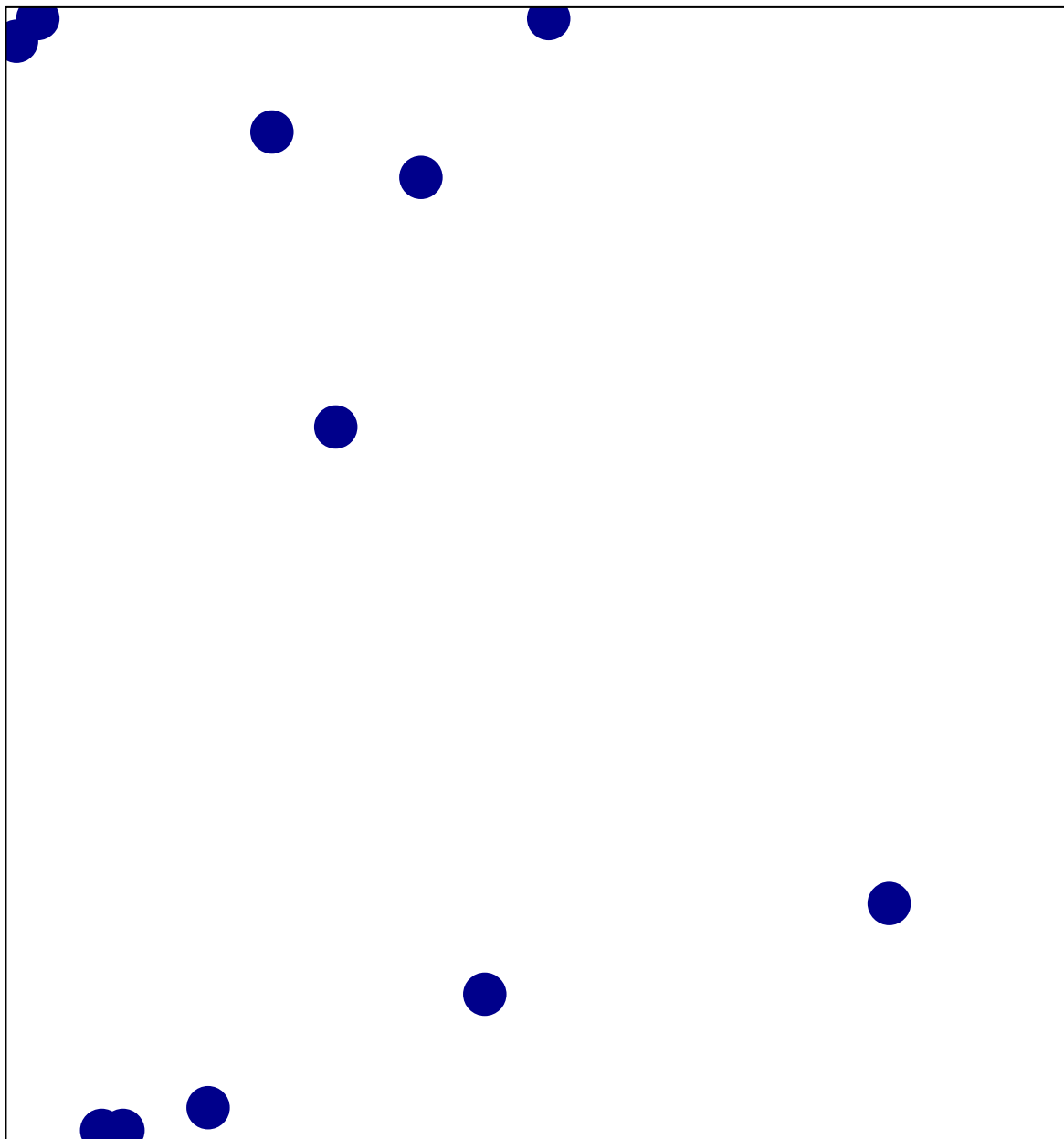


negative regulation of transcription regulatory region DNA binding



features = 11
chi-square p = 0.84

ve regulation of transcription regulatory region DNA



features = 11 , max = 1