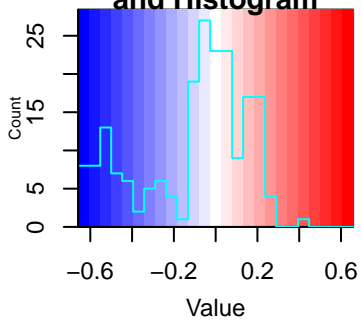
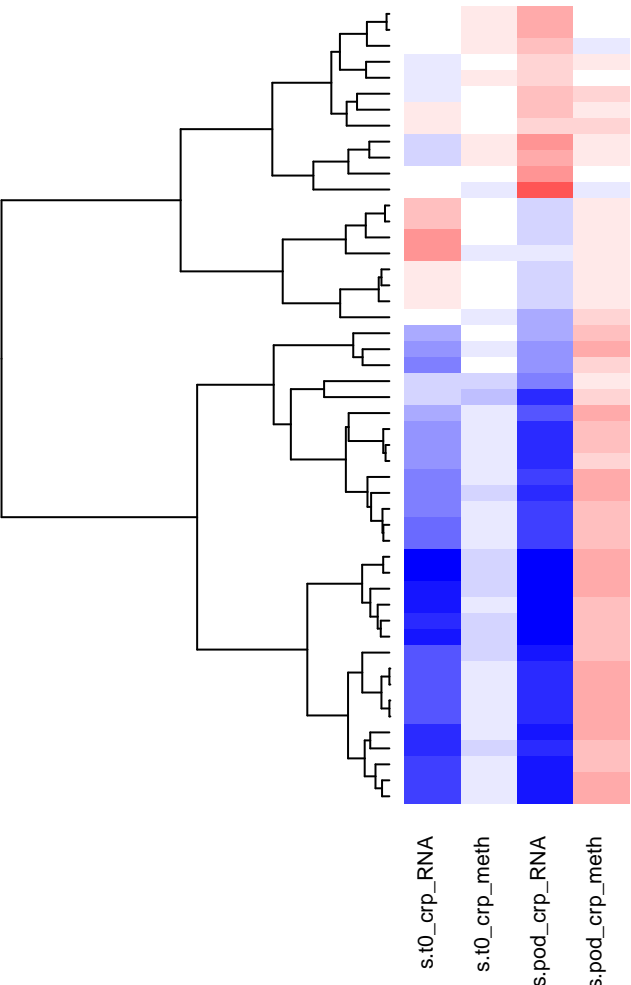


Color Key and Histogram



4way



- Vesicle-mediated transport
- Membrane Trafficking
- Metabolism of lipids
- Signal Transduction
- Metabolism
- Signaling by Interleukins
- Immune System
- Cytokine Signaling in Immune system
- Platelet activation, signaling and aggregation
- Signaling by Receptor Tyrosine Kinases
- Innate Immune System
- Neutrophil degranulation
- Cell Cycle, Mitotic
- Cell Cycle
- Cell Cycle Checkpoints
- Interferon Signaling
- RNA Polymerase II Transcription
- Gene expression (Transcription)
- Generic Transcription Pathway
- Metabolism of RNA
- Signaling by ROBO receptors
- Regulation of expression of SLITs and ROBOs
- Cellular response to starvation
- Translation
- rRNA modification in the nucleus and cytosol
- Influenza Infection
- Major pathway of rRNA processing in the nucleolus and cytosol
- rRNA processing in the nucleus and cytosol
- rRNA processing
- Influenza Viral RNA Transcription and Replication
- SRP-dependent cotranslational protein targeting to membrane
- Ribosomal scanning and start codon recognition
- Activation of the mRNA upon binding of the cap-binding complex and eIFs, and subsequent binding to 43S
- Translation initiation complex formation
- Peptide chain elongation
- Eukaryotic Translation Elongation
- Viral mRNA Translation
- Selenocysteine synthesis
- Formation of a pool of free 40S subunits
- Eukaryotic Translation Termination
- Formation of the ternary complex, and subsequently, the 43S complex
- Nonsense-Mediated Decay (NMD)
- Nonsense Mediated Decay (NMD) enhanced by the Exon Junction Complex (EJC)
- Eukaryotic Translation Initiation
- Cap-dependent Translation Initiation
- Nonsense Mediated Decay (NMD) independent of the Exon Junction Complex (EJC)
- Selenoamino acid metabolism
- Response of EIF2AK4 (GCN2) to amino acid deficiency
- GTP hydrolysis and joining of the 60S ribosomal subunit
- L13a-mediated translational silencing of Ceruloplasmin expression

s.t0_crp_RNA
s.t0_crp_meth
s.pod_crp_RNA
s.pod_crp_meth