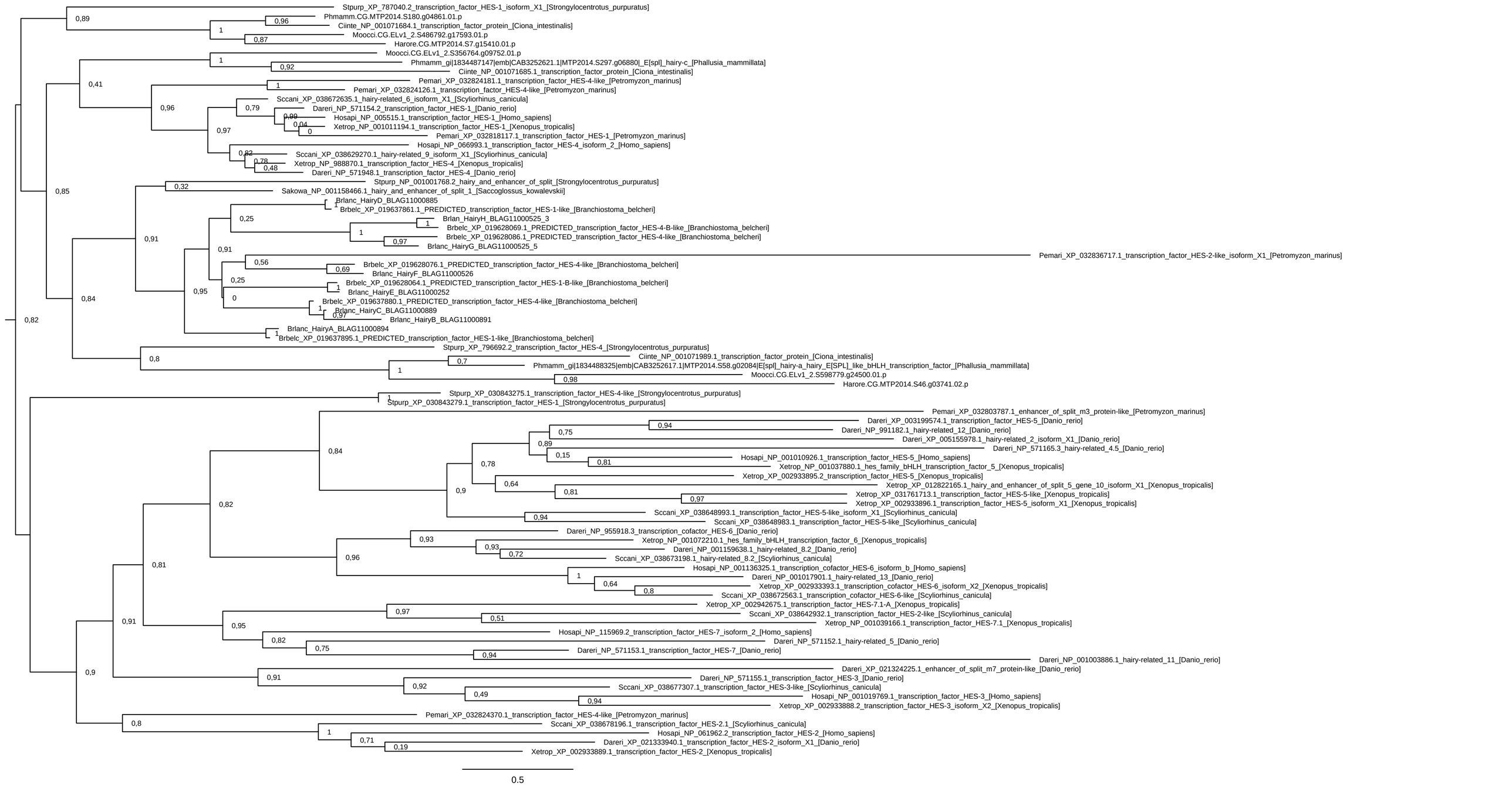
Stpurp_XP_787040.2_transcription_factor_HES-1_isoform_X1_[Strongylocentrotus_purpuratus]
Phmamm.CG.MTP2014.S180.g04861.01.p
Ciinte_NP_001071684.1_transcription_factor_protein_[Ciona_intestinalis]
Moocci.CG.ELv1_2.S486792.g17593.01.p
Harore.CG.MTP2014.S7.g15410.01.p
Moocci.CG.ELv1_2.S356764.g09752.01.p
Phmamm_gi|1834487147|emb|CAB3252621.1|MTP2014.S297.g06880|_E[spl]_hairy-c_[Phallusia_mammillata]
Ciinte_NP_001071685.1_transcription_factor_protein_[Ciona_intestinalis]
Pemari_XP_032824181.1_transcription_factor_HES-4-like_[Petromyzon_marinus]
Pemari_XP_032824126.1_transcription_factor_HES-4-like_[Petromyzon_marinus]
Sccani_XP_038672635.1_hairy-related_6_isoform_X1_[Scyliorhinus_canicula]
Dareri_NP_571154.2_transcription_factor_HES-1_[Danio_rerio]
Hosapi_NP_005515.1_transcription_factor_HES-1_[Homo_sapiens]
Xetrop_NP_001011194.1_transcription_factor_HES-1_[Xenopus_tropicalis]
Pemari_XP_032818117.1_transcription_factor_HES-1_[Petromyzon_marinus]
Hosapi_NP_066993.1_transcription_factor_HES-4_isoform_2_[Homo_sapiens]
Sccani_XP_038629270.1_hairy-related_9_isoform_X1_[Scyliorhinus_canicula]
Xetrop_NP_988870.1_transcription_factor_HES-4_[Xenopus_tropicalis]
Dareri_NP_571948.1_transcription_factor_HES-4_[Danio_rerio]
Stpurp_NP_001001768.2_hairy_and_enhancer_of_split_[Strongylocentrotus_purpuratus]
Sakowa_NP_001158466.1_hairy_and_enhancer_of_split_1_[Saccoglossus_kowalevskii]
Brlanc_HairyD_BLAG11000885
Brbelc_XP_019637861.1_PREDICTED_transcription_factor_HES-1-like_[Branchiostoma_belcheri]
Brlan_HairyH_BLAG11000525_3
Brbelc_XP_019628069.1_PREDICTED_transcription_factor_HES-4-B-like_[Branchiostoma_belcheri]
Brbelc_XP_019628086.1_PREDICTED_transcription_factor_HES-4-like_[Branchiostoma_belcheri]
Brlanc_HairyG_BLAG11000525_5
Pemari_XP_032836717.1_transcription_factor_HES-2-like_isoform_X1_[Petromyzon_marinus]
Brbelc_XP_019628076.1_PREDICTED_transcription_factor_HES-4-like_[Branchiostoma_belcheri]
Brlanc_HairyF_BLAG11000526
Brbelc_XP_019628064.1_PREDICTED_transcription_factor_HES-1-B-like_[Branchiostoma_belcheri]
Brlanc_HairyE_BLAG11000252
Brbelc_XP_019637880.1_PREDICTED_transcription_factor_HES-4-like_[Branchiostoma_belcheri]
Brlanc_HairyC_BLAG11000889
Brlanc_HairyB_BLAG11000891
Brlanc_HairyA_BLAG11000894
Brbelc_XP_019637895.1_PREDICTED_transcription_factor_HES-1-like_[Branchiostoma_belcheri]
Stpurp_XP_796692.2_transcription_factor_HES-4_[Strongylocentrotus_purpuratus]
Ciinte_NP_001071989.1_transcription_factor_protein_[Ciona_intestinalis]
Phmamm_gi|1834488325|emb|CAB3252617.1|MTP2014.S58.g02084|E[spl]_hairy-a_hairy_E[SPL]_like_bHLH_transcription_factor_[Phallusia_mammillata]
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Stpurp_XP_030843275.1_transcription_factor_HES-4-like_[Strongylocentrotus_purpuratus]
Stpurp_XP_030843279.1_transcription_factor_HES-1_[Strongylocentrotus_purpuratus]
Pemari_XP_032803787.1_enhancer_of_split_m3_protein-like_[Petromyzon_marinus]
Dareri_XP_003199574.1_transcription_factor_HES-5_[Danio_rerio]
Dareri_NP_991182.1_hairy-related_12_[Danio_rerio]
Dareri_XP_005155978.1_hairy-related_2_isoform_X1_[Danio_rerio]
Dareri_NP_571165.3_hairy-related_4.5_[Danio_rerio]
Hosapi_NP_001010926.1_transcription_factor_HES-5_[Homo_sapiens]
Xetrop_NP_001037880.1_hes_family_bHLH_transcription_factor_5_[Xenopus_tropicalis]
Xetrop_XP_002933895.2_transcription_factor_HES-5_[Xenopus_tropicalis]
Xetrop_XP_012822165.1_hairy_and_enhancer_of_split_5_gene_10_isoform_X1_[Xenopus_tropicalis]
Xetrop_XP_031761713.1_transcription_factor_HES-5-like_[Xenopus_tropicalis]
Xetrop_XP_002933896.1_transcription_factor_HES-5_isoform_X1_[Xenopus_tropicalis]
Sccani_XP_038648993.1_transcription_factor_HES-5-like_isoform_X1_[Scyliorhinus_canicula]
Sccani_XP_038648983.1_transcription_factor_HES-5-like_[Scyliorhinus_canicula]
Dareri_NP_955918.3_transcription_cofactor_HES-6_[Danio_rerio]
Xetrop_NP_001072210.1_hes_family_bHLH_transcription_factor_6_[Xenopus_tropicalis]
Dareri_NP_001159638.1_hairy-related_8.2_[Danio_rerio]
Sccani_XP_038673198.1_hairy-related_8.2_[Scyliorhinus_canicula]
Hosapi_NP_001136325.1_transcription_cofactor_HES-6_isoform_b_[Homo_sapiens]
Dareri_NP_001017901.1_hairy-related_13_[Danio_rerio]
Xetrop_XP_002933393.1_transcription_cofactor_HES-6_isoform_X2_[Xenopus_tropicalis]
Sccani_XP_038672563.1_transcription_cofactor_HES-6-like_[Scyliorhinus_canicula]
Xetrop_XP_002942675.1_transcription_factor_HES-7.1-A_[Xenopus_tropicalis]
Sccani_XP_038642932.1_transcription_factor_HES-2-like_[Scyliorhinus_canicula]
Xetrop_NP_001039166.1_transcription_factor_HES-7.1_[Xenopus_tropicalis]
Hosapi_NP_115969.2_transcription_factor_HES-7_isoform_2_[Homo_sapiens]
Dareri_NP_571152.1_hairy-related_5_[Danio_rerio]
Dareri_NP_571153.1_transcription_factor_HES-7_[Danio_rerio]
Dareri_NP_001003886.1_hairy-related_11_[Danio_rerio]
Dareri_XP_021324225.1_enhancer_of_split_m7_protein-like_[Danio_rerio]
Dareri_NP_571155.1_transcription_factor_HES-3_[Danio_rerio]
Sccani_XP_038677307.1_transcription_factor_HES-3-like_[Scyliorhinus_canicula]
Hosapi_NP_001019769.1_transcription_factor_HES-3_[Homo_sapiens]
Xetrop_XP_002933888.2_transcription_factor_HES-3_isoform_X2_[Xenopus_tropicalis]
Pemari_XP_032824370.1_transcription_factor_HES-4-like_[Petromyzon_marinus]
Sccani_XP_038678196.1_transcription_factor_HES-2.1_[Scyliorhinus_canicula]
Hosapi_NP_061962.2_transcription_factor_HES-2_[Homo_sapiens]
Dareri_XP_021333940.1_transcription_factor_HES-2_isoform_X1_[Danio_rerio]
Xetrop_XP_002933889.1_transcription_factor_HES-2_[Xenopus_tropicalis]
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