*Muribaculaceae* genomes assembled from metagenomes suggest genetic drivers of differential response to acarbose treatment in mice

Byron J. Smitha,\*

Richard A. Millerb

Thomas M. Schmidta,c,#

aDepartment of Ecology & Evolutionary Biology, University of Michigan, Ann Arbor, MI, USA.

bDepartment of Pathology and Geriatrics Center, University of Michigan, Ann Arbor, MI, USA.

cDepartment of Internal Medicine, University of Michigan, Ann Arbor, MI, USA.

\*Present address: J. David Gladstone Institutes, San Francisco, CA, USA.

#Correspondence: [schmidti@umich.edu](mailto:schmidti@umich.edu)

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