



Reputation at Risk: How Animal Welfare, Antimicrobial Resistance and Social Responsibility Are Shaping Consumer Perception

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Antimicrobial resistance and aquaculture

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Power of Collaboration

AMR is a global issue

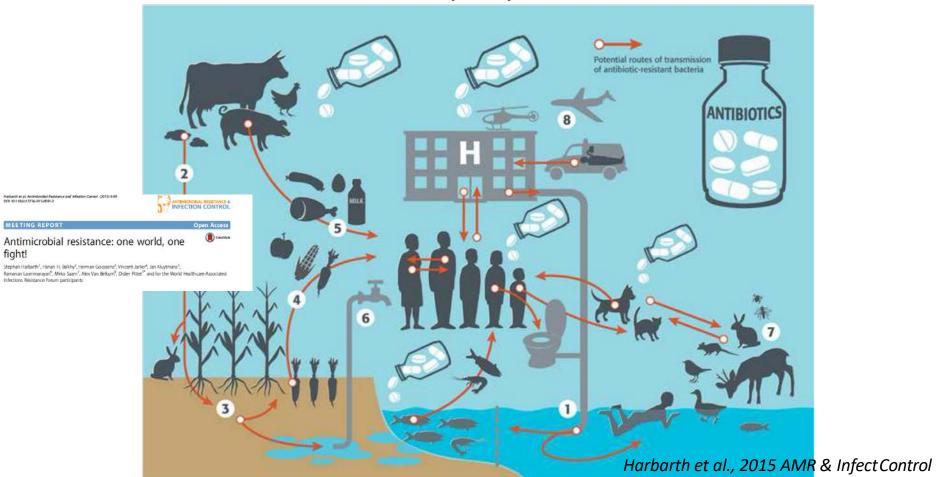




TACKLING ANTIMICROBIAL RESISTANCE:

Supporting national measures to address infection prevention and control and water, sanitation and hygiene in health care settings Antimicrobial resistance (AMR) presents a significant threat to human health. World leaders have agreed that tackling AMR will require addressing both health and agriculture concerns with a focus on prevention. Improving infection prevention and control (IPC) and water, sanitation, and hygiene (WASH) is one of the five objectives in the World Health Organization's (WHO) AMR Global Action Plan. Nowhere is reducing infection more important than in health care facilities. Joint, immediate action to address IPC and WASH is essential.

One health perspective



Characteristics of antibiotic use (ABU) in aquaculture

- Aquaculture systems are complex and dynamic, with many factors driving the (mis)use of antibiotics
- ABU in aquaculture is different to that in humans and livestock, since it is administered directly in to the aquatic environment usually through medicated feed
- Antibiotics and feed are often supplied separately in aquaculture. This
 means that farmers have to mix the antibiotics with the feed, resulting
 in a high risk of occupational exposure
- Fish do not metabolise antibiotics effectively, and it has been estimated that 75% of the antibiotics fed to fish will be excreted back in their active form in to the aquatic environment through faeces
- Aquaculture systems are often linked to the natural water environment, facilitating contamination of the environment and waterways, and having the potential to lead to the emergence and dissemination of ABR further than terrestrial livestock systems

Growth of food animal production > thirty years from 1980 and aquaculture sector in 2014

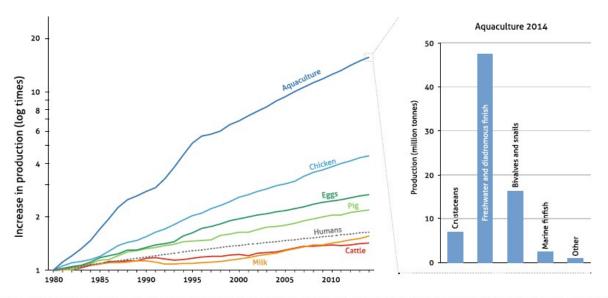
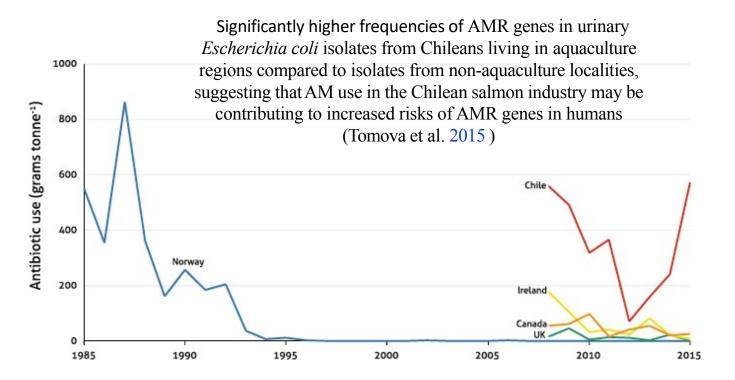


Fig. 1 Proportionate growth of the human population, aquaculture, poultry, pigs, and cattle between 1980 and 2013, and the present composition of the aquaculture sector *Source* (FAO 2016a; The World Bank 2017; FAO 2016b)



Antibiotic use in Atlantic salmon farming- major producers



Elevated AMR in bacteria from farmed fish

Briver September 10 intl1 intl2 blaSHV blaOXA aac(6')-lb-cr blaCTX-M

Rivers and lakes in Pune, India

ARG biomarkers farming

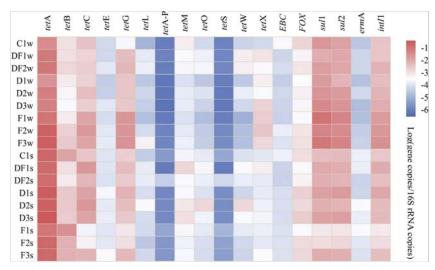


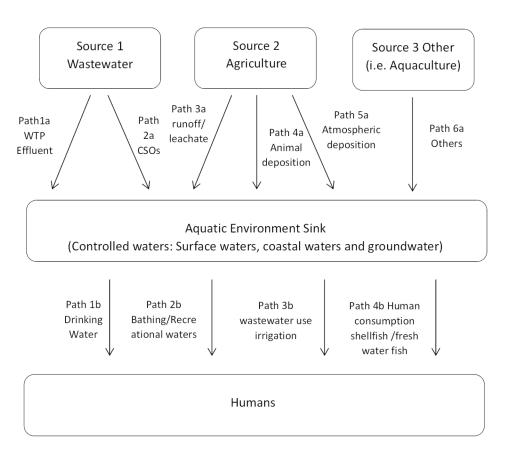
Fig. 3. A "heat map" of resistance genes in eighteen samples. Sample names were abbreviated with letters representing pond and sample type: capital C, DF, D and F (control pond, duck-fish pond, duck pond and fish pond) and lower case w and s (water sample and sediment sample). The different colored cells represent the Log (relative abundance) values of genes. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

Pearl river delta, South China Huang et al., Chemosphere 2018

Marathe et al. J. Biosci. 41(3), September 2016, 341–346 Indian Academy of Sciences 341

Report for Defra UK Gov

Systems Map AMR/ARGs Flows



James et al., 2018 draft report commissioned by Defra UKGov

Thames Investigation

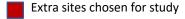
- -69 sites located along the Thames catchment were chosen for sampling.
- -Sampling begins at the source of each tributary and continues along the tributary where potential sources of AMR (eg. WWTPs and fish farms) are located.



	qPCR Targets
Gene Target	Resistance to/Marker
16S rDNA	bacterial marker
Intl1	class 1 integron
QacE	quaternary ammonium compounds
TetM	Tetracycline
RuBac	Ruminant assoc. Bacteroides
HuBac	Human assoc. Bacteroides
QnrS	fluoroquinolones
ErmF	erythromycin (macrolide)
CTX-M (Group 1)	CTX-M-1 and 15 (beta lactams)
E. coli	faecal contamination
Enterococci	faecal contamination

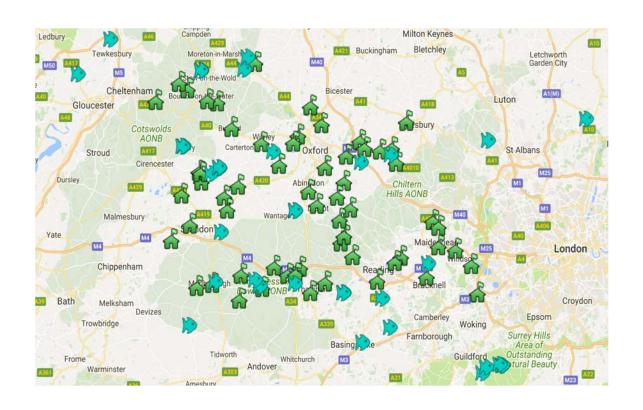
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Routinely monitored by CEH



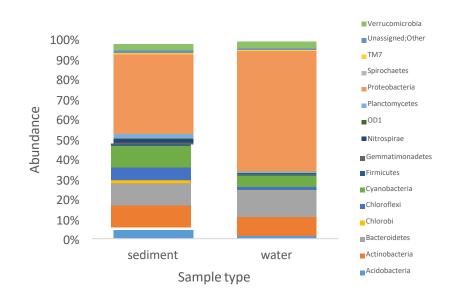
- -DNA was purified from triplicate samples and qPCR was used to determine resistance loads at each site.
- -16s rDNA amplicon sequencing on all sites
- -Metagenomes on 20 core sites

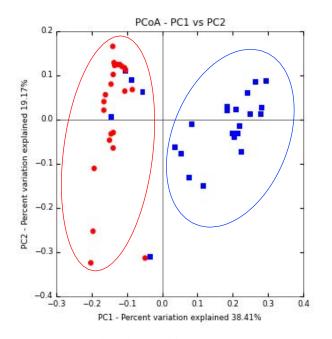
Location of fish farms and wastewater treatment plants



Sampling Planktonic and Benthic Phases

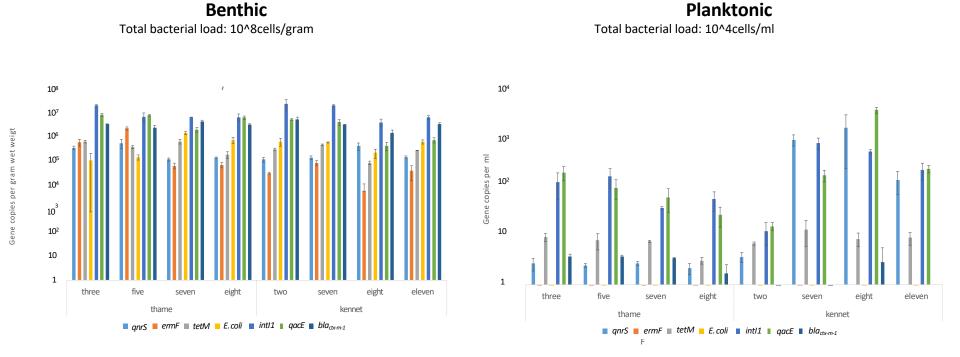
- -Planktonic and benthic phases differed significantly in phyla abundance.
- -Benthic phases displayed higher bacterial diversity compared to planktonic phase





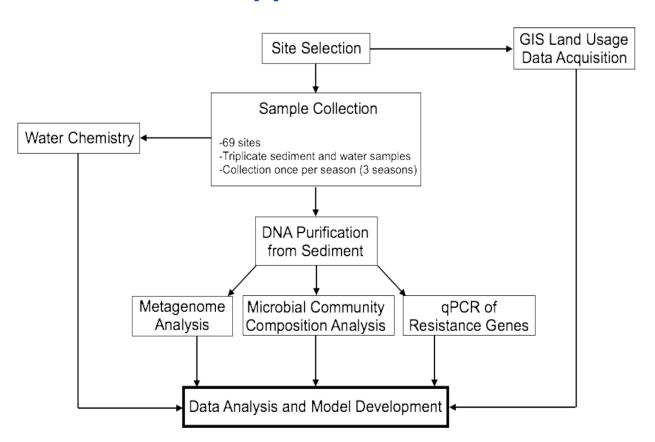
Red=sediment; blue= water

Benthic phase contains higher bacterial loads



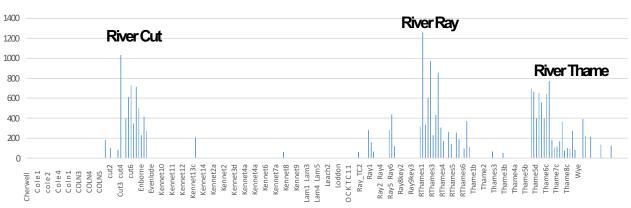
Benthic phase constitutes a greater reservoir for ARGs than planktonic phase

Approach

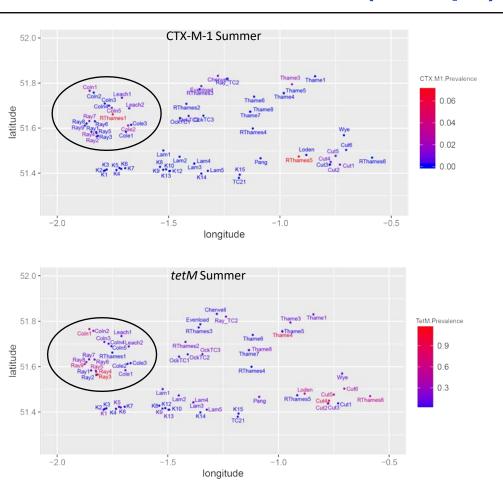


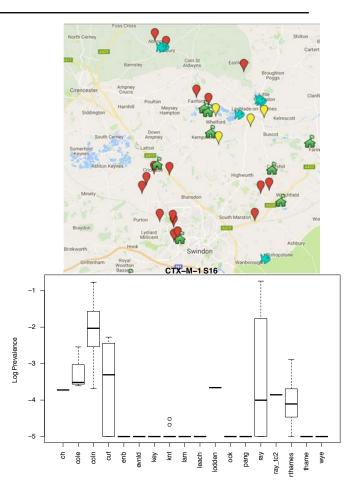
Summary of abundance of organic pollutant analytes Three seasons

Organic Pollutant Analytes

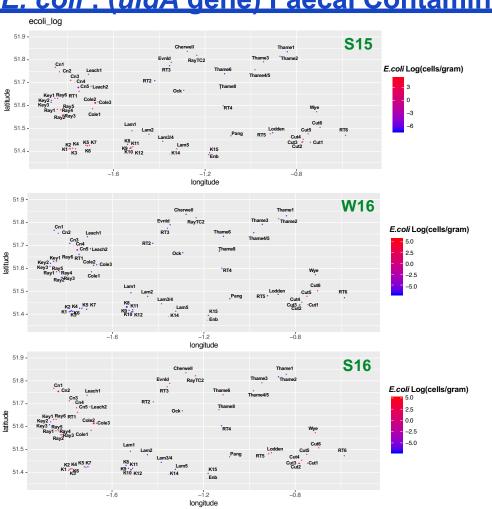


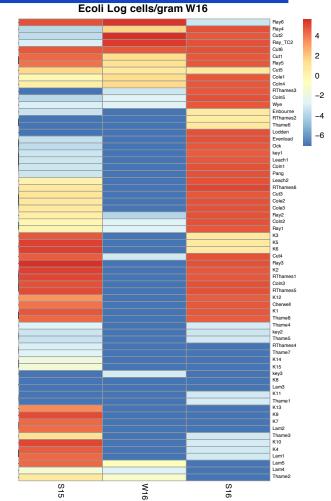
Prevalence of CTX-M-1 (Group1) / tetM

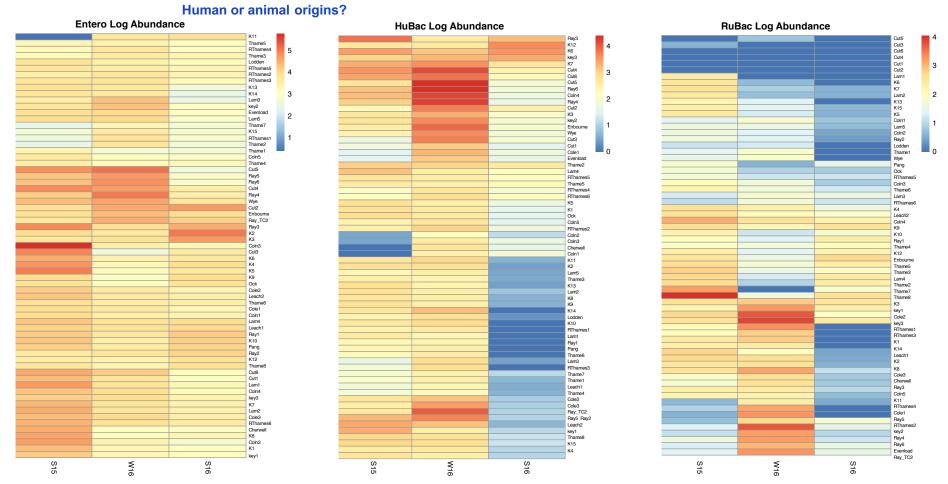




E. coli: (uidA gene) Faecal Contamination Indicator

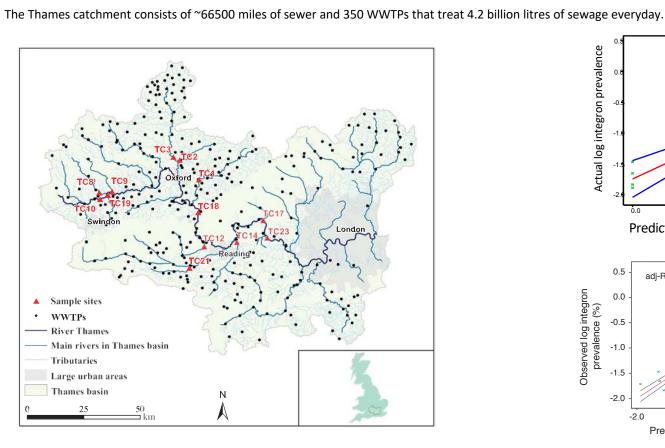


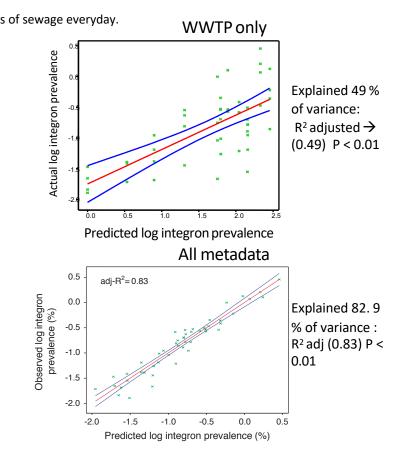




HuBaC and RuBac: Seeing differences between abundance of each within various tributaries

Previous model work on the Environmental Resistome: Thames Catchment





Amos et al., 2015. The ISME Journal, 9, 1467–1476

In conclusion

Alternatives to ABU

Vaccines*
Probiotics*
Improved microbiomes
Animal husbandry
Improved sanitation
Identification of drivers of AMR

*Shrimp farming, viral diseases white spot syndrome virus (WSSV) use GMO *Bacillus subtilis* spores that display the VP28 capsid protein of WSSV and when administered in feed appears to protect against white spot disease. Protective mechanism unclear; shrimp are not thought to produce antibodies, but presentation of the viral antigens does produce some level of specific immunity. *Hoelzer et al., Vet Research 2018*

Improving the natural defences:

initiate innate and subsequent adaptive immune responses, e.g. triggering the host's pattern recognition receptors

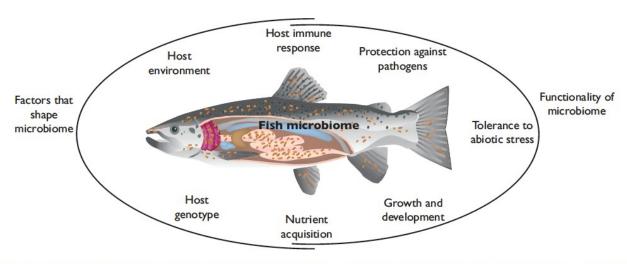
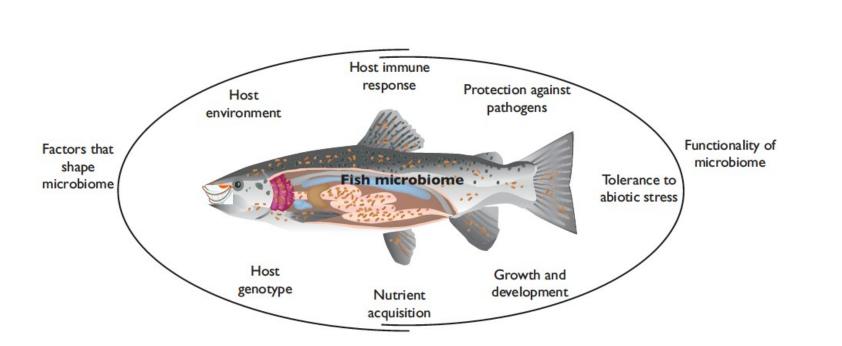


Figure 1. Schematic overview of different life-support functions of the fish microbiome and the impact of the host and ambient environment on the microbiome diversity, assembly and functions.



RESERVOIRS



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