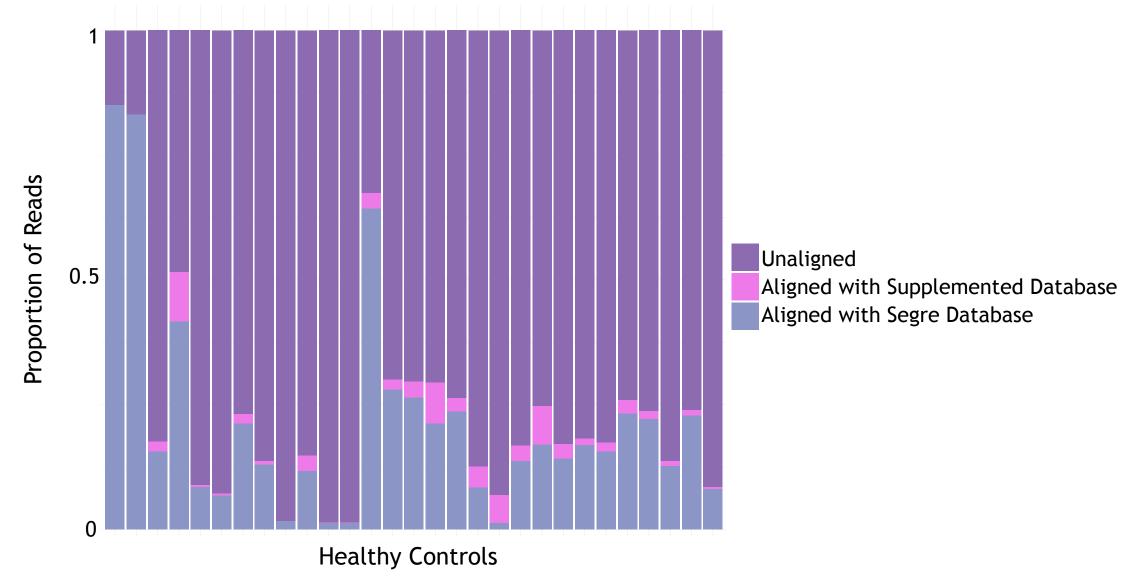
## More reads are mapped with addition of genomes



## Questions and Goals

• In our hands, how much of the skin microbiome is dark matter?

 What are the genus and species-level holes in the database that we can fill with publicly available reference genomes?

 What are the strain level holes in the database that we need to fill by culturing and sequencing?