Painting courtesy of Matt Cotten @mlcotten13



Genomic epidemiology and population genetics of emerging RNA viruses

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eLife 2018;7:e31257



Coronavirus discovered in 2012

Coronavirus discovered in 2012 Severe morbidity/mortality*



Coronavirus discovered in 2012 Severe morbidity/mortality*

Parallels with SARS-CoV inevitable



NPR (blog)



SARS veterans tackle coronavirus Nature.com - Oct 3. 2012 Last week, the researchers reported the genome sequence of the new coronavirus and the first diagnostic tests to screen for it - two major advances that will ...

When New Diseases Emerge, Experts Are Faster On The Uptake North Country Public Radio - Oct 3, 2012

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Coronavirus-Patient ohne besondere Schutzmaßnahmen behandelt Spiegel Online - Nov 23, 2012 Anspannung bei den Gesundheitsbehörden: Ein Mann aus Katar hatte sich mit dem

Coronavirus discovered in 2012 Severe morbidity/mortality* Parallels with SARS-CoV inevitable "Mysterious epidemiology"

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Coronavirus discovered in 2012 Severe morbidity/mortality* Parallels with SARS-CoV inevitable "Mysterious epidemiology" But MERS-CoV restricted to Arabian peninsula



Coronavirus discovered in 2012 Severe morbidity/mortality* Parallels with SARS-CoV inevitable "Mysterious epidemiology" But MERS-CoV restricted to Arabian peninsula No evidence of widespread infection in humans Very high seroprevalence in camels



Spoiler alert

MERS is a recurring zoonosis in humans



The Questions

How frequent is spillover into humans?

How well does MERS-CoV do in humans?

Which host(s) responsible for virus maintenance? How asymmetric is virus transfer between hosts?





Data so far

Epidemiological

Sequence

Andrew "Phylogenetics Rambo" Rambaut's curated case list MERS-CoV genomes from humans and camels



~300 genomes ~2:1 human to camel ratio

Studies into camel-human interface so far

Traditional epi

Phylodynamics



Problems

Difficult to link cases into clusters

regions."

Sequence sampling skewed, models used inappropriate







Not caused by prior





How good (bad) is MERS-CoV in humans?



How good (bad) is MERS-CoV in humans?



Webstation adaptation/mis-assigned recombinants?



Conclusions

Target camels to stop MERS-CoV



Conclusions

Target camels to stop MERS-CoV Continue genomic surveillance



Conclusions

Target camels to stop MERS-CoV Continue genomic surveillance Example of sequence to epidemiology



Thank you

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MERS-CoV spillover at the camel-human interface. eLife 2018; 7:e31257









