

# CURRENT GENOMICS

**Guest Editor:**

Current Genomics, Volume 21, Number 1, 2020

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## Computational RNA Epigenetics

### CONTRIBUTORS

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***A Brief Survey for MicroRNA Precursor Identification Using Machine Learning Methods***  
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***Extremely-randomized-tree-based Prediction of N6-methyladenosine Sites in Saccharomyces cerevisiae***  
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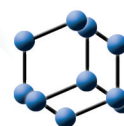
[www.benthamscience.com/journals/cg](http://www.benthamscience.com/journals/cg)

ISSN: 1875-5488 (Online)  
ISSN: 1389-2029 (Print)



**Editor-in-Chief:**  
Current Genomics

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