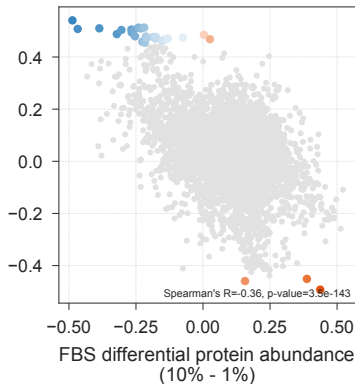


PCs enrichment scores (NES)



- GO_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GO_CRISTAE_FORMATION
- GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX
- REACTOME_CRISTAE_FORMATION
- GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE
- GO_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
- GO_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT
- GO_MITOCHONDRIAL_PROTEIN_COMPLEX
- GO_INNER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX
- GO_ORGANELLE_INNER_MEMBRANE
- GO_ORGANELLAR_RIBOSOME
- GO_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT
- REACTOME_RESPIRATORY_ELECTRON_TRANSPORT_ATP_SYNTHESIS_BY_CHEMIOSMOTIC_COUPLING_AND_HEAT_PRODUCTION_BY_UNCOUPLING_PROTEINS
- GO_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION
- GO_MITOCHONDRIAL_GENE_EXPRESSION
- GO_MITOCHONDRIAL_TRANSLATION
- REACTOME_MITOCHONDRIAL_TRANSLATION
- GO_RESPIRATORY_CHAIN_COMPLEX
- GO_NADH_DEHYDROGENASE_COMPLEX
- GO_RESPIRASOME
- GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY
- GO_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY
- GO_NADH_DEHYDROGENASE_ACTIVITY
- GO_NUCLEAR_INNER_MEMBRANE
- GO_SMALL_SUBUNIT_PROCESSOME
- GO_90S_PRERIBOSOME
- GO_PROTEASOME_ACCESSORY_COMPLEX
- GO_CHAPERONE_COMPLEX
- REACTOME_COOPERATION_OF_PREFOLDIN_AND_TRIC_CCT_IN_ACTIN_AND_TUBULIN_FOLDING