

Pig-borne disease most likely jumped into humans when rearing practices changed

March 31 2015



A domestic pig on an organic farm in Solothurn, Switzerland. Image: Wikimedia Commons

Almost every pig carries harmless strains of the *S. suis* bacterium - such strains are known as 'commensal' strains. However, a more virulent group of strains of the bacteria also exist, which cause disease in pigs worldwide and are a major driver of antibiotic use for prevention. Increasingly, this group of strains is also implicated in serious human diseases such as meningitis and septicaemia.

In order to understand the genetic basis of disease in *S. suis*, an international study, led by the Bacterial Respiratory Diseases of Pigs-1 Technology (BRaDP1T) consortium, examined the genomes of 375 clinical samples from pigs and humans from the UK and Vietnam and combined these with data already published on 15 *S. suis* [genome](#)

sequences and draft genomes from around the world. They found that the commensal strains and disease-causing strains differed genetically; in particular, the disease-causing strains have between 50 and 100 fewer genes than the commensal strains.

Dr Lucy Weinert from the Department of Veterinary Medicine at the University of Cambridge, first author of the study, says: "It seems that the loss of genes is associated with causing disease. This is something we see quite often in [bacteria](#), but for reasons that are unclear. One possibility is that the missing genes are those that hinder the function of newly-acquired virulence factors in the genomes."

By examining the *S. suis*'s 'tree of life' - which looks at how the bacteria have evolved over time - the researchers were able to show that the emergence of a group of strains causing meningitis in pigs and the human form of the disease dates back to the 1920s, when pig production was intensified with the introduction of wide-scale indoor rearing of meat-producing pigs in larger groups, supported by government schemes that favoured larger producers with regular throughput. However, despite having jumped the species barrier from pig to human, the bacteria do not appear to have adapted to infect humans.

"A group of more [virulent strains](#) seem to have emerged at around the time the pig industry changed, and it is these [strains](#) that mostly cause disease in [pigs](#) and humans," says Professor Duncan Maskell, Head of the School of the Biological Sciences at Cambridge.

"Human *S. suis* [disease](#) in the West is extremely rare, but is seen more frequently in south east Asia. It is most likely spread to humans through poor food hygiene practices or other environmental factors. This emphasises the importance of monitoring such practices and putting policies in place to reduce the risk of the spread of infections between species."

More information: Weinert, LA et al. Genomic signatures of human and animal disease in the zoonotic pathogen *Streptococcus suis*. *Nature Communications*; 31 March 2015.

Provided by University of Cambridge

Citation: Pig-borne disease most likely jumped into humans when rearing practices changed (2015, March 31) retrieved 10 April 2024 from <https://phys.org/news/2015-03-pig-borne-disease-humans-rearing.html>

<p>This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.</p>
--