

**LATION\_OF\_TRAFFICKING, REACTOME\_RAB\_REGULATION\_OF\_TRAFFICKING**

- REACTOME\_RAB\_GEF5\_EXCHANGE\_GTP\_FOR\_GDP\_ON\_RABS, REACTOME\_RAB\_GEF5\_EXCHANGE\_GTP\_FOR\_GDP\_ON\_RABS
- REACTOME\_TP53\_REGULATES\_METABOLIC\_GENES, REACTOME\_TP53\_REGULATES\_METABOLIC\_GENES
- REACTOME\_RETROGRADE\_TRANSPORT\_AT\_THE\_TRANS\_GOLGI\_NETWORK, REACTOME\_RETROGRADE\_TRANSPORT\_AT\_THE\_TRANS\_GOLGI\_NETWORK
- KEGG\_ENDOCYTOSIS, KEGG\_ENDOCYTOSIS
- HOLLEMAN\_ASPARAGINASE\_RESISTANCE\_ALL\_DN, HOLLEMAN\_ASPARAGINASE\_RESISTANCE\_ALL\_DN
- PID\_AJDISS\_2PATHWAY, PID\_AJDISS\_2PATHWAY
- WU\_HBX\_TARGETS\_2\_UP, WU\_HBX\_TARGETS\_2\_UP
- WP\_AMPACTIVATED\_PROTEIN\_KINASE\_AMPK\_SIGNALING, WP\_AMPACTIVATED\_PROTEIN\_KINASE\_AMPK\_SIGNALING
- REACTOME\_SYNTHESIS\_OF\_PIPS\_AT\_THE\_PLASMA\_MEMBRANE, REACTOME\_SYNTHESIS\_OF\_PIPS\_AT\_THE\_PLASMA\_MEMBRANE
- PID\_IGF1\_PATHWAY, PID\_IGF1\_PATHWAY
- WP\_TRANSLATION\_INHIBITORS\_IN\_CHRONICALLY\_ACTIVATED\_PDGFRA\_CELLS, WP\_TRANSLATION\_INHIBITORS\_IN\_CHRONICALLY\_ACTIVATED\_PDGFRA\_CELLS
- YAGI\_AML\_RELAPSE\_PROGNOSIS, YAGI\_AML\_RELAPSE\_PROGNOSIS
- KEGG\_MTOR\_SIGNALING\_PATHWAY, KEGG\_MTOR\_SIGNALING\_PATHWAY
- REACTOME\_REGULATION\_OF\_GENE\_EXPRESSION\_IN\_BETA\_CELLS, REACTOME\_REGULATION\_OF\_GENE\_EXPRESSION\_IN\_BETA\_CELLS
- WP\_CELL\_MIGRATION\_AND\_INVASION\_THROUGH\_P75NTR, WP\_CELL\_MIGRATION\_AND\_INVASION\_THROUGH\_P75NTR
- MULLIGHAN\_NPM1\_MUTATED\_SIGNATURE\_1\_DN, MULLIGHAN\_NPM1\_MUTATED\_SIGNATURE\_1\_DN
- KEGG\_REGULATION\_OF\_AUTOPHAGY, KEGG\_REGULATION\_OF\_AUTOPHAGY