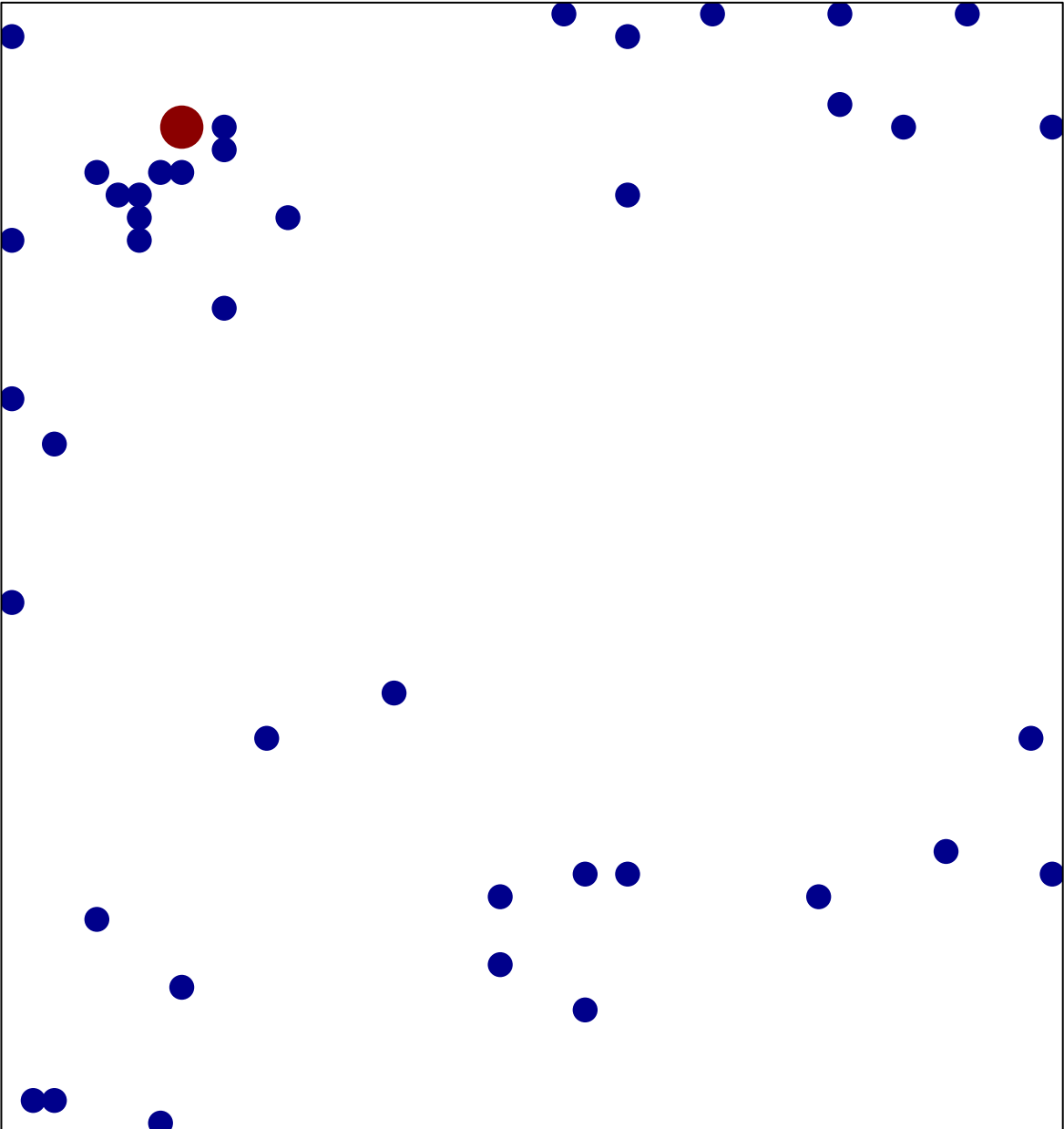


REACTOME_CLEAVAGE_OF_GROWING_TRANSCRIPT_IN_THE_TERMINATION_REGION



features = 43
chi-square p = 0.81

AVERAGE_OF_GROWING_TRANSCRIPT_IN_THE_TERM



features = 43 , max = 2