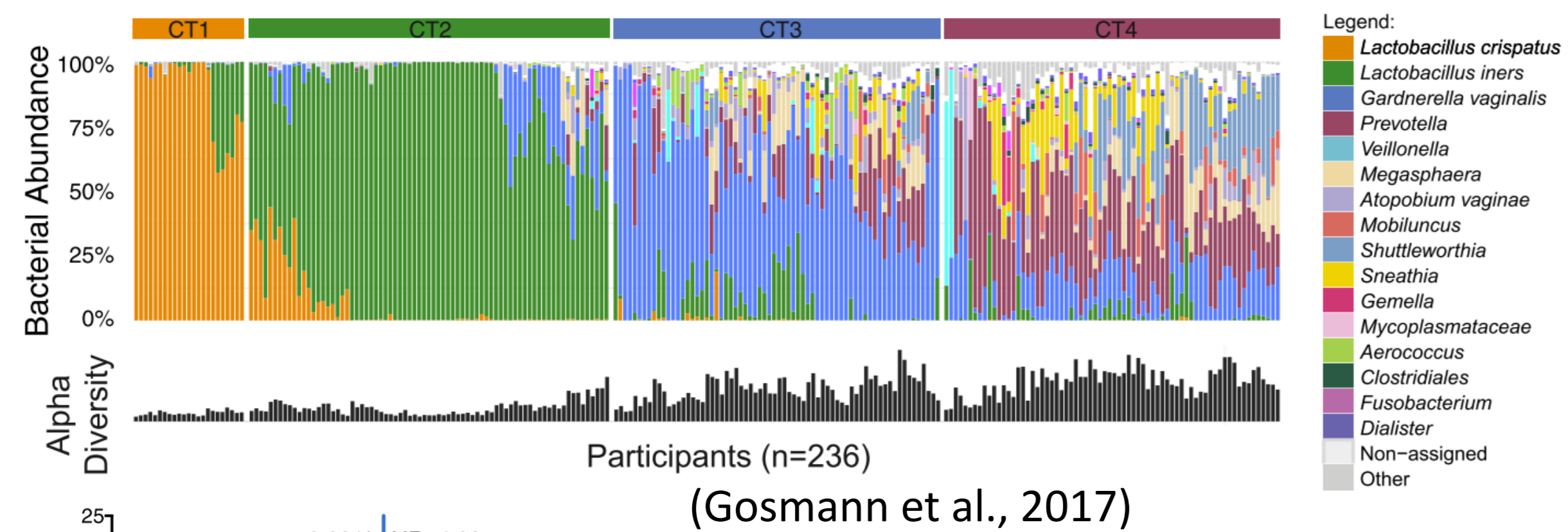


Combining standard bacterial vaginosis treatment with cystine uptake inhibitors to block growth of *Lactobacillus iners* is a potential target for shifting the cervicovaginal microbiota towards health-associated *Lactobacillus crispatus*-dominant communities

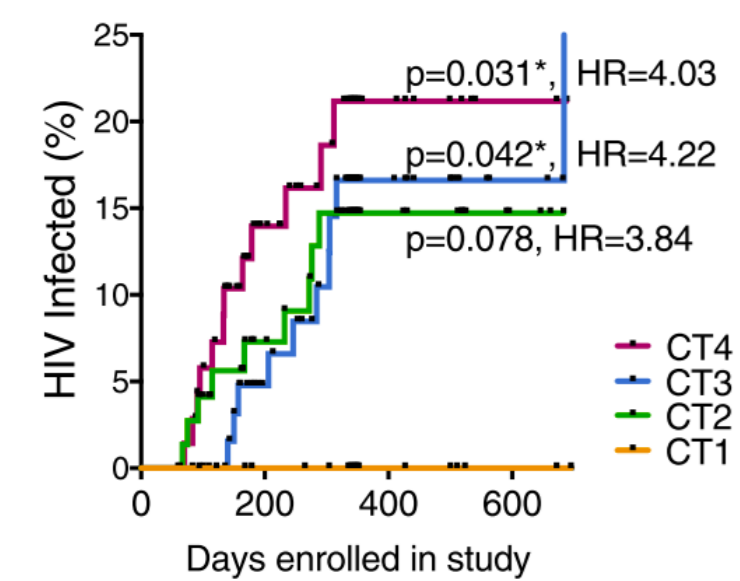
Seth M. Bloom*, Nomfuneko A. Mafunda, Benjamin M. Woolston, Matthew R. Hayward, Josephine F. Frempong, Jiawu Xu, Alissa Mitchell, Xavier Westergaard, Justin K. Rice, Namit Choksi, Emily P. Balskus, Caroline M. Mitchell, Douglas S. Kwon

Introduction:

- Cervicovaginal microbiome domination by *Lactobacillus crispatus* is associated with protection from HIV and with other beneficial outcomes.
- By contrast, *L. iners*-dominance is associated with adverse outcomes and higher risk of bacterial vaginosis (BV) and HIV (Fig).



(Gosmann et al., 2017)

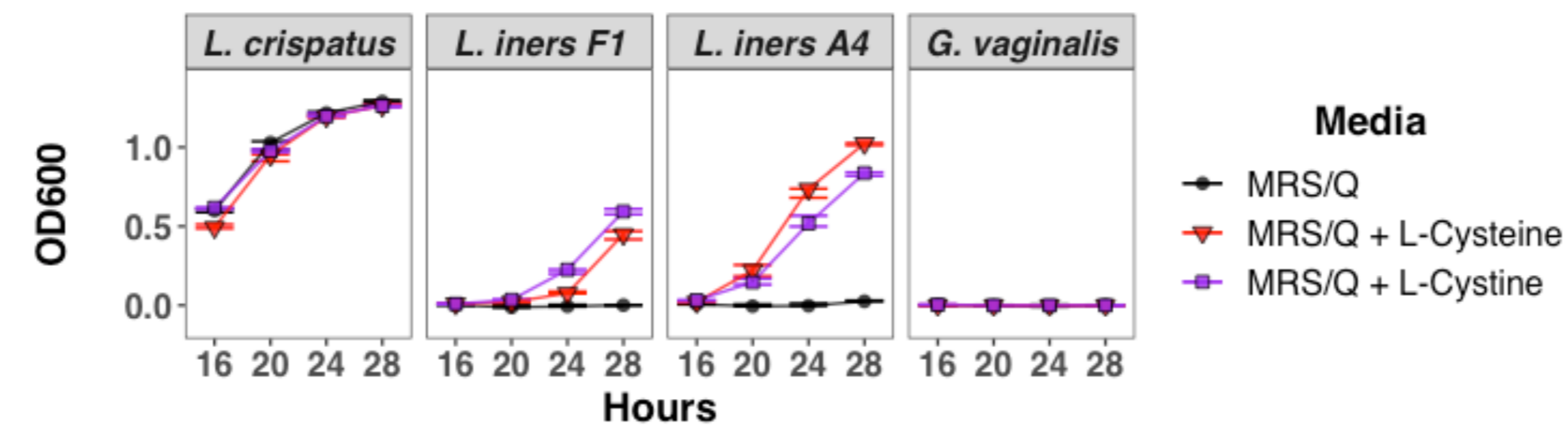


- BV treatment with metronidazole (MTZ), to which lactobacilli are resistant, typically produces *L. iners*-dominant (not *L. crispatus*-dominant) communities, and post-treatment relapse rates are high.
- Selectively inhibiting *L. iners* during BV treatment could mitigate HIV risk by favoring shifts to *L. crispatus*-dominant communities, but *L. iners* has been difficult to phenotypically characterize *in vitro* because it fails to grow in standard *Lactobacillus* MRS media unlike other cervicovaginal lactobacilli.
- We investigated *L. iners* growth characteristics to identify the basis for its unique growth requirements and to develop strategies for inhibiting it in favor of *L. crispatus*.**

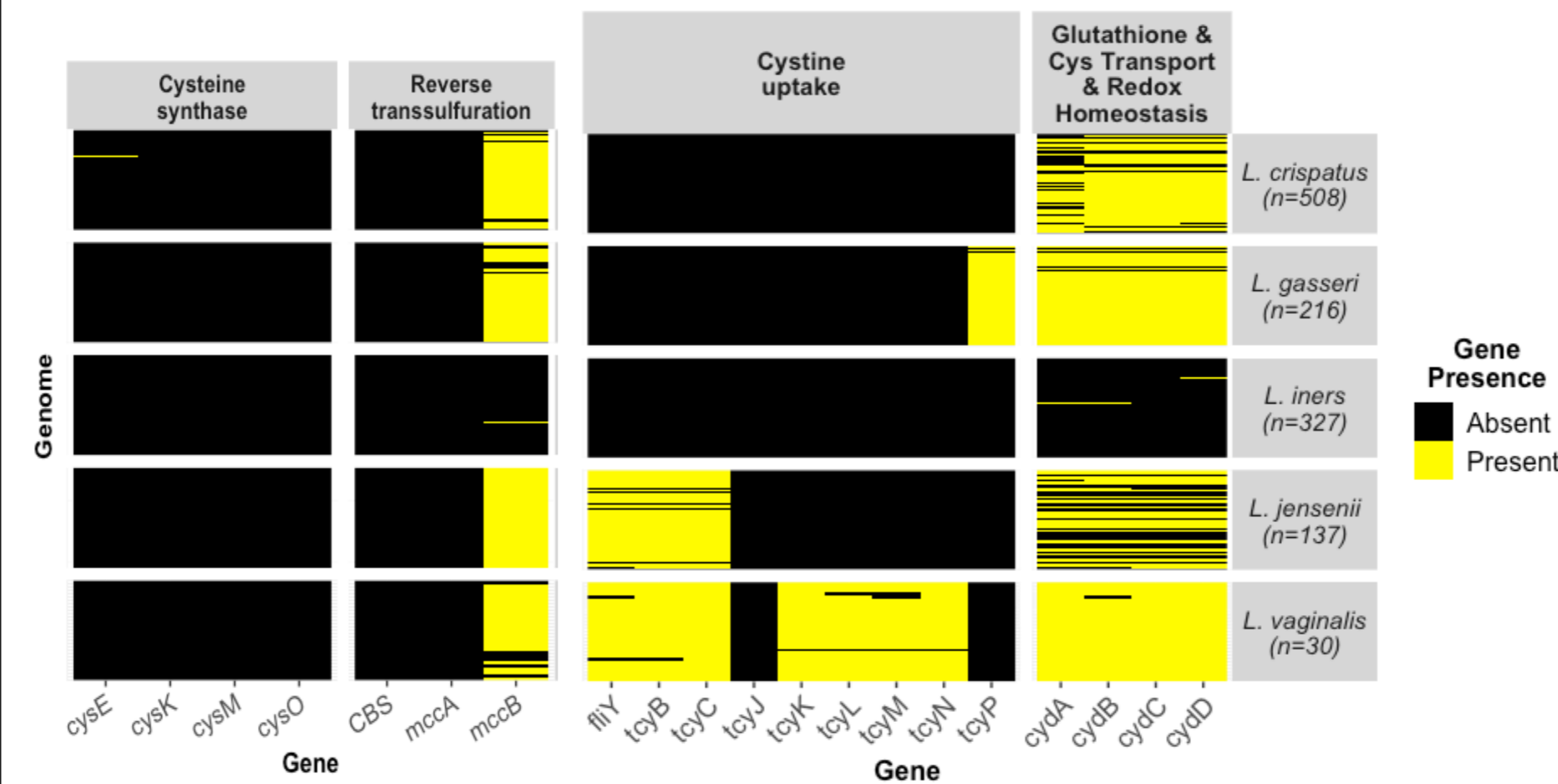
Methods:

- Bacteria were cultured anaerobically. Mono-culture growth in broth +/- supplements or inhibitors was quantified by optical density.
- Comprehensive *Lactobacillus* genome collections assembled from novel and published isolate genomes and from metagenomically assembled genomes were annotated using Prokka, Roary, and eggNOG.
- Competitive mixed cultures employing mock microbial communities +/- inhibitors were quantified via 16S rRNA gene sequencing.

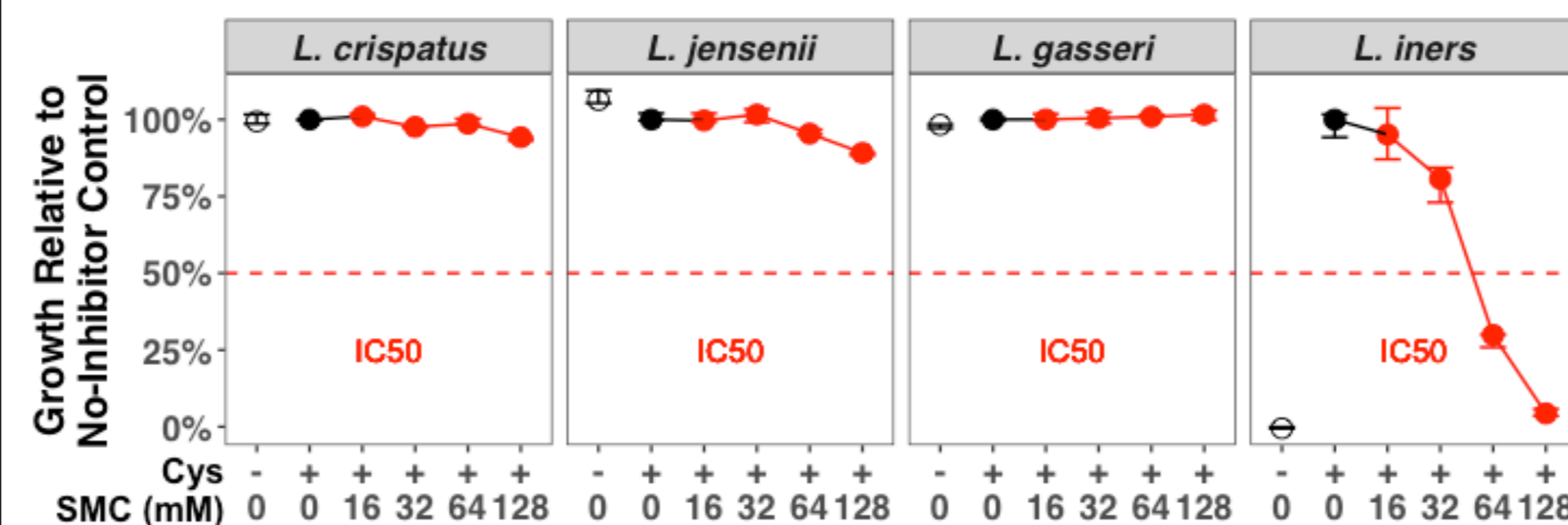
Cysteine supplementation supports *L. iners* growth in MRS media



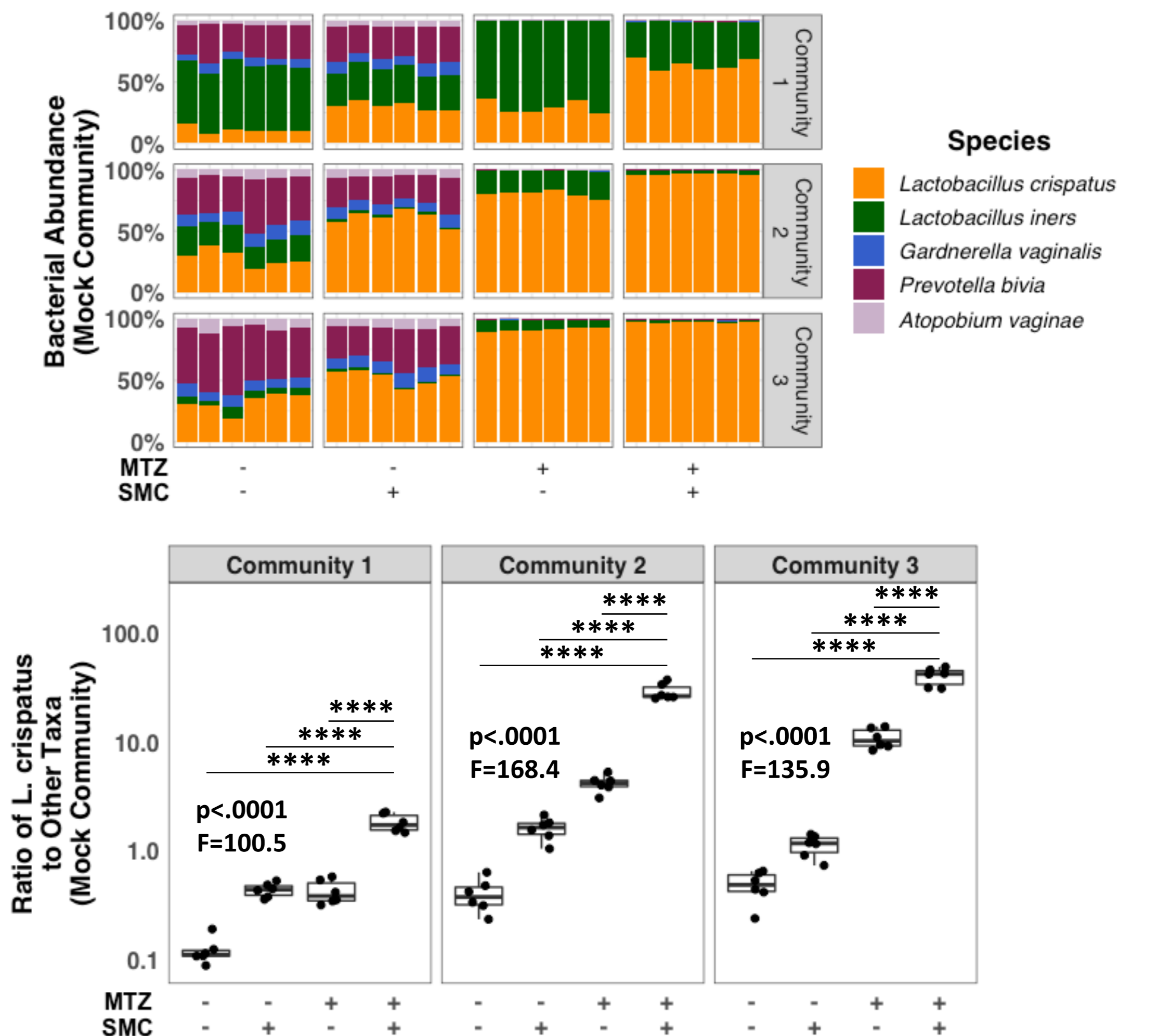
Major cervicovaginal *Lactobacilli* all lack canonical cysteine biosynthesis pathways, but *L. iners* uniquely lacks the *cydABCD* cysteine/glutathione transport operon



A cystine uptake inhibitor (SMC) selectively blocks growth of *L. iners*



MTZ combined with SMC shifts mock BV-like communities to *L. crispatus* dominance more effectively than MTZ alone



Conclusions:

- L. iners* has unique nutritional requirements related to uptake of exogenous cysteine/cystine that can be selectively inhibited.
- Targeting cysteine uptake to inhibit *L. iners* is a candidate strategy to shift the cervicovaginal microbiota toward *L. crispatus*-dominated communities associated with low HIV risk and other beneficial health outcomes.

Support:

Harvard University CFAR/NIAID 5P30AI060354-15 (2019-present)
NIH T32 AI007387 (2016-2018)
smbloom@mgh.harvard.edu