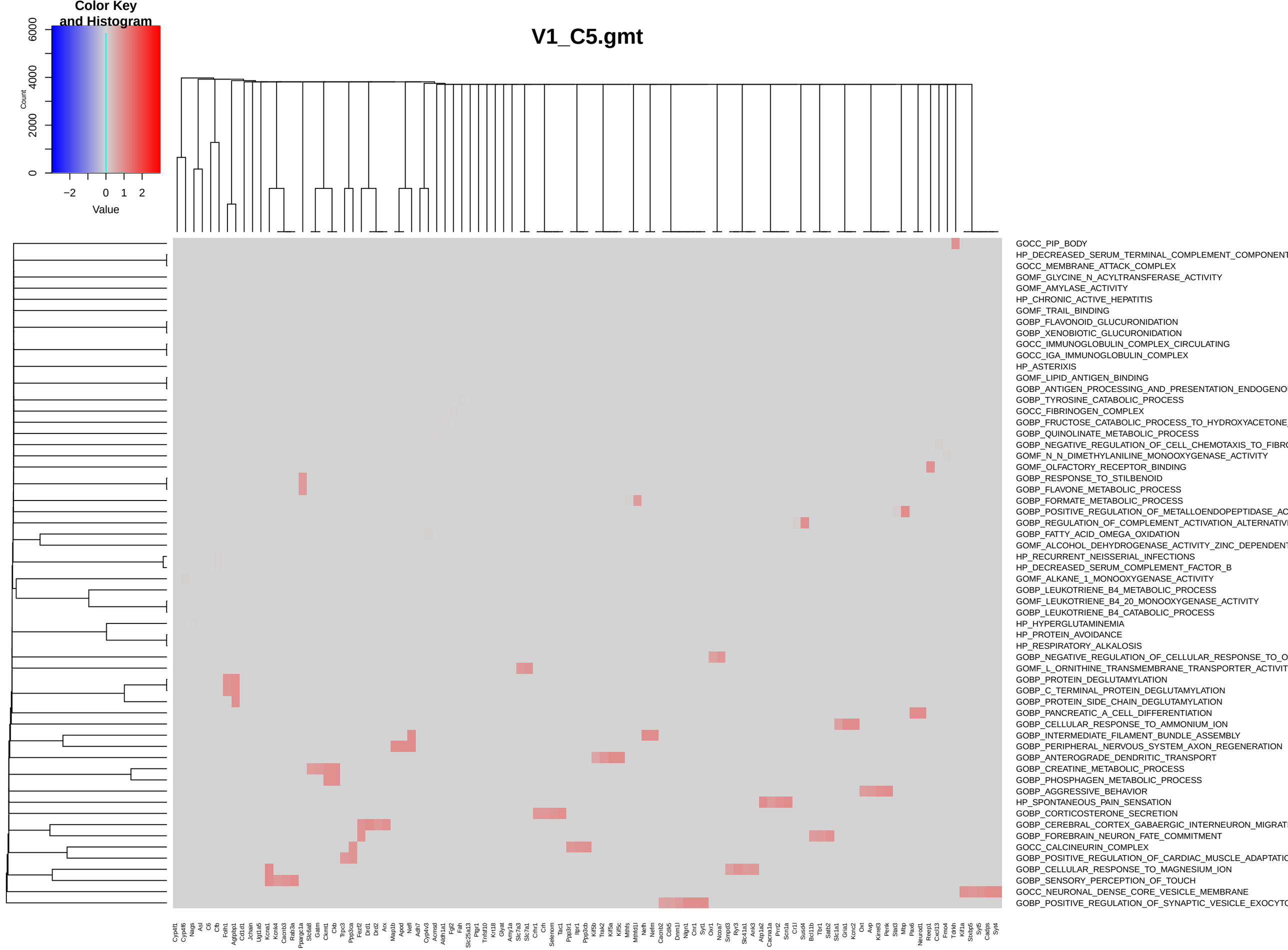
Color Key
and Histogram
Count
0
2000
4000
6000
−2
0
1
2
Value
V1_C5.gmt
GOCC_PIP_BODY
HP_DECREASED_SERUM_TERMINAL_COMPLEMENT_COMPONENT
GOCC_MEMBRANE_ATTACK_COMPLEX
GOMF_GLYCINE_N_ACYLTRANSFERASE_ACTIVITY
GOMF_AMYLASE_ACTIVITY
HP_CHRONIC_ACTIVE_HEPATITIS
GOMF_TRAIL_BINDING
GOBP_FLAVONOID_GLUCURONIDATION
GOBP_XENOBIOTIC_GLUCURONIDATION
GOCC_IMMUNOGLOBULIN_COMPLEX_CIRCULATING
GOCC_IGA_IMMUNOGLOBULIN_COMPLEX
HP_ASTERIXIS
GOMF_LIPID_ANTIGEN_BINDING
GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_ENDOGENO
GOBP_TYROSINE_CATABOLIC_PROCESS
GOCC_FIBRINOGEN_COMPLEX
GOBP_FRUCTOSE_CATABOLIC_PROCESS_TO_HYDROXYACETONE
GOBP_QUINOLINATE_METABOLIC_PROCESS
GOBP_NEGATIVE_REGULATION_OF_CELL_CHEMOTAXIS_TO_FIBRO
GOMF_N_N_DIMETHYLANILINE_MONOOXYGENASE_ACTIVITY
GOMF_OLFACTORY_RECEPTOR_BINDING
GOBP_RESPONSE_TO_STILBENOID
GOBP_FLAVONE_METABOLIC_PROCESS
GOBP_FORMATE_METABOLIC_PROCESS
GOBP_POSITIVE_REGULATION_OF_METALLOENDOPEPTIDASE_AC
GOBP_REGULATION_OF_COMPLEMENT_ACTIVATION_ALTERNATIVE
GOBP_FATTY_ACID_OMEGA_OXIDATION
GOMF_ALCOHOL_DEHYDROGENASE_ACTIVITY_ZINC_DEPENDENT
HP_RECURRENT_NEISSERIAL_INFECTIONS
HP_DECREASED_SERUM_COMPLEMENT_FACTOR_B
GOMF_ALKANE_1_MONOOXYGENASE_ACTIVITY
GOBP_LEUKOTRIENE_B4_METABOLIC_PROCESS
GOMF_LEUKOTRIENE_B4_20_MONOOXYGENASE_ACTIVITY
GOBP_LEUKOTRIENE_B4_CATABOLIC_PROCESS
HP_HYPERGLUTAMINEMIA
HP_PROTEIN_AVOIDANCE
HP_RESPIRATORY_ALKALOSIS
GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_O
GOMF_L_ORNITHINE_TRANSMEMBRANE_TRANSPORTER_ACTIVIT
GOBP_PROTEIN_DEGLUTAMYLATION
GOBP_C_TERMINAL_PROTEIN_DEGLUTAMYLATION
GOBP_PROTEIN_SIDE_CHAIN_DEGLUTAMYLATION
GOBP_PANCREATIC_A_CELL_DIFFERENTIATION
GOBP_CELLULAR_RESPONSE_TO_AMMONIUM_ION
GOBP_INTERMEDIATE_FILAMENT_BUNDLE_ASSEMBLY
GOBP_PERIPHERAL_NERVOUS_SYSTEM_AXON_REGENERATION
GOBP_ANTEROGRADE_DENDRITIC_TRANSPORT
GOBP_CREATINE_METABOLIC_PROCESS
GOBP_PHOSPHAGEN_METABOLIC_PROCESS
GOBP_AGGRESSIVE_BEHAVIOR
HP_SPONTANEOUS_PAIN_SENSATION
GOBP_CORTICOSTERONE_SECRETION
GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_MIGRAT
GOBP_FOREBRAIN_NEURON_FATE_COMMITMENT
GOCC_CALCINEURIN_COMPLEX
GOBP_POSITIVE_REGULATION_OF_CARDIAC_MUSCLE_ADAPTATIO
GOBP_CELLULAR_RESPONSE_TO_MAGNESIUM_ION
GOBP_SENSORY_PERCEPTION_OF_TOUCH
GOCC_NEURONAL_DENSE_CORE_VESICLE_MEMBRANE
GOBP_POSITIVE_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTO
Cyp4f1
Cyp4f6
Nags
Asl
C6
Cfb
Folh1
Agtpbp1
Cd1d1
Jchain
Ugt1a5
Kcna1
Kcnk4
Cacnb3
Rab3a
Ppargc1a
Slc6a8
Gatm
Ckmt1
Ckb
Trpc3
Ppp3ca
Fezf2
Drd1
Drd2
Aox
Map1b
Apod
Nefl
Adh7
Cyp4v3
Acmsd
Aldh1a1
Fgl2
Fah
Slc25a13
Ptgr1
Tnfsf10
Krt18
Glyat
Amy1a
Slc7a3
Slc7a1
Crhr1
Crh
Selenom
Tac1
Ppp3r1
Itpr1
Ppp3cb
Klf5b
Trak2
Klf5a
Klf5c
Mtrfs
Mthfd1l
Nefh
Nefm
Cacnb2
Cdk5
Dnm1l
Nlgn1
Cnr1
Syt1
Oxt
Ncoa7
Smpd3
Ryr3
Slc41a1
Ank3
Atp1a2
Cacna1a
Prrt2
Scn1a
Cr1l
Sus4
Bcl11b
Tbr1
Satb2
Slc1a1
Gria1
Kcnc2
Oxt
Avp
Kirrel3
Penk
Stat3
Mtap
Pax6
Neurod1
Reep1
Cxcl13
Fmo4
Tdrkh
Klf1a
Stxbp5
Syt5
Cadps
Syt4