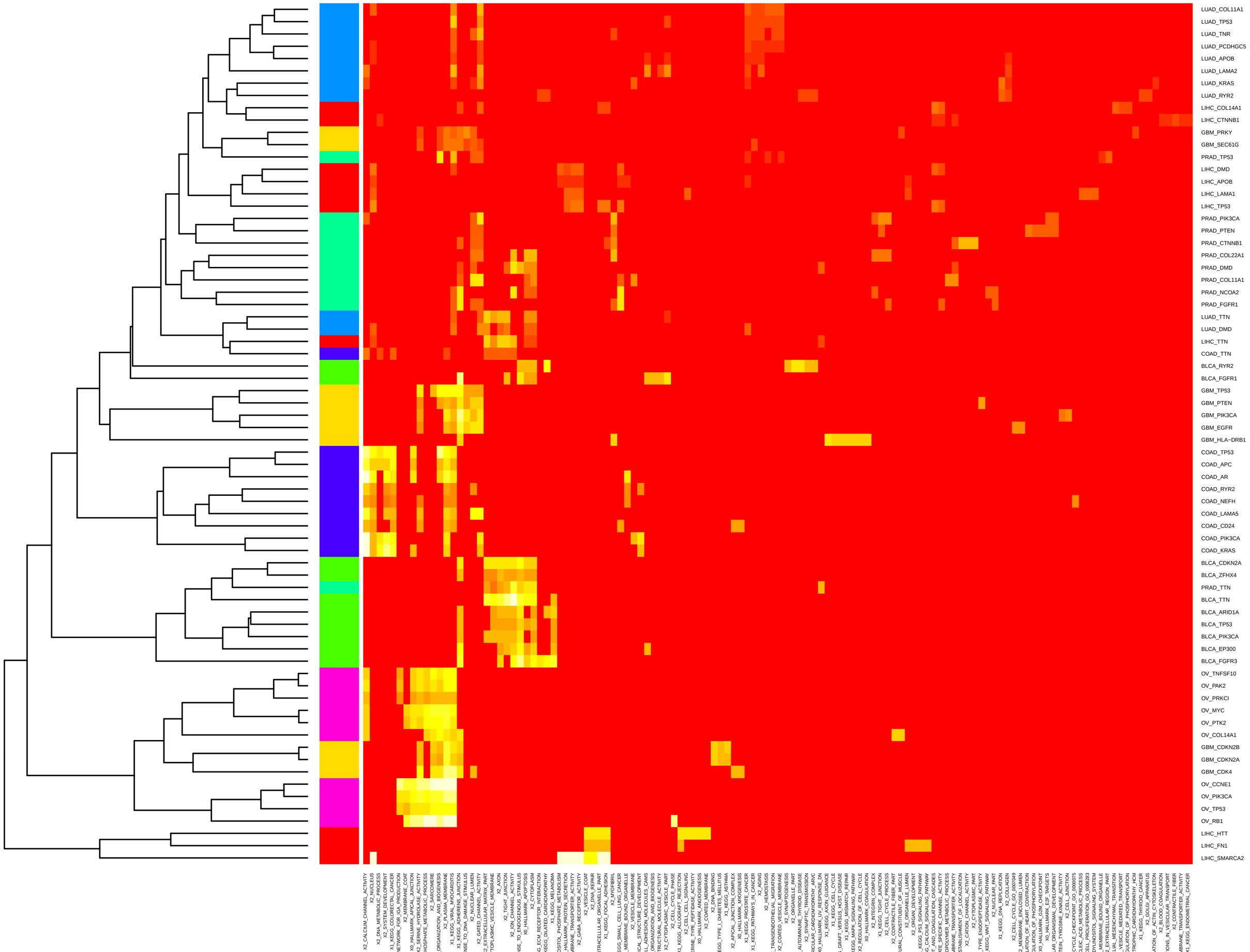
LUAD_COL11A1
LUAD_TP53
LUAD_TNR
LUAD_PCDHGC5
LUAD_APOB
LUAD_LAMA2
LUAD_KRAS
LUAD_RYR2
LIHC_COL14A1
LIHC_CTNNB1
GBM_PRKY
GBM_SEC61G
PRAD_TP53
LIHC_DMD
LIHC_APOB
LIHC_LAMA1
LIHC_TP53
PRAD_PIK3CA
PRAD_PTEN
PRAD_CTNNB1
PRAD_COL22A1
PRAD_DMD
PRAD_COL11A1
PRAD_NCOA2
PRAD_FGFR1
LUAD_TTN
LUAD_DMD
LIHC_TTN
COAD_TTN
BLCA_RYR2
BLCA_FGFR1
GBM_TP53
GBM_PTEN
GBM_PIK3CA
GBM_EGFR
GBM_HLA−DRB1
COAD_TP53
COAD_APC
COAD_AR
COAD_RYR2
COAD_NEFH
COAD_LAMA5
COAD_CD24
COAD_PIK3CA
COAD_KRAS
BLCA_CDKN2A
BLCA_ZFHX4
PRAD_TTN
BLCA_TTN
BLCA_ARID1A
BLCA_TP53
BLCA_PIK3CA
BLCA_EP300
BLCA_FGFR3
OV_TNFSF10
OV_PAK2
OV_PRKCI
OV_MYC
OV_PTK2
OV_COL14A1
GBM_CDKN2B
GBM_CDKN2A
GBM_CDK4
OV_CCNE1
OV_PIK3CA
OV_TP53
OV_RB1
LIHC_HTT
LIHC_FN1
LIHC_SMARCA2
X2_CALCIUM_CHANNEL_ACTIVITY
X2_NUCLEUS
X2_DNA_METABOLIC_PROCESS
X2_SYSTEM_DEVELOPMENT
X1_KEGG_COLORECTAL_CANCER
X2_NETWORK_FOR_IGA_PRODUCTION
X2_MEMBRANE_COAT
X0_HALLMARK_APOPTOSIS
X2_SERINE-HYDROLASE_ACTIVITY
X2_PHOSPHATE_METABOLIC_PROCESS
X2_SARCOMERE
X2_ORGANIZATION_AND_BIOGENESIS
X2_PLASMA_MEMBRANE
X1_KEGG_VIRAL_MYOCARDITIS
X1_KEGG_ADHERENS_JUNCTION
X2_RESPONSE_TO_DNA_DAMAGE_STIMULUS
X2_NUCLEAR_LUMEN
X2_GATED_CHANNEL_ACTIVITY
X2_EXTRACELLULAR_MATRIX_PART
X2_CYTOPLASMIC_VESICLE_MEMBRANE
X2_AXON
X2_TIGHT_JUNCTION
X2_ION_CHANNEL_ACTIVITY
X0_HALLMARK_APOPTOSIS
X2_CYTOPLASM
X2_ECM_RECEPTOR_INTERACTION
X1_KEGG_DILATED_CARDIOMYOPATHY
X1_KEGG_MELANOMA
X2_PHOSPHOINOSITOL_PHOSPHATE_METABOLISM
X0_HALLMARK_PROTEIN_SECRETION
X2_MEMBRANE_TRANSPORTER_ACTIVITY
X2_GABA_RECEPTOR_ACTIVITY
X2_VESICLE_COAT
X2_EXTRACELLULAR_ORGANELLE_PART
X1_KEGG_FOCAL_ADHESION
X2_MYOFIBRIL
X1_KEGG_SMALL_CELL_LUNG_CANCER
X2_MEMBRANE_BOUND_ORGANELLE
X2_VESICLE_MEMBRANE
X2_CHEMICAL_STRUCTURE_DEVELOPMENT
X2_CELL_ADHESION_MOLECULES_CAMS
X2_ORGANIZATION_AND_BIOGENESIS
X2_STRUCTURAL_MOLECULE_ACTIVITY
X2_CYTOPLASMIC_VESICLE_PART
X2_CELL_CYCLE_PHASE
X2_CELL_CYCLE_ALIGNMENT_REDUCTION
X1_KEGG_ALLOGRAFT_REJECTION
X2_CELL_CELL_SIGNALING
X2_SERINE_TYPE_PEPTIDASE_ACTIVITY
X0_HALLMARK_ANGIOGENESIS
X2_COATED_MEMBRANE
X2_DNA_BINDING
X1_KEGG_TYPE_I_DIABETES_MELLITUS
X1_KEGG_ASTHMA
X2_APICAL_JUNCTION_COMPLEX
X0_HALLMARK_MYOGENESIS
X1_KEGG_PROSTATE_CANCER
X1_KEGG_PATHWAYS_IN_CANCER
X2_AGING
X2_HEMOSTASIS
X2_TRANSENDOTHELIAL_MIGRATION
X2_COATED_VESICLE_MEMBRANE
X2_SYNAPTOGENESIS
X2_ORGANELLE_PART
X2_AUTOIMMUNE_THYROID_DISEASE
X2_SYNAPTIC_TRANSMISSION
X0_HALLMARK_CARDIOMYOPATHY_ARVC
X0_HALLMARK_UV_RESPONSE_DN
X1_KEGG_AXON_GUIDANCE
X1_KEGG_CELL_CYCLE
X1_KEGG_GRAFT_VERSUS_HOST_DISEASE
X1_KEGG_MISMATCH_REPAIR
X1_KEGG_ALANINE_ASPARTATE_AND_GLUTAMATE_METABOLISM
X2_REGULATION_OF_CELL_CYCLE
X0_HALLMARK_COAGULATION
X2_INTEGRIN_COMPLEX
X1_KEGG_TIGHT_JUNCTION
X2_CELL_CYCLE_PROCESS
X2_CONTRACTILE_FIBER_PART
X2_STRUCTURAL_CONSTITUENT_OF_MUSCLE
X2_ORGANELLE_LUMEN
X2_ORGAN_DEVELOPMENT
X1_KEGG_P53_SIGNALING_PATHWAY
X2_CALCIUM_SIGNALING_PATHWAY
X2_COAGULATION_CASCADES
X2_VOLTAGE-GATED_CHANNEL_ACTIVITY
X2_ENZYME_BINDING_ACTIVITY
X2_MEMBRANE_TRANSPORTER_ACTIVITY
X2_ESTABLISHMENT_OF_LOCALIZATION
X2_CATION_CHANNEL_ACTIVITY
X2_CYTOPLASMIC_PART
X1_KEGG_TYPE_ENDOPEPTIDASE_ACTIVITY
X1_KEGG_WNT_SIGNALING_PATHWAY
X2_NUCLEAR_PART
X1_KEGG_DNA_REPLICATION
X2_COLLAGEN
X2_CELL_CYCLE_GO_0007049
X2_MEMBRANE_ENCLOSED_LUMEN
X2_ACTIVATION_OF_PROTEIN_KINASE_ACTIVITY
X2_REGULATION_OF_PHOSPHORYLATION
X0_HALLMARK_G2M_CHECKPOINT
X0_HALLMARK_EPE_TARGETS
X2_CELLULAR_ORGANISMAL_DEVELOPMENT
X2_CELL_JUNCTION
X1_KEGG_PROTEIN_TYROSINE_KINASE_ACTIVITY
X2_CELL_CYCLE_CHECKPOINT_GO_0000075
X2_ELEC_ACID_METABOLIC_PROCESS
X2_CELL_PROLIFERATION_GO_0008283
X2_PHOSPHOINOSITOL_SIGNALING_SYSTEM
X2_MEMBRANE_BOUND_ORGANELLE
X2_EXTRACELLULAR_REGION_PART
X2_ACTIVATION_OF_MAPKKK_ACTIVITY
X2_MEMBRANE_TRANSPORT
X2_VESICLE_MEDIATED_TRANSPORT
X2_REGULATION_OF_PHOSPHORYLATION
X1_KEGG_THYROID_CANCER
X2_GOLGI_APPARATUS
X2_REGULATION_OF_ACTIN_CYTOSKELETON
X2_BLOOD_COAGULATION
X2_CONTRACTILE_FIBER
X2_PROTEINS_IN_VESICULAR_TRANSPORT
X2_MEMBRANE_TRANSPORTER_ACTIVITY
X1_KEGG_ENDOMETRIAL_CANCER