R−SMAD binding
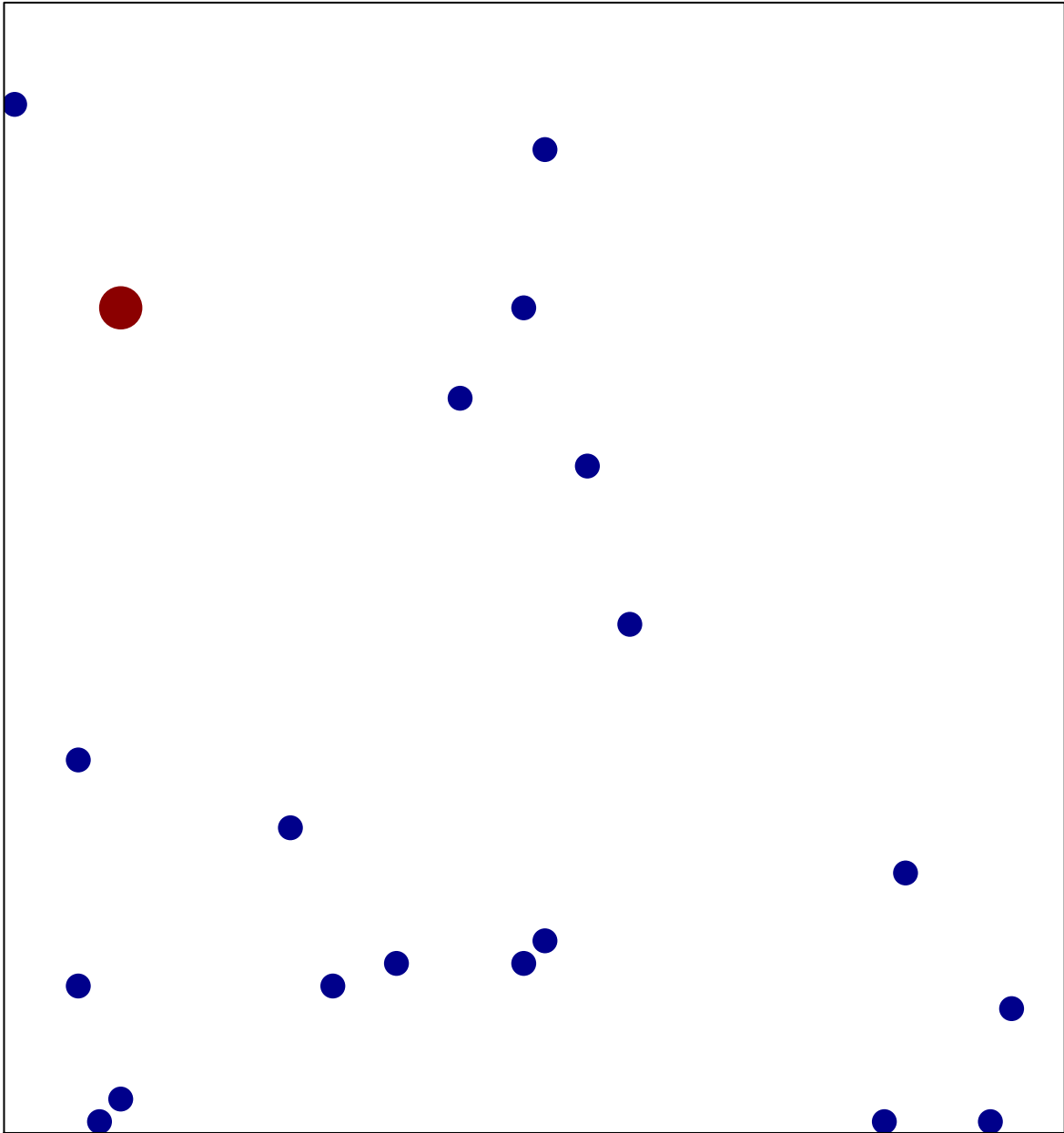
features = 21
chi−square p = 0.82

R–SMAD binding

features = 21 , max = 2