Motif-based method for genome-wide prediction of eukaryotic gene clusters

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Details

Beteiligte Forschungseinheiten

Microbiome Dynamics Gianni Panagiotou Mehr erfahren

International Leibniz Research School Peter F. Zipfel Mehr erfahren

Leibniz-HKI-Autor*innen



Ekaterina Shelest

Details



Vladimir Shelest

Details



Thomas Wolf

Details

Themenfelder

Vorhersage genregulatorischer Elemente und Gencluster in Pilzen