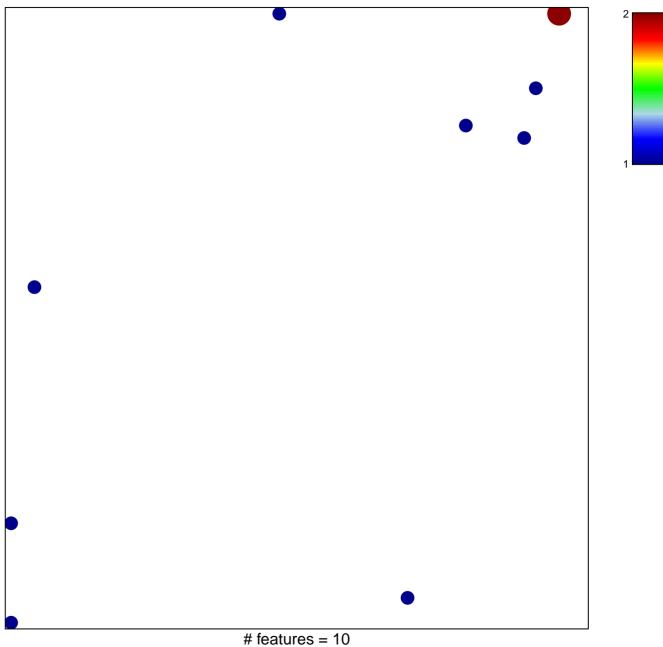
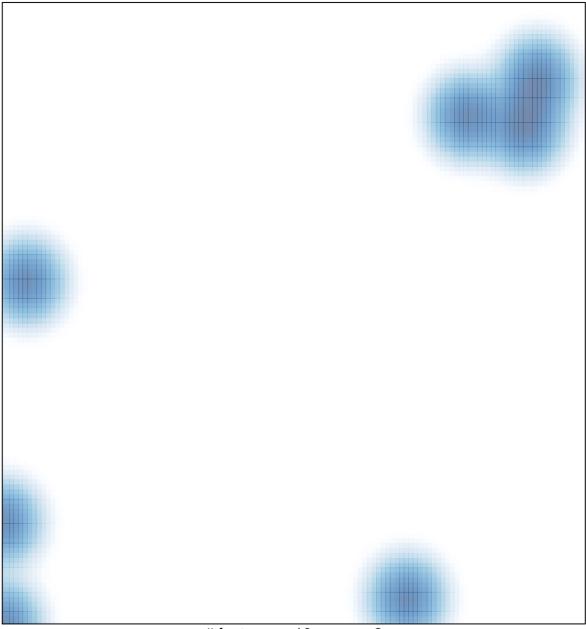
ribosomal small subunit binding



chi-square p = 0.82

## ribosomal small subunit binding



# features = 10, max = 2