Orthogroup usage across cts and gene modules HoiH23|lipophil_2 Hhon|lipophil_ Tadh|İipophil TrH2|lipophil Hhon|lipophil_1 HoiH23|lipophil_1 Hhon|unknown_1 TrH2|unknown_1 HoiH23|unknown_1 Tadh|fibre TrH2|fibre Hhon|fibre HoiH23|fibre_1 HoiH23|fibre_2 Tadh|unknown_2.2 Tadhjunkhown_2.2
TrH2|unknown_2.2
Tadhjunknown_2.1
TrH2|unknown_2.1
TrH2|unknown_2.4
TrH2|unknown_2.3
TrH2|epithelia_gland_like
Trdb|epithelia_gland_like Tadh|epithelia_gland_like Tadh|gland_1 Hhon|epithelia_gland_like Hhon|gland_1 Hhon|gland_1_like HoiH23|epithelia_gland_like HoiH23|gland_1 Hhon|epithelia_unknown Tadh|epithelia_unknown_3 Tadh epithelia_unknown_ Tadh epithelia_ventral TrH2|epithelia_ventral Hhon|epithelia_ventral HoiH23|epithelia_ventral Tadh|epithelia_dorsal TrH2|epithelia_dorsal Hhon|epithelia_dorsal TrH2|epithelia_dorsal_like HoiH23|epithelia_dorsal HoiH23|epithelia_unkanwn Todklepithelia_unkanwn Tadh|epithelia_unknown_1 TrH2|epithelia_unknown_1
Tadh|peptidergic_delta TrH2|peptidergic_delta Hhon|peptidergic_delta Tadh|peptidergic_beta frac TrH2|peptidergic_beta Hhon|peptidergic_beta HoiH23|peptidergic_beta HoiH23|peptidergic_gamma 8.0 0.6 0.4 Hhon|peptidergic_gamma
Tadh|peptidergic_gamma
TrH2|peptidergic_gamma
TrH2|peptidergic_lambda
TrH2|peptidergic_lambda
HoiH23|peptidergic_lambda
Hon|peptidergic_lambda 0.2 0 Hhon|peptidergic_lambda Tadh|peptidergic_epsilon TrH2|peptidergic_epsilon HoiH23|peptidergic_epsilon Hhon|peptidergic_other_2 Tadh|peptidergic_omicron TrH2|peptidergic_omicron Hhon|peptidergic_omicron Hhon|peptidergic_other_1 Tadh|peptidergic_zeta TrH2|peptidergic_zeta Hhon|peptidergic_zeta HoiH23|peptidergic_zeta Tadh|peptidergic_theta TrH2|peptidergic_theta HoiH23|peptidergic_theta Hhon|peptidergic_theta Tadh|peptidergic_iota TrH2[peptidergic_iota Hhon|peptidergic_iota HoiH23|peptidergic_iota Tadh|peptidergic_eta TrH2|peptidergic_eta Hhon peptidergic_eta HoiH23 peptidergic_eta Tadh|peptidergic_nu TrH2|peptidergic_nu HoiH23|peptidergic_nu Tadh|peptidergic_mu
TrH2|peptidergic_mu
HoiH23|peptidergic_mu Hhon|peptidergic_mu
Tadh|peptidergic_kappa
TrH2|peptidergic_kappa
Hhon|peptidergic_kappa Hhon|peptidergic_kappa Tadh|peptidergic_alpha TrH2|peptidergic_alpha Hhon|peptidergic_alpha HoiH23|peptidergic_alpha

Hhon|peptidergic_progenitors Tadh|peptidergic_progenitors HoiH23|peptidergic_progenitors TrH2|peptidergic_progenitors

102 row

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peptidergic_gamma_like gc003a peptidergic_gamma gc017 peptidergic_lambda gc006 peptidergic_lepsilon gc020 peptidergic_omicron gc014 peptidergic_zeta gc018 peptidergic_alpha gc008 peptidergic_progenitors gc027 6000b gc030 peptidergic_theta gc026 | peptidergic_theta gc015 | peptidergic_tota gc012 ပ gc028 epithelia_unknown gc005 peptidergic_delta gc019 peptidergic_beta gc010 panpeptidergic gc004 peptidergic_eta gc016 peptidergic_nu gc011a peptidergic_mu_iota_like gc003b peptidergic_mu gc013 peptidergic_kappa gc023 peptidergic_alpha gc008 ပ lipophil_2 gc002b lipophil gc002a unknown_1 gc029 cell_cycle_1 gc007b meiosis gc007a epithelia_like gc032 epithelia_ventral gc001a fibre gc024 gland gc022 unknown 2.1 gc009 cell_cycle_2 gc021 cilia dorsal epithelia_dorsal epithelia

0.6

0.2

Orthogroup usage across cts and gene modules