

**red module  
consensus 99**

Term

Transcription regulator activity

Dna binding transcription factor activity

Sequence specific dna binding

Chromatin

Cis regulatory region sequence specific dna binding

Intermediate filament cytoskeleton

Dna binding transcription activator activity

Chromosome

Phagophore assembly site

0 1 2 3 4

$-\log_{10}(\text{FDR})$

DB

CC

MF

93

68

78

55

56

12

24

71

6