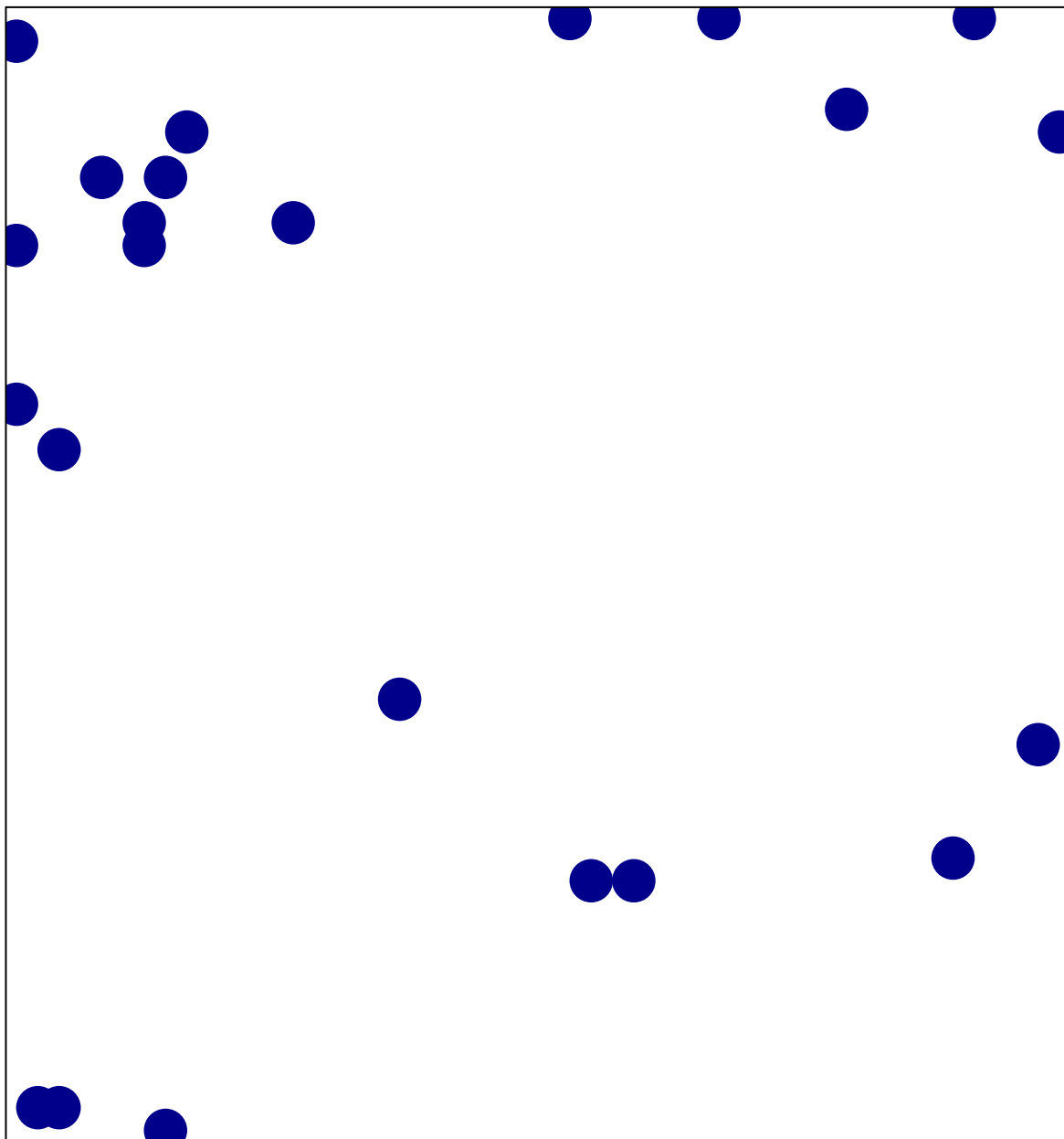


REACTOME_PROCESSING_OF_CAPPED_INTRONLESS_PRE_MRNA



features = 23
chi-square p = 0.83

SOME_PROCESSING_OF_CAPPED_INTRONLESS_PR



features = 23 , max = 1