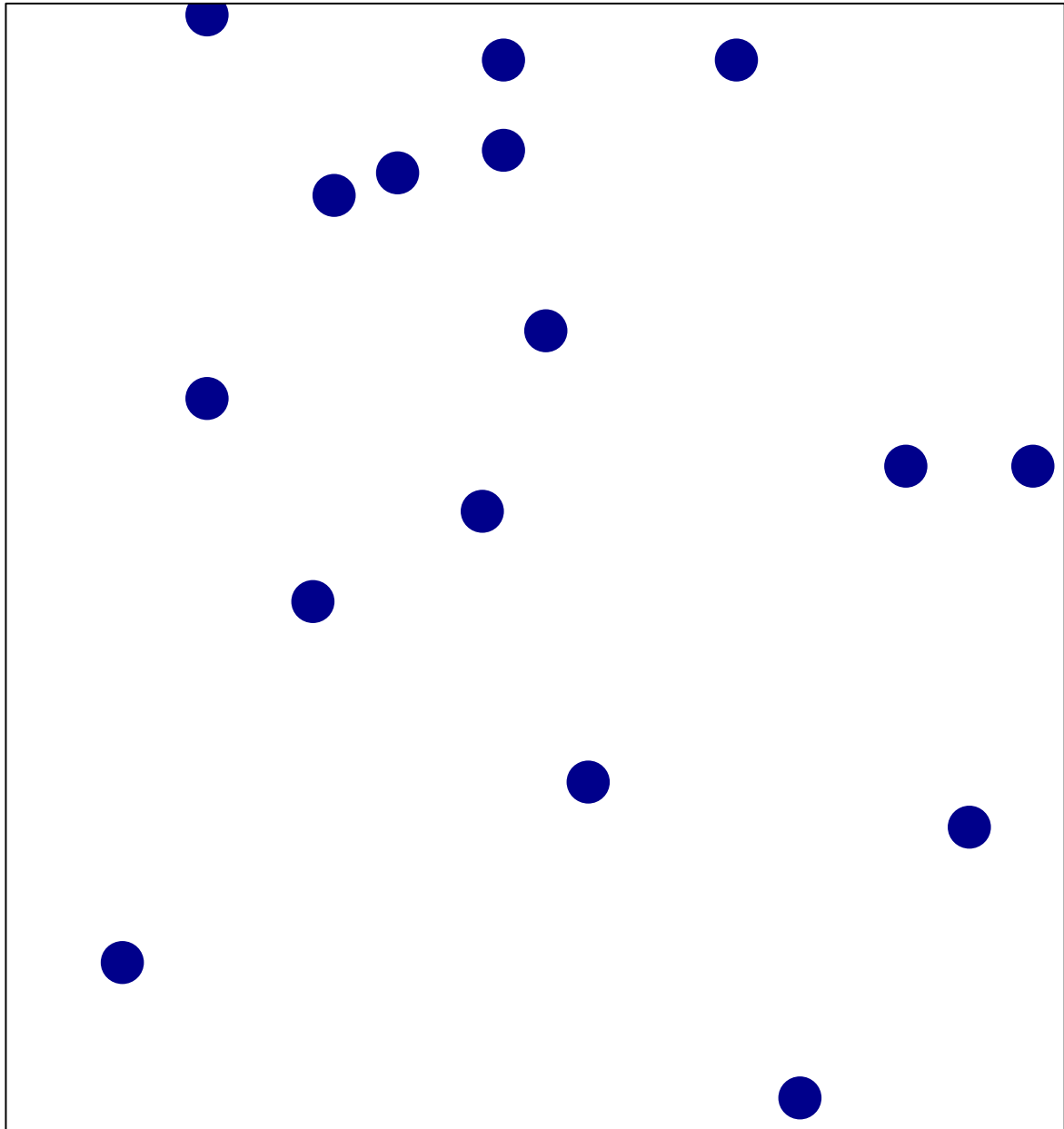
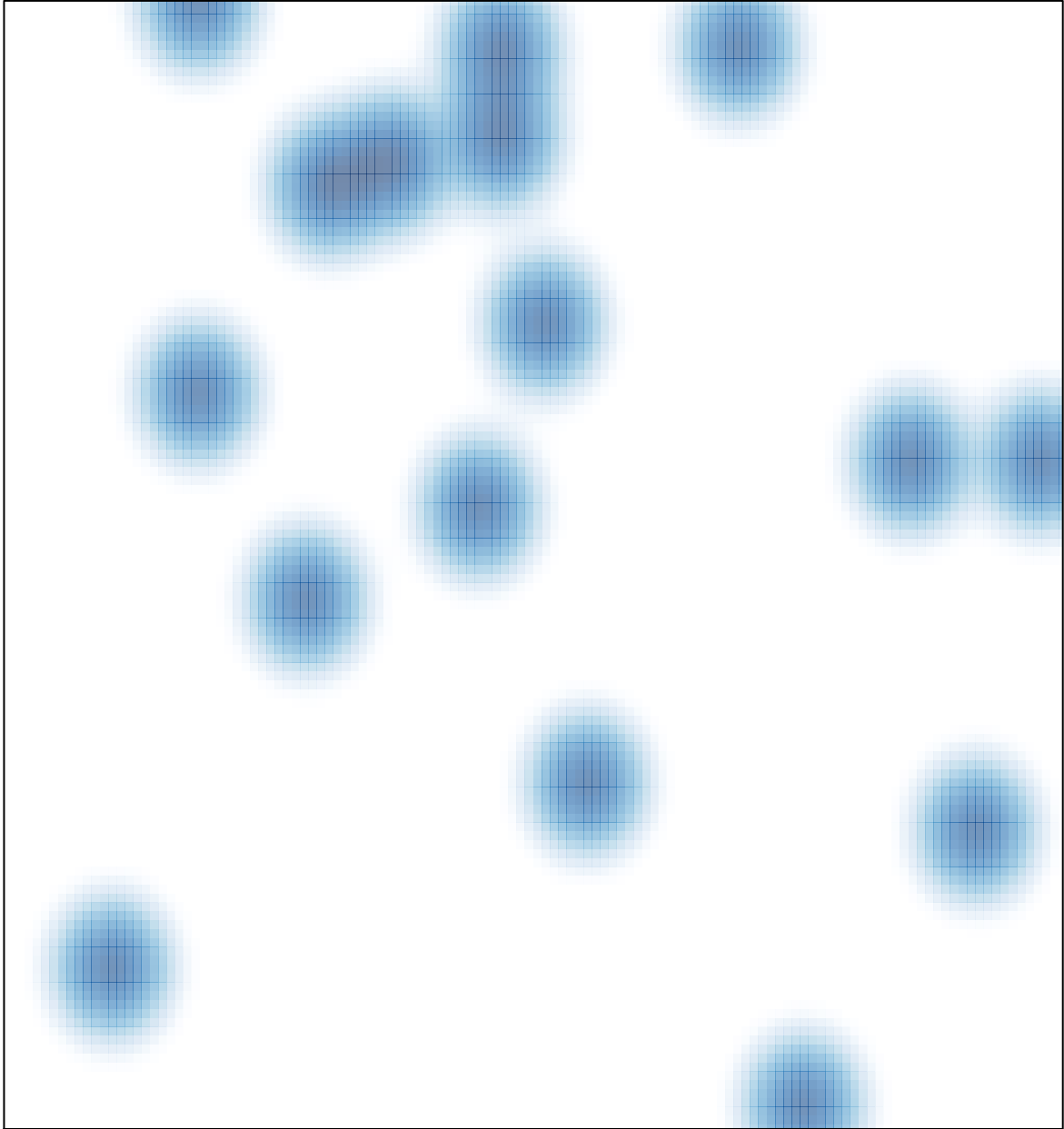


# melanosome organization



# features = 16  
chi-square p = 0.85

# melanosome organization



# features = 16 , max = 1