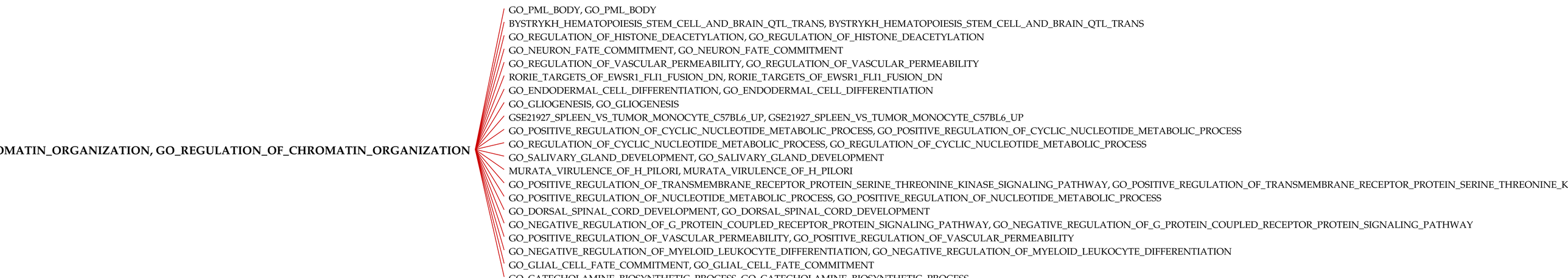
OMATIN_ORGANIZATION, GO_REGULATION_OF_CHROMATIN_ORGANIZATION
GO_PML_BODY, GO_PML_BODY
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_TRANS
GO_REGULATION_OF_HISTONE_DEACETYLATION, GO_REGULATION_OF_HISTONE_DEACETYLATION
GO_NEURON_FATE_COMMITMENT, GO_NEURON_FATE_COMMITMENT
GO_REGULATION_OF_VASCULAR_PERMEABILITY, GO_REGULATION_OF_VASCULAR_PERMEABILITY
RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_DN, RORIE_TARGETS_OF_EWSR1_FLI1_FUSION_DN
GO_ENDODERMAL_CELL_DIFFERENTIATION, GO_ENDODERMAL_CELL_DIFFERENTIATION
GO_GLIOGENESIS, GO_GLIOGENESIS
GSE21927_SPLEEN_VS_TUMOR_MONOCYTE_C57BL6_UP, GSE21927_SPLEEN_VS_TUMOR_MONOCYTE_C57BL6_UP
GO_POSITIVE_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS
GO_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS, GO_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS
GO_SALIVARY_GLAND_DEVELOPMENT, GO_SALIVARY_GLAND_DEVELOPMENT
MURATA_VIRULENCE_OF_H_PILORI, MURATA_VIRULENCE_OF_H_PILORI
GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_K
GO_POSITIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS
GO_DORSAL_SPINAL_CORD_DEVELOPMENT, GO_DORSAL_SPINAL_CORD_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_PROTEIN_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_PROTEIN_SIGNALING_PATHWAY
GO_POSITIVE_REGULATION_OF_VASCULAR_PERMEABILITY, GO_POSITIVE_REGULATION_OF_VASCULAR_PERMEABILITY
GO_NEGATIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION
GO_GLIAL_CELL_FATE_COMMITMENT, GO_GLIAL_CELL_FATE_COMMITMENT
GO_CATECHOLAMINE_BIOSYNTHETIC_PROCESS, GO_CATECHOLAMINE_BIOSYNTHETIC_PROCESS