Sequencing data (N=3) shows Wuhan coronavirus integration in bacteria (*Prevotella* mostly). Sequencing artifact - or is the virus infecting both bacterial and human cells?

Sandeep Chakraborty

Letter

RETRACTION NOTE

I retract this paper - initially I thought that the bacteria and the virus had integrations, but these turned out to be chimeric reads, a sequencing artifact. Please follow my other research thread that SARS-Cov2 is enabling anaerobic bacteria to colonize the lungs at <https://osf.io/usztn/>.

References