

GENETICS

# VIRUS REACTIVATION

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A recent [Nature study](#) showed Ebola reactivation in a previously infected patient:

The 2021 lineage shows considerably lower divergence than would be expected during sustained human-to-human transmission, which suggests a persistent infection with reduced replication or a period of latency.

The most recent viral genome shared 10 substitutions that evolved during the previous epidemic making it unlikely that there was a new animal spillover event. I always wondered how second & third wave of COVID-19 started in Germany. Was it really a new spread or just a reactivation? A [Frontiers review](#) concluded from the existing literature that

our study, consisting more than a total of 113,715 patients, indicates that the RP-SARS-CoV-2 scenario occurs plausibly due to reactivation, reinfection, viral shedding, or testing errors.

So far, there are 240 documented COVID-19 cases of reinfection reported worldwide [according to the reinfection tracker](#). In the case of Ebola there is a known viral persistence in semen while transmission through milk and cervicovaginal fluid is also possible (similar for [COVID-19](#) although [neuronal persistence](#) seems to be more relevant). Maybe we need more immunological studies [particular in long COVID](#) if there is a continuous or intermittent antigenic stimulation due to persistence of an antigenic reservoir.

The fourth wave in Germany is caused by the delta variant, reactivation of alpha is certainly not a major factor. So we will only know in the next few years if reactivation is responsible for small regional outbreaks in unvaccinated communities.

