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Advances in the development of genome engineering tools — such as zinc finger proteins, TALEs, and more recently the CRISPR/Cas9 system — are allowing us to study the genomes of diverse organisms at the functional level with unprecedented ease and accuracy. This collection of 20 articles highlights this young, dynamic field with a special issue covering the applications of genome editing technologies in all kingdoms of life, both wet-lab and computational methodological advances, novel insights into the biology of naturally occurring CRISPR/Cas systems, discussions on the challenges and opportunities of these new techniques, and more.

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