

2010 : April 2010 - Fast Breaking Papers : Patrik Nosil on Population Genomics

fast breaking papers - 2010

April 2010



Patrik Nosil talks with *ScienceWatch.com* and answers a few questions about this month's Fast Breaking Paper Paper in the field of Environment & Ecology.



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Article Title: Divergent selection and heterogeneous genomic divergence

Authors: **Nosil, P**; Funk, DJ; Ortiz-Barrientos, D

Journal: MOL ECOL, Volume: 18, Issue: 3, Page: 375-402, Year: FEB 2009

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SW: Why do you think your paper is highly cited?

This article is likely highly cited because it provides both a conceptual and empirical review of a newly emerged field of study—"population genomics."

SW: Would you summarize the significance of your paper in layman's terms?

The article synthesizes our knowledge concerning how natural selection impacts the genome, and especially how natural selection creates differences in levels of genetic divergence across the genome.

SW: How did you become involved in this research, and were there any problems along the way?

This review article emerged from empirical "population genomic" studies conducted in two groups of insects. In both studies, my coauthors, Daniel Funk and Scott Egan of Vanderbilt University, and I identified genomic regions subject to natural selection. These empirical findings naturally then led into a review of the overall literature on genomic divergence.

SW: Where do you see your research leading in the future?

Recent advances in molecular biology ("next-generation sequencing") and computational approaches

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will allow the effects of natural selection on the genome to be examined in more precise, as well as new and creative, ways.

SW: Do you foresee any social or political implications for your research?

Understanding how natural selection affects the genome might increase our understanding of the origin and maintenance of genetic diversity.

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KEYWORDS: WHITEFISH COREGONUS-CLUPEAFORMIS; ADAPTIVE POPULATION DIVERGENCE; QUANTITATIVE TRAIT LOCI; BEBBIANAE LEAF BEETLES; DETECT CANDIDATE LOCI; HOST-PLANT ADAPTATION; DNA-SEQUENCE DATA; GENE FLOW; NATURAL-SELECTION; REPRODUCTIVE ISOLATION.

Additional information:

[Scott Eagan](#) | [Daniel Funk](#)

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