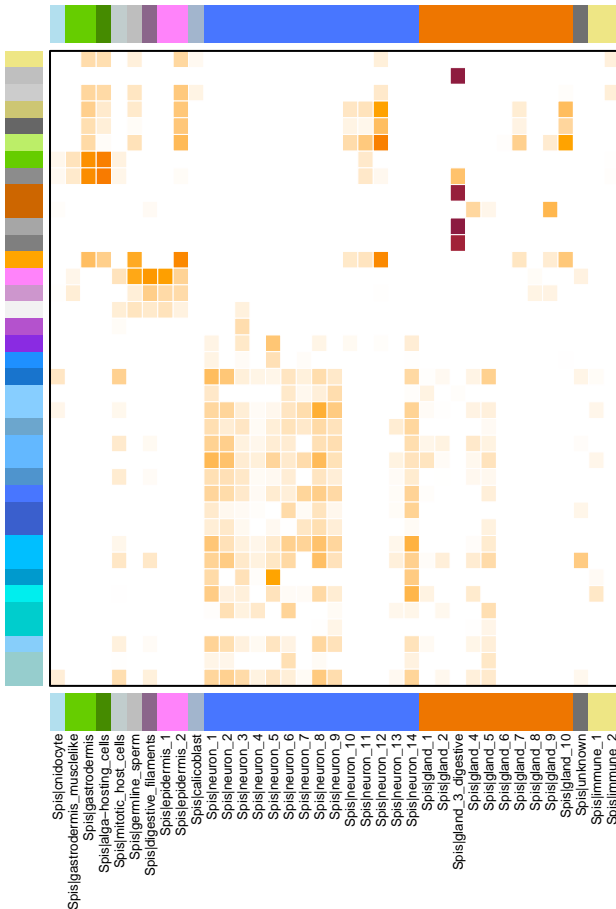


Spis (2686 markers, FC>=1.50), n = 37

Hhon, n = 38

Hhon|lipophil_1
Hhon|trans_lipophil_1_gland
Hhon|trans_lipophil_1_epithelia
Hhon|lipophil_2
Hhon|trans_lipophil_2_epithelia
Hhon|unknown_1
Hhon|fibre
Hhon|trans_fibre_gland
Hhon|gland_1
Hhon|gland_1_like
Hhon|trans_gland_epithelia_ventral
Hhon|trans_gland_epithelia_dorsal
Hhon|epithelia_gland_like
Hhon|epithelia_ventral
Hhon|epithelia_unknown
Hhon|trans_epithelia_dorsal_ventral
Hhon|epithelia_dorsal
Hhon|peptidergic_progenitors
Hhon|peptidergic_alpha
Hhon|peptidergic_beta
Hhon|peptidergic_gamma
Hhon|peptidergic_gamma_like
Hhon|peptidergic_delta
Hhon|peptidergic_epsilon
Hhon|peptidergic_epsilon_like
Hhon|peptidergic_zeta
Hhon|peptidergic_eta
Hhon|peptidergic_theta
Hhon|peptidergic_theta_like
Hhon|peptidergic_iota
Hhon|peptidergic_iota_like
Hhon|peptidergic_kappa
Hhon|peptidergic_lambda
Hhon|peptidergic_mu
Hhon|peptidergic_mu_like
Hhon|peptidergic_omicron
Hhon|peptidergic_other_1
Hhon|peptidergic_other_2



Hhon|lipophil_1
Hhon|trans_lipophil_1_gland
Hhon|trans_lipophil_1_epithelia
Hhon|lipophil_2
Hhon|trans_lipophil_2_epithelia
Hhon|unknown_1
Hhon|fibre
Hhon|trans_fibre_gland
Hhon|gland_1
Hhon|gland_1_like
Hhon|trans_gland_epithelia_ventral
Hhon|trans_gland_epithelia_dorsal
Hhon|epithelia_gland_like
Hhon|epithelia_ventral
Hhon|epithelia_unknown
Hhon|trans_epithelia_dorsal_ventral
Hhon|epithelia_dorsal
Hhon|peptidergic_progenitors
Hhon|peptidergic_alpha
Hhon|peptidergic_beta
Hhon|peptidergic_gamma
Hhon|peptidergic_gamma_like
Hhon|peptidergic_delta
Hhon|peptidergic_epsilon
Hhon|peptidergic_epsilon_like
Hhon|peptidergic_zeta
Hhon|peptidergic_eta
Hhon|peptidergic_theta
Hhon|peptidergic_theta_like
Hhon|peptidergic_iota
Hhon|peptidergic_iota_like
Hhon|peptidergic_kappa
Hhon|peptidergic_lambda
Hhon|peptidergic_mu
Hhon|peptidergic_mu_like
Hhon|peptidergic_omicron
Hhon|peptidergic_other_1
Hhon|peptidergic_other_2

wpearson

0.8
0.6
0.4
0.2
0