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Recent Work

Title

MicroPheno: predicting environments and host phenotypes from 16S rRNA gene sequencing using a k-mer based representation of shallow sub-samples.

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Authors

Asgari, Ehsaneddin
Garakani, Kiavash
McHardy, Alice C
et al.

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Corrigendum

MicroPheno: predicting environments and host phenotypes from 16S rRNA gene sequencing using a k-mer based representation of shallow sub-samples

Ehsaneddin Asgari^{1,2}, Kiavash Garakani¹, Alice C. McHardy²,
Mohammad R.K. Mofrad^{1,3,*}

¹Molecular Cell Biomechanics Laboratory, Departments of Bioengineering and Mechanical Engineering, University of California, Berkeley, CA, 94720, USA, ²Computational Biology of Infection Research, Helmholtz Center for Infection Research, 38124 Braunschweig, Germany and ³Molecular Biophysics and Integrated Bioimaging, Lawrence Berkeley National Lab, Berkeley, CA, 94720, USA

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The author wishes to inform readers that the affiliations for the authors were incorrect in the original paper. They appear correctly above.

The paper has been corrected online.