Identifying possibly novel sources of antimicrobial resistance in uncultivated bacterial and archeal metagenome-assembled genomes

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Authors

- Jackie Purdue
 - © 0000-0003-3131-0688 · jackiepurdue · У jackiepurdue

 Department of Computer Science, Dalhousie University · Funded by Grant XXXXXXXXX
- Jane Roe

Department of Something, University of Whatever; Department of Whatever, University of Something

References