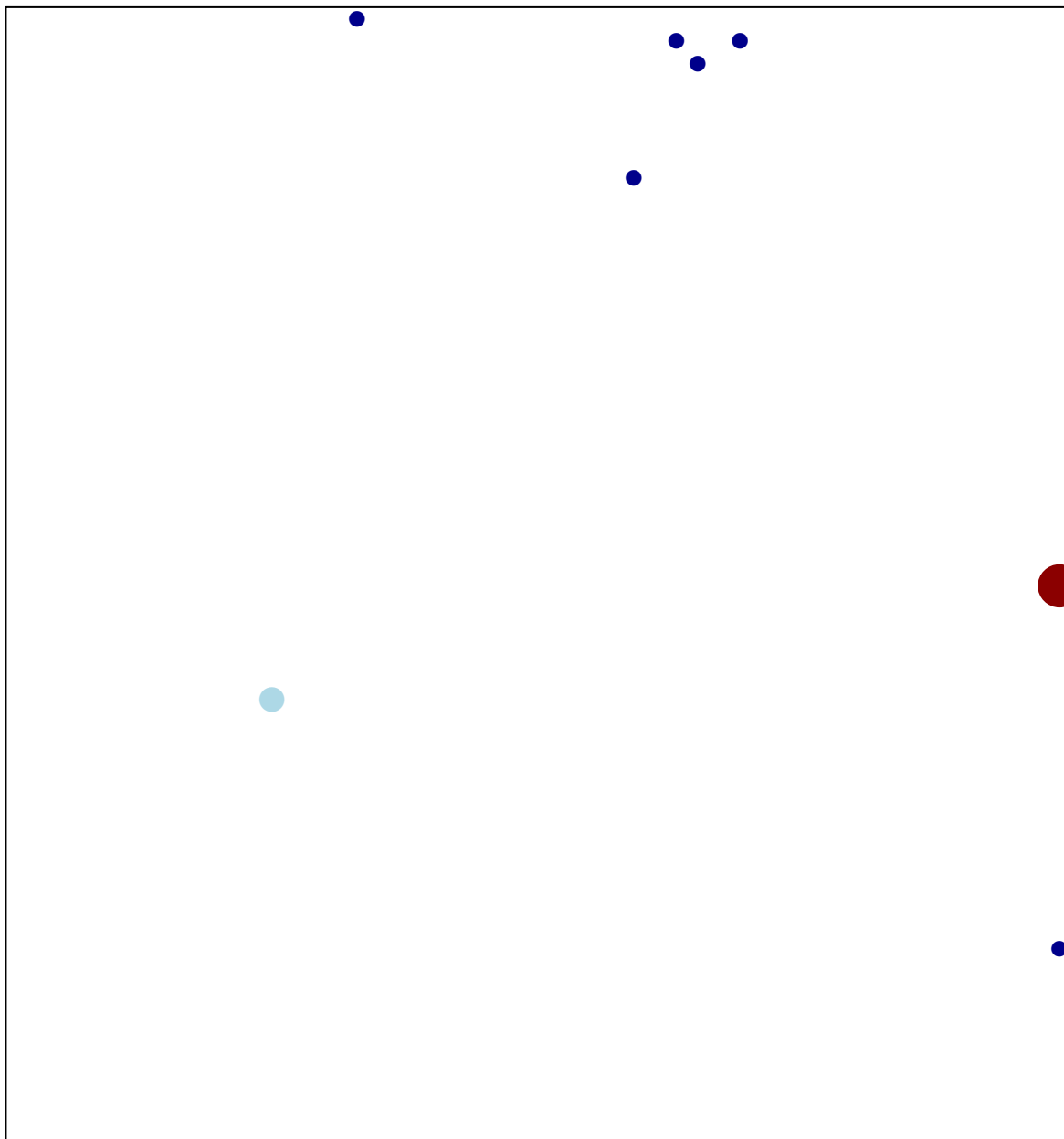


REACTOME_TRANSLATION_INITIATION_COMPLEX_FORMATION



features = 12
chi-square p = 0.76

TOME_TRANSLATION_INITIATION_COMPLEX_FORM



features = 12 , max = 4