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Articles selected by Faculty *of* **1000**: comparative genomics of fruit ripening; gene co-regulation during evolution; histone acetylation during plant development; microRNA gene identification; ChIP-on-chip analysis in *E. coli* 

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#### Summary

A selection of evaluations from Faculty *of* **1000** covering fruit ripening comparative genomics; gene co-regulation during evolution; histone acetylation during plant development; microRNA gene identification; ChIP-on-chip analysis in *E. coli*.

# Comparative genomics of fruit ripening

**Comprehensive EST analysis of tomato and comparative genomics of fruit ripening.** Fei Z, Tang X, Alba RM, White JA, Ronning CM, Martin GB, Tanksley SD, Giovannoni JJ. *Plant J* 2004, **40:**47-59.

For the Faculty of 1000 evaluation of this article please see: http://genomebiology.com/reports/F1000/gb-2004-5-11-356.asp#Fei

# Gene co-regulation during evolution

Gene co-regulation is highly conserved in the evolution of eukaryotes and prokaryotes. Snel B, van Noort V, Huynen MA. *Nucleic Acids Res* 2004, **32:**4725-4731.

For the Faculty of 1000 evaluation of this article please see: http://genomebiology.com/reports/F1000/gb-2004-5-11-356.asp#Snel

## Histone acetylation during plant development

**Reversible histone acetylation and deacetylation mediate genome-wide, promoter-dependent, and locus-specific changes in gene expression during plant development.** Tian L, Fong MP, Wang JJ, Wei NE, Jiang H, Doerge RW, Chen ZJ. *Genetics* 2004, September 15.

For the Faculty of 1000 evaluation of this article please see: http://genomebiology.com/reports/F1000/gb-2004-5-11-356.asp#Tian

### MicroRNA gene identification

Patterns of flanking sequence conservation and a characteristic upstream motif for microRNA gene identifications. Ohler U, Yekta S, Lim LP, Bartel DP, Burge CB. *RNA* 2004, **10**:1309-1322.

For the Faculty of 1000 evaluation of this article please see: http://genomebiology.com/reports/F1000/gb-2004-5-11-356.asp#Ohler

#### ChIP-on-chip analysis in E. coli

Genomic studies with *Escherichia coli* MelR protein: applications of chromatin immunoprecipitation and microarrays. Grainger DC, Overton TW, Reppas N, Wade JT, Tamai E, Hobman JL, Constantinidou C, Struhl K, Church G, Busby SJ. *J Bacteriol* 2004, **186**:6938-6943.

For the Faculty of 1000 evaluation of this article please see: http://genomebiology.com/reports/F1000/gb-2004-5-11-356.asp#Grainger