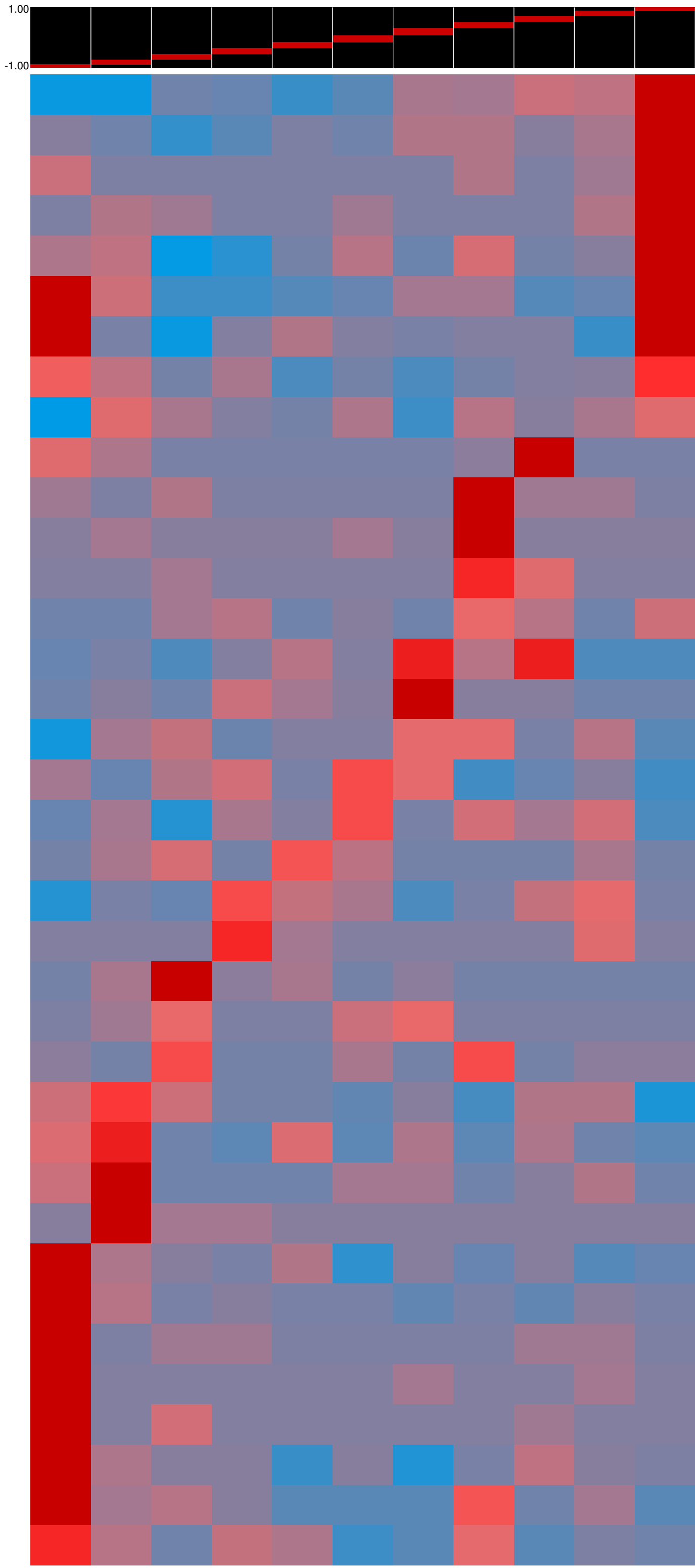


3 Over-representation

Under-representation -3



- GOBP_MITOCHONDRIAL_GENE_EXPRESSION
- GOBP_THIOESTER_BIOSYNTHETIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_CYTOPLASMIC_TRANSLATION
- GOCC_MITOTIC_SPINDLE_MICROTUBULE
- GOBP_NCRNA_TRANSCRIPTION
- GOCC_PROTEIN_ACETYLTRANSFERASE_COMPLEX
- GOBP_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME
- GOBP_POSITIVE_REGULATION_OF_ATPASE_ACTIVITY
- GOBP_RESPONSE_TO_OSMOTIC_STRESS
- HP_ANTERIOR_OPEN_BITE_MALOCCLUSION
- HP_ABNORMAL_BLOOD_CHLORIDE_CONCENTRATION
- HP_HYPOPLASTIC_FRONTAL_SINUSES
- GOCC_PARANODAL_JUNCTION
- GOBP_NODAL_SIGNALING_PATHWAY
- HP_ABNORMAL_EPIPHYSEAL_OSSIFICATION
- GOMF_PHOSPHOLIPASE_A2_ACTIVITY_CONSUMING_1_2_DIPAL
- GOCC_PLATELET_ALPHA_GRANULE_LUMEN
- HP_ATHETOSIS
- HP_ABNORMALITY_OF_BLADDER_MORPHOLOGY
- GOMF_METALLOENDOPEPTIDASE_INHIBITOR_ACTIVITY
- GOBP_INTERLEUKIN_10_PRODUCTION
- HP_ACANTHOLYSIS
- GOBP_CELL_PROLIFERATION_INVOLVED_IN_HEART_MORPHOGENESIS
- GOMF_PHOSPHOLIPASE_INHIBITOR_ACTIVITY
- HP_INCREASED_CIRCULATING_IGG_LEVEL
- GOBP_PROTEIN_LOCALIZATION_TO_CILIUM
- GOBP_REGULATION_OF_CELLULAR_AMINE_METABOLIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR
- GOBP_MIRNA_LOADING_ONTO_RISC_INVOLVED_IN_GENE_SILENCING
- GOBP_RECOMBINATIONAL_REPAIR
- GOBP_PROTEIN_K11_LINKED_UBIQUITINATION
- GOCC_ISWI_TYPE_COMPLEX
- GOCC_CORE_MEDIATOR_COMPLEX
- GOBP_REGULATION_OF_MICROTUBULE_BINDING
- GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLISM
- GOMF_METAL_CLUSTER_BINDING
- GOBP_MULTI_ORGANISM_LOCALIZATION