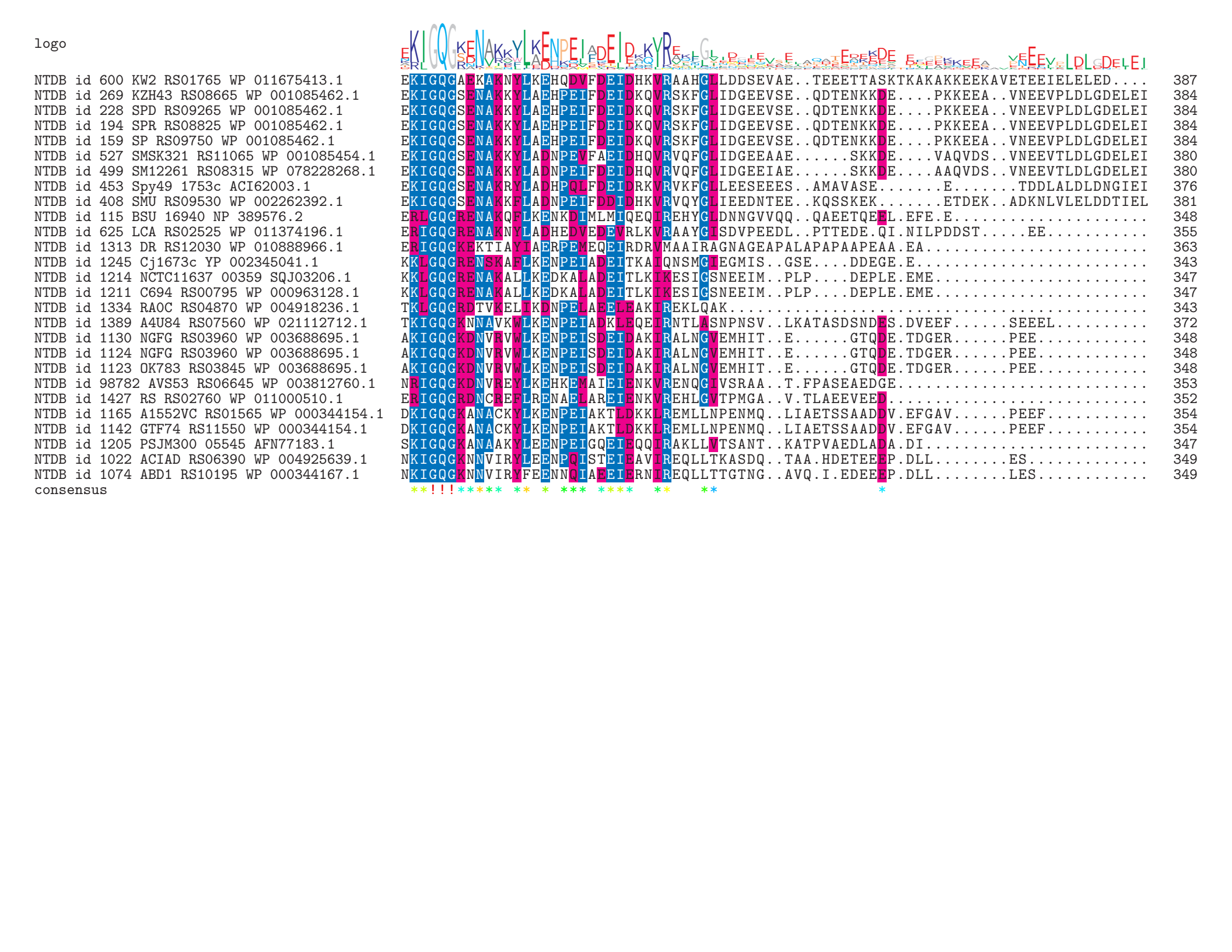

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consensus


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NTDB id 499 SM12261 RS08315 WP 078228268.1 EKIGQGSENAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAE......SKKDE....AAQVDS..VNEEVTLDLGDELEI 380
NTDB id 453 Spy49 1753c ACI62003.1 EKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEES..AMAVASE.......E.......TDDLALDLDNGIEI 376
NTDB id 408 SMU RS09530 WP 002262392.1 EKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEE..KQSSKEK.......ETDEK..ADKNLVLELDDTIEL 381
NTDB id 115 BSU 16940 NP 389576.2 ERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQ..QAEETQEEL.EFE.E..... 348
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NTDB id 1124 NGFG RS03960 WP 003688695.1 AKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEMHIT..E......GTQDE.TDGER......PEE..... 348
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consensus

X non conserved
X similar
X $\geq$ 50% conserved