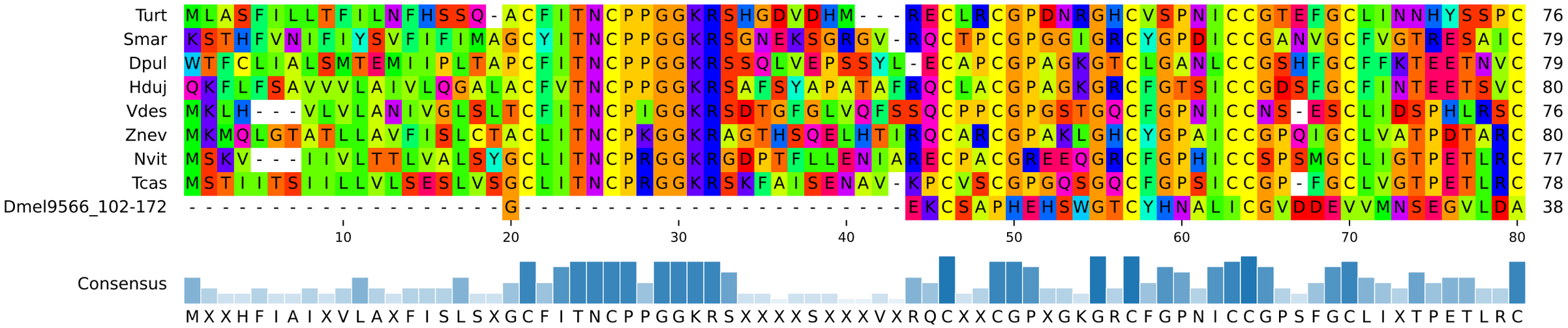
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Dpul WTFCLIALSMTEMIIPLTAPCFITNCPPGGKRSSQLVEPSSYL-ECAPCGPAGKGTCLGANLCCGSHFGCFFKTEETNVC 79
Hduj QKFLFSAVVVLAIVLQGALACFVTNCPPGGKRSAFSYAPATAFRQCLACGPAGKGRCFGTSICCGDSFGCFINTEETSVC 80
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Dmel9566_102-172 ----------G----------EKCSAPHEHSWGTCYHNALICGVDDEVVMNSEGVLDA 38
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Consensus
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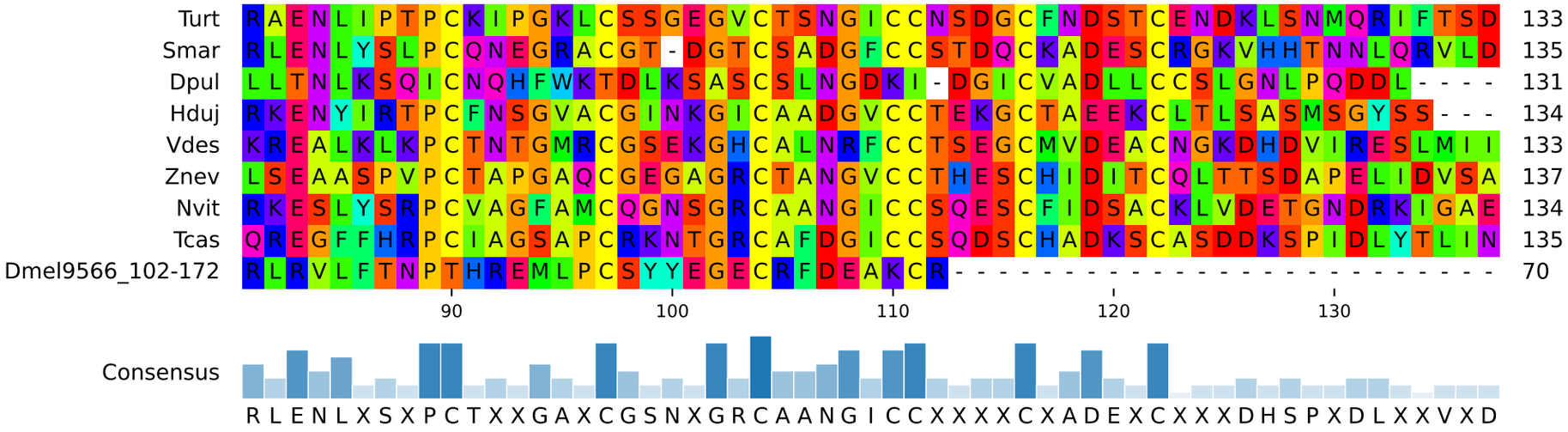
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Hduj RKENYIRTPCFNSGVACGINKGICAADGVCCTEKGCTAEEKCLTLSASMSGYSS--- 134
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90 100 110 120 130
Consensus
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