

CORRECTION

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# Correction to: An application of slow feature analysis to the genetic sequences of coronaviruses and influenza viruses

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**Correction to: Hum Genomics 15, 26 (2021)**

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Following publication of the original article [1], the authors would like to change all the word DNA to nucleotide or genetic.

The viruses that we analysed have an RNA genome, however the sequences obtained from the repository (NCBI) are reported as DNA. In this manuscript a neutral term "nucleotide" sequence(s) or "genetic" structure should be used instead of DNA sequences or DNA structure

The original article [1] has been updated.

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1. Tsonis AA, et al. An application of slow feature analysis to the genetic sequences of coronaviruses and influenza viruses. *Hum Genomics*. 2021;15:26.

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