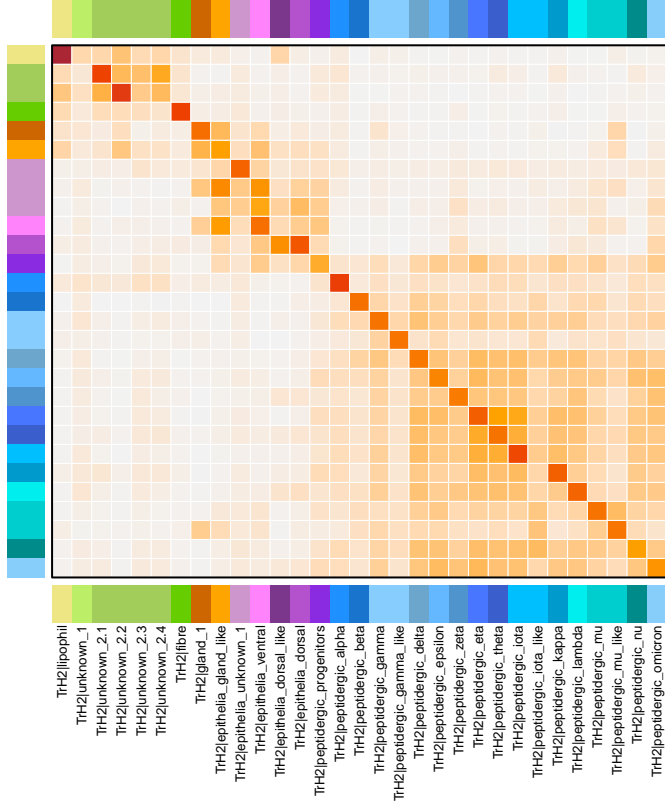


Tadh, n = 28

Tadh|lipophil
Tadh|unknown_2.1
Tadh|unknown_2.2
Tadh|fibre
Tadh|gland_1
Tadh|epithelia_gland_like
Tadh|epithelia_unknown_1
Tadh|epithelia_unknown_2
Tadh|epithelia_unknown_3
Tadh|epithelia_ventral
Tadh|epithelia_dorsal
Tadh|peptidergic_progenitors
Tadh|peptidergic_alpha
Tadh|peptidergic_beta
Tadh|peptidergic_gamma
Tadh|peptidergic_gamma_like
Tadh|peptidergic_delta
Tadh|peptidergic_epsilon
Tadh|peptidergic_zeta
Tadh|peptidergic_eta
Tadh|peptidergic_theta
Tadh|peptidergic_iota
Tadh|peptidergic_kappa
Tadh|peptidergic_lambda
Tadh|peptidergic_mu
Tadh|peptidergic_mu_like
Tadh|peptidergic_nu
Tadh|peptidergic_omicron



TrH2 (2795 markers, FC>=2.00), n = 31

TrH2|lipophil
TrH2|unknown_1
TrH2|unknown_2.1
TrH2|unknown_2.2
TrH2|unknown_2.3
TrH2|unknown_2.4
TrH2|fibre
TrH2|gland_1
TrH2|epithelia_gland_like
TrH2|epithelia_unknown_1
TrH2|epithelia_ventral
TrH2|epithelia_dorsal_like
TrH2|epithelia_dorsal
TrH2|peptidergic_progenitors
TrH2|peptidergic_alpha
TrH2|peptidergic_beta
TrH2|peptidergic_gamma
TrH2|peptidergic_gamma_like
TrH2|peptidergic_delta
TrH2|peptidergic_epsilon
TrH2|peptidergic_zeta
TrH2|peptidergic_eta
TrH2|peptidergic_theta
TrH2|peptidergic_iota
TrH2|peptidergic_iota_like
TrH2|peptidergic_kappa
TrH2|peptidergic_lambda
TrH2|peptidergic_mu
TrH2|peptidergic_mu_like
TrH2|peptidergic_nu
TrH2|peptidergic_omicron

Tadh|lipophil
Tadh|unknown_2.1
Tadh|unknown_2.2
Tadh|fibre
Tadh|gland_1
Tadh|epithelia_gland_like
Tadh|epithelia_unknown_1
Tadh|epithelia_unknown_2
Tadh|epithelia_unknown_3
Tadh|epithelia_ventral
Tadh|epithelia_dorsal
Tadh|peptidergic_progenitors
Tadh|peptidergic_alpha
Tadh|peptidergic_beta
Tadh|peptidergic_gamma
Tadh|peptidergic_gamma_like
Tadh|peptidergic_delta
Tadh|peptidergic_epsilon
Tadh|peptidergic_zeta
Tadh|peptidergic_eta
Tadh|peptidergic_theta
Tadh|peptidergic_iota
Tadh|peptidergic_kappa
Tadh|peptidergic_lambda
Tadh|peptidergic_mu
Tadh|peptidergic_mu_like
Tadh|peptidergic_nu
Tadh|peptidergic_omicron

