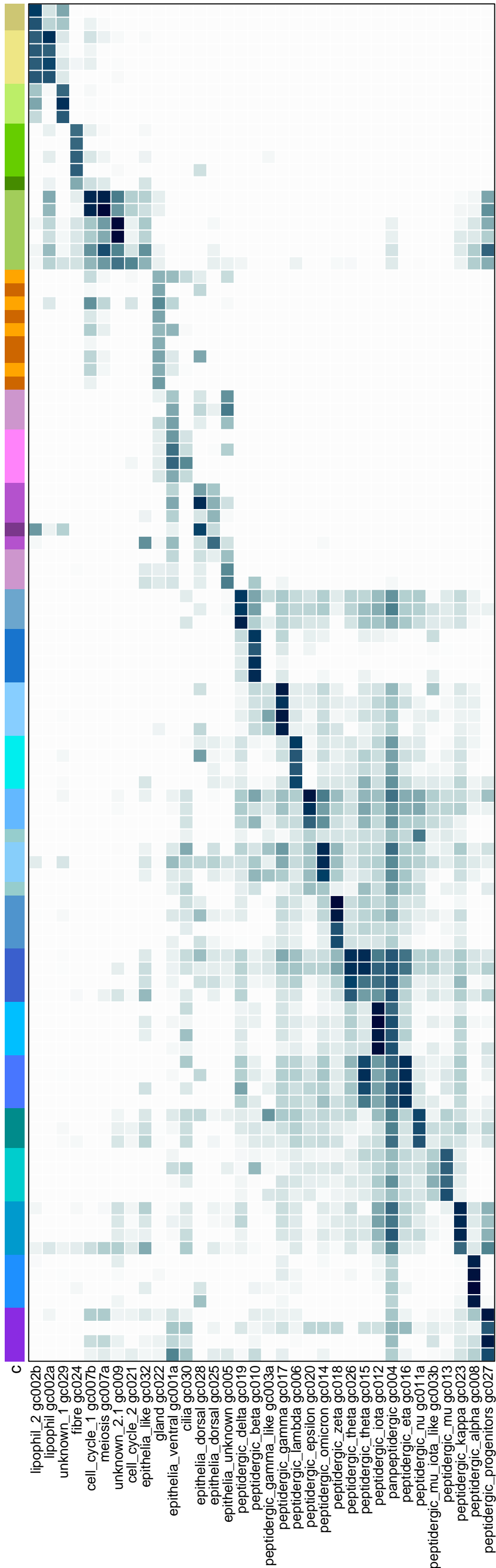


Orthogroup usage across cts and gene modules

n = 102 rows



- HoiH23|lipophil_2
- Hhon|lipophil_2
- Tadh|lipophil
- 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
- TrH2|unknown_2.2
- Tadh|unknown_2.1
- TrH2|unknown_2.1
- TrH2|unknown_2.4
- TrH2|unknown_2.3
- TrH2|epithelia_gland_like
- TrH2|gland_1
- 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_2
- 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_unknown
- Tadh|epithelia_unknown_1
- TrH2|epithelia_unknown_1
- Tadh|peptidergic_delta
- TrH2|peptidergic_delta
- Hhon|peptidergic_delta
- Tadh|peptidergic_beta
- TrH2|peptidergic_beta
- Hhon|peptidergic_beta
- HoiH23|peptidergic_beta
- HoiH23|peptidergic_gamma
- Hhon|peptidergic_gamma
- Tadh|peptidergic_gamma
- TrH2|peptidergic_gamma
- Tadh|peptidergic_lambda
- TrH2|peptidergic_lambda
- HoiH23|peptidergic_lambda
- 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
- HoiH23|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

frac

0.8

0.6

0.4

0.2

0

Orthogroup usage across cts and gene modules

n = 144 rows

