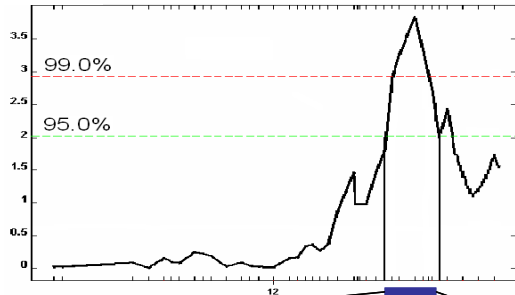


A

Location: 95% confidence

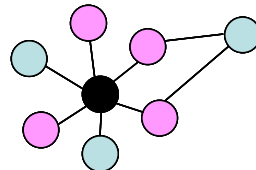


C

Function:

i. Assign putative function

Protein interactions & Networks



ii. GO enrichment

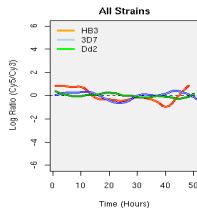
B

Gene Content:

Sequence Polymorphisms

Dd2: aatagacttca

HB3: aatagactaca

TranscriptomeExpression correlations