NEW APPLICATIONS

DEVELOPING COST-EFFECTIVE GENOMIC TOOLS TO TRACK AND UNDERSTAND FUNGAL CROP DISEASES

DISCOVERY OF NEW VIRUSES ENABLED BY PROXIMITY LIGATION TECHNOLOGY AND NEW FUNDING

NEW FEATURES
NEW PUBLICATIONS

Differential Expression of α, β, and γ Protocadherin Isoforms During Differentiation, Aging, and Cancer

Genome sequence of Monilinia vaccinii-corymbosi sheds light on mummy berry disease infection of blueberry and mating type
Gene flux and acid-imposed selection are the main drivers of antimicrobial resistance in broiler chicks infected with *Salmonella*.

Chromosome-scale genome assembly of the sea louse *Caligus rogercresseyi* by SMRT sequencing and Hi-C analysis.

Improving Spirodela polyrhiza genome and proteomic analyses reveal a conserved chromosomal structure.

Chromosome-level genome assembly of Ophiirhiza pumila reveals the evolution of camptothecin biosynthesis.

Tracking Reservoirs of Antimicrobial Resistance Genes in a Complex Microbial Community Using Metagenomic Hi-C.

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