

Fatal human encephalitis with new virus and characterized by standardized and validated Shotgun metagenomic approach

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Fatal human encephalitis with new virus and characterized by standardized and validated Shotgun metagenomic approach

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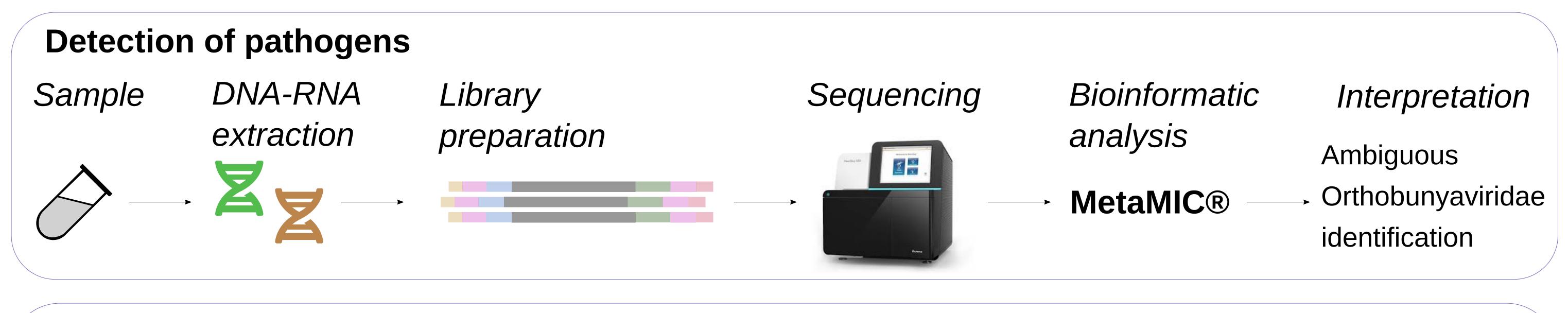
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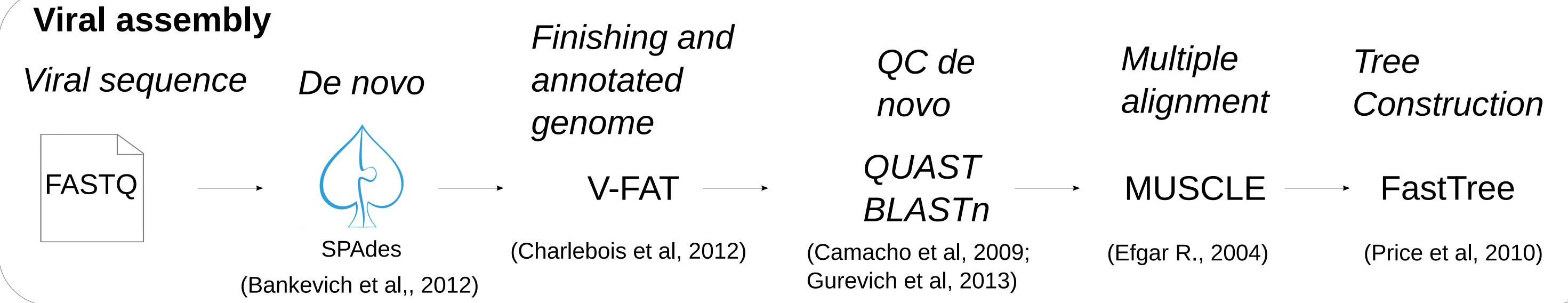
Context

Shotgun metagenomics have demonstrated high potential to detect all pathogens in undignosed syndromic infectious disease. Thus, we have developed an accredited (ISO EN NF 15189) SMg technology (MetaMIC®) to explore atypical and severe clinical cases.

We report a case of a 58year-old woman, admitted to the emergency department for neurological syndrome, treated as epileptic disease. Despite intensive symptomatic treatment therapy, the patient falls into a coma and subsequently died with encephalitis symptoms. MRIs concluded to probable severe status epilepticus and all conventional microbiology testing assays remained negative on clinical samples including sequential cerebrospinal fluids and cerebral biopsy which were tested by SMg.

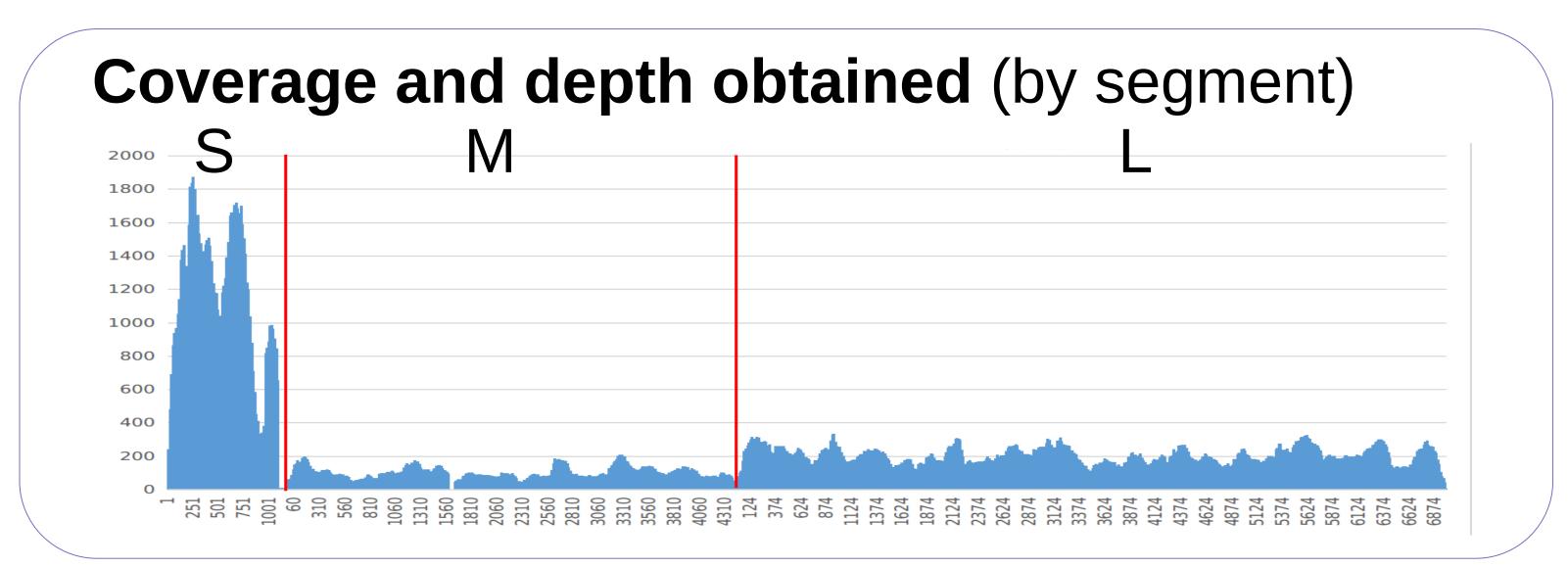
Materials and Methods





Results

Viral phylogeny of Orthobunyaviridae Koongol virus - KP792667 Umbre virus - KP792685 0.21 0 S Segment Witwatersrand virus - KP792688 0.04 M'Poko virus - AM711133 0.09 Umbre virus - KP792686 M Segment Koongol virus - KP792668 Witwatersrand virus - KP792689 L Segment Witwatersrand virus - KP792690 Koongol virus - KP792669 Umbre virus - KP792687 Little Sussex virus - KT720483



Conclusion

- The patient harbored a encephalitis infection with new virus impossible to detect with conventional diagnostic tools.
- This case highlighted the major advantage to have a reliable SMg tool to diagnose new pathogens in a world with frequent emerging infectious diseases













