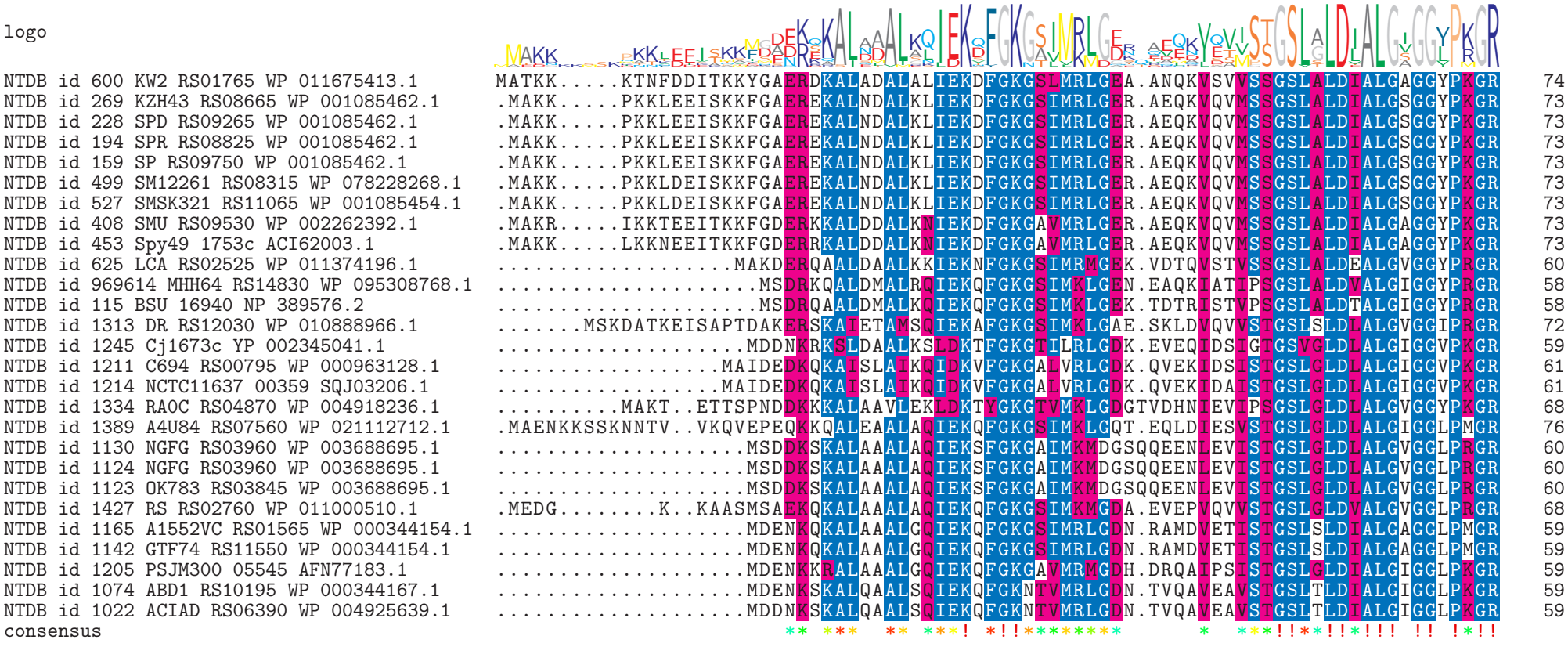
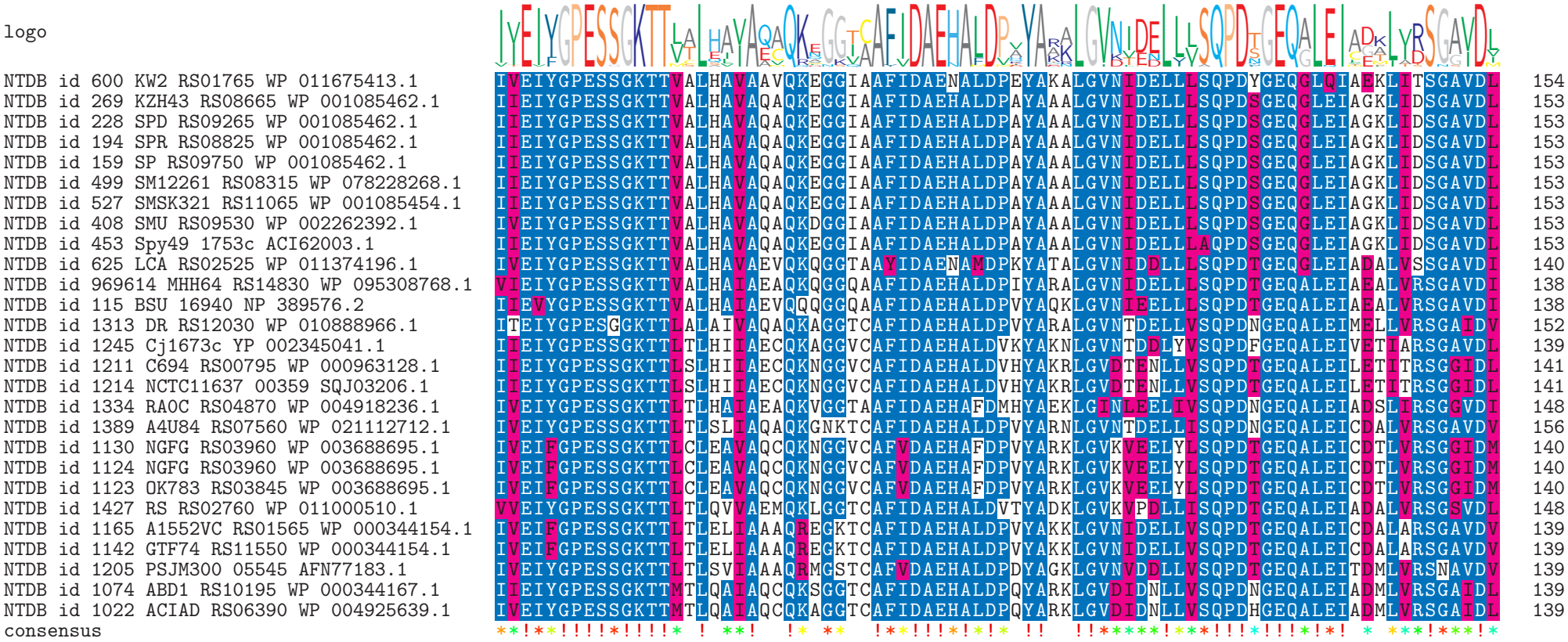
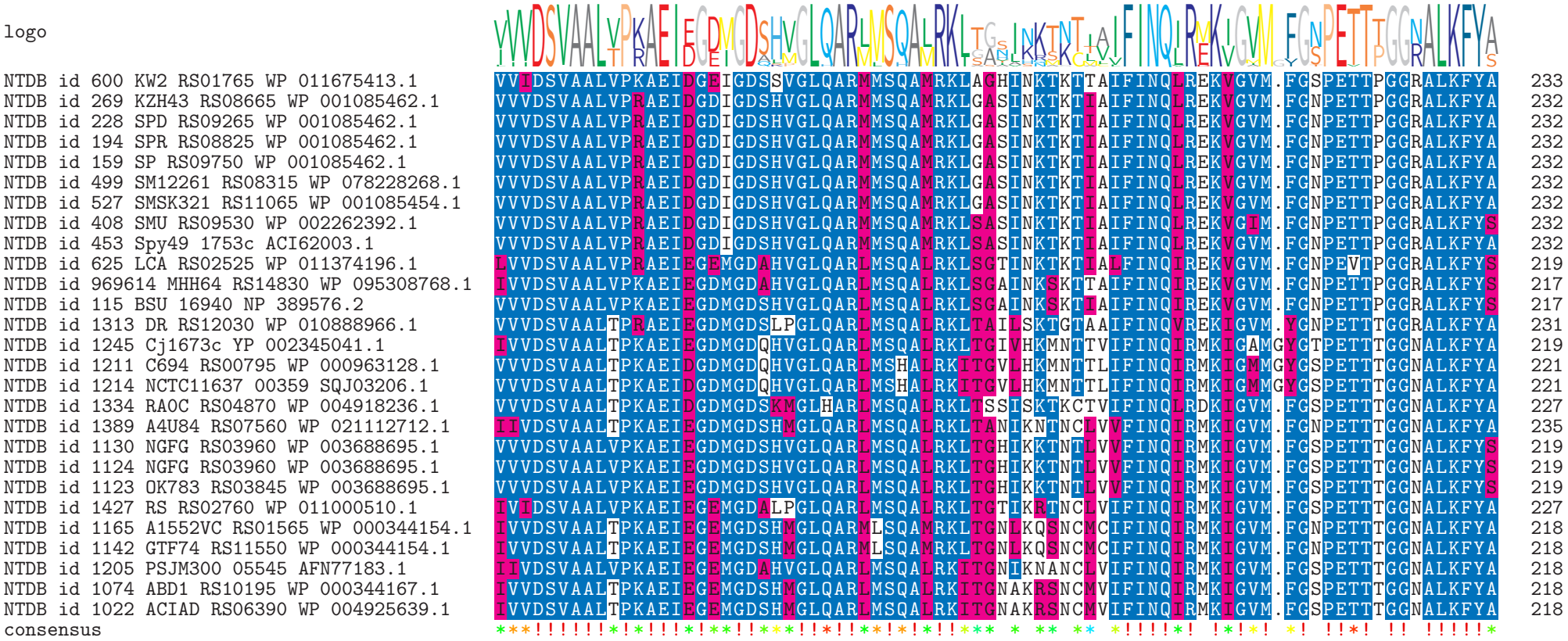
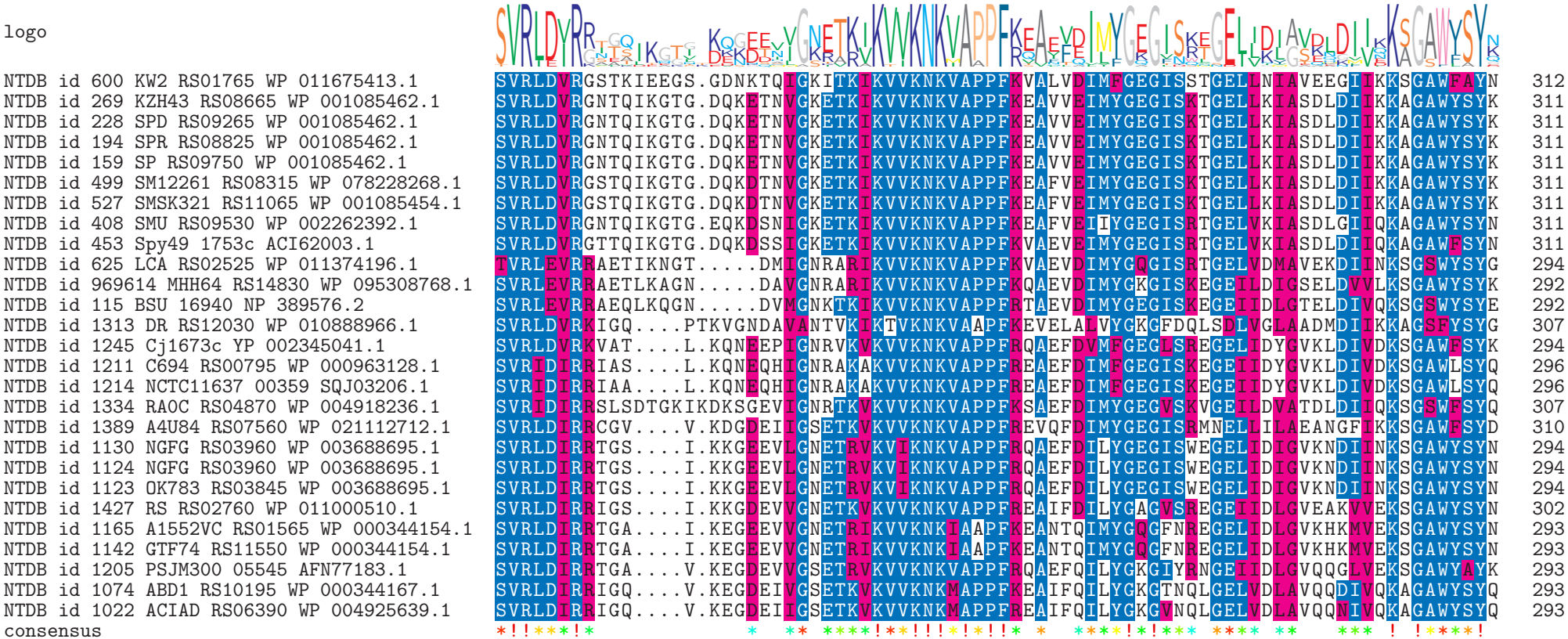
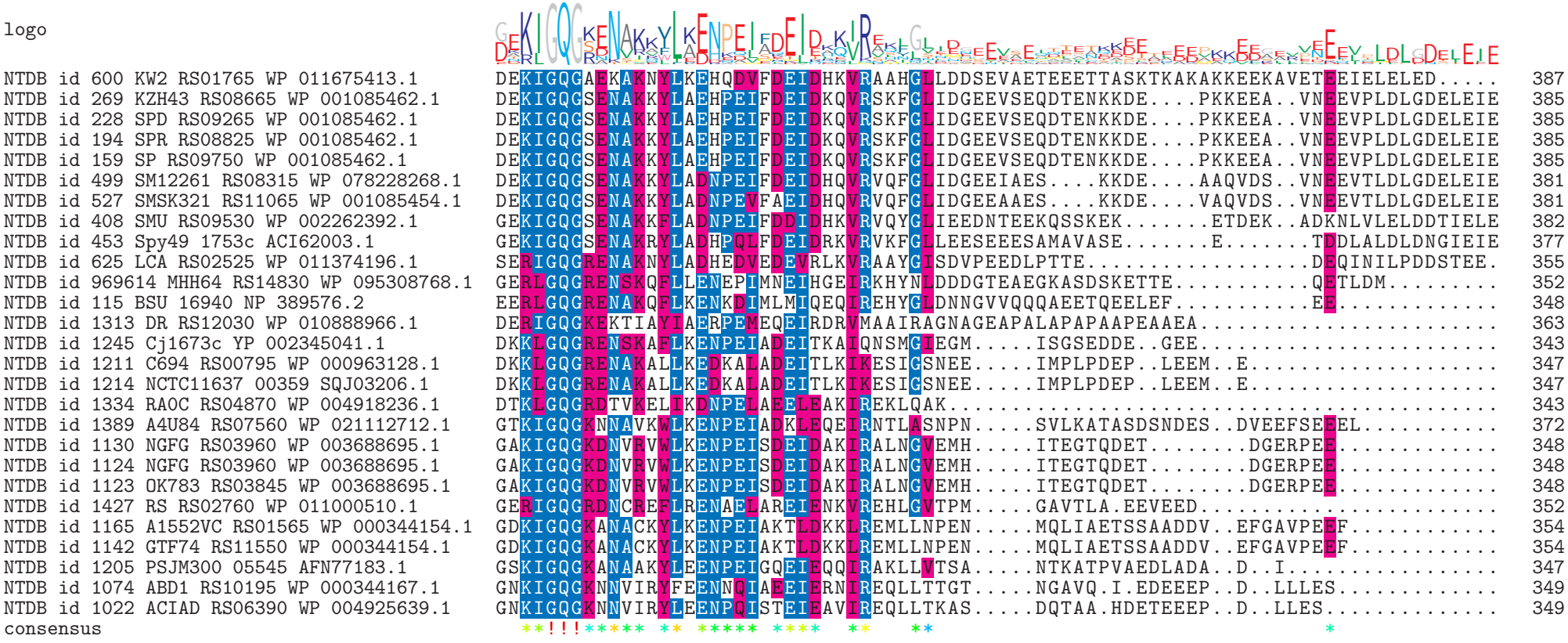
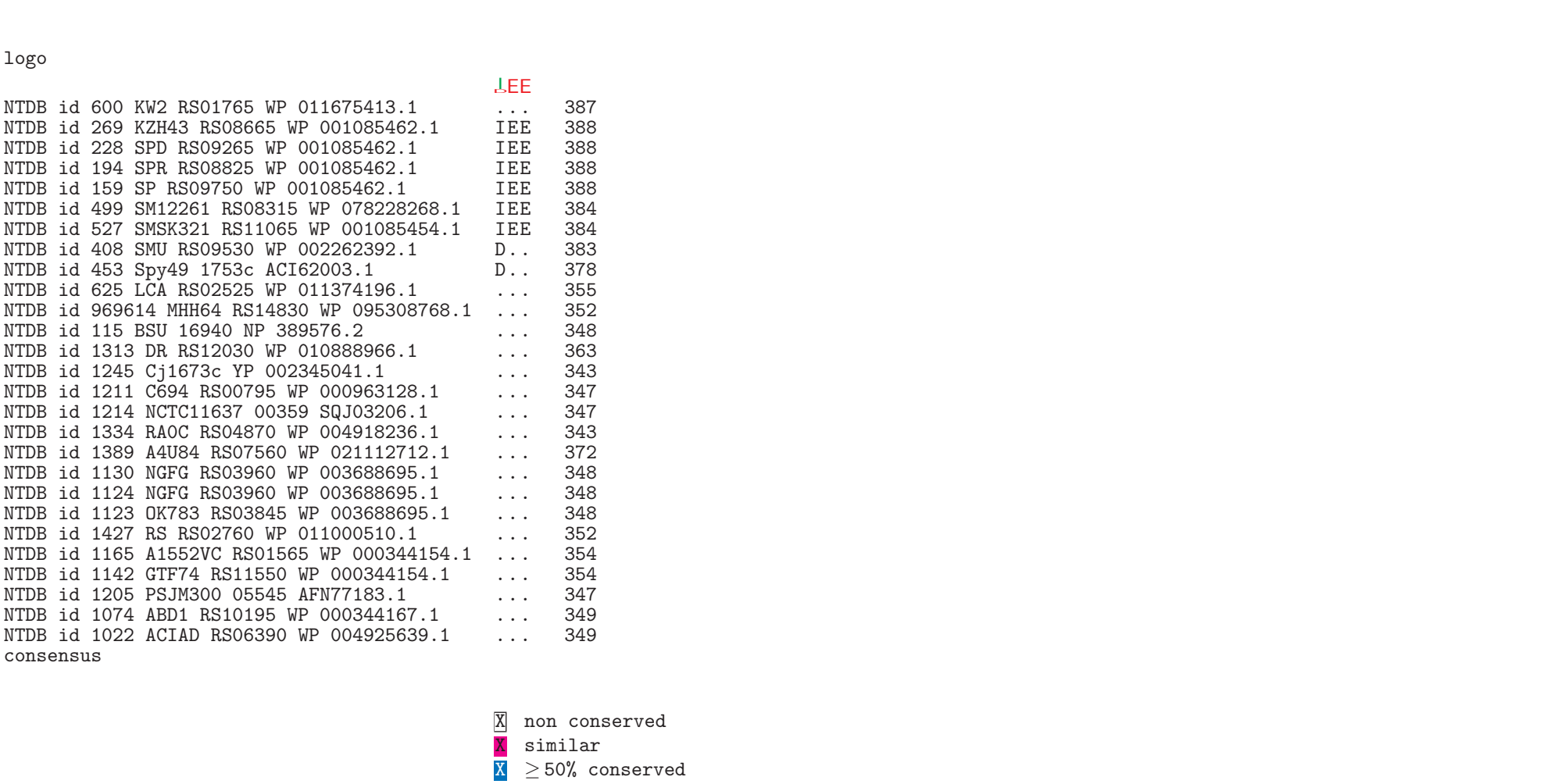

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NTDB id 1142 GTF74 RS11550 WP 000344154.1 SVRLDIRRTGA....I.KEGEEVVGNETRIKVVKNKIAAPFKEANTQIMYGQGFNREGELIDLGVKHKMVEKSGAWYSYN 293
NTDB id 1205 PSJM300 05545 AFN77183.1 SVRLDIRRTGA....V.KEGDEVVGSETRVKVVKNKVAPPFRQAEFQILYGKGIYRNGEIIDLGVQQGLVEKSGAWYAYK 293
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NTDB id 527 SMSK321 RS11065 WP 001085454.1 DEKIGQGSENAKKYLADNPEVFAEIDHQVRVQFGLIDGEEAAES....KKDE....VAQVDS..VNEEVTLDLGDELEIE 381
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NTDB id 969614 MHH64 RS14830 WP 095308768.1 GERLGQGRENSKQFLLENEPIMNEIHGEIRKHYNLDDDGTEAEGKASDSKETTE.....QETLDM..... 352
NTDB id 115 BSU 16940 NP 389576.2 EERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQAEETQEELEF.....EE..... 348
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NTDB id 1142 GTF74 RS11550 WP 000344154.1 GDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPEN.....MQLIAETSSAADDV..EFGAVPEEF..... 354
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consensus
X non conserved
X similar
X ≥ 50% conserved