

<https://www.coincoin.fr.eu.org/?Sans-titre-34054>



Fil Threadreaderapp

- 6- Webographie -

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In new study led by @bdadonaite, we measure how all mutations to H5 influenza HA affect four molecular phenotypes relevant to pandemic risk : Results can inform surveillance of ongoing evolution of H5N1. [biorxiv.org/co...](https://www.biorxiv.org/content/10.1101/2020.03.10.334441v1)