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Articles selected by Faculty *of 1000*: linkage disequilibrium across the human genome; identifying low abundance proteins; bacterial phylogenetics; *trans*-acting regulatory variation in yeast; promoter detection in *Arabidopsis*

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Summary

A selection of evaluations from Faculty of **1000** covering the distribution of linkage disequilibrium across the human genome, identifying low abundance and membrane proteins, bacterial phylogenetics, *trans*-acting regulatory variation in yeast and promoter detection in *Arabidopsis*.

Linkage disequilibrium across the human genome

Linkage disequilibrium and inference of ancestral recombination in 538 single-nucleotide polymorphism clusters across the human genome. Clark AG, Nielsen R, Signorovitch J, Matise TC, Glanowski S, Heil J, Winn-Deen ES, Holden AL, Lai E. *Am J Hum Genet* 2003, **73**:285-300.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2003-4-9-339.asp#Clark>

Identifying low abundance proteins

Unseen proteome: mining below the tip of the iceberg to find low abundance and membrane proteins. Pedersen SK, Harry JL, Sebastian L, Baker J, Traini MD, McCarthy JT, Manoharan A, Wilkins MR, Gooley AA, Righetti PG, *et al.* *J Proteome Res* 2003, **2**:303-311.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2003-4-9-339.asp#Pedersen>

Bacterial phylogenetics

Phylogenetics and the cohesion of bacterial genomes. Daubin V, Moran NA, Ochman H. *Science* 2003, **301**:829-832.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2003-4-9-339.asp#Daubin>

Trans-acting regulatory variation in yeast

***Trans* -acting regulatory variation in *Saccharomyces cerevisiae* and the role of transcription factors.** Yvert G, Brem RB, Whittle J, Akey JM, Foss E, Smith EN, Mackelprang R, Kruglyak L. *Nat Genet* 2003, Aug 3.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2003-4-9-339.asp#Yvert>

Promoter detection in *Arabidopsis*

Computational approaches to identify promoters and *cis* -regulatory elements in plant genomes. Rombauts S, Florquin K, Lescot M, Marchal K, Rouzé P, van de Peer Y. *Plant Physiol* 2003, **132**:1162-1176.

For the Faculty of 1000 evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2003-4-9-339.asp#Rombauts>