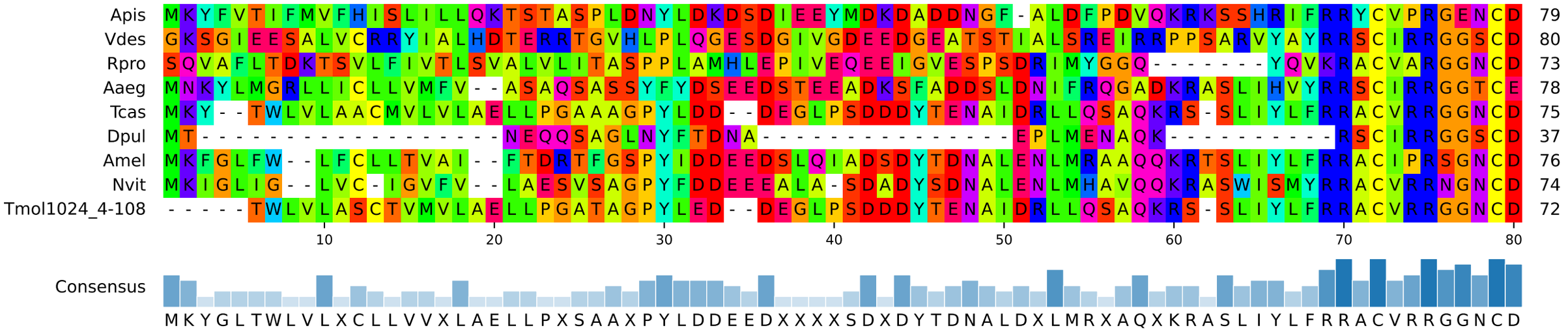
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Rpro SQVAFLTDKTSVLFIVTLSVALVLITASPPLAMHLEPIVEQEEIGVESPSDRIMYGGQ-------YQVKRACVARGGNCD 73
Aaeg MNKYLMGRLLICLLVMFV--ASAQSASSYFYDSEEDSTEEADKSFADDSLDNIFRQGADKRASLIHVYRRSCIRRGGTCE 78
Tcas MKY--TWLVLAACMVLVLAELLPGAAAGPYLDD--DEGLPSDDDYTENAIDRLLQSAQKRS-SLIYLFRRACVRRGGNCD 75
Dpul MT----------NEQQSAGLNYFTDNA----------EPLMENAQK----------RSCIRRGGSCD 37
Amel MKFGLFW--LFCLLTVAI--FTDRTFGSPYIDDEEDSLQIADSDYTDNALENLMRAAQQKRTSLIYLFRRACIPRSGNCD 76
Nvit MKIGLIG--LVC-IGVFV--LAESVSAGPYFDDEEEALA-SDADYSDNALENLMHAVQQKRASWISMYRRACVRRNGNCD 74
Tmol1024_4-108 -----TWLVLASCTVMVLAELLPGATAGPYLED--DEGLPSDDDYTENAIDRLLQSAQKRS-SLIYLFRRACVRRGGNCD 72
10
20
30
40
50
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Consensus
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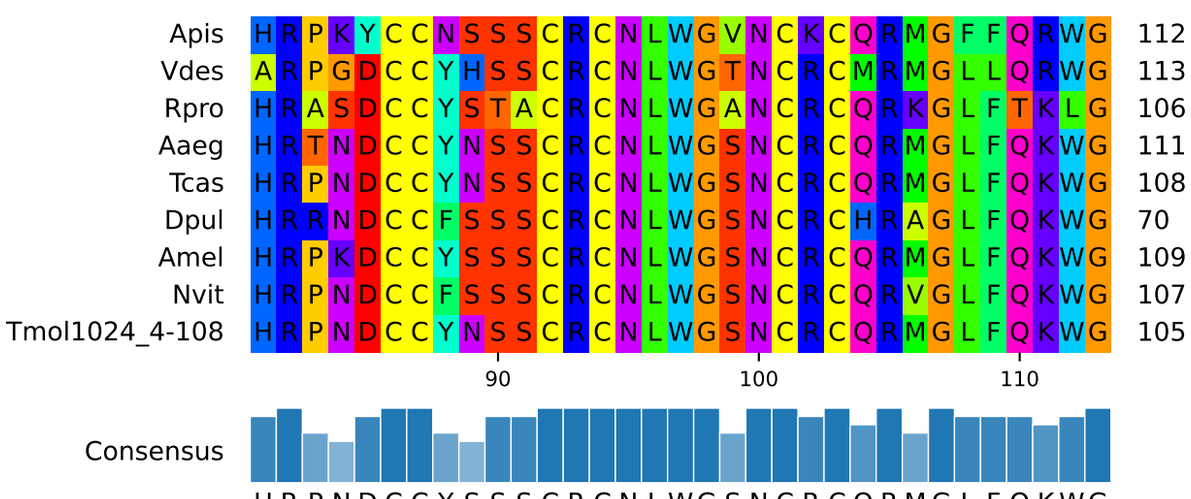
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Vdes ARPGDCCYHSSCRCNLWGTNCRCMRMGLLQRWG 113
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Aaeg HRTNDCCYNSSCRCNLWGSNCRCQRMGLFQKWG 111
Tcas HRPNDCCYNSSCRCNLWGSNCRCQRMGLFQKWG 108
Dpul HRRNDCCFSSSCRCNLWGSNCRCHRAGLFQKWG 70
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Nvit HRPNDCCFSSSCRCNLWGSNCRCQRVGLFQKWG 107
Tmol1024_4-108 HRPNDCCYNSSCRCNLWGSNCRCQRMGLFQKWG 105
90
100
110
Consensus
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