

Red mark syndrome in rainbow trout



Jacob G. Schmidt

Red mark syndrome in rainbow trout

DTU



Future

Problem

Background

Project

Model

Agent

Response



Red mark syndrome in rainbow trout



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Starting c. 2010 Danish fish farmers started reporting red marks

2015 questionnaire: 1 of 3 farms with RMS. Increasing.

Now most important disease in DK aquaculture (alongside BCWD)

The fish looks sick

RMS is very difficult to manage

“Slow disease”



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Likely a disease endemic to western North America, where it is known as strawberry disease and affects *Oncorhynchus* sp. First reports are from middle of 20th century.

One report from France in mid-80s, but apart from this the first European RMS reports are from the UK c. 2003.

The disease has since spread to many European countries as well as Iran and Chile.



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The disease has been proposed to be associated with an adenovirus, *Flavobacterium psychrophilum* and a Rickettsia-like organism (RLO).

The first association with an RLO was made in 2008. This has since been considered the most likely candidate for a causative agent but a definite connection has not been made.

The RLO is now known to be more closely related to the recently described family Midichloriaceae, and is thus now referred to as Midichloria-like organism (MLO).



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- **Firmly establish what causes RMS**

- Developing an infection model
- Diagnostics
- NGS

- **Describe RMS symptom development**

- Histology: H&E, IHC, ISH
- Images

- **Manipulating the disease**

- Testing different factors such as antibiotics and glucocorticoids



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- **We first aimed at a cohabitation model of infection.**
- RMS-affected fish brought in from Møborg fish farm (~170 g).
- 3 fish were screened for vira, bacteria and parasites.
 - Vira: None (on cell culture)
 - Bacteria: Few (on BA, TYA and CHAB). *F. psychrophilum*.
 - Parasites: Few. Only *Ichthyobodo necator* detected.
- Cohabitation with own SPF fish (~90 g)
- All fish were PIT-tagged prior to cohabitation.



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- We first aimed at a cohabitation model of infection.

Control

Cohabitation

Untreated

Formalin treated

Untreated

~12°C

1X

2X

2X

30 SPF fish

15 SPF fish

15 SPF fish



10 Seeder fish



10 Seeder fish



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- **Infection facilities.**



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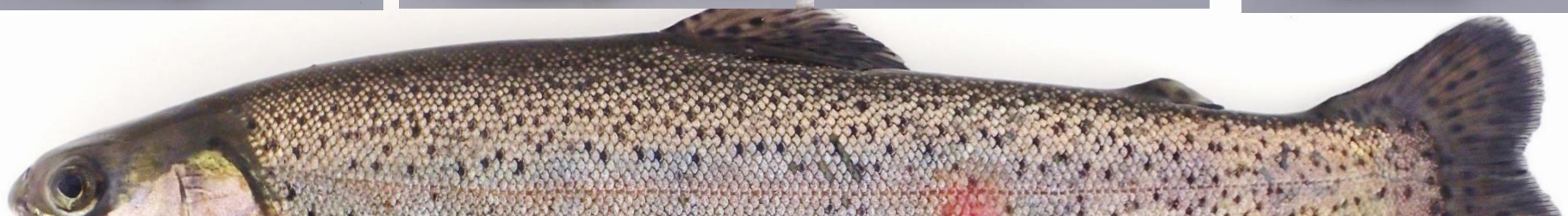
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• Seeders



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• Problems

- After 16 days of cohabitation SPF fish started to die. Over the next two weeks 11 out of 60 cohabitants died or were killed.
 - Probably a mix of mostly bacterial factors – sometimes infection of tagging site.
 - *F. psychrophilum* was the only bacterial fish pathogen detected from deceased or moribund fish – and only in small numbers.
 - No virus detected.
 - Two fish had fungal infection.



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- **More problems**

- Just as the fish started to recover, Ich infection was detected.
- Salt treatment
- 4 fish were killed due to heavy infection (several hundred trophonts).
- No further mortality.



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- **Eureka moment**

- Just as the Ich infection was wearing off the fish developed lesions consistent with early RMS symptoms

(~550 degree days)



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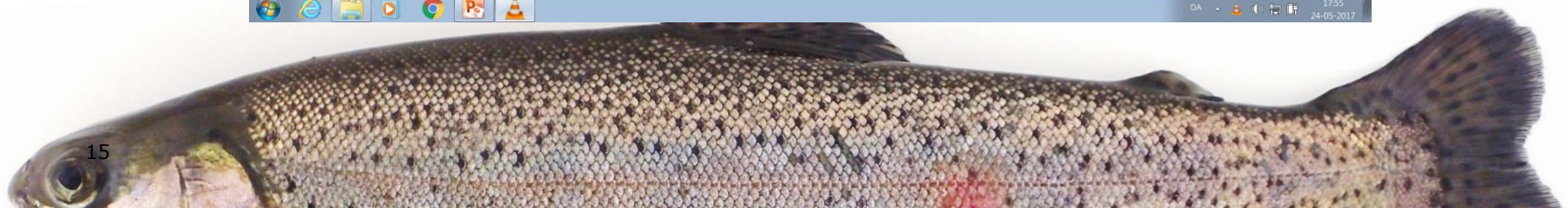
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- Symptoms developing nicely



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- Cohabitants after ~900 degree days at 12 °C



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- **Cohabitant 518999**

W0/0 dg



W7/420 dg



W9/740 dg



W10/850 dg



W12/970 dg



W14/1150 dg



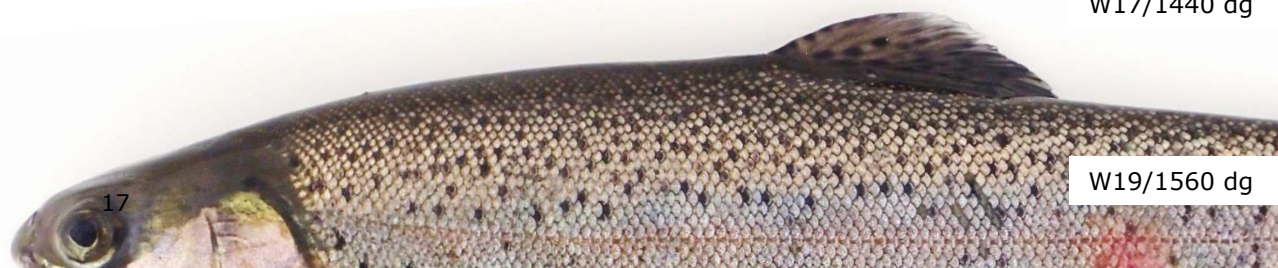
W16/1350 dg



W17/1440 dg



W19/1560 dg



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- **Cohabitant 518999**



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- **Cohabitation model established**

- Over the past year we have successfully transferred RMS to 5 passages of new cohabitants by using the old cohabitants as seeders for new SPF cohabitants
- Ich is eradicated
- *F. psychrophilum* is still in the system although it causes only very occasional mortality now.



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• Other infection approaches:

- Injection of tissue homogenate (organs and skin)
 - Hoping to avoid *F. psychrophilum*
- Limited success. Only few symptoms, and mostly at injection sites.
- Still *F. psychrophilum*.



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- **Other infection approaches:**

- Indirect cohabitation
- Also successful, but fewer lesions, longer incubation time.
- Actually seems to resemble farm situation somewhat better
- For experimental purposes, direct cohabitation more applicable



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• **Diagnostics:**

- No virus detected
(no CPE on RTG, BF-2 and EPC)
- No obvious pathogenic bacteria detected on agar plates
(BA, TYA, CHAB) – apart from *F. psychrophilum*
- MLO detected with specific qPCR and with 16S rDNA NGS
 - Correlation with disease



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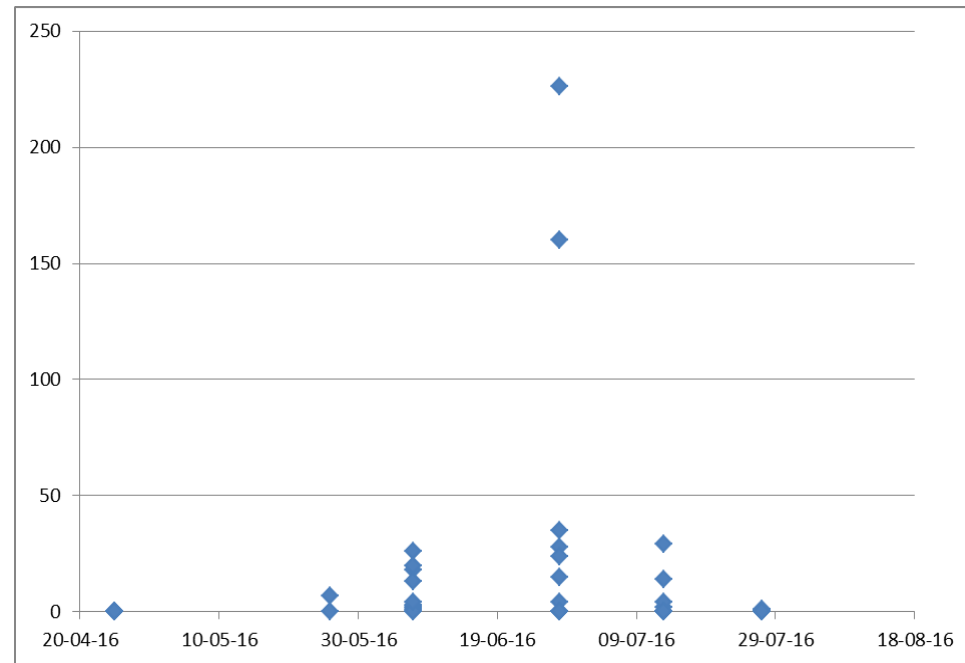
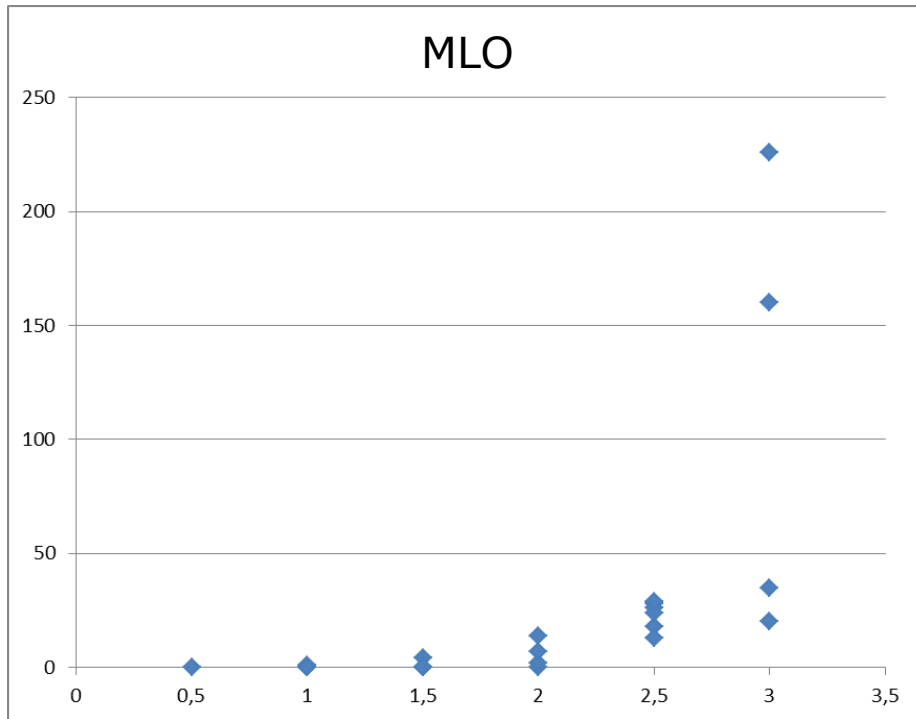
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• Diagnostics:

MLO



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- **Diagnostics:**

- Next generation sequencing of bacterial 16S rDNA further shows that MLO is the only bacterial group that correlates with RMS lesions



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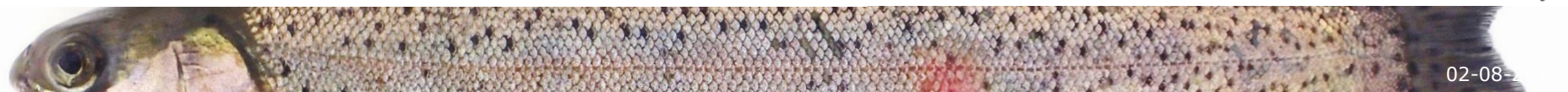
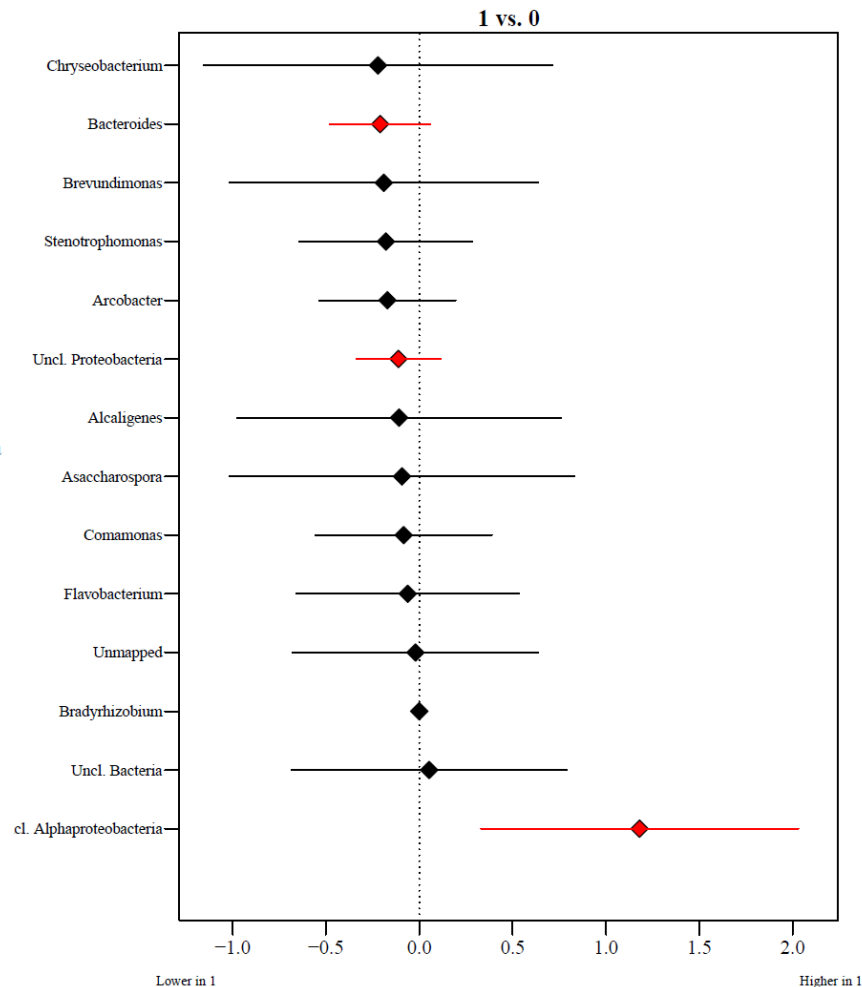
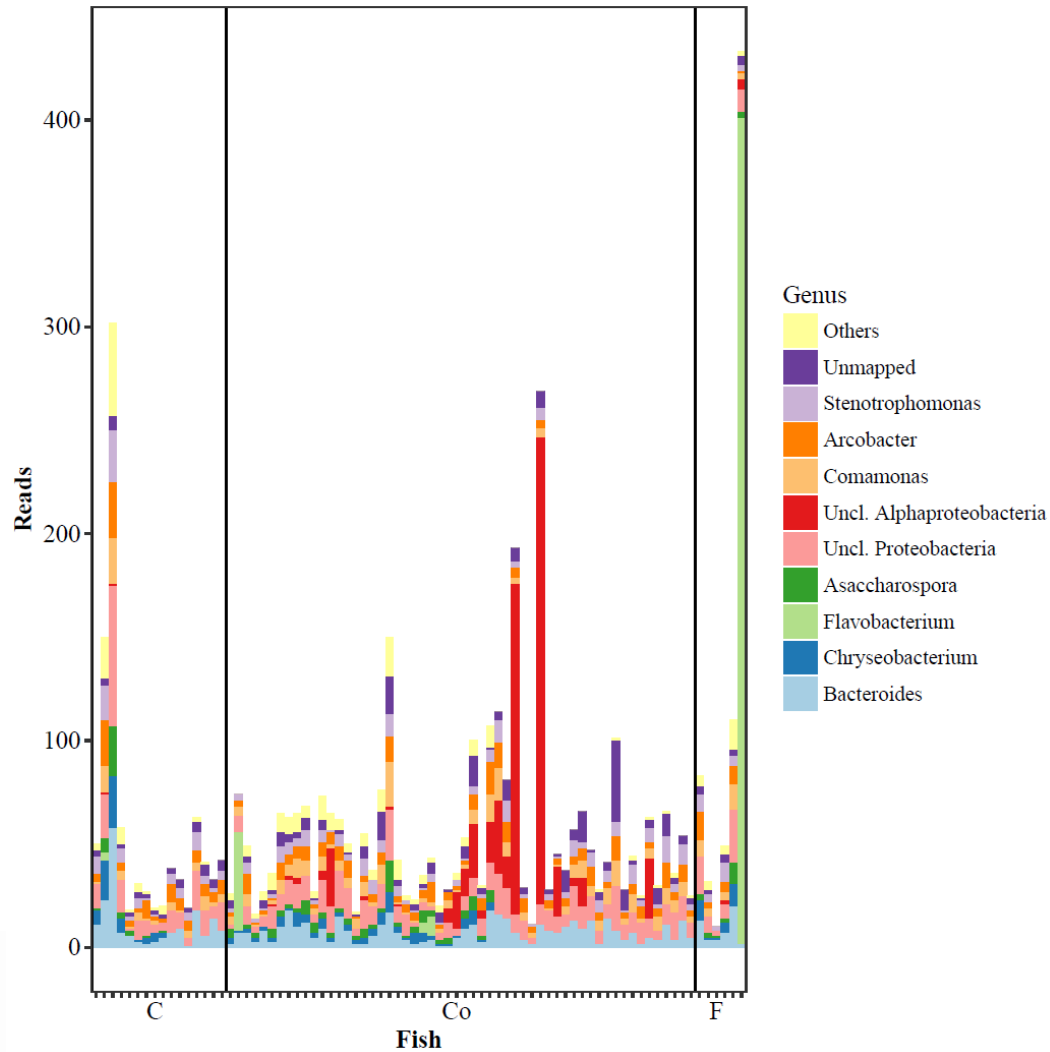
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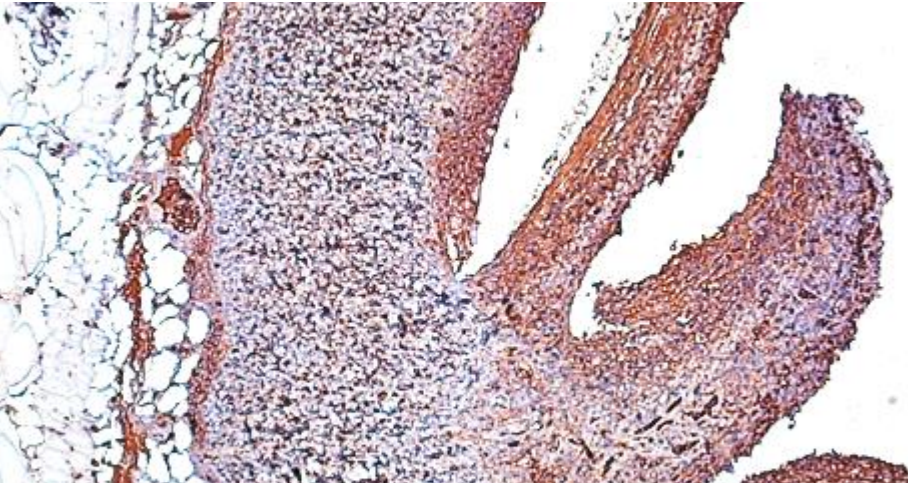
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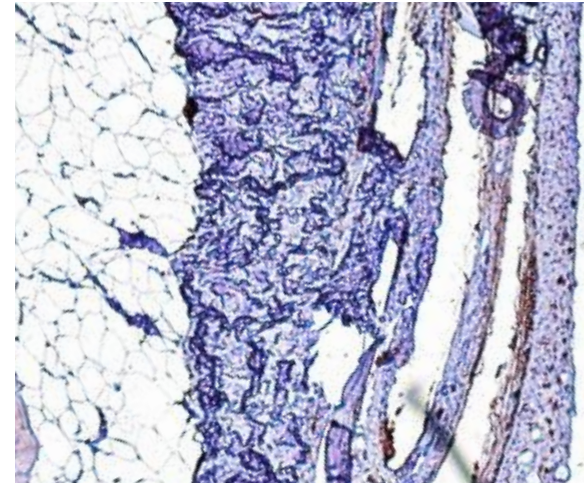
Future

- **Immunohistochemistry: IgM**

RMS



Control



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- **In this project:**

- Fish farms:

- NGS
- Questionnaires

- At DTU:

- Continuous optimization of the infection model
- RMS experiments with the model
 - Antibiotics
 - Glucocorticoids
 - Fish size
 - Disease vectors



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- **Future RMS projects:**

- WGS
- Vaccine development

- ETC!



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The end

