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Articles selected by Faculty of 1000 Biology: genetically identical SNPs; detailed histone modification mapping; plant gene-expression diversity; photosynthesis gene evolution; ϵ -Proteobacteria diversity.

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Summary

A selection of evaluations from Faculty of **1000** Biology Genetically identical SNPs; detailed histone modification mapping; plant gene-expression diversity; photosynthesis gene evolution; ε-Proteobacteria diversity.

Genetically identical SNPs

Genetically indistinguishable SNPs and their influence on inferring the location of disease-associated variants. Lawrence R, Evans DM, Morris AP, Ke X, Hunt S, Paolucci M, Ragoussis J, Deloukas P, Bentley D, Cardon LR. *Genome Res* 2005, **15**:1503-1510.

For the Faculty of 1000 Biology evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-12-362.asp#Lawrence>

Detailed histone modification mapping

Single-nucleosome mapping of histone modifications in *S. cerevisiae*. Liu CL, Kaplan T, Kim M, Buratowski S, Schreiber SL, Friedman N, Rando OJ. *PLoS Biol* 2005, **3**:e328.

For the Faculty of 1000 Biology evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-12-362.asp#Liu>

Plant gene-expression diversity

Genomic survey of gene expression diversity in *Arabidopsis thaliana*. Kliebenstein DJ, West MA, van Leeuwen H, Kim K, Doerge RW, Michelmore RW, St Clair DA. *Genetics* 2005, October 3.

For the Faculty of 1000 Biology evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-12-362.asp#Kliebenstein>

Photosynthesis gene evolutione

Potential photosynthesis gene recombination between *Prochlorococcus* and *Synechococcus* via viral intermediates. Zeidner G, Bielawski JP, Shmoish M, Scanlan DJ, Sabehi G, Béjà O. *Environ Microbiol* 2005, 7:1505-1513.

For the Faculty of 1000 Biology evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-12-362.asp#Zeidner>

ε -Proteobacteria diversity

Distribution, phylogenetic diversity and physiological characteristics of ε -Proteobacteria in a deep-sea hydrothermal field. Nakagawa S, Takai K, Inagaki F, Hirayama H, Nunoura T, Horikoshi K, Sako Y. *Environ Microbiol* 2005, 7:1619-1632.

For the Faculty of 1000 Biology evaluation of this article please see: <http://genomebiology.com/reports/F1000/gb-2005-6-12-362.asp#Nakagawa>