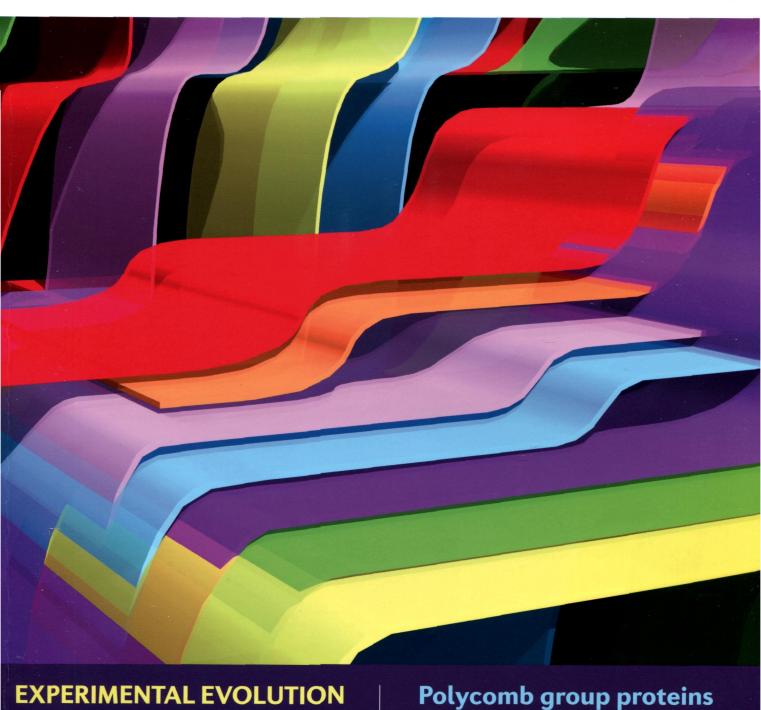


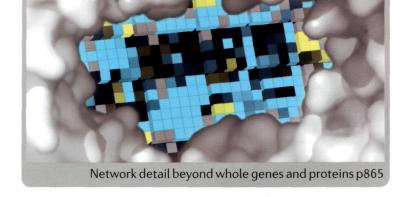
Fresh insights into genome dynamics

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GENETICS



Complexity of complexes



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Jeffrey E. Barrick and Richard E. Lenski This Review describes how whole-genome sequencing of experimental populations of microorganisms and other species is yielding new insights into evolutionary processes. These details include variation in mutation rates, diverse genetic routes to adaptation and dynamic interactions within and between populations.

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Evolution of crop species: genetics of domestication and diversification

Rachel S. Meyer and Michael D. Purugganan The recent improvement in technologies to identify genetic variants linked with quantitative traits has allowed the identification of variants that are associated with crop domestication. This Review considers these approaches and their application to the study of crop domestication.

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A new world of Polycombs: unexpected partnerships and emerging functions

Yuri B. Schwartz and Vincenzo Pirrotta Polycomb group (PcG) proteins have long been known to have repressive functions. Recently, novel complexes of PcG proteins have been identified and have been shown to have novel molecular and biological functions, which are discussed in this Review.

High-resolution network biology: connecting sequence with function

Colm J. Ryan, Peter Cimermančič, Zachary A. Szpiech, Andrei Sali, Rvan D. Hernandez and Nevan I. Krogan Network analyses of protein-protein, genetic and drug-gene interactions have typically been limited to analyses at the whole-protein and whole-gene levels. This Review discusses how high-resolution network analyses at the domain and residue levels are providing valuable insights into diverse research fields that range from structural biology to human disease.

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Gene regulation by antisense transcription Vicent Pelechano and Lars M. Steinmetz

Antisense transcription is increasingly being recognized as an important regulator of gene expression across all kingdoms of life and through a range of regulatory modes. Antisense transcripts are also emerging as facilitators of rapid evolution of gene regulation.



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