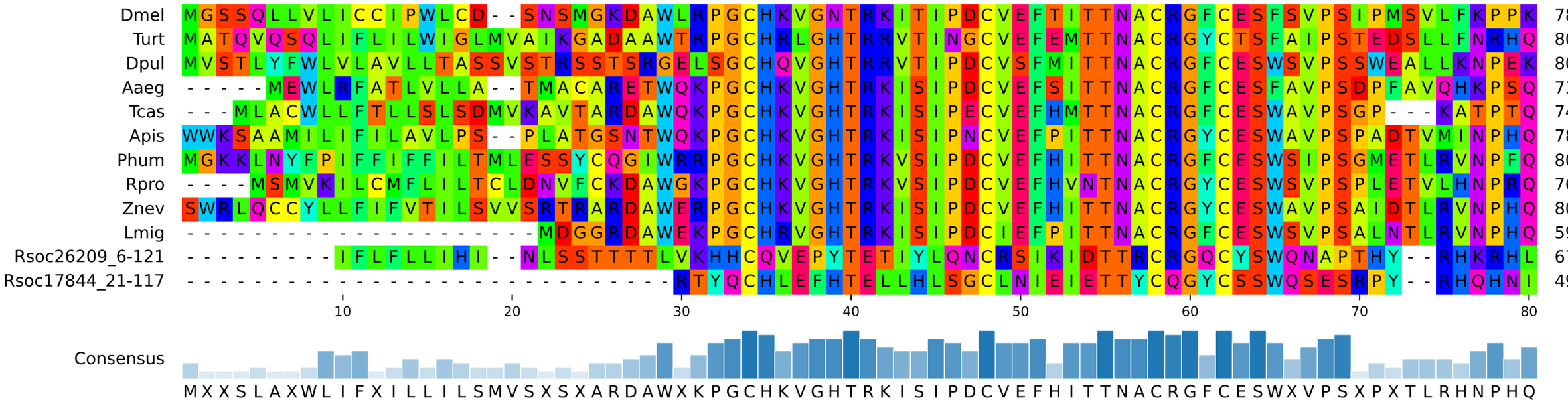
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Aaeg -----MEWLRFATLVLLA--TMACARETWQKPGCHKVGHTRKISIPDCVEFSITTNACRGFCESFAVPSDPFAVQHKPSQ 73
Tcas ---MLACWLLFTLLSLSDMVKAVTARDAWQKPGCHKVGHTRKISIPECVEFHMTTNACRGFCESWAVPSGP---KATPTQ 74
Apis WWKSAAMILIFILAVLPS--PLATGSNTWQKPGCHKVGHTRKISIPNCVEFPITTNACRGYCESWAVPSPADTVMINPHQ 78
Phum MGKKLNYFPIFFIFFILTMLESSYCQGIWRRPGCHKVGHTRKVSIPDCVEFHITTNACRGFCESWSIPSGMETLRVNPFQ 80
Rpro ----MSMVKILCMFLILTCLDNVFCKDAWGKPGCHKVGHTRKVSIPDCVEFHVNTNACRGYCESWSVPSPLETVLHNPRQ 76
Znev SWRLQCCYLLFIFVTILSVVSRTRARDAWERPGCHKVGHTRKISIPDCVEFHITTNACRGYCESWAVPSAIDTLRVNPHQ 80
Lmig ----------MDGGRDAWEKPGCHRVGHTRKISIPDCIEFPITTNACRGFCESWSVPSALNTLRVNPHQ 59
Rsoc26209_6-121 ---------IFLFLLIHI--NLSSTTTTLVKHHCQVEPYTETIYLQNCRSIKIDTTRCRGQCYSWQNAPTHY--RHKRHL 67
Rsoc17844_21-117 ----------RTYQCHLEFHTELLHLSGCLNIEIETTYCQGYCSSWQSESRPY--RHQHNI 49
10 20 30 40 50 60 70 80
Consensus
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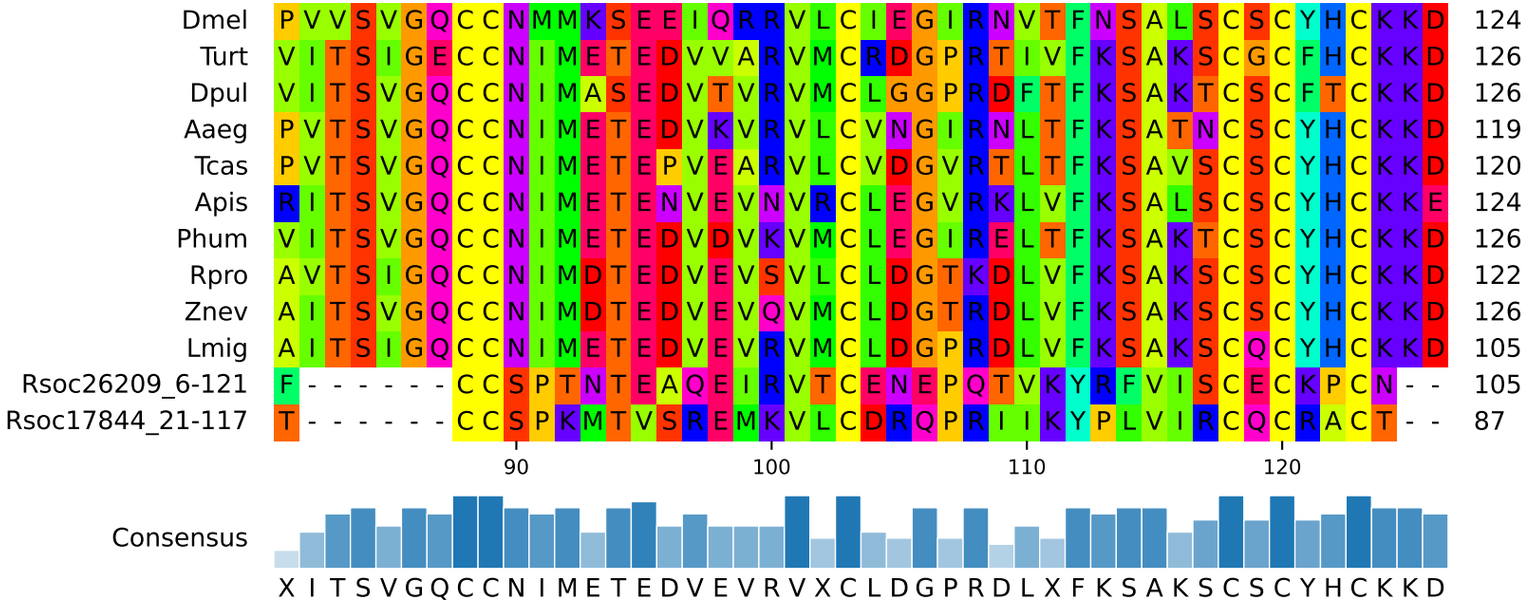
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Tcas PVTSVGQCCNIMETEPVEARVLCVDGVRTLTFKSAVSCSCYHCKKD 120
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Rsoc26209_6-121 F------CCSPTNTEAQEIRVTCENEPQTVKYRFVISCECKPCN-- 105
Rsoc17844_21-117 T------CCSPKMTVSREMKVLCDRQPRIIKYPLVIRCQCRACT-- 87
90 100 110 120
Consensus
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