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African Swine Fever Virus

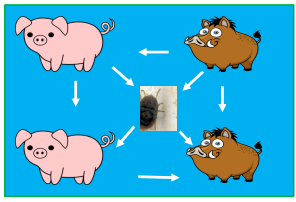
- African swine fever virus: Asfarviridae family
 - dsDNA genome 170-190kb
 - 150+ genes
 - Very few have been studied in detail ~20
- Highly lethal (100%) to subclinical infection
- Hemorrhage, edema, ascites and shock
- Wild Boar, Warthogs, domestic pigs
- Virus/Host factors responsible for different outcomes
 - Remain poorly understood
 - Different strains have different virulence
 - Varying host species symptoms/lethality
- Long-term persistent/latent infection
- Rapid and efficient transmission among pigs



AFV (100x) (100000x)
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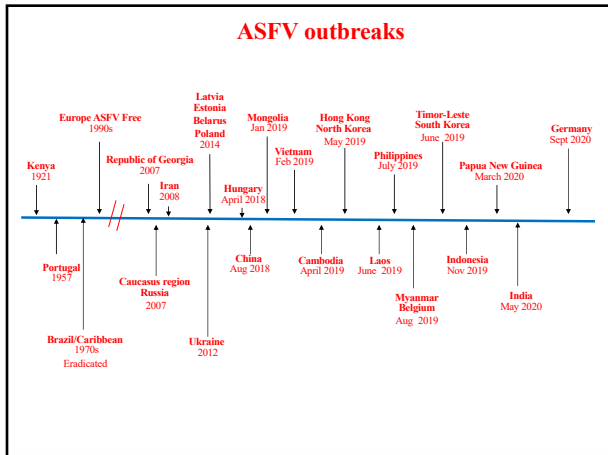
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African swine fever virus: transmission

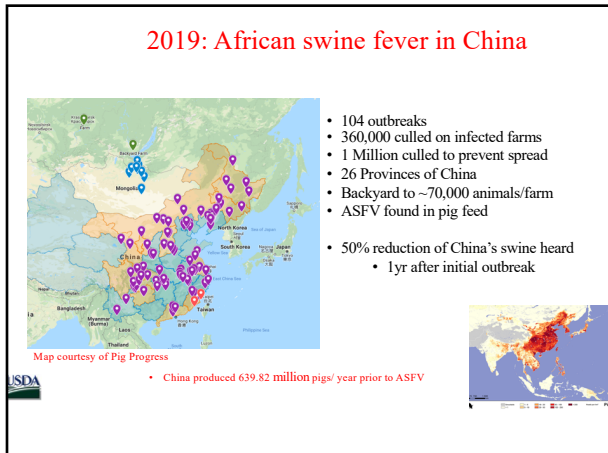


- Several days in feces
- Months in contaminated pens
- Years in frozen carcasses
- Pork products: Found over 140 days in salted/dried hams
- Feeding swill, garbage, waste

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Live attenuated vaccines (LAV) are effective preventing homologous challenge

Attenuated isolates were produced by:

- Low virulence field isolates
- Viruses attenuated by tissue culture passages
- Viruses with genetically engineered deletions

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Live attenuated vaccines (LAV) are effective preventing homologous challenge

Attenuated isolates were produced by:

• Low

In this study we investigated whether deletion of genes from the ASFV low virulence isolate OUR T88/3 would reduce adverse clinical reactions post-immunisation but maintain induction of high levels of protection post-challenge with virulent isolate OUR T88/1. We took advantage of the adjacent genomic location of

Immunisation with OUR T88/3ΔDP2 did not appear to significantly reduce the adverse clinical reactions observed in pigs compared to parental OUR T88/3 since two pigs in each of groups 1 and 2 had transient joint swelling. Although it is possible that fever virus isolate OUR T88/3 decreases its ability to protect against

deletion virus OUR T88/3ΔDP2 compared to 100% protection with the parental virus OUR T88/3. Thus the deletion of the two genes DP71L and DP96R from OUR T88/3 strain reduced its ability to protect pigs against challenge with virulent virus.

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Live attenuated vaccines (LAV) are effective preventing homologous challenge

Journal of General Virology (2001), 82, 513-523. Printed in Great Britain

In the current study, we investigated the effect of deletion of genes (A238L [28-30]; A224L [31,32]; EP153R [33,34] and A276R [35]), involved in virus-host interaction and immune system control on protection and virulence of the naturally-attenuated ASFV strain NH/P68.

Vaccine challenge. Pigs immunized with 10⁶ TCID₅₀ of either vaccine candidates produced in COS7 cells or parental NH/P68 (produced in PAM), developed mild clinical signs (slightly raised body temperatures, necrotic skin areas and joint swelling, associated with chronic ASFV infection [25,27,45,46]) or remained asymptomatic.

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Live attenuated vaccines (LAV) are effective preventing homologous challenge

Attenuated isolates were produced by:

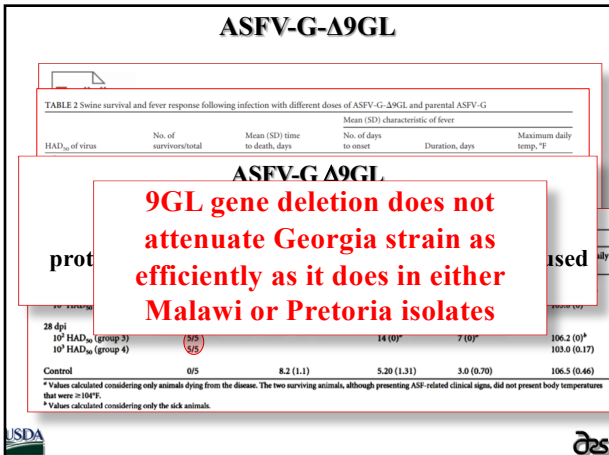
• Low

Low virulent field isolates retain residual virulence

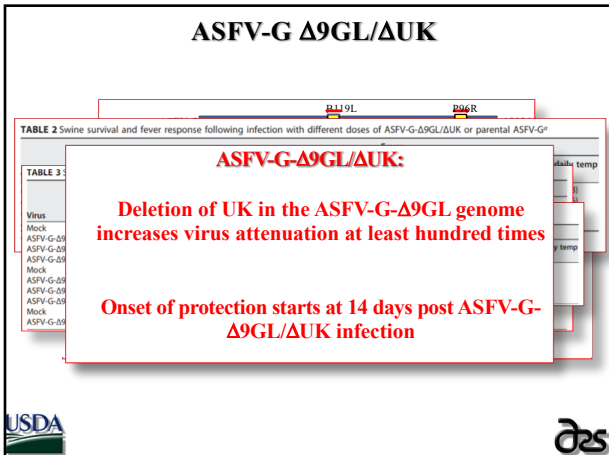
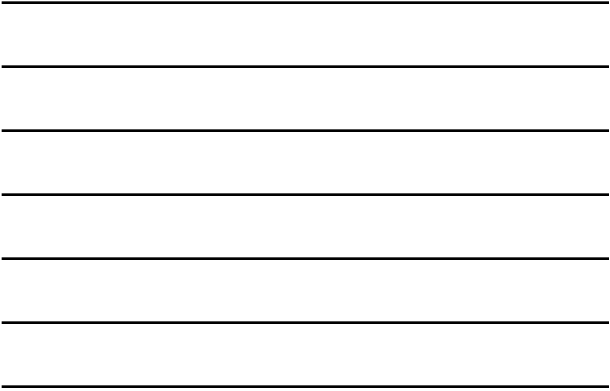
inoculated pig (PW17) showed weak peaks of fever (40.3-40.7°C) from 8 to 12 dpi accompanied by the appearance of cyanosis in

intermittent weak viraemia (CI > 35) was detected in PW15 from 22 to 94 dpi, while PW18 remained aviraemic on all analysis days (up to 101 dpi). The remaining in-contact animals (PW14 and PW16) presented mild transitory clinical signs, specifically joint swelling, from 7 to 16 dpi, particularly in PW16. In PW14, the viraemia lasted for

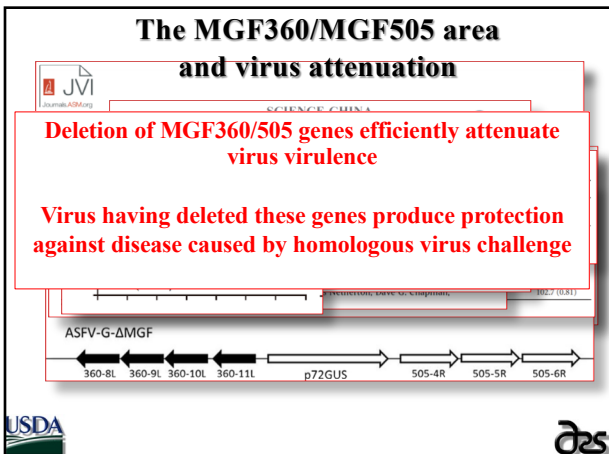
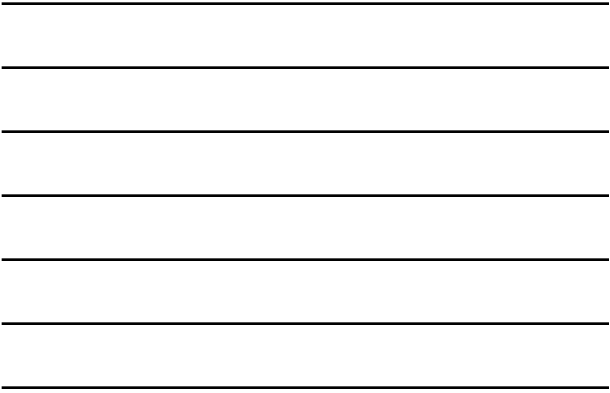
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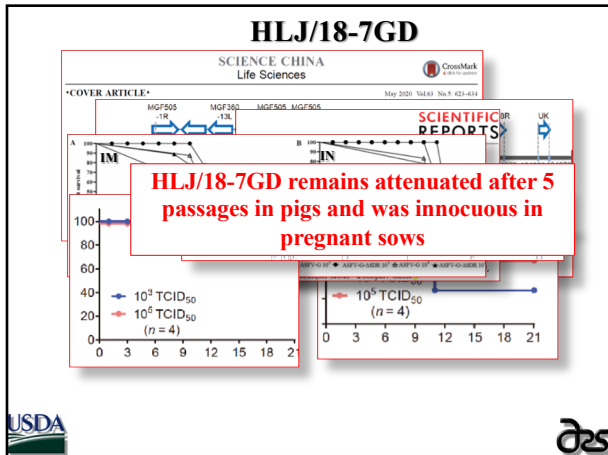


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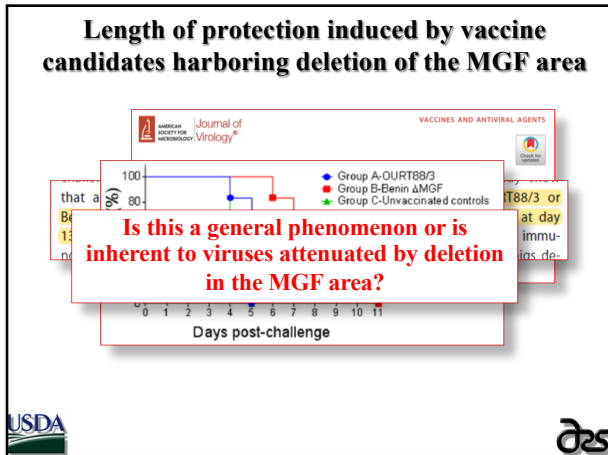


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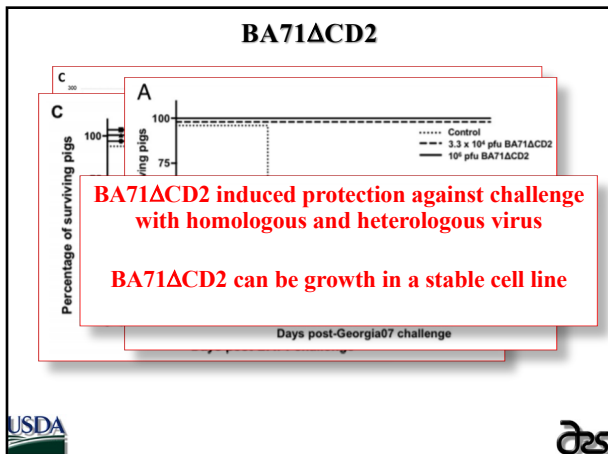




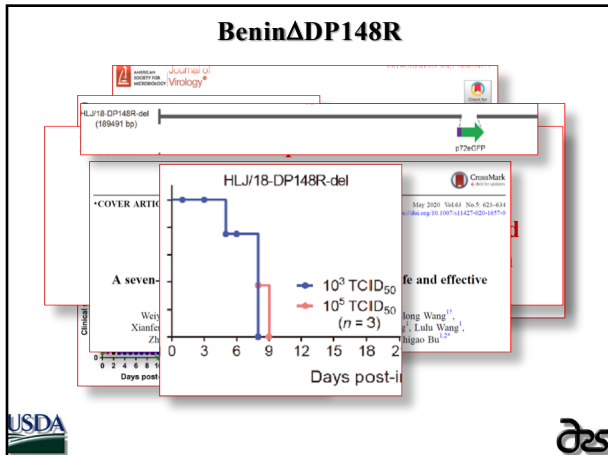
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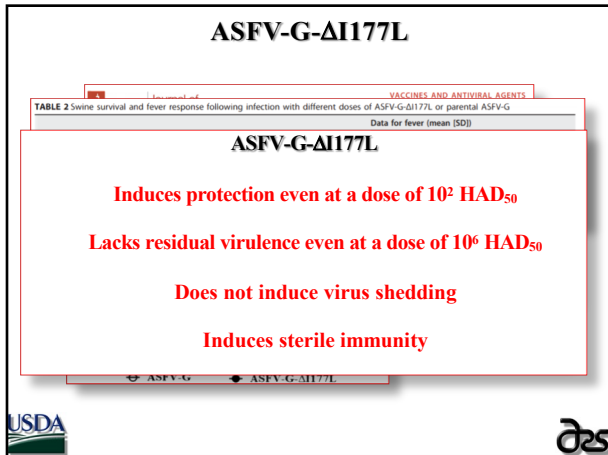
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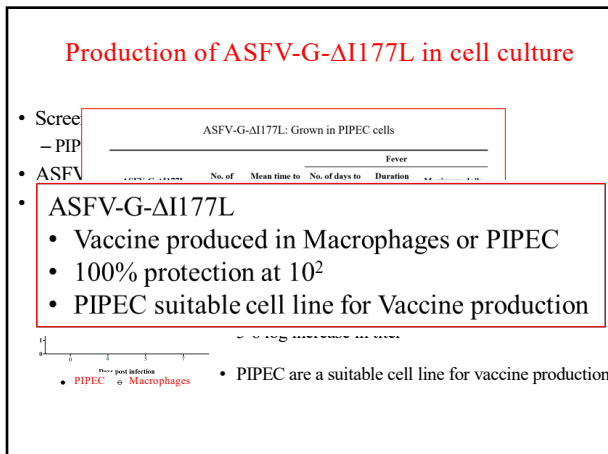
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Gaps

Study critical aspects of vaccine candidates (onset and duration of immunity, genetic stability, attenuation stability, minimal protection doses, possible routes of inoculation)

Standardization of methodologies to test efficacy of vaccine candidates

Develop vaccine candidates with DIVA capability

Investigate the bases of the differential effect in virus virulence of deleting the same genes among different virus isolates (9GL, CD2, NL, UK, TK, DP148R)

Continue the discovery and characterization of virulence associate genes

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ARS vaccine candidates

ASFV-G-Δ9GL
ASFV-G-ASFV-G-Δ9GL/ΔUK
ASFV-G-ΔMGF
ASFV-G-ΔI177L
ASFV-G-ΔI177L/ΔLVR

All these candidates have been patented and are being in the process of being licensed by veterinary pharmaceutical companies

Thank you

Questions?

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