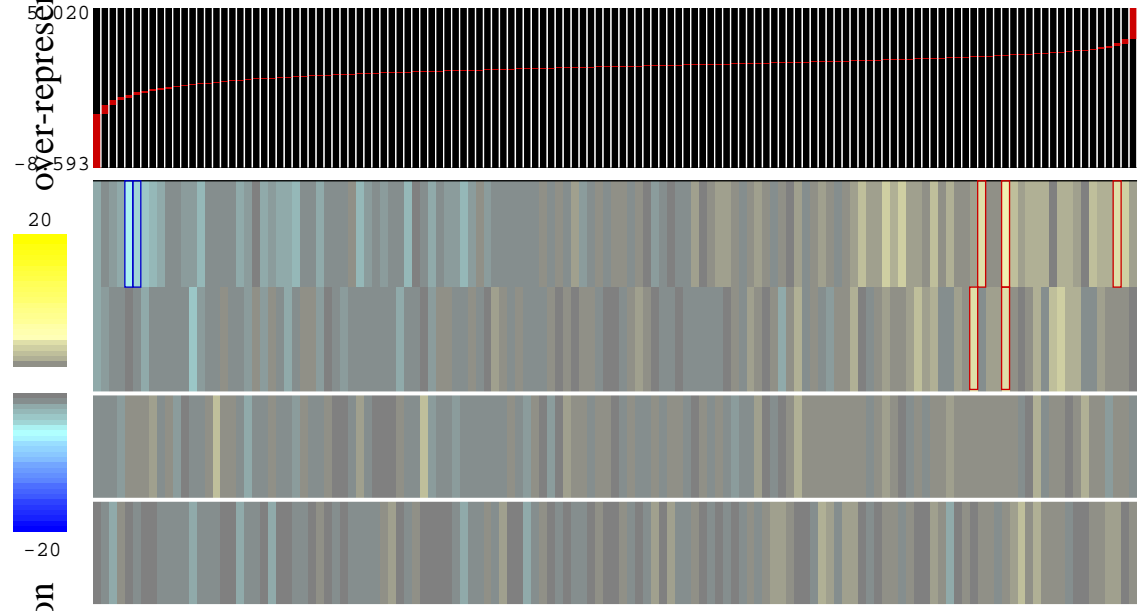


over-representation

representation



STRUCTURAL molecule activity; P<0.001

coenzyme binding, P<0.01

cytoplasmic part, P<0.01

regulation of development, P<0.01

oxidoreductase activity, P<0.01

