Enrichment analysis of signatures

Legend:
- Sign 0
- Sign 1
- Sign 2
- Sign 3
- Sign 4
- Sign 5
- Sign 6
- Sign 7
- Sign 8
- Sign 9

Categories (top to bottom):
- Outer membrane
- Endoplasmic reticulum tubular network
- Mitochondrion
- Mitochondrial envelope
- Envelope
- Carboxylic acid binding
- Organic acid binding
- 2 oxoglutarate dependent dioxygenase activity
- Oxidoreductase activity acting on paired donors with [...]
- Endoplasmic reticulum lumen
- Supramolecular complex
- Microtubule
- Structural constituent of cytoskeleton
- Supramolecular polymer
- Polymeric cytoskeletal fiber
- Ion transmembrane transporter activity
- Intrinsic component of organelle membrane
- Secondary active transmembrane transporter activity
- Active ion transmembrane transporter activity
- Active transmembrane transporter activity
- Nuclear periphery
- Nucleolus
- Single stranded dna binding
- Mrna binding
- Rna binding
- Inner mitochondrial membrane protein complex
- Mitochondrial envelope
- Mitochondrial protein containing complex
- Mitochondrion
- Organelle inner membrane
- Intracellular protein containing complex
- Proteasome core complex
- Peptidase complex
- Endopeptidase complex
- Proteasome complex
- Vacuole
- Vacuolar lumen
- Secretory vesicle
- Secretory granule
- Azurophil granule
- Rna binding
- Catalytic step 2 spliceosome
- Mrna binding
- Spliceosomal complex
- Ribonucleoprotein complex
- Intrinsic component of organelle membrane
- Intrinsic component of endoplasmic reticulum membrane
- Organelle subcompartment
- Nuclear outer membrane endoplasmic reticulum membrane network
- Endoplasmic reticulum

X-axis: Sum of -log10(p-value)