

CLUSTAL 2.1 multiple sequence alignment

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RdpA      MHAALSPLSQRFERIAVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
FT_5      MHAALSPLSQRFERIAVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
FT_C      MHAALTPLTNKYRFIDVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
FT_R      MHAALTPLTNKYRFIDVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
FT_T      MHAALTPLTNKYRFIDVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
FT_Tv3    MHAALTPLTNKYRFIDVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
FT_Tv7    MHAALTPLTNKYRFIDVQPLTGVLGAEITGVDLREPLDDSTWNEILDAFHTYQVIYFPGQ
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RdpA      AITNEQHIAFSRRFGPVDVPVLLKSIEGYPEVQMIRREANESGRVIGDDWHTDSTFLDAP
FT_5      AITNEQHIAFSRRFGPVDVPVLLKSIEGYPEVQMIRREANESGRVIGENWHTDSTFLDAP
FT_C      AITNEQHIAFSRRFGPVDVPVLLKSIEGYPEVQMIRREANESGRVIGDDWHTDSTFLDAP
FT_R      AITNEQHIAFSRRFGPVDVPVLLKSIEGYPEVQMIRREANESSRVIGDDWHSdstFLDAP
FT_T      AITNEQHIAFSRRFGPVDVPVILKSIEGYPEVQMIRREANESSRFIGDDWHTDSTFLDAP
FT_Tv3    AITNEQHIAFSRRFGPVDVPVILKSIEGYPEVQMIRREANESSRYIGDDWHADSTFLDAP
FT_Tv7    AITNEQHIAFSRRFGPVDVPVILKSIEGYPEVQMIRREANESSRYIGDDWHADSTFLDAP
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RdpA      PAAVVMRAIDVPEHGGDTGFLSMYTAWETLSPTMQATIEGLNVVHSATRVFGSLYQAQNR
FT_5      PAAVVMYAKEIPPYGGDTLFTSMYTAWETLSPTMQATIEGLNVVHSATRVFGSLYQAQNR
FT_C      PAAVVMRAIEVPEYGGDTGFLSMYSAWETLSPTMQATIEGLNVVHSATKVFGSLYQATNW
FT_R      PAAVVMRAIEVPEYGGDTGFLSMYSAWETLSPTMQATIEGLNVVHSATKVFGSLYQATNW
FT_T      PAAVVMRAIEVPEYGGDTGFLSMYSAWETLSPTMQATIEGLNVVHSATKVFGSLYQATNW
FT_Tv3    PAAVVMRAIEVPEYGGDTGFLSMYSAWETLSPTMQATIEGLNVVHSATKVFGSLYQATNW
FT_Tv7    PAAVVMRAIEVPEYGGDTGFLSMYSAWETLSPTMQATIEGLNVVHSATKVFGSLYQATNW
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RdpA      RFSNTSVKVMVDVDAGDRET VHPLVVTHPGSGRKGlyVNQVYCQRIEGMTDAESKPLLQFL
FT_5      RFSNTSVKVMVDVDAGDRET VHPLVVTHPETGRKGlyVNQVYCQRIEGMSEKESEPLLSFL
FT_C      RFSNTSVKVMVDVDAGDRET VHPLVVTHPVTGRRALYCNQVYCQKIQGMTDAESKSLQFL
FT_R      RFSNTSVKVMVDVDAGDRET VHPLVVTHPVTGRRALYCNQVYCQKIQGMTDAESKSLQFL
FT_T      RFSNTSVKVMVDVDAGDRET VHPLVVTHPVTGRRALYCNQVYCQKIQGMTDAESKSLQFL
FT_Tv3    RFSNTSVKVMVDVDAGDRET VHPLVVTHPVTGRRALYCNQVYCQKIQGMTDAESKSLQFL
FT_Tv7    RFSNTSVKVMVDVDAGDRET VHPLVVTHPVTGRRALYCNQVYCQKIQGMTDAESKSLQFL
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RdpA      YEHAtrFDfTCrVRWKKDQVLVWDNLCTMhRAVPDYAGKFRYLTrTTVGGVrPAR
FT_5      FAHATKPEfTCrVRWQEGDVLVWDNLCTQHYAVPDYAGKFRYLTrTTVGGVrPAR
FT_C      YEHAtrFDfTCrVRWKKDQVLVWDNLCTMhRAVPDYAGKFRYLTrTTVAGDKPSR
FT_R      YEHAtrFDfTCrVRWKKDQVLVWDNLCTMhRAAPDYAGKFRYLTrTTVAGDKPSR
FT_T      YEHAtrFDfTCrVRWKKDQVLVWDNLCTMhRAVPDYAGKFRYLTrTTVAGDKPSR
FT_Tv3    YEHAtrFDfTCrVRWKKDQVLVWDNLCTMhRAVPDYAGKFRYLTrTTVAGDKPSR
FT_Tv7    YEHAtrFDfTCrVRWKKDQVLVWDNLCTMhRAVPDYAGKFRYLTrTTVAGDKPSR
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