**Access and citation statistics, current as of June 2011**

* ***A novel sensitive microarray approach for differential screening using probes labeled with two different radio-elements*** by Salin H, Vujasinovic T, Mazurie A, Maitrejean S, Menini C, Mallet J and Dumas S. Published in *Nucleic Acids Research*, Feb 2002; 30(4):e17
* Number of accesses (downloads from the journal’s website): data not available
* Number of citations in scholarly journals publications: **22** (sources: ISI Web of Knowledge[[1]](#footnote-1) and CrossRef[[2]](#footnote-2)); complete list in appendix A (p. 3).
* ***Gene networks inference using dynamic bayesian networks*** by Perrin BE, Ralaivola L, Mazurie A, Bottani S, Mallet J and d'Alché-Buc F. Published in *Bioinformatics*, Oct 2003; 19(2):II138-II148
* Number of accesses (downloads from the journal’s website): data not available
* Number of citations in scholarly journals’ publications: **128** (source: Google Scholar[[3]](#footnote-3)); complete list in appendix B (p. 5).
* ***An evolutionary and functional assessment of regulatory network motifs*** by Mazurie A, Bottani S and Vergassola M. Published in *Genome Biology*, Mar 2005; 6(4):R35
* Number of accesses (downloads from the journal’s website): **6961** (source: BioMed Central[[4]](#footnote-4))
* Number of citations in scholarly journals’ publications: **49** (sources: Google Scholar[[5]](#footnote-5), ISI Web of Knowledge[[6]](#footnote-6), PubMed[[7]](#footnote-7)); complete list in appendix C (p. 16).
* Tagged as ‘**must read**’ by the Faculty of 1000[[8]](#footnote-8)
* ***Integrating genome-scale data for gene essentiality prediction*** by Roberts SB, Mazurie A and Buck GA. Published in *Chemistry and Biodiversity*, 2007; 4(11):2618-2630
* Number of accesses (downloads from the journal’s website): data not available
* Number of citations in scholarly journals’ publications: **2** (source: Google Scholar[[9]](#footnote-9)); complete list in appendix D (p. 20).
* ***Phylogenetic distances are encoded in networks of interacting pathways*** by Mazurie A, Bonchev D, Schwikowski B and Buck GA. Published in *Bioinformatics*, Nov 2008; 24(22):2579-2585
* Number of accesses (downloads from the journal’s website): **1698** (source: Bioinformatics[[10]](#footnote-10))
* Number of citations in scholarly journals’ publications: **16** (sources: ISI Web of Knowledge[[11]](#footnote-11) and Google Scholar[[12]](#footnote-12)); complete list in appendix E (p. 21).
* ***Path-a-Way: a strategy for network analysis of microarray data*** by Arasappan D, Mazurie A, Alves J, Bonchev D and Buck GA. Presented at *International Conference on BioMedical Engineering and Informatics* and published in proceedings, p. 432-436 (2008)
* Number of accesses (downloads from the journal’s website): data not available
* Number of citations in scholarly journals’ publications: 0.
* ***Evolution of metabolic network organization*** by Mazurie A, Bonchev D, Schwikowski B and Buck GA. Published in *BMC Systems Biology*, May 2010; 4(1):59
* Number of accesses (downloads from the journal’s website): **2670** (source: BioMed Central[[13]](#footnote-13))
* Number of citations in scholarly journals’ publications: **3** (sources: ISI Web of Knowledge[[14]](#footnote-14) and Google Scholar[[15]](#footnote-15)); complete list in appendix F (p. 23)
* Tagged as ‘**Highly accessed**’ (source: BMC Systems Biology[[16]](#footnote-16))
* ***Physiology of Pseudomonas aeruginosa in biofilms as revealed by transcriptome analysis*** by Folsom JP, Richards L, Pitts B, Roe F, Ehrlich GD, Parker A, Mazurie A and Stewart PS. Published in *BMC Microbiology*, Nov 2010; 10:294
* Number of accesses (downloads from the journal’s website): **3008** (source: BioMed Central[[17]](#footnote-17))
* Number of citations in scholarly journals’ publications: 0.
* Tagged as ‘**Highly accessed**’ (source: BMC Microbiology[[18]](#footnote-18))

**Note**: the two latest publications in BMC Systems Biology and BMC Microbiology are too recent (2010) to have been significantly cited in any other publication, due to the average length for a publication to be peer-reviewed and published (about a year). However, their tagging as ‘Highly accessed’ by the journal’s website demonstrate the interest shown by scientists in the field for the content of this work.

Appendix A**.** Publications in scholarly journals citing the manuscript ‘*A novel sensitive microarray approach for differential screening using probes labeled with two different radio-elements’*

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Appendix D. Publications in scholarly journals citing the manuscript ‘*Integrating genome-scale data for gene essentiality prediction’*

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**Appendix E**. Publications in scholarly journals citing the manuscript ‘[*Phylogenetic distances are encoded in networks of interacting pathways*](http://apps.isiknowledge.com/CitedFullRecord.do?product=WOS&SID=3F1c568fnp43F@n1pBc&search_mode=CitedFullRecord&isickref=174670337&db_id=WOS&colname=WOS)’

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**Appendix F**. Publications in scholarly journals citing the manuscript ‘*Evolution of metabolic network organization*’

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7. http://goo.gl/swssY [↑](#footnote-ref-7)
8. See document #5 provided as initial evidence under section 2.A.3 [↑](#footnote-ref-8)
9. http://goo.gl/GpzMs [↑](#footnote-ref-9)
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16. http://goo.gl/DxFsq [↑](#footnote-ref-16)
17. http://goo.gl/O4Lmh [↑](#footnote-ref-17)
18. http://goo.gl/clzxr [↑](#footnote-ref-18)