

PublisherInfo		
PublisherName	:	BioMed Central
PublisherLocation	:	London
PublisherImprintName	:	BioMed Central

Articles selected by Faculty *of 1000*: muscle differentiation regulatory network; the rules of transcriptional regulation; rapid experimental determination of DNA-binding protein motifs; polymerase poised and ready to go; bacterial tree accounting for lateral transfer

ArticleInfo		
ArticleID	:	3512
ArticleDOI	:	10.1186/gb-2005-6-6-327
ArticleCitationID	:	327
ArticleSequenceNumber	:	17
ArticleCategory	:	Paper report
ArticleFirstPage	:	1
ArticleLastPage	:	3
ArticleHistory	:	RegistrationDate : 2005-5-9

		OnlineDate : 2005-5-9
ArticleCopyright	:	BioMed Central Ltd2005
ArticleGrants	:	
ArticleContext	:	130596666

Summary

A selection of evaluations from Faculty of 1000 covering muscle differentiation regulatory network; the rules of transcriptional regulation; rapid experimental determination of DNA-binding protein motifs; polymerase poised and ready to go; bacterial tree accounting for lateral transfer.

Muscle differentiation regulatory network

An initial blueprint for myogenic differentiation. Blais A, Tsikitis M, Acosta-Alvear D, Sharan R, Kluger Y, Dynlacht BD. *Genes Dev* 2005, **19**:553-569.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-6-327.asp#Blais>

The rules of transcriptional regulation

Cis -regulatory logic of short-range transcriptional repression in *Drosophila melanogaster*. Kulkarni MM, Arnosti DN. *Mol Cell Biol* 2005, **25**:3411-3420.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-6-327.asp#Kulkarni>

Rapid experimental determination of DNA-binding protein motifs

DIP-chip: rapid and accurate determination of DNA-binding specificity. Liu X, Noll DM, Lieb JD, Clarke ND. *Genome Res* 2005, **15**:421-427.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-6-327.asp#Liu>

Polymerase poised and ready to go

Genome-wide analyses reveal RNA polymerase II located upstream of genes poised for rapid response upon *S. cerevisiae* stationary phase exit. Radonjic M, Andrau JC, Lijnzaad P, Kemmeren P, Kockelkorn TT, van Leenen D, van Berkum NL, Holstege FC. *Mol Cell* 2005, **18**:171-183.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-6-327.asp#Radonjic>

Bacterial tree accounting for lateral transfer

Evolutionary origins of genomic repertoires in bacteria. Lerat E, Daubin V, Ochman H, Moran NA. *PLoS Biol* 2005, **3**:e130.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-6-327.asp#Lerat>