

20. Gout JF, Thomas WK, Smith Z, Okamoto K, Lynch M (2013) Large-scale detection of in vivo transcription errors. *Proc Natl Acad Sci USA* 110(46):18584–18589.
21. Acevedo A, Brodsky L, Andino R (2014) Mutational and fitness landscapes of an RNA virus revealed through population sequencing. *Nature* 505(7485):686–690.
22. Acevedo A, Andino R (2014) Library preparation for highly accurate population sequencing of RNA viruses. *Nat Protoc* 9(7):1760–1769.