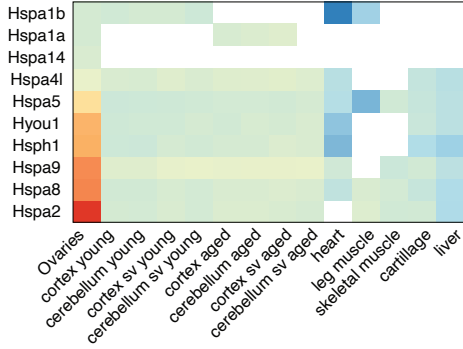
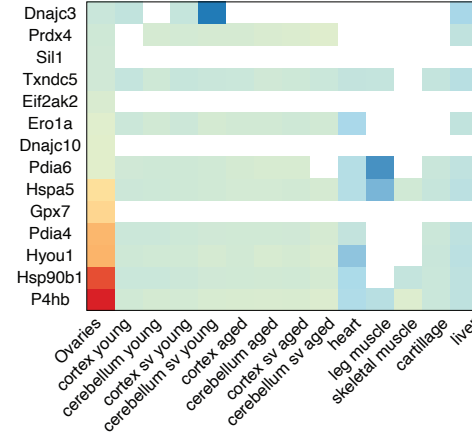


enriched terms

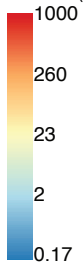
Heat shock protein 70 family



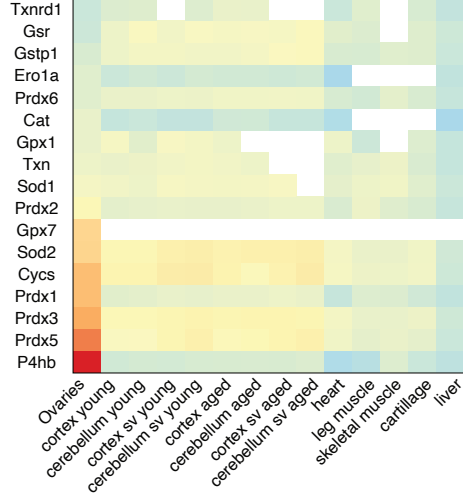
Endoplasmic reticulum chaperone complex, and unfolded protein response



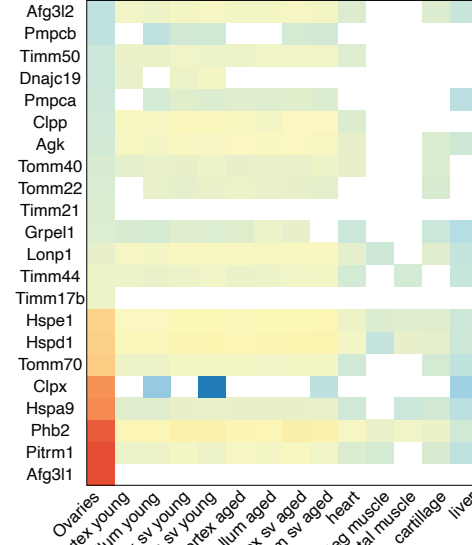
half-life (days)



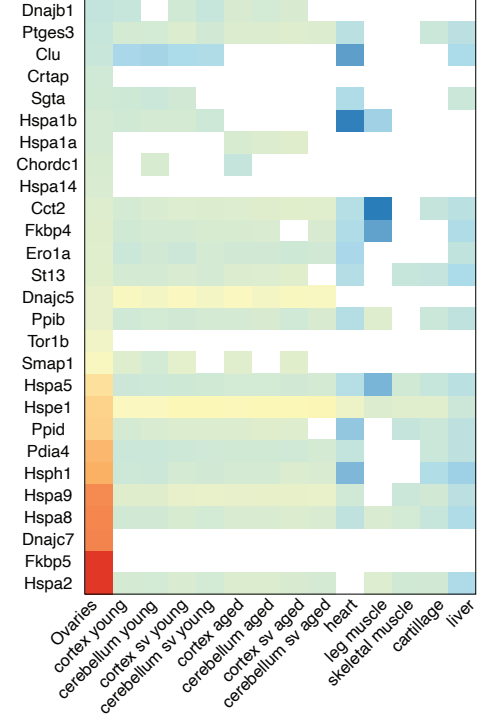
Detoxification of Reactive Oxygen Species



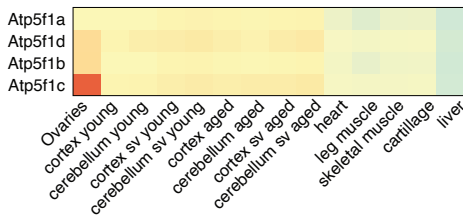
Protein targeting to mitochondrion, and m-aaa complex



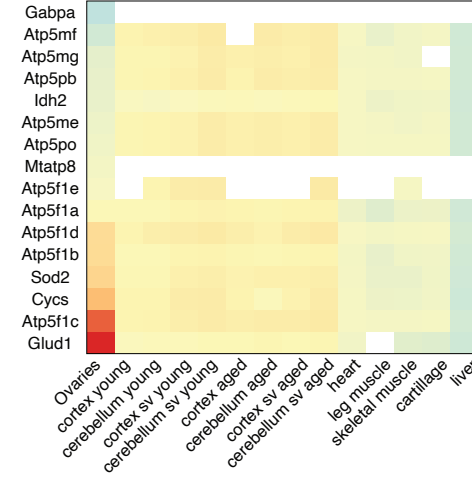
Chaperone-mediated protein folding



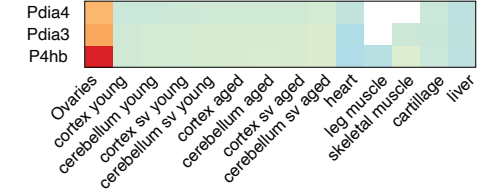
Mitochondrial proton-transporting atp synthase complex, catalytic sector f(1)



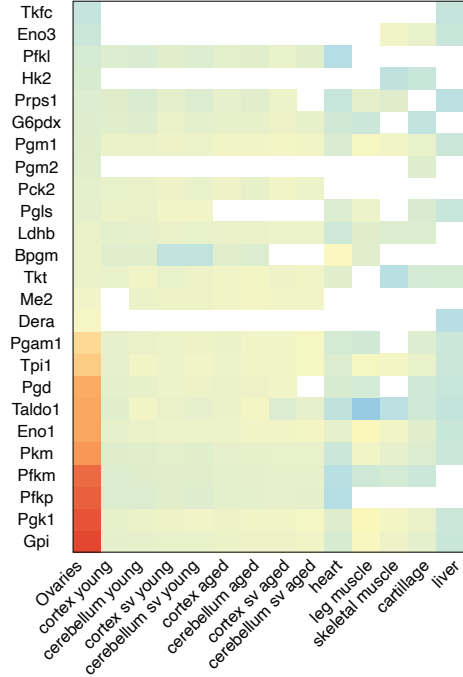
Mitochondrial biogenesis



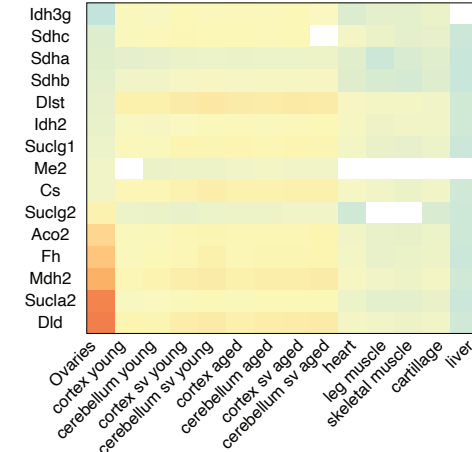
Protein disulphide isomerase



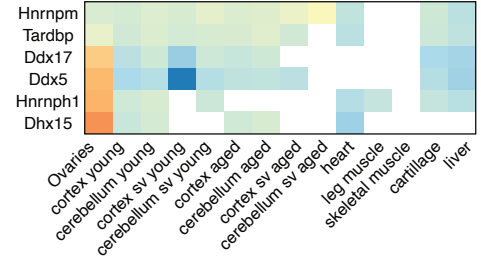
Pentose phosphate pathway, and pyruvate metabolic process



Citric acid cycle (TCA cycle)

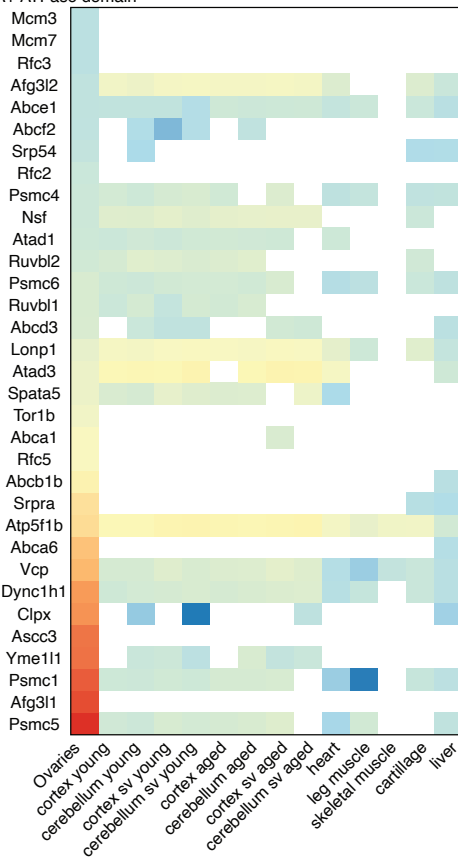


Drosha complex

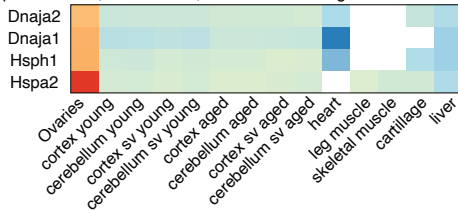


enriched terms (p2)

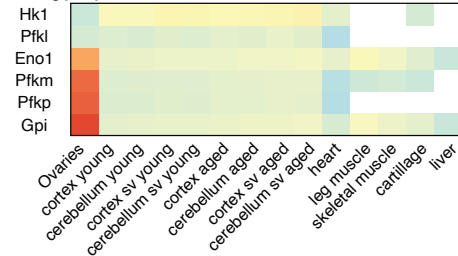
AAA+ ATPase domain



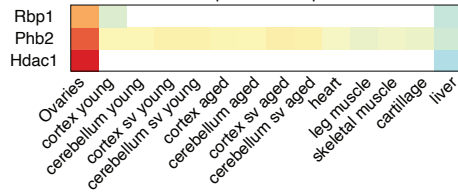
Chaperone DnaJ, and HSPH1, nucleotide-binding domain



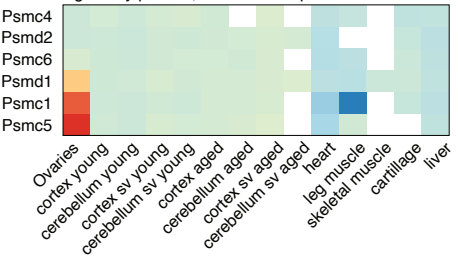
Canonical glycolysis



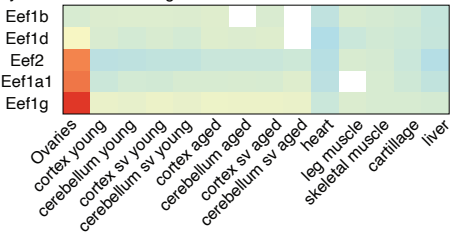
Hdac1-Ino80-Kdm1a-Phb2-Rbp1-Taf5 complex



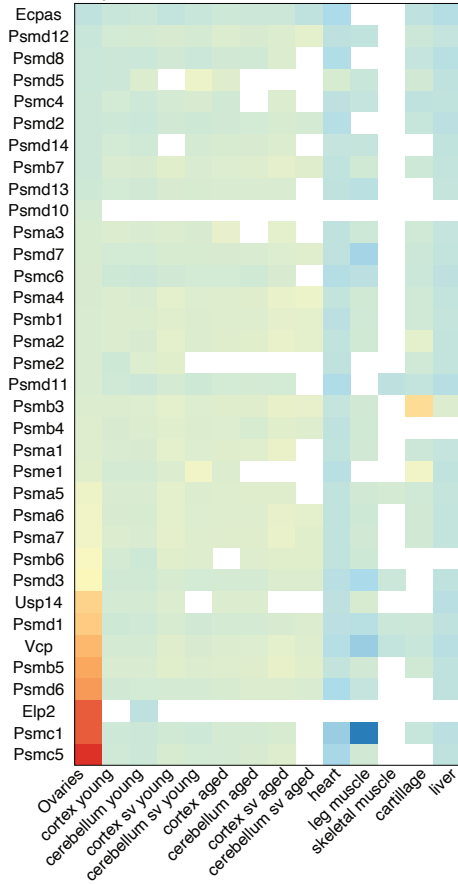
Proteasome regulatory particle, base subcomplex



Eukaryotic Translation Elongation



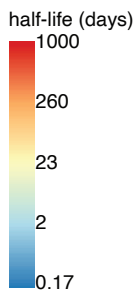
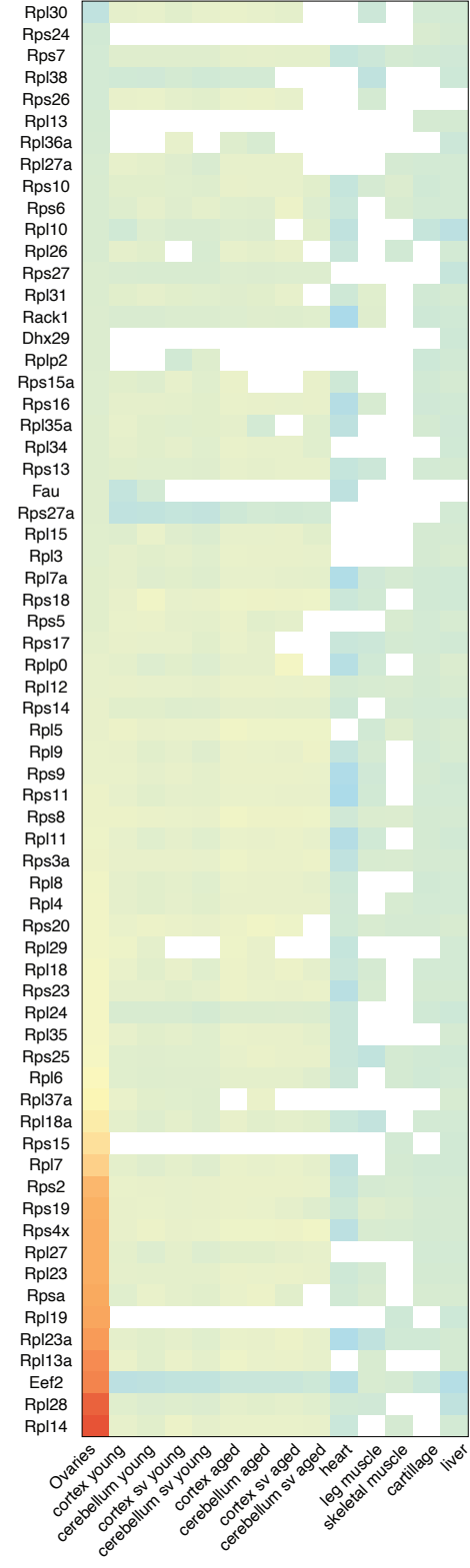
Proteasome complex



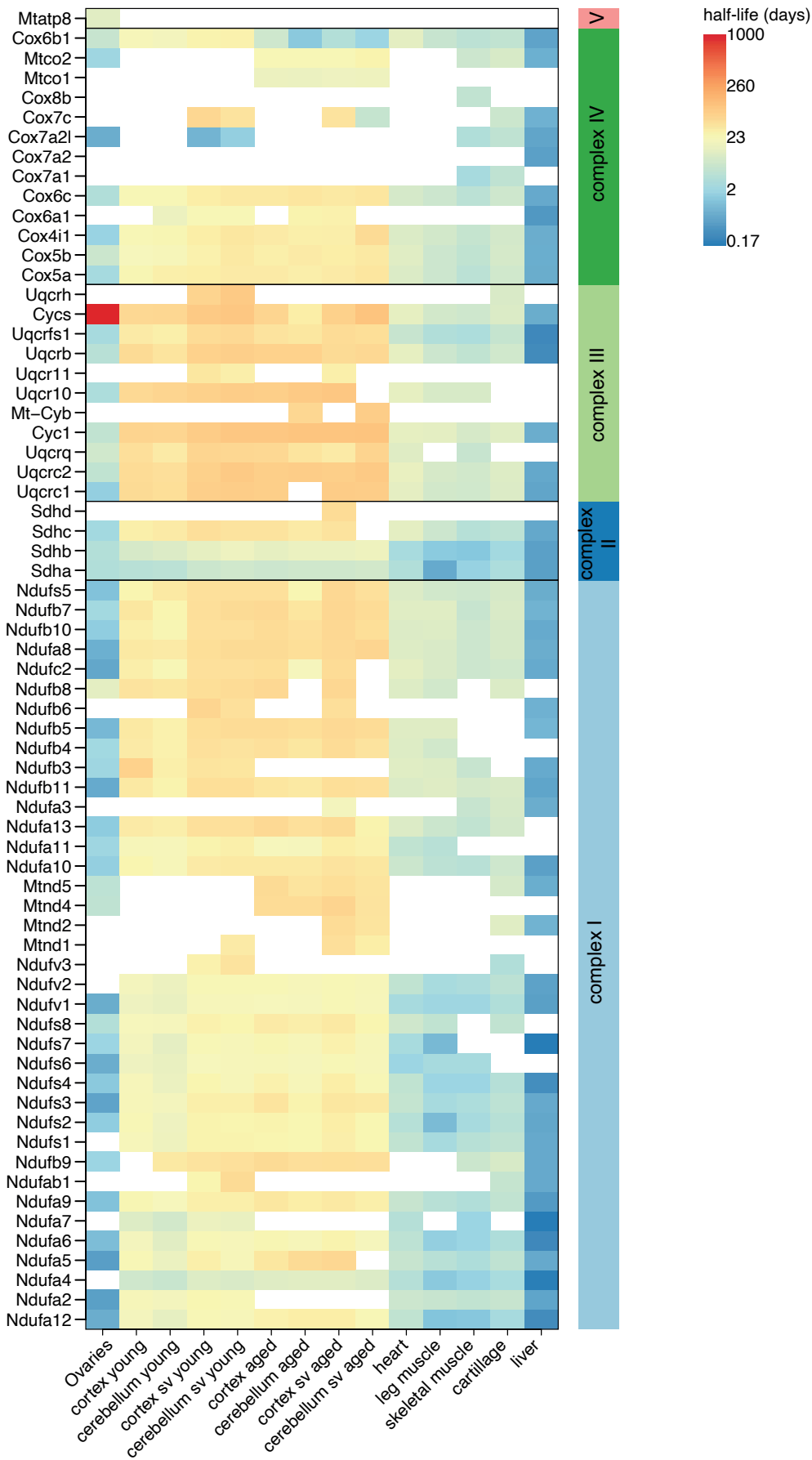
Cytosolic large ribosomal subunit



Cytosolic ribosome

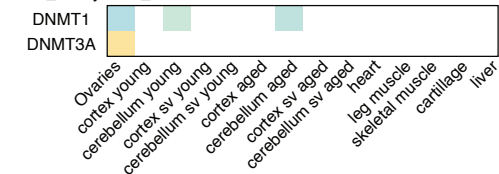


Respiratory chain complex

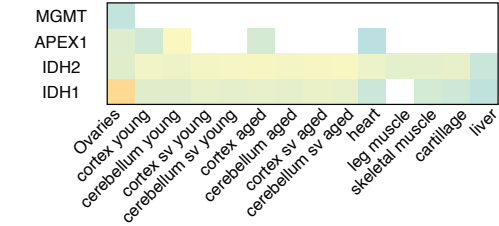


Epigenetic regulators

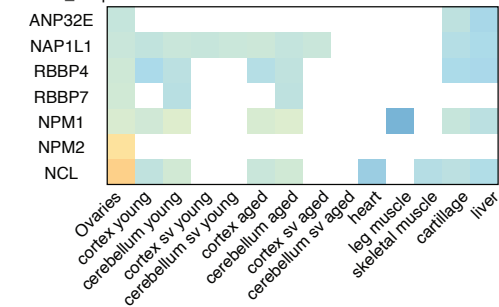
DNA_methylation_writer



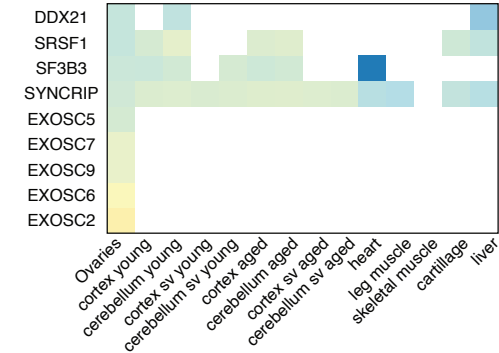
DNA_methylation_eraser



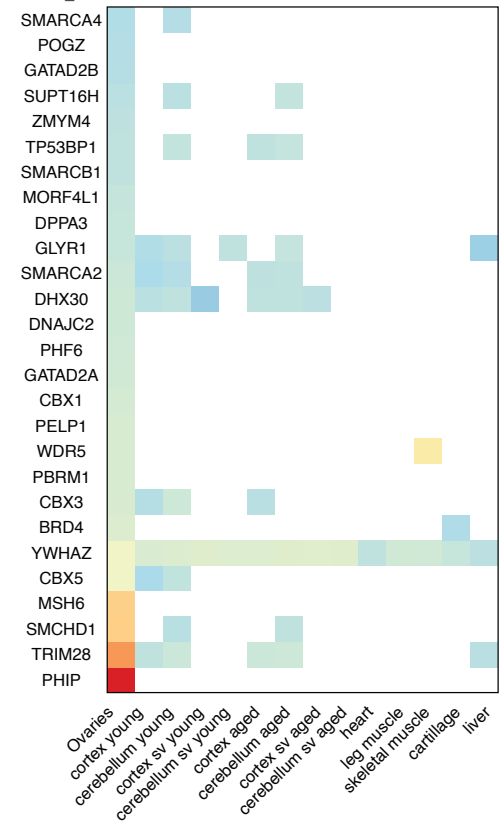
histone_chaperone



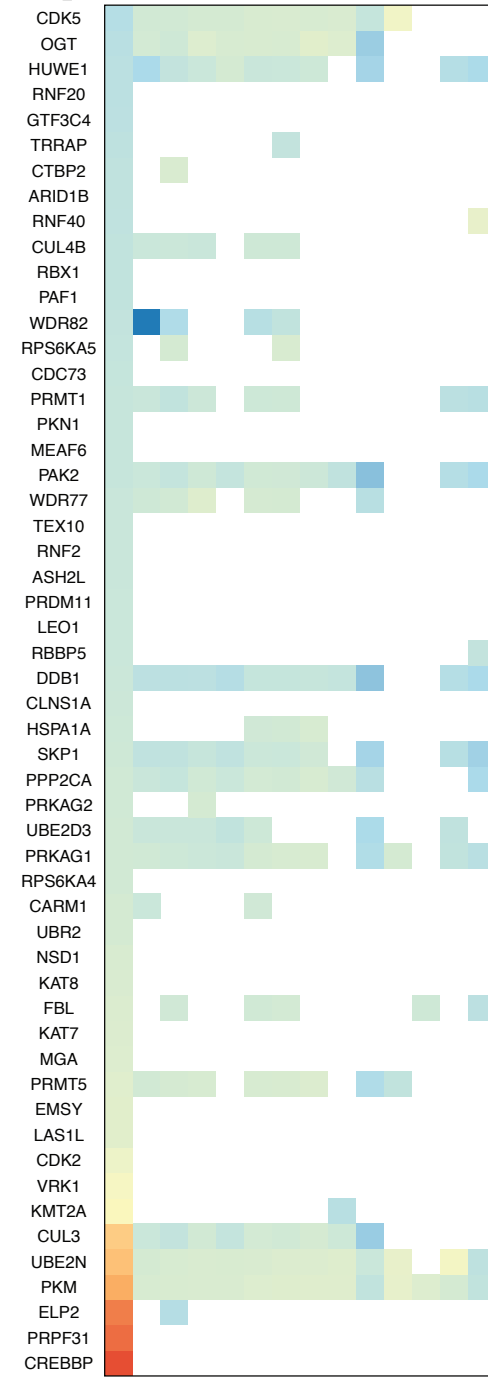
RNA_modification



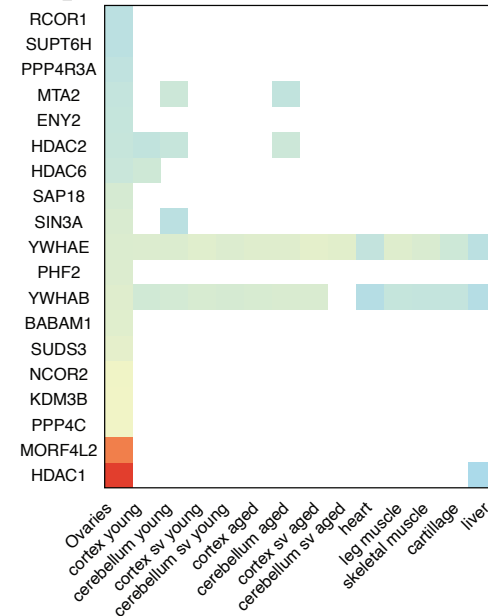
histone_reader



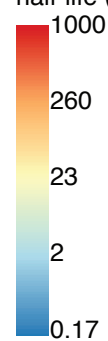
histone_writer



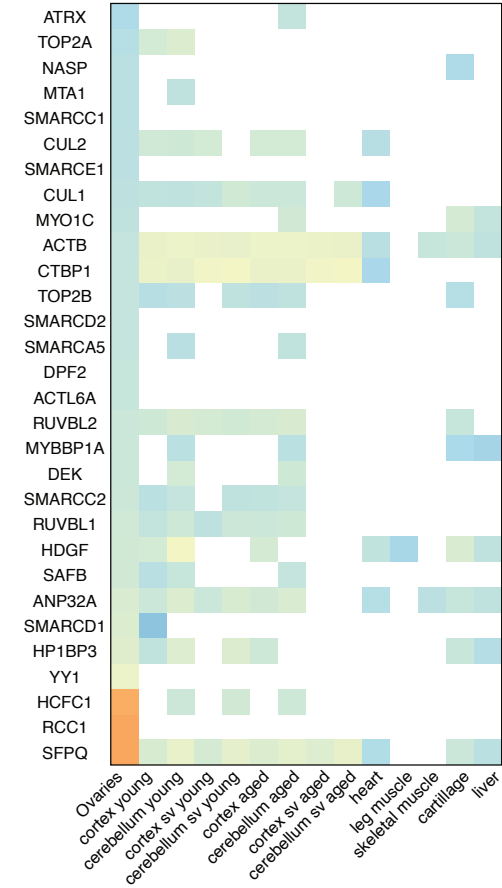
histone_eraser



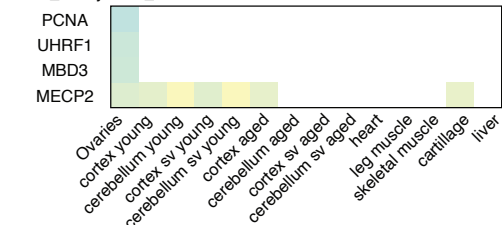
half-life (days)



chromatin_remover



DNA_methylation_reader



Chaperones

