Imogene beta.1-77-73c9-dirty

[Run]   [Reset]

Genomewide identification of cisregulatory motifs and modules

* Execution mode [Choice ▼]

⊟ General options

Family of species to consider [Choice ▼]

Width of the motif [10]

Threshold used for motif scanning [13.0]

Extent of the motif search within an alignment [20]

⊞ Motgen options

⊞ Scangen options

References :Genomewide identification of cis regulatory motifs and modules underlying gene coregulation using statistics and phylogeny, National Academy of Sciences of the United States of America. August 17, 2010 vol. 107 no. 33 14615-14620

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