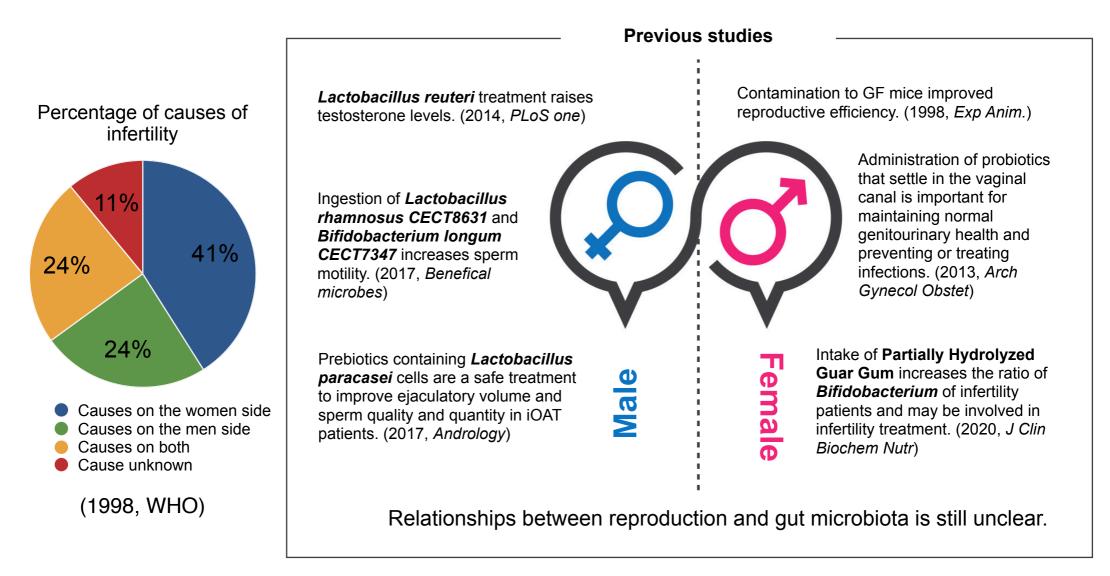
Elucidation of the mechanisms of gut microbiota-sex hormone interaction

82024683, BI program, Haruno Takahashi

Background / Aim of study

[Problem] About **186 million** people around the world are currently suffering from **infertility**.



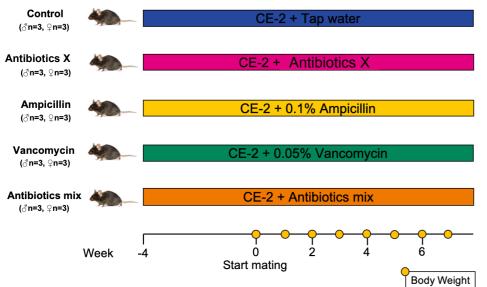


Clarify the effect of gut microbiota on reproductive function.

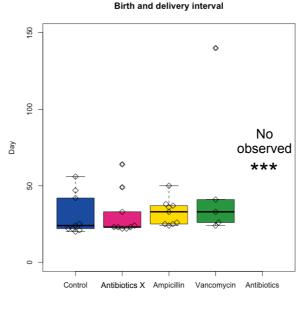
Antibiotic treatment delayed the birth-delivery interval

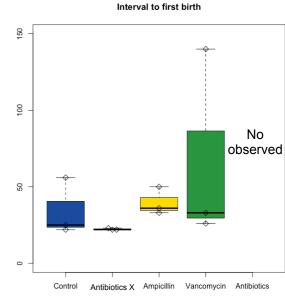
Reproductive efficiency evaluation test

@Tsuruoka CNV; free intake of CE-2 (n=3 pairs)

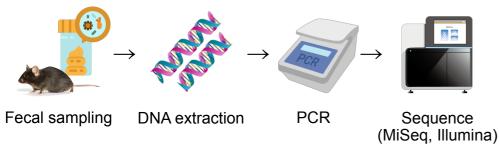


Birth and delivery intervals

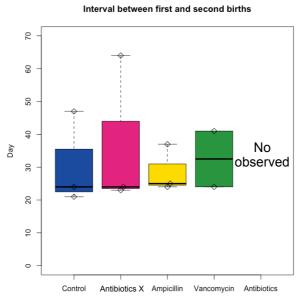


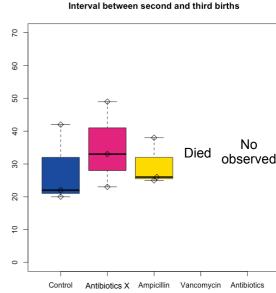


16S rRNA gene analysis

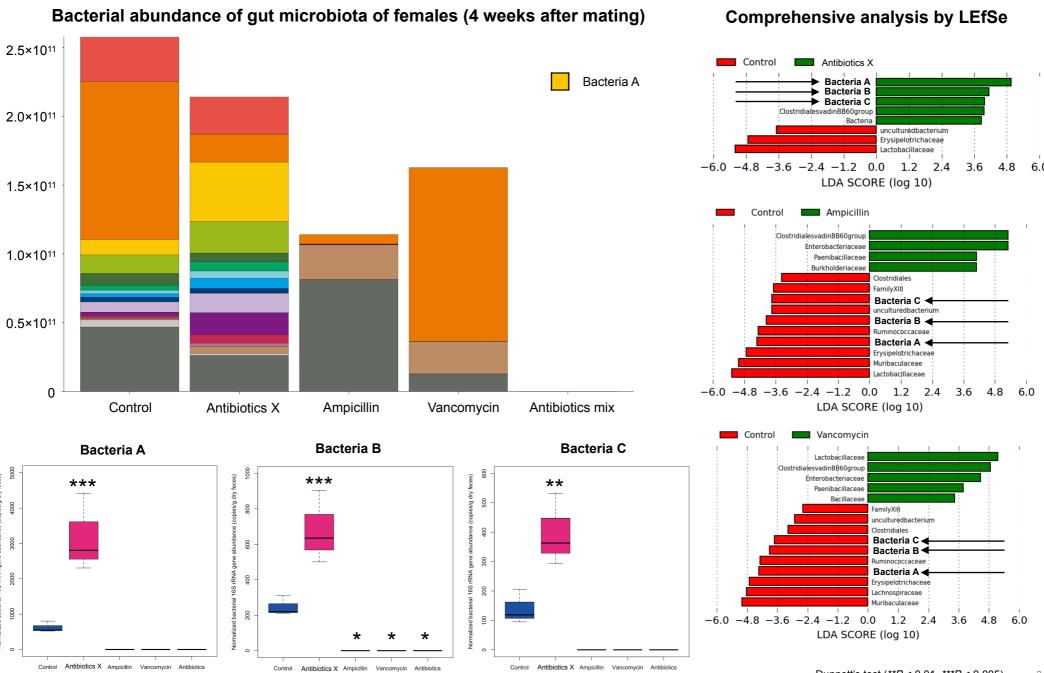


Searched for gut bacterial candidates associated with reproductive efficiency.

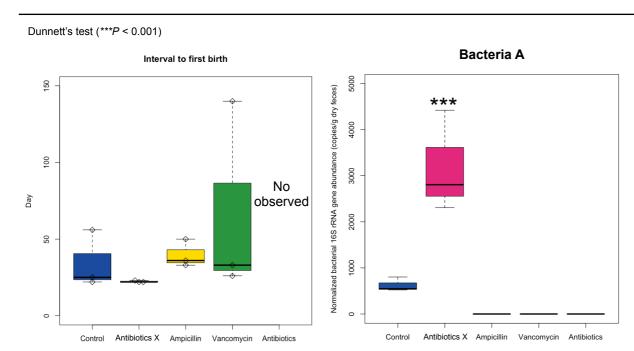




Administration of antibiotics X significantly increased bacteria A



Bacteria A may promote pregnancy in mice



Suggestion baed on the results

- ✔ Antibiotics treatment delayed the birth-delivery interval.
 - → Dysbiosis of gut microbiota leads infertility in mice.
 - → Pregnancy may be accelerated by administration of antibiotics X.
- ✓ Administration of antibiotics X significantly increased bacteria A, B and C.
 - → The amount of these bacteria may be related to the reproductive function.

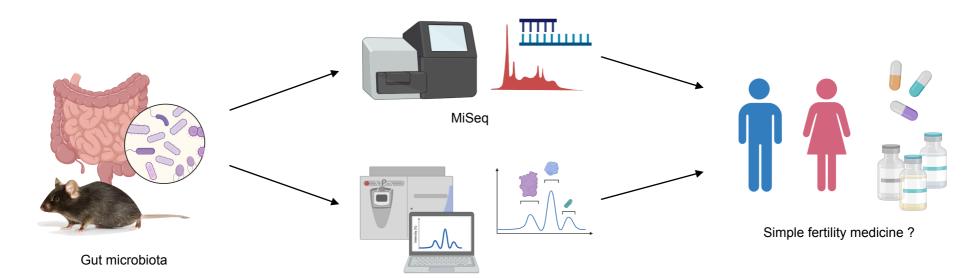


Track more detailed effects of gut microbiota on reproductive function.

Future plans

Biomarker analysis Aged mice antibiotics X treatment experiment ✓ Gut microbiota analysis Control CE-2 + Tap water ✓ Metabolome analysis (♂n=5, ♀n=5) **Antibiotics X** CE-2 + Antibiotics X (♂n=5, ♀n=5) ✓ Measurement of sex hormone Antibiotics treatment Week -2 2 -3 Gene expression analysis Start mating

Expected outcome



Capillary electrophoresis-mass spectrometry

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