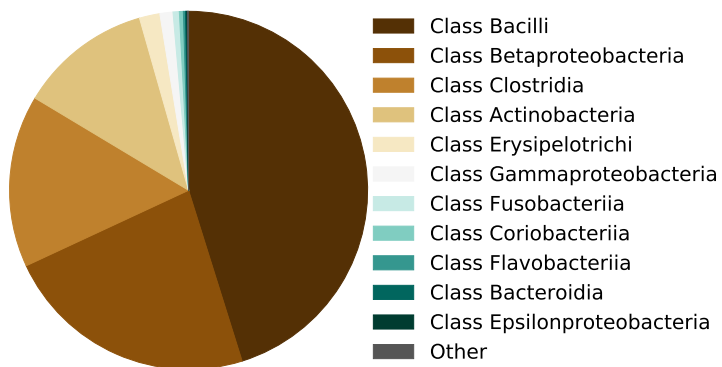


CHARLES WARDEN

What's in your oral sample?



Your most abundant microbes:

Taxonomy	Sample
Genus <i>Streptococcus</i>	27.8%
Family Neisseriaceae	22.6%
Genus <i>Granulicatella</i>	15.1%
Genus <i>Actinomyces</i>	10.6%

Your most enriched microbes:

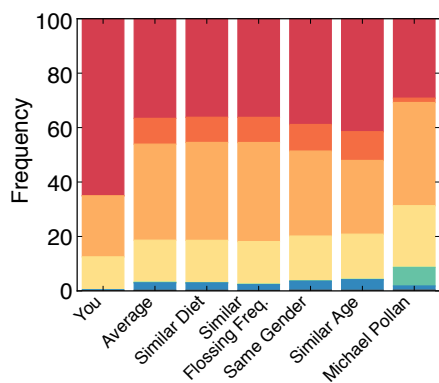
Taxonomy	Sample	Population	Fold
Genus <i>Actinomyces</i>	10.55%	2.44%	4x
Genus <i>Vagococcus</i>	0.12%	0.03%	4x
Genus <i>Butyrivibrio</i>	0.38%	0.01%	53x
Family Neisseriaceae	22.56%	6.34%	4x

Your sample contained 9 rare taxa, including the following: Genus *Paenibacillus*, Genus *Solibacillus*, Genus *Salinicoccus*, Genus *Trichococcus*, Genus *Devosia*.

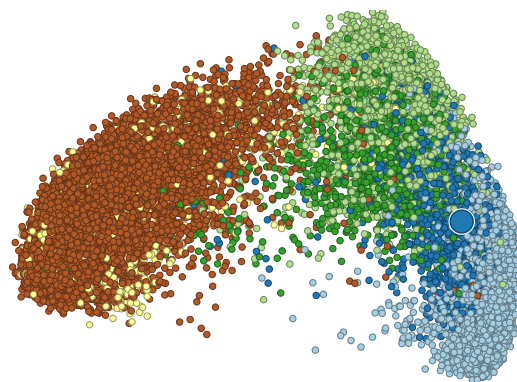
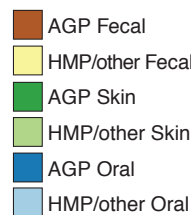
This sample was registered on unknown at 21:35.

How do your oral microbes compare to others?

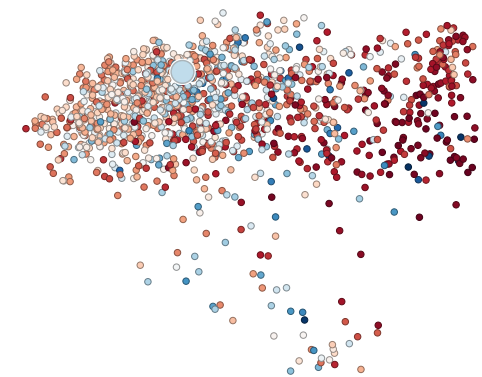
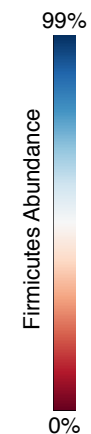
● You ● Others ● Missing metadata



Different Subpopulations



Different Body Sites



The American Gut Population