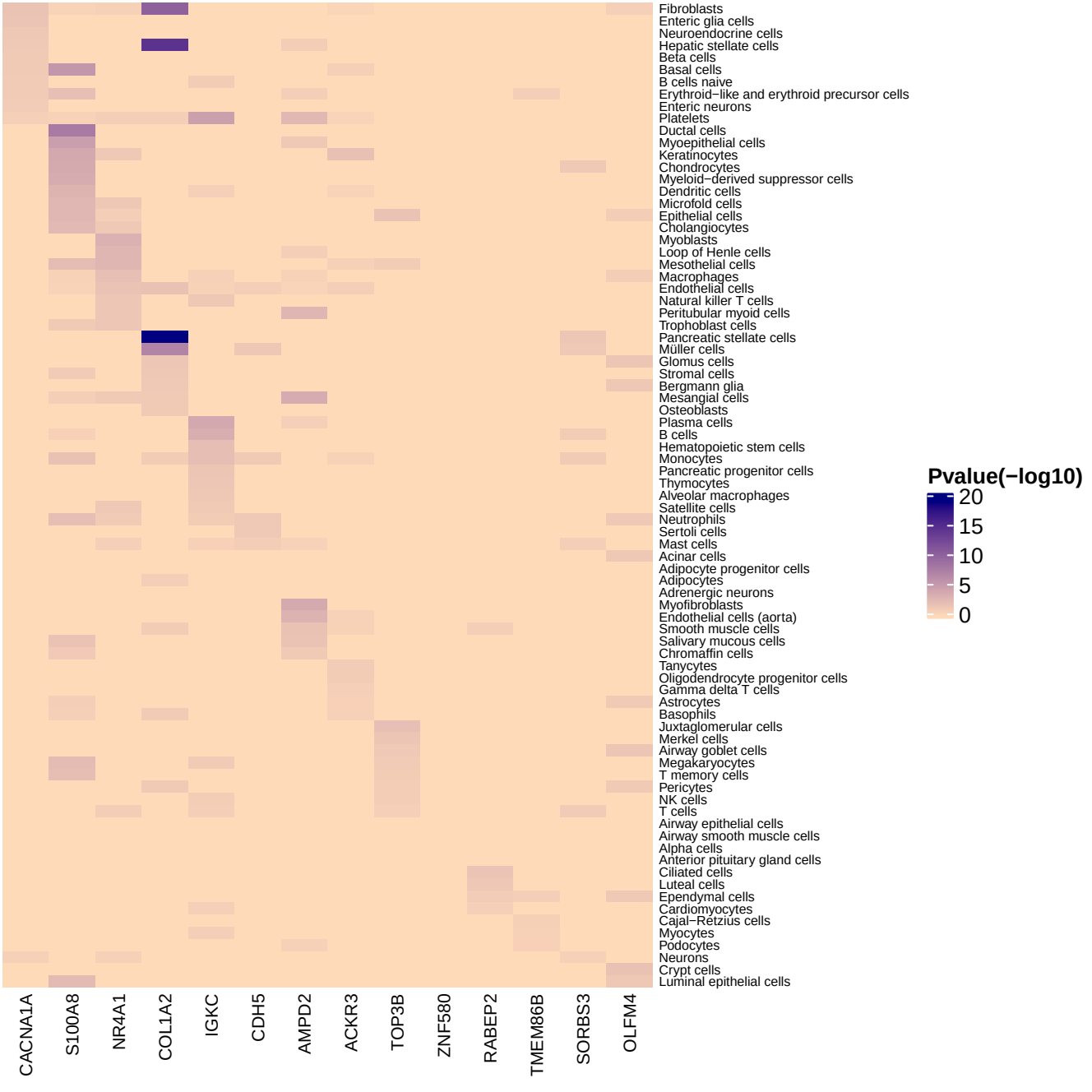
Fibroblasts
Enteric glia cells
Neuroendocrine cells
Hepatic stellate cells
Beta cells
Basal cells
B cells naive
Erythroid−like and erythroid precursor cells
Enteric neurons
Platelets
Ductal cells
Myoepithelial cells
Keratinocytes
Chondrocytes
Myeloid−derived suppressor cells
Dendritic cells
Microfold cells
Epithelial cells
Cholangiocytes
Myoblasts
Loop of Henle cells
Mesothelial cells
Macrophages
Endothelial cells
Natural killer T cells
Peritubular myoid cells
Trophoblast cells
Pancreatic stellate cells
Müller cells
Glomus cells
Stromal cells
Bergmann glia
Mesangial cells
Osteoblasts
Plasma cells
B cells
Hematopoietic stem cells
Monocytes
Pancreatic progenitor cells
Thymocytes
Alveolar macrophages
Satellite cells
Neutrophils
Sertoli cells
Mast cells
Acinar cells
Adipocyte progenitor cells
Adipocytes
Adrenergic neurons
Myofibroblasts
Endothelial cells (aorta)
Smooth muscle cells
Salivary mucous cells
Chromaffin cells
Tanycytes
Oligodendrocyte progenitor cells
Gamma delta T cells
Astrocytes
Basophils
Juxtaglomerular cells
Merkel cells
Airway goblet cells
Megakaryocytes
T memory cells
Pericytes
NK cells
T cells
Airway epithelial cells
Airway smooth muscle cells
Alpha cells
Anterior pituitary gland cells
Ciliated cells
Luteal cells
Ependymal cells
Cardiomyocytes
Cajal−Retzius cells
Myocytes
Podocytes
Neurons
Crypt cells
Luminal epithelial cells
CACNA1A
S100A8
NR4A1
COL1A2
IGKC
CDH5
AMPD2
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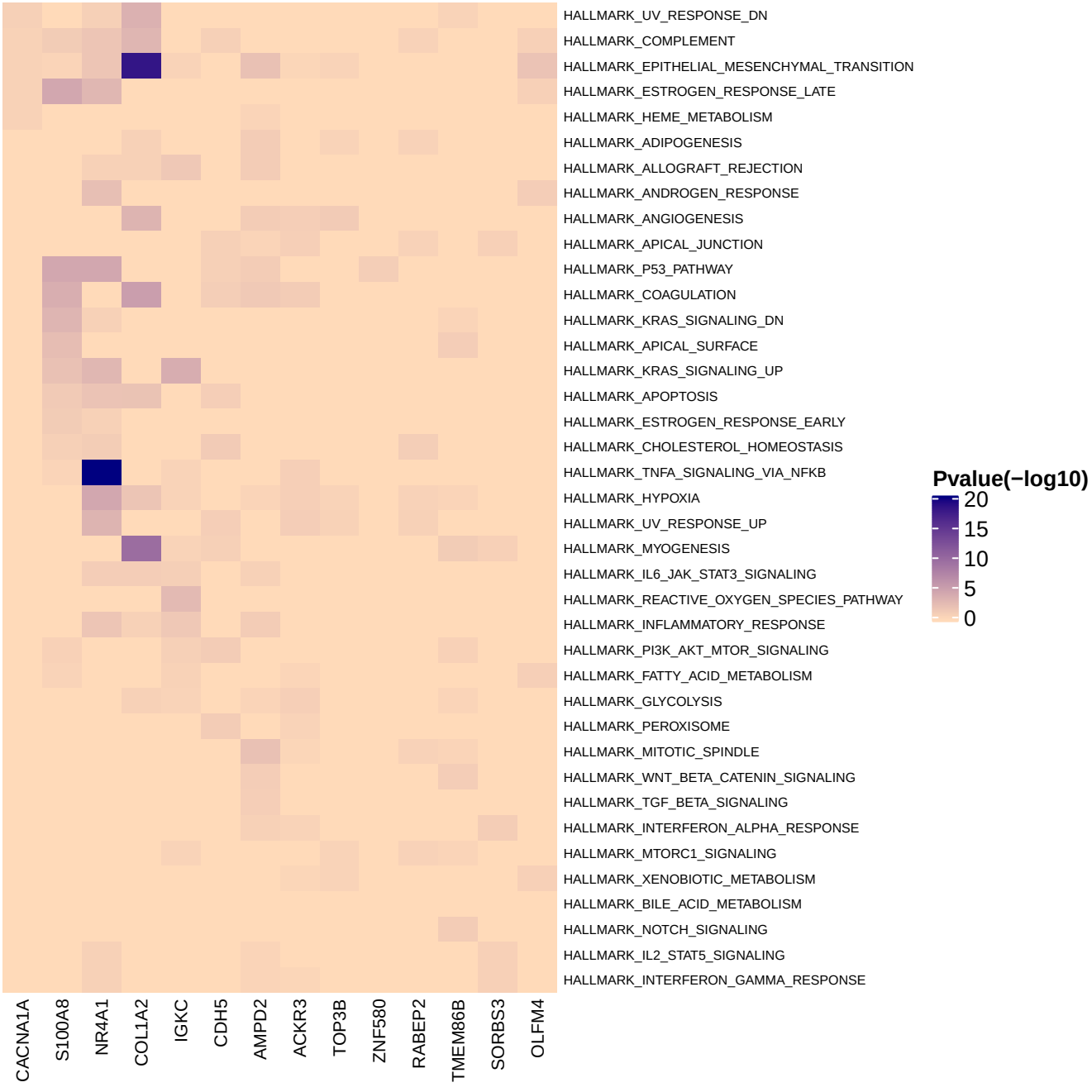
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HALLMARK_COMPLEMENT
HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION
HALLMARK_ESTROGEN_RESPONSE_LATE
HALLMARK_HEME_METABOLISM
HALLMARK_ADIPOGENESIS
HALLMARK_ALLOGRAFT_REJECTION
HALLMARK_ANDROGEN_RESPONSE
HALLMARK_ANGIOGENESIS
HALLMARK_APICAL_JUNCTION
HALLMARK_P53_PATHWAY
HALLMARK_COAGULATION
HALLMARK_KRAS_SIGNALING_DN
HALLMARK_APICAL_SURFACE
HALLMARK_KRAS_SIGNALING_UP
HALLMARK_APOPTOSIS
HALLMARK_ESTROGEN_RESPONSE_EARLY
HALLMARK_CHOLESTEROL_HOMEOSTASIS
HALLMARK_TNFA_SIGNALING_VIA_NFKB
HALLMARK_HYPOXIA
HALLMARK_UV_RESPONSE_UP
HALLMARK_MYOGENESIS
HALLMARK_IL6_JAK_STAT3_SIGNALING
HALLMARK_REACTIVE_OXYGEN_SPECIES_PATHWAY
HALLMARK_INFLAMMATORY_RESPONSE
HALLMARK_PI3K_AKT_MTOR_SIGNALING
HALLMARK_FATTY_ACID_METABOLISM
HALLMARK_GLYCOLYSIS
HALLMARK_PEROXISOME
HALLMARK_MITOTIC_SPINDLE
HALLMARK_WNT_BETA_CATENIN_SIGNALING
HALLMARK_TGF_BETA_SIGNALING
HALLMARK_INTERFERON_ALPHA_RESPONSE
HALLMARK_MTORC1_SIGNALING
HALLMARK_XENOBIOTIC_METABOLISM
HALLMARK_BILE_ACID_METABOLISM
HALLMARK_NOTCH_SIGNALING
HALLMARK_IL2_STAT5_SIGNALING
HALLMARK_INTERFERON_GAMMA_RESPONSE
CACNA1A
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NR4A1
COL1A2
IGKC
CDH5
AMPD2
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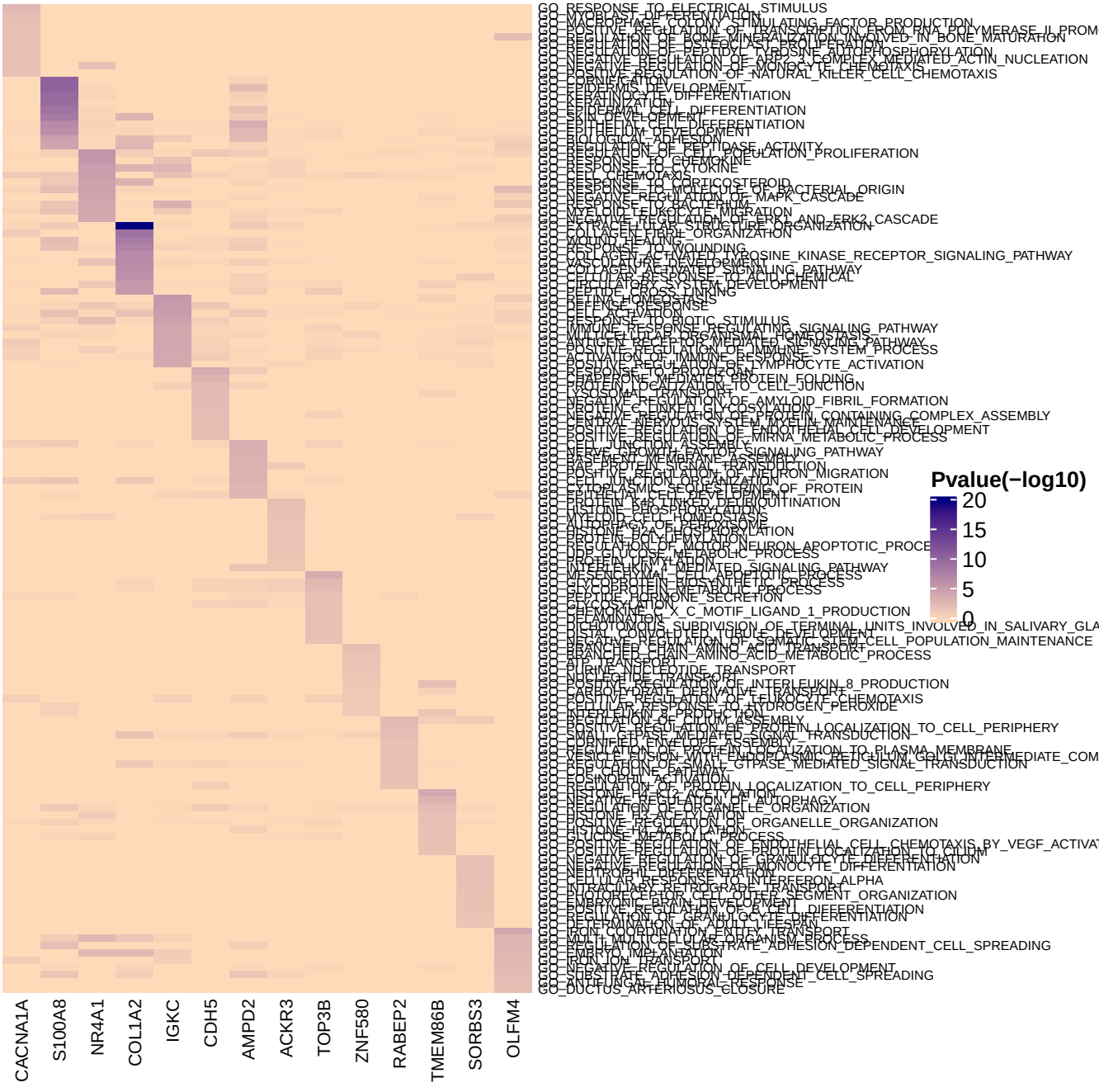
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OLFM4
GO_RESPONSE_TO_ELECTRICAL_STIMULUS
GO_MACROPHAGE_COLONY_STIMULATING_FACTOR_PRODUCTION
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROM
GO_REGULATION_OF_BONE_MINERALIZATION_INVOLVED_IN_BONE_MATURATION
GO_REGULATION_OF_OSTEOCLAST_PROLIFERATION
GO_REGULATION_OF_PEPTIDYL_TYROSINE_AUTOPHOSPHORYLATION
GO_POSITIVE_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
GO_NEGATIVE_REGULATION_OF_MONOCYTE_CHEMOTAXIS
GO_NEGATIVE_REGULATION_OF_NATURAL_KILLER_CELL_CHEMOTAXIS
GO_CORNIFICATION
GO_EPIDERMIS_DEVELOPMENT
GO_KERATINOCYTE_DIFFERENTIATION
GO_KERATINIZATION
GO_EPIDERMAL_CELL_DIFFERENTIATION
GO_SKIN_DEVELOPMENT
GO_EPITHELIAL_CELL_DIFFERENTIATION
GO_EPITHELIUM_DEVELOPMENT
GO_BIOLOGICAL_ADHESION
GO_REGULATION_OF_PEPTIDASE_ACTIVITY
GO_REGULATION_OF_CELL_POPULATION_PROLIFERATION
GO_RESPONSE_TO_CHEMOKINE
GO_RESPONSE_TO_CYTOKINE
GO_CELL_CHEMOTAXIS
GO_RESPONSE_TO_CORTICOSTEROID
GO_RESPONSE_TO_MOLECULE_OF_BACTERIAL_ORIGIN
GO_NEGATIVE_REGULATION_OF_MAPK_CASCADE
GO_RESPONSE_TO_BACTERIUM
GO_MYELOID_LEUKOCYTE_MIGRATION
GO_NEGATIVE_REGULATION_OF_ERK1_AND_ERK2_CASCADE
GO_EXTRACELLULAR_STRUCTURE_ORGANIZATION
GO_COLLAGEN_FIBRIL_ORGANIZATION
GO_WOUND_HEALING
GO_RESPONSE_TO_WOUNDING
GO_PLATELET_ACTIVATION
GO_VASCULATURE_DEVELOPMENT
GO_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_TYROSINE_KINASE_RECEPTOR_SIGNALING_PATHWAY
GO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_SIGNALING_PATHWAY
GO_CELLULAR_RESPONSE_TO_ANGIOTENSIN
GO_CIRCULATORY_SYSTEM_DEVELOPMENT
GO_PEPTIDE_CROSS_LINKING
GO_RETINAL_HOMEOSTASIS
GO_DEFENSE_RESPONSE
GO_PLATELET_ACTIVATION
GO_RESPONSE_TO_BIOTIC_STIMULUS
GO_IMMUNE_RESPONSE_-_REGULATING_SIGNALING_PATHWAY
GO_MUSCLE_CELLULAR_ORGANISMAL_HOMEOSTASIS
GO_ANTIGEN_RECEPTOR_MEDIATED_SIGNALING_PATHWAY
GO_POSITIVE_REGULATION_OF_IMMUNE_SYSTEM_PROCESS
GO_ACTIVATION_OF_IMMUNE_RESPONSE
GO_POSITIVE_REGULATION_OF_LYMPHOCYTE_ACTIVATION
GO_RESPONSE_TO_PROTOZOAN
GO_CHAPERONE_MEDIATED_PROTEIN_FOLDING
GO_PROTEIN_LOCALIZATION_TO_CELL_JUNCTION
GO_LYSOSOMAL_TRANSPORT
GO_NEGATIVE_REGULATION_OF_AMYLOID_FIBRIL_FORMATION
GO_PEPTIDE_CROSS_LINKING_VIA_CYSTEINE
GO_CENTRAL_NERVOUS_SYSTEM_MYELIN_MAINTENANCE
GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_MIRNA_METABOLIC_PROCESS
GO_CELL_JUNCTION_ASSEMBLY
GO_NERVE_GROWTH_FACTOR_SIGNALING_PATHWAY
GO_BASEMENT_MEMBRANE_ASSEMBLY
GO_ANTIGEN_PROCESSING_AND_PRESENTATION
GO_BASOPHIL_REGULATION_OF_NEURON_MIGRATION
GO_CELL_JUNCTION_ORGANIZATION
GO_CELLULAR_MACROMOLECULE_LOCALIZATION
GO_EPITHELIAL_CELL_DEVELOPMENT
GO_PROTEIN_SUMOylation
GO_PROTEIN_K48_LINKED_UBIQUITINATION
GO_HISTONE_PHOSPHORYLATION
GO_MYELOID_CELL_HOMEOSTASIS
GO_MITOPHAGY_-_ROTTENHOUSE
GO_HISTONE_H2A_PHOSPHORYLATION
GO_PROTEIN_POLYUBIQUITINATION
GO_REGULATION_OF_MOTOR_NEURON_APOPTOTIC_PROCE
GO_DDE_GLUCOSE_METABOLIC_PROCESS
GO_PROTEIN_UBIQUITINATION
GO_INTERLEUKIN_4_MEDIATED_SIGNALING_PATHWAY
GO_MESENCHYMAL_CELL_APOPTOTIC_PROCESS
GO_GLYCOPROTEIN_BIOSYNTHETIC_PROCESS
GO_PEPTIDE_HORMONE_METABOLIC_PROCESS
GO_PEPTIDE_HORMONE_SECRETION
GO_GLYCOSYLATION
GO_CHEMOKINE_(C_X_C_MOTIF)_LIGAND_1_PRODUCTION
GO_DEAMINATION
GO_DICHOTOMOUS_SUBDIVISION_OF_TERMINAL_UNITS_INVOLVED_IN_SALIVARY_GLA
GO_DISTAL_CONVOLUTED_TUBULE_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_SOMATIC_STEM_CELL_POPULATION_MAINTENANCE
GO_BRANCHED_CHAIN_AMINO_ACID_TRANSPORT
GO_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS
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GO_PURINE_NUCLEOTIDE_TRANSPORT
GO_NUCLEOTIDE_TRANSPORT
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GO_INTERLEUKIN_8_PRODUCTION
GO_REGULATION_OF_PODOSOME_ASSEMBLY
GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY
GO_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION
GO_PONTIFICAL_LYMPHOPEESTASIS
GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_PLASMA_MEMBRANE
GO_VESICLE_FUSION_WITH_ENDOPLASMIC_RETICULUM_GOLGI_INTERMEDIATE_COM
GO_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION
GO_CDP_CHOLINE_PATHWAY
GO_EOSINOPHIL_ACTIVATION
GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CELL_PERIPHERY
GO_HISTONE_H4K12_ACETYLATION
GO_NEGATIVE_REGULATION_OF_AUTOPHAGY
GO_REGULATION_OF_ORGANELLE_ORGANIZATION
GO_HISTONE_H2_ACETYLATION
GO_POSITIVE_REGULATION_OF_ORGANELLE_ORGANIZATION
GO_HISTONE_H4_ACETYLATION
GO_GLUCOSE_METABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS_BY_VEGF_ACTIVAT
GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CILIUM
GO_NEGATIVE_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION
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GO_NEUTROPHIL_DIFFERENTIATION
GO_CELLULAR_RESPONSE_TO_INTERFERON_ALPHA
GO_INTRACILIARY_RETROGRADE_TRANSPORT
GO_PHOTORECEPTOR_CELL_OUTER_SEGMENT_ORGANIZATION
GO_RHOMBIC_LIP_DORSAL_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_B_CELL_DIFFERENTIATION
GO_REGULATION_OF_GRANULOCYTE_DIFFERENTIATION
GO_DEFERMENTATION_OF_ADULT_LITTERSPIN
GO_IRON_COORDINATION_ENTITY_TRANSPORT
GO_MULTIMERIC_COMPLEX_DISASSEMBLY
GO_REGULATION_OF_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING
GO_EMBRYO_IMPLANTATION
GO_IRON_ION_TRANSPORT
GO_NEGATIVE_REGULATION_OF_CELL_DEVELOPMENT
GO_SUBSTRATE_ADHESION_DEPENDENT_CELL_SPREADING
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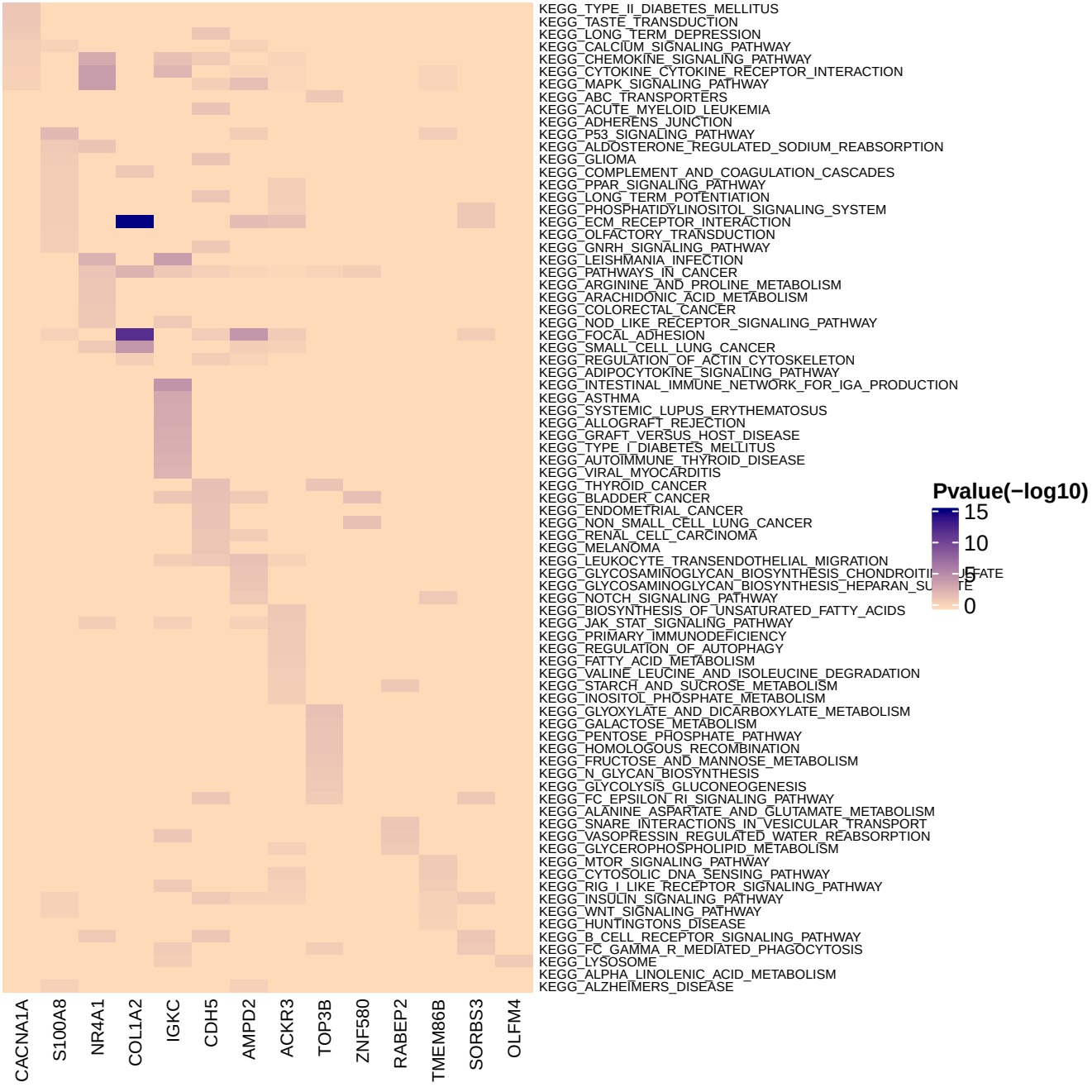
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KEGG_CYTOKINE_CYTOKINE_RECEPTOR_INTERACTION
KEGG_MAPK_SIGNALING_PATHWAY
KEGG_ABC_TRANSPORTERS
KEGG_ACUTE_MYELOID_LEUKEMIA
KEGG_ADHERENS_JUNCTION
KEGG_P53_SIGNALING_PATHWAY
KEGG_ALDOSTERONE_REGULATED_SODIUM_REABSORPTION
KEGG_GLIOMA
KEGG_COMPLEMENT_AND_COAGULATION_CASCADES
KEGG_PPAR_SIGNALING_PATHWAY
KEGG_LONG_TERM_POTENTIATION
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KEGG_ECM_RECEPTOR_INTERACTION
KEGG_OLFACTORY_TRANSDUCTION
KEGG_GNRH_SIGNALING_PATHWAY
KEGG_LEISHMANIA_INFECTION
KEGG_PATHWAYS_IN_CANCER
KEGG_ARGININE_AND_PROLINE_METABOLISM
KEGG_ARACHIDONIC_ACID_METABOLISM
KEGG_COLORECTAL_CANCER
KEGG_NOD_LIKE_RECEPTOR_SIGNALING_PATHWAY
KEGG_FOCAL_ADHESION
KEGG_SMALL_CELL_LUNG_CANCER
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KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY
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KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS
KEGG_ALLOGRAFT_REJECTION
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KEGG_TYPE_I_DIABETES_MELLITUS
KEGG_AUTOIMMUNE_THYROID_DISEASE
KEGG_VIRAL_MYOCARDITIS
KEGG_THYROID_CANCER
KEGG_BLADDER_CANCER
KEGG_ENDOMETRIAL_CANCER
KEGG_NON_SMALL_CELL_LUNG_CANCER
KEGG_RENAL_CELL_CARCINOMA
KEGG_MELANOMA
KEGG_LEUKOCYTE_TRANSENDOTHELIAL_MIGRATION
KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_CHONDROITIN_SULFATE
KEGG_GLYCOSAMINOGLYCAN_BIOSYNTHESIS_HEPARAN_SULFATE
KEGG_NOTCH_SIGNALING_PATHWAY
KEGG_BIOSYNTHESIS_OF_UNSATURATED_FATTY_ACIDS
KEGG_JAK_STAT_SIGNALING_PATHWAY
KEGG_PRIMARY_IMMUNODEFICIENCY
KEGG_REGULATION_OF_AUTOPHAGY
KEGG_FATTY_ACID_METABOLISM
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION
KEGG_STARCH_AND_SUCROSE_METABOLISM
KEGG_INOSITOL_PHOSPHATE_METABOLISM
KEGG_GLYOXYLATE_AND_DICARBOXYLATE_METABOLISM
KEGG_GALACTOSE_METABOLISM
KEGG_PENTOSE_PHOSPHATE_PATHWAY
KEGG_HOMOLOGOUS_RECOMBINATION
KEGG_FRUCTOSE_AND_MANNOSE_METABOLISM
KEGG_N_GLYCAN_BIOSYNTHESIS
KEGG_GLYCOLYSIS_GLUCONEOGENESIS
KEGG_FC_EPSILON_RI_SIGNALING_PATHWAY
KEGG_ALANINE_ASPARTATE_AND_GLUTAMATE_METABOLISM
KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT
KEGG_VASOPRESSIN_REGULATED_WATER_REABSORPTION
KEGG_GLYCEROPHOSPHOLIPID_METABOLISM
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KEGG_CYTOSOLIC_DNA_SENSING_PATHWAY
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COL1A2
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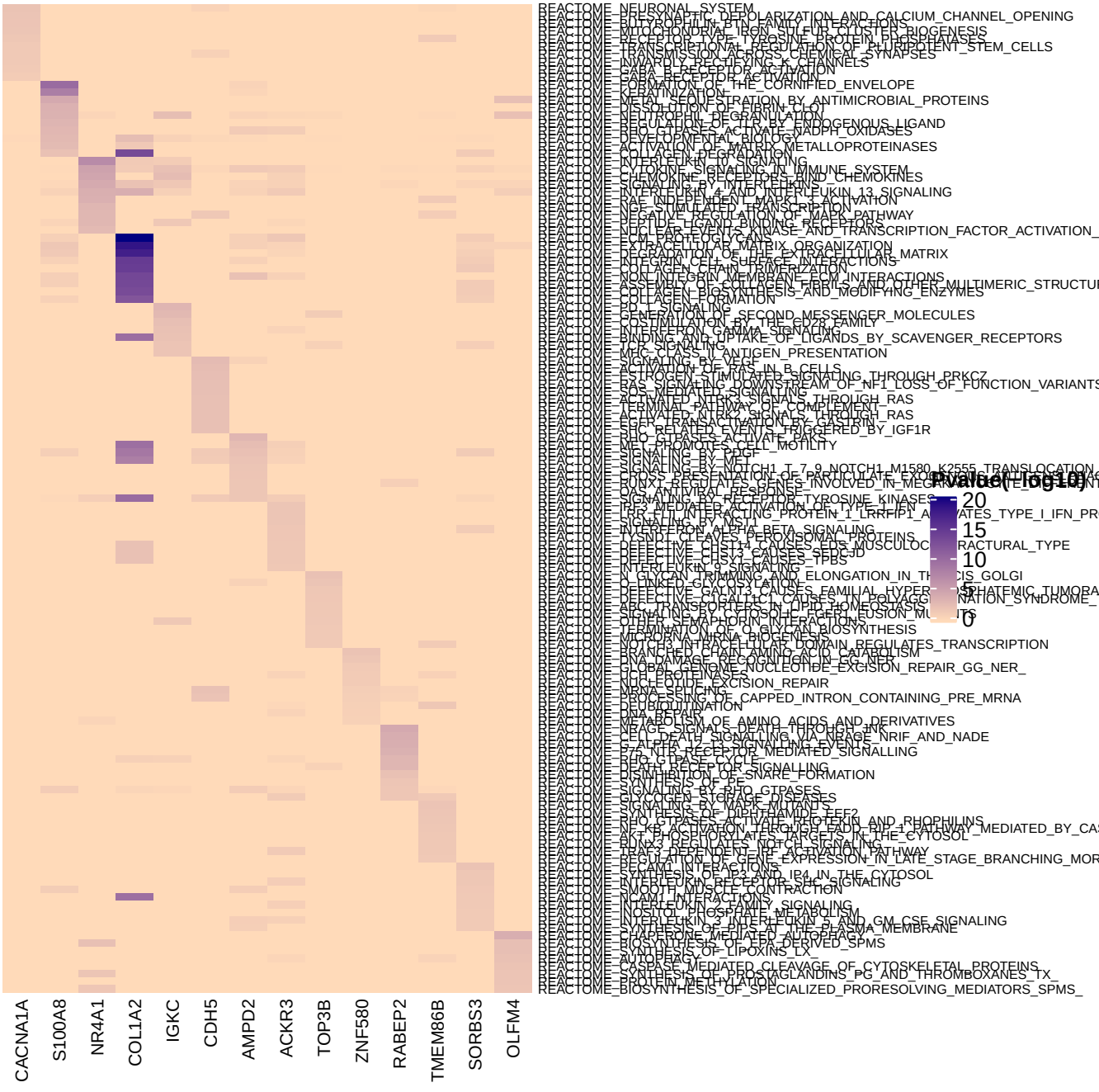
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TMEM86B
SORBS3
OLFM4
REACTOME_NEURONAL_SYSTEM
REACTOME_DUTYSHPIN_DEPOLARIZATION_AND_CALCIUM_CHANNEL_OPENING
REACTOME_BIOTIN_TRANSPORT_AND_METABOLISM
REACTOME_MITOCHONDRIAL_IRON_SULFUR_CLUSTER_BIOGENESIS
REACTOME_RECEPTOR_TYPE_TYROSINE_PROTEIN_PHOSPHATASES
REACTOME_TRANSCRIPTIONAL_REGULATION_OF_PLURIPOTENT_STEM_CELLS
REACTOME_TRANSMISSION_ACROSS_CHEMICAL_SYNAPSES
REACTOME_INWARDLY_RECTIFYING_K+_CHANNELS
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REACTOME_GABA_RECEPTOR_ACTIVATION
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REACTOME_KERATINIZATION
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REACTOME_DISSOLUTION_OF_FIBRIN_CLOT
REACTOME_NEUTROPHIL_DEGRANULATION
REACTOME_REGULATION_OF_TLR_BY_ENDOGENOUS_LIGAND
REACTOME_NADPH_OXIDASE_(ACTIVATE_NADPH_OXIDASES
REACTOME_DEVELOPMENTAL_BIOLOGY
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REACTOME_CELL_CELL_REGRADATION
REACTOME_INTERLEUKIN_10_SIGNALING
REACTOME_CYTOKINE_SIGNALING_IN_IMMUNE_SYSTEM
REACTOME_CHEMOKINE_RECEPTORS_BIND_CHEMOKINES
REACTOME_SIGNALING_BY_INTERLEUKINS
REACTOME_INTERLEUKIN_4_AND_INTERLEUKIN_13_SIGNALING
REACTOME_NAD_DEPENDENT_DEACETYLASE_ACTIVATION
REACTOME_NOTCH_HES_MEDIATED_TRANSCRIPTION
REACTOME_NEGATIVE_REGULATION_OF_MAPK_PATHWAY
REACTOME_CELL_CELL_COMMUNICATION_BINDING_RECEPTORS
REACTOME_NUCLEAR_EVENTS_KINASE_AND_TRANSCRIPTION_FACTOR_ACTIVATION_
REACTOME_ECM_PROTEOGLYCANS
REACTOME_EXTRACELLULAR_MATRIX_ORGANIZATION
REACTOME_DEGRADATION_OF_THE_EXTRACELLULAR_MATRIX
REACTOME_INTEGRIN_CELL_SURFACE_INTERACTIONS
REACTOME_LAMININ_CHAIN_MEMBRANE_ACTIVATION
REACTOME_NON_INTEGRIN_MEMBRANE_ECM_INTERACTIONS
REACTOME_ASSEMBLY_OF_COLLAGEN_FIBRILS_AND_OTHER_MULTIMERIC_STRUCTUI
REACTOME_COLLAGEN_BIOSYNTHESIS_AND_MODIFYING_ENZYMES
REACTOME_COLLAGEN_FORMATION
REACTOME_PD-1_SIGNALING
REACTOME_GENERATION_OF_SECOND_MESSENGER_MOLECULES
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REACTOME_SIGNALING_BY_VEGF
REACTOME_ACTIVATION_OF_RAS_IN_B_CELLS
REACTOME_ESTROGEN_STIMULATED_SIGNALING_THROUGH_PRKCZ
REACTOME_RAS_SIGNALING_DOWNSTREAM_OF_NF1_LOSS_OF_FUNCTION_VARIANTS
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REACTOME_OAS_ANTIVIRAL_RESPONSE_
REACTOME_SIGNALING_BY_RECEPTOR_TYROSINE_KINASES_
REACTOME_IRF3_MEDIATED_ACTIVATION_OF_TYPE_I_IFN_ATES_TYPE_I_IFN_PR
REACTOME_TRP_FTL_INTERACTING_PROTEIN_1_LRRFIP1_A
REACTOME_SIGNALING_BY_MSTR1_
REACTOME_INTERFERON_ALPHA_BETA_SIGNALING_
REACTOME_TYSNDT_CLEAVES_PEROXISOMAL_PROTEINS_
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REACTOME_DEFECTIVE_CHST3_CAUSES_SEDCJD_
REACTOME_DEFECTIVE_CHSY1_CAUSES_TPBS_
REACTOME_INTERLEUKIN_9_SIGNALING_
REACTOME_N_GLYCAN_TRIMMING_AND_ELONGATION_IN_THIS_GOLGI
REACTOME_O-LINKED_GLYCOSYLATION_
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REACTOME_DEFECTIVE_GALNT3_CAUSES_HYPERPHOSPHATEMIA_TUMORAL_CALCINOSIS_
REACTOME_ABC_TRANSPORTERS_IN_LIPID_HOMEOSTASIS_
REACTOME_SIGNALING_BY_CYTOSOLIC_FGFR_FUSION_MUTANTS_
REACTOME_OTHER_SEMAPHORIN_INTERACTIONS_
REACTOME_TERMINATION_OF_O_GLYCAN_BIOSYNTHESIS_
REACTOME_MICRORNA_(MIRNA)_BIOGENESIS_
REACTOME_NOTCH3_INTRACELLULAR_DOMAIN_REGULATES_TRANSCRIPTION_
REACTOME_BRANCHED_CHAIN_AMINO_ACID_CATABOLISM_
REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER_
REACTOME_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR_GG_NER_
REACTOME_VCP_P97_PROTEIN_ESCRT_
REACTOME_NUCLEOTIDE_EXCISION_REPAIR_
REACTOME_MRNA_SPLICING_
REACTOME_PROCESSING_OF_CAPPED_INTRON_CONTAINING_PRE_MRNA_
REACTOME_DEUBIQUITINATION_
REACTOME_DNA_REPAIR_
REACTOME_METABOLISM_OF_AMINO_ACIDS_AND_DERIVATIVES_
REACTOME_MAPK_SIGNALS_DEATH_THROUGH_JNK_
REACTOME_CELL_DEATH_SIGNALLING_VIA_NRAGE_NRIF_AND_NADE_
REACTOME_ERBB4_11_SIGNALING_EVENTS_
REACTOME_P75_NTR_RECEPTOR_MEDIATED_SIGNALLING_
REACTOME_RHO_GTPASE_CYCLE_
REACTOME_DEATH_RECEPTOR_SIGNALING_
REACTOME_DISINHIBITION_OF_SNARE_FORMATION_
REACTOME_SYNTHESIS_OF_PE_
REACTOME_SIGNALING_BY_RHO_GTPASES_
REACTOME_GLYCOGEN_STORAGE_DISEASES_
REACTOME_SIGNALING_BY_MAPK_MUTANTS_
REACTOME_SYNTHESIS_OF_DIPHTHAMIDE_EEF2_
REACTOME_RHO_GTPASES_ACTIVATE_KTN_KATANIN_AND_RHOPHILINS_
REACTOME_NF-KB_ACTIVATION_THROUGH_FADD_RIP-1_PATHWAY_MEDIATED_BY_CA
REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_CYTOSOL_
REACTOME_RUNX3_REGULATES_NOTCH_SIGNALING_
REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY_
REACTOME_REGULATION_OF_GENE_EXPRESSION_IN_LATE_STAGE_BRANCHING_MOR
REACTOME_CELL_CELL_INTERACTIONS_
REACTOME_SYNTHESIS_OF_IP3_AND_IP4_IN_THE_CYTOSOL_
REACTOME_INTERLEUKIN_RECEPTOR_SHC_SIGNALING_
REACTOME_SMOOTH_MUSCLE_CONTRACTION_
REACTOME_NCAM1_INTERACTIONS_
REACTOME_INTERLEUKIN_4_FAMILY_SIGNALING_
REACTOME_INOSITOL_PHOSPHATE_METABOLISM_
REACTOME_INTERLEUKIN_3_INTERLEUKIN_5_AND_GM_CSF_SIGNALING_
REACTOME_SYNTHESIS_OF_PIPS_IN_THE_PLASMA_MEMBRANE_
REACTOME_CHAPERONE_MEDIATED_AUTOPHAGY_
REACTOME_BIOSYNTHESIS_OF_EPA_DERIVED_SPMS_
REACTOME_CONVERSION_OF_LIPOXINS_LX_
REACTOME_AUTOPHAGY_
REACTOME_CASPASE_MEDIATED_CLEAVAGE_OF_CYTOSKELETAL_PROTEINS_
REACTOME_SYNTHESIS_OF_PROSTAGLANDINS_PG_AND_THROMBOXANES_TX_
REACTOME_PRIMARY_HEMOSTASIS_
REACTOME_BIOSYNTHESIS_OF_SPECIALIZED_PRORESOLVING_MEDIATORS_SPMS_
pvalue(-log10)
20
15
10
5
0