



Antiviral defense systems drive persistence of antimicrobial-resistant bacteria but limit the transfer of antimicrobial resistance genes in anaerobic digestion

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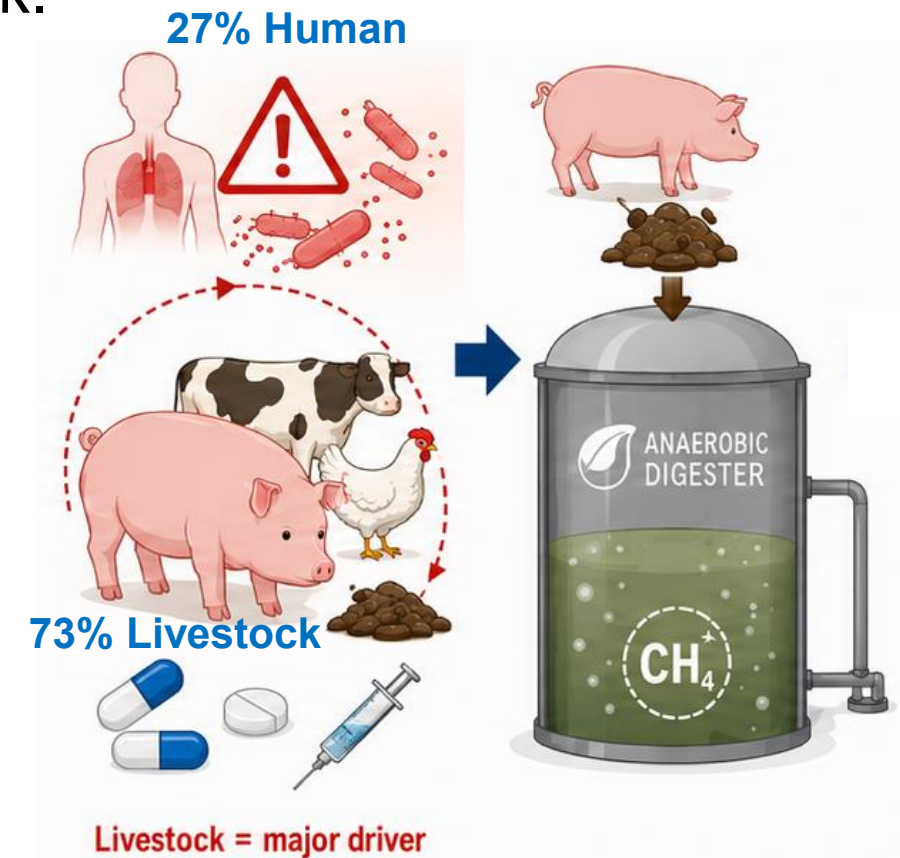
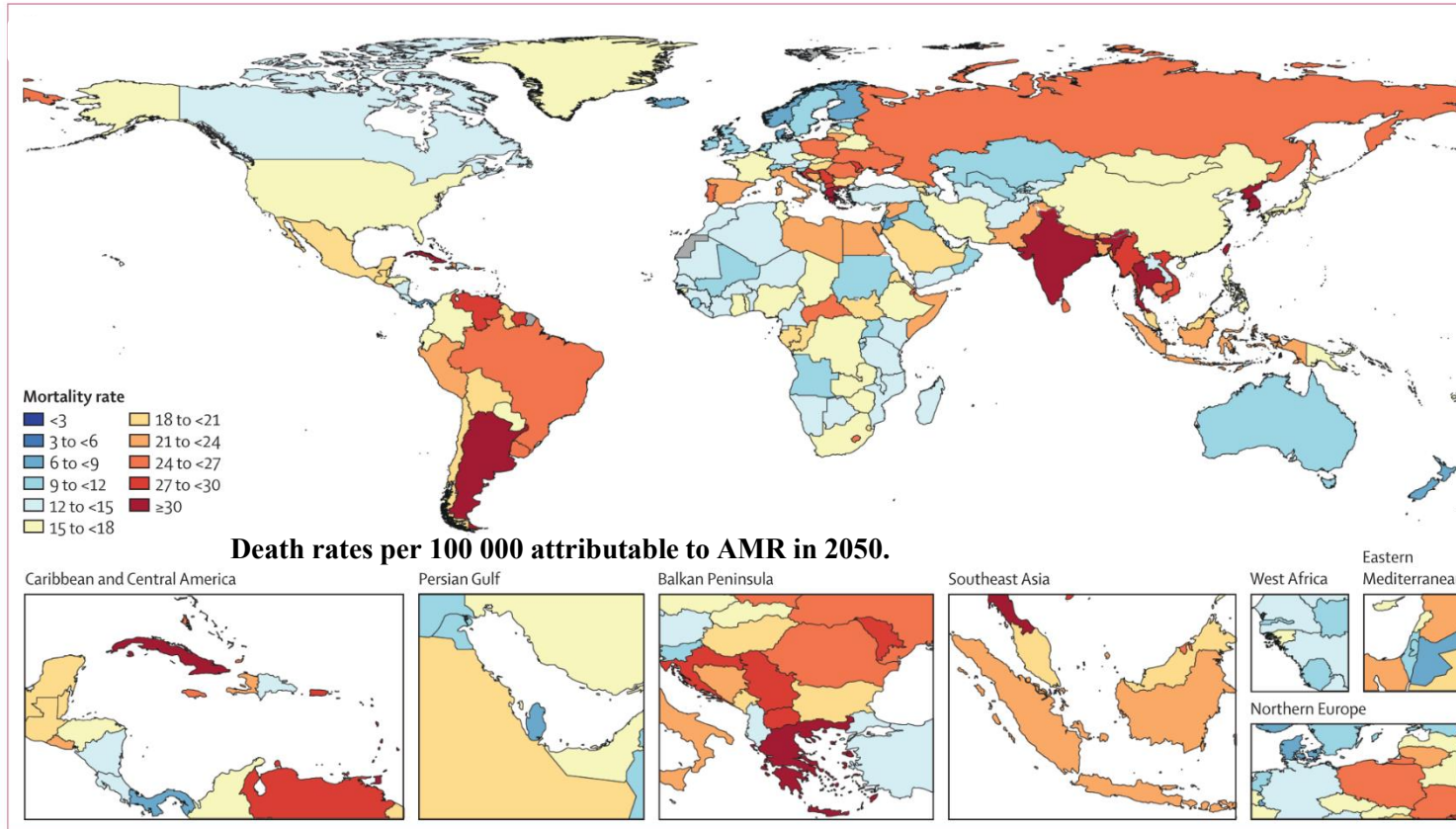


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Background

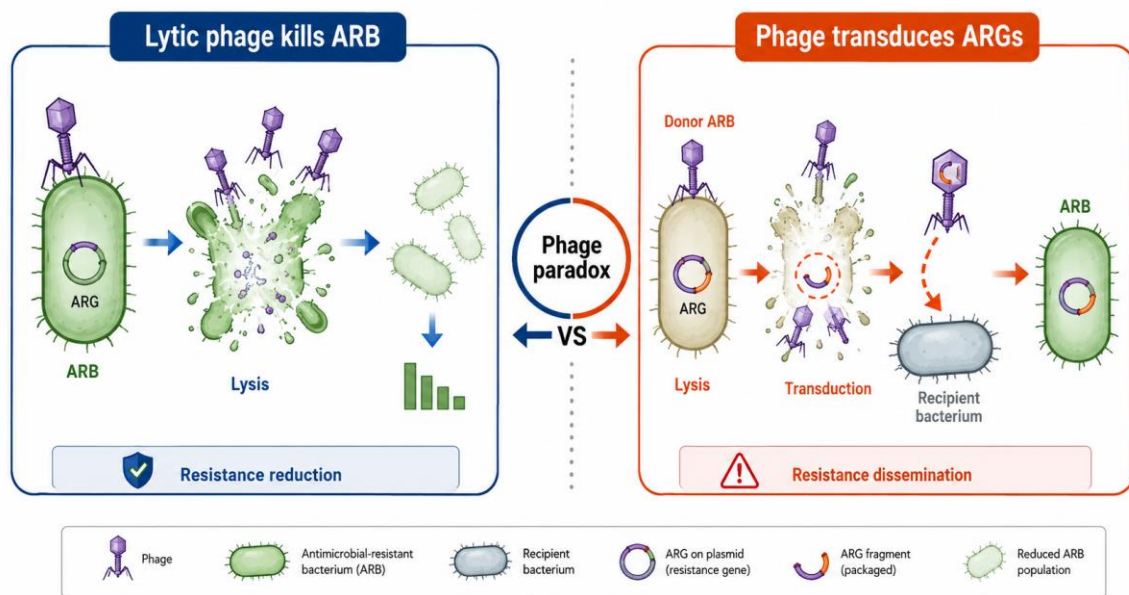
- **Antimicrobial resistance (AMR) is a severe "One Health" crisis**, associated with approximately 4.71 million deaths globally in 2021.
- Approximately **73% of global antibiotics are used in livestock farming**, playing a crucial role in the One Health framework.





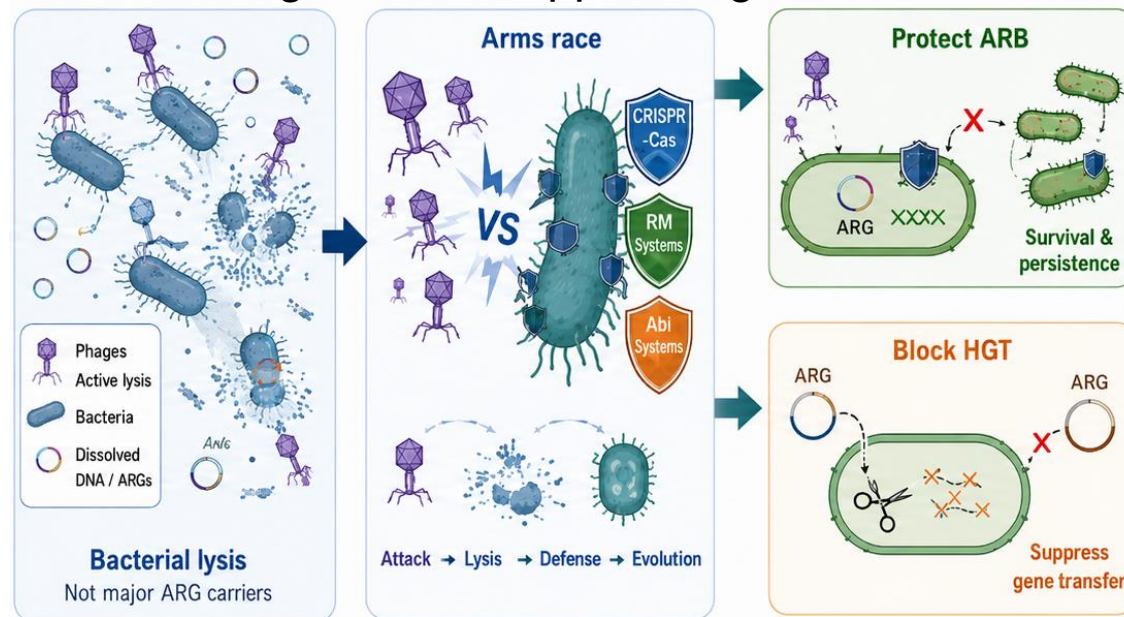
Background

□ The paradox of phages in AMR transmission: Lysis vs. Transduction



Which one dominates?

□ The duality of ADS in AMR transmission: Protecting ARB vs Suppressing HGT

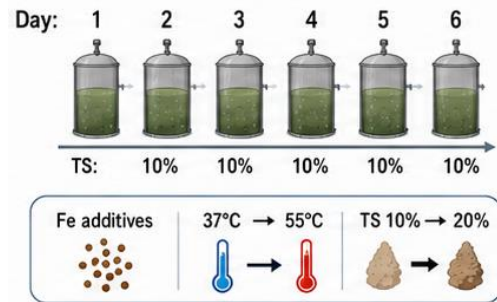


To be verified

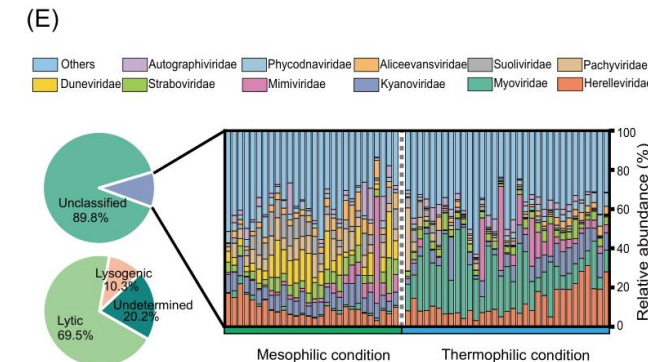
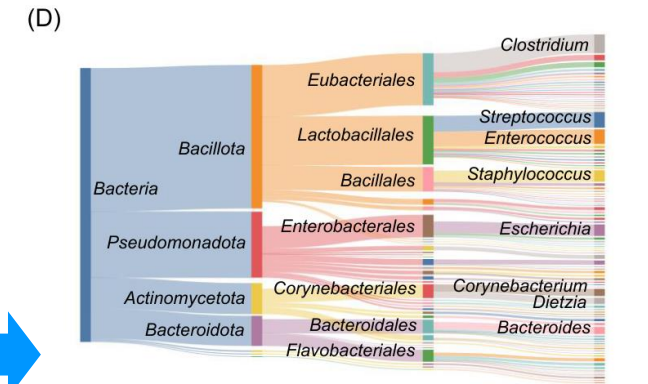
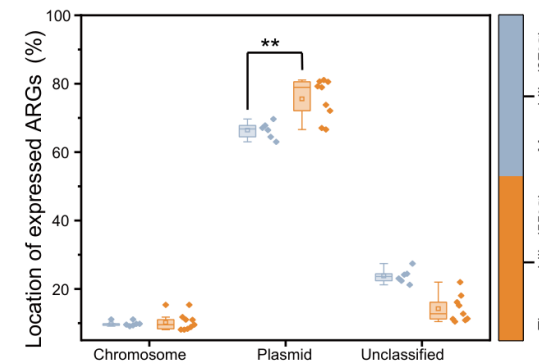
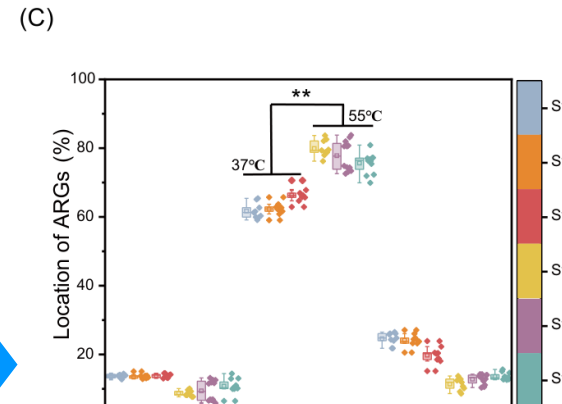
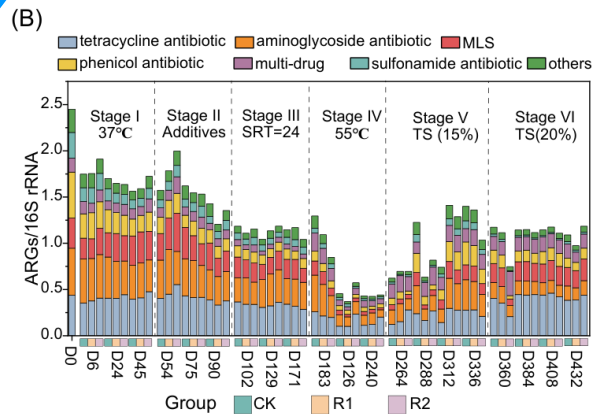
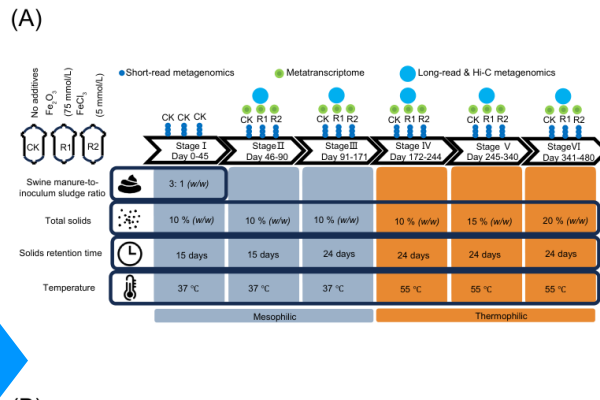
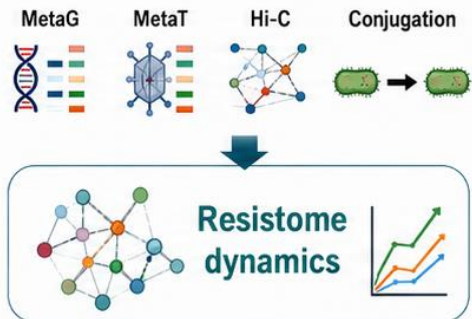


Lysis overwhelmingly outweighs transduction

440-day semi-continuous swine manure AD experiment



Multi-omics & validation



- 440-day semi-continuous anaerobic digestion
- Multi-omics and Hi-C
- Validation via qPCR and conjugation experiments
- Significant changes in ARGs with operational variations.
- Thermophilic conditions enhance ARG reduction, but increased TS leads to enrichment.

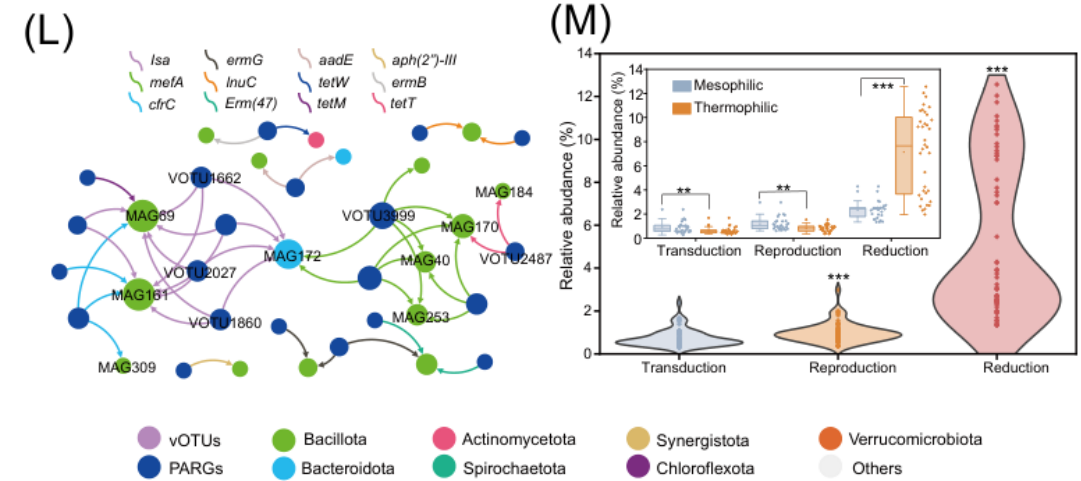
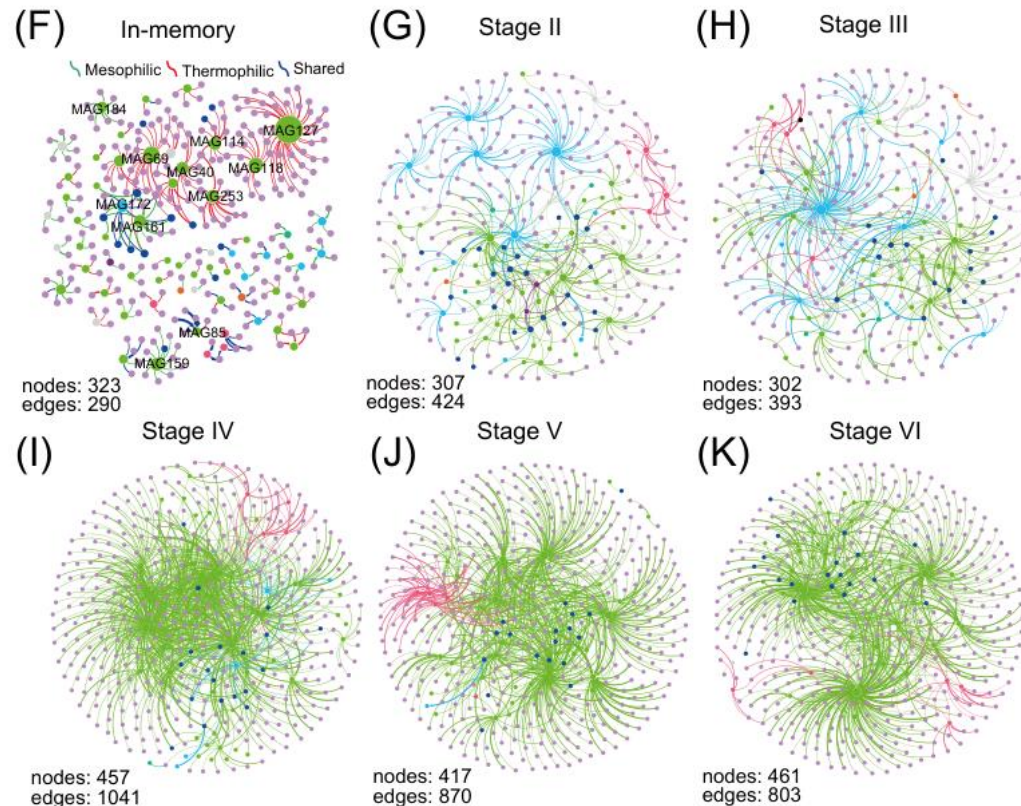
- 70.5%–77.6% of ARGs are located on plasmids.
- Most ARGs are silent; expressed ARGs are primarily plasmid-borne.

- Bacillota* are the main ARG hosts, but actively expressed ARGs are concentrated in *Pseudomonadota*.
- Diverse unknown viral communities exist, with 69.5% being lytic phages.

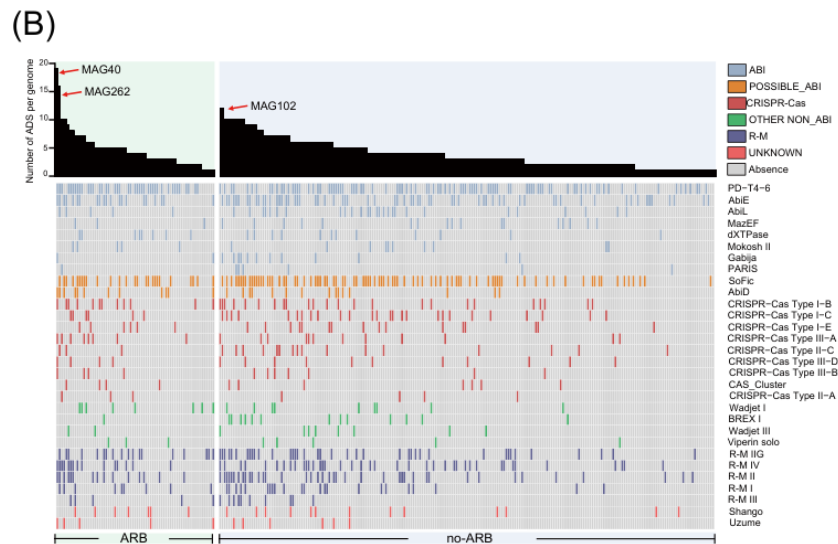


Lysis overwhelmingly outweighs transduction

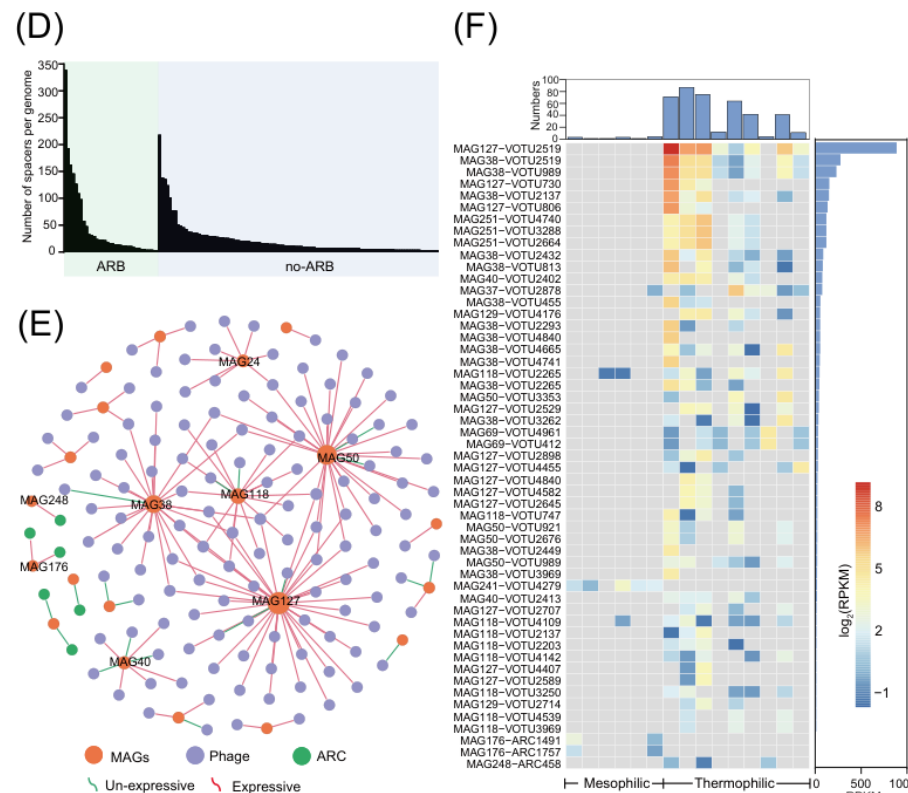
- Reconstructed historical (In-memory) phage-host interactions based on CRISPR-Cas, tRNA, and sequence homology.
- Reconstructed on-going phage-host interactions at different stages based on Hi-C.



- Identified 26 vOTUs carrying ARGs, accounting for $0.55 \pm 0.32\%$ of total ARGs and $2.56 \pm 1.04\%$ of the viral community.
- Detected 45 transduction events, including cross-phylum transfers.
- The abundance of phages driving reduction ($5.16 \pm 3.54\%$) was significantly higher than those involved in transduction ($0.74 \pm 0.39\%$) and reproduction ($0.99 \pm 0.45\%$) ($p < 0.001$).
- Thermophilic anaerobic digestion significantly weakened the transduction and reproduction capabilities of phages for ARGs, but remarkably enhanced their reduction effect.



- On average, **each ARB carries 4.5 ADS, significantly higher than the 3.7 in non-ARB** ($p < 0.05$).



- Meanwhile, CRISPR-Cas also targets ARG-carrying contigs, **indicating its inhibitory role in the horizontal transfer of ARGs.**

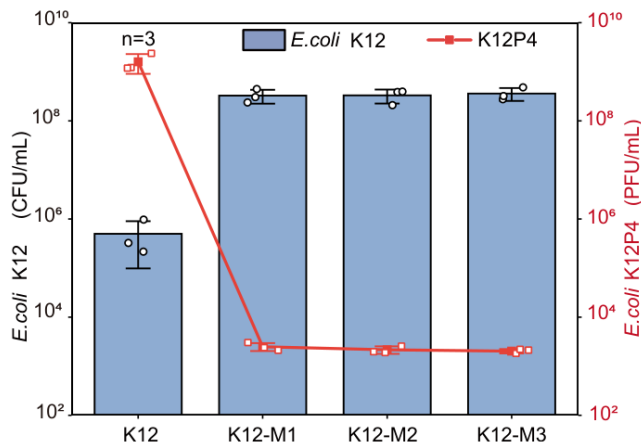


The dual role of antiviral defense systems

Donor:

Verification of antiviral defense *E. coli* K12 strains

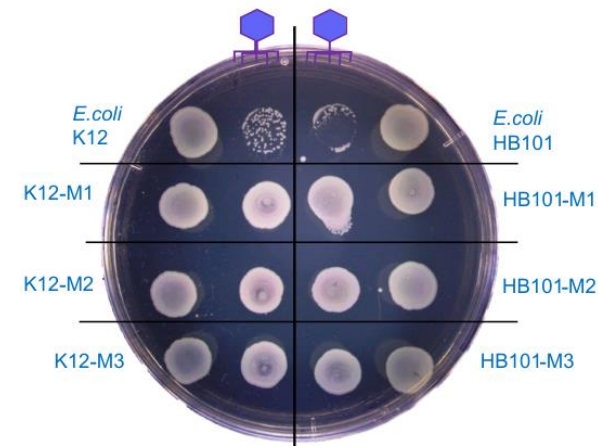
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Phenotypic verification of

antiviral defense strains

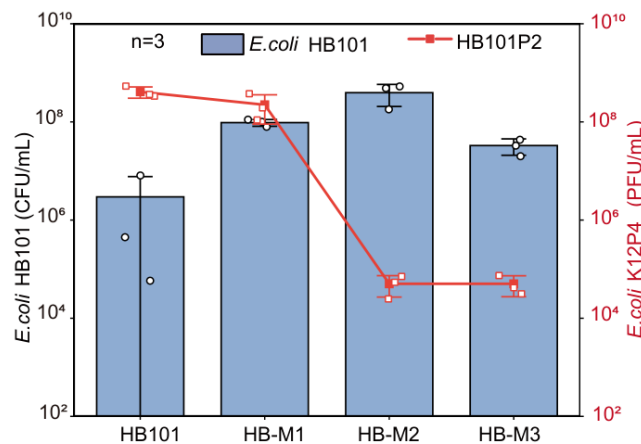
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Recipient:

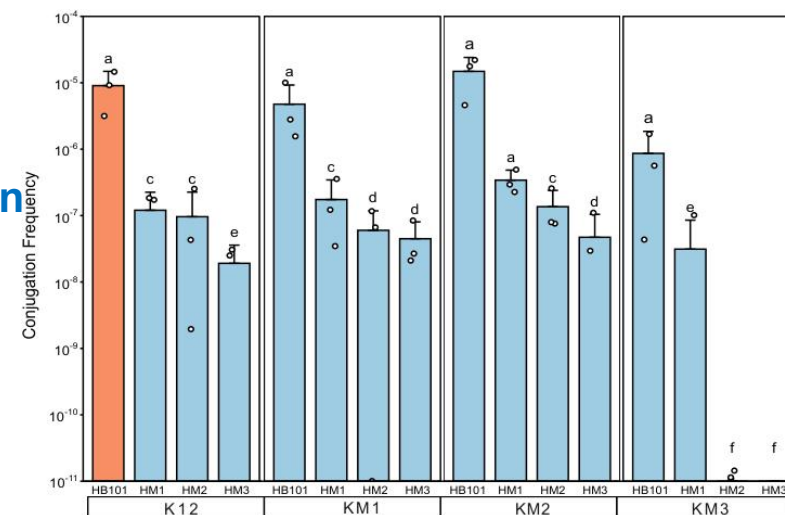
Verification of antiviral defense *E. coli* HB101 strains

(H)



Emergence of antiviral defense **significantly decreased conjugation frequency**

(J)



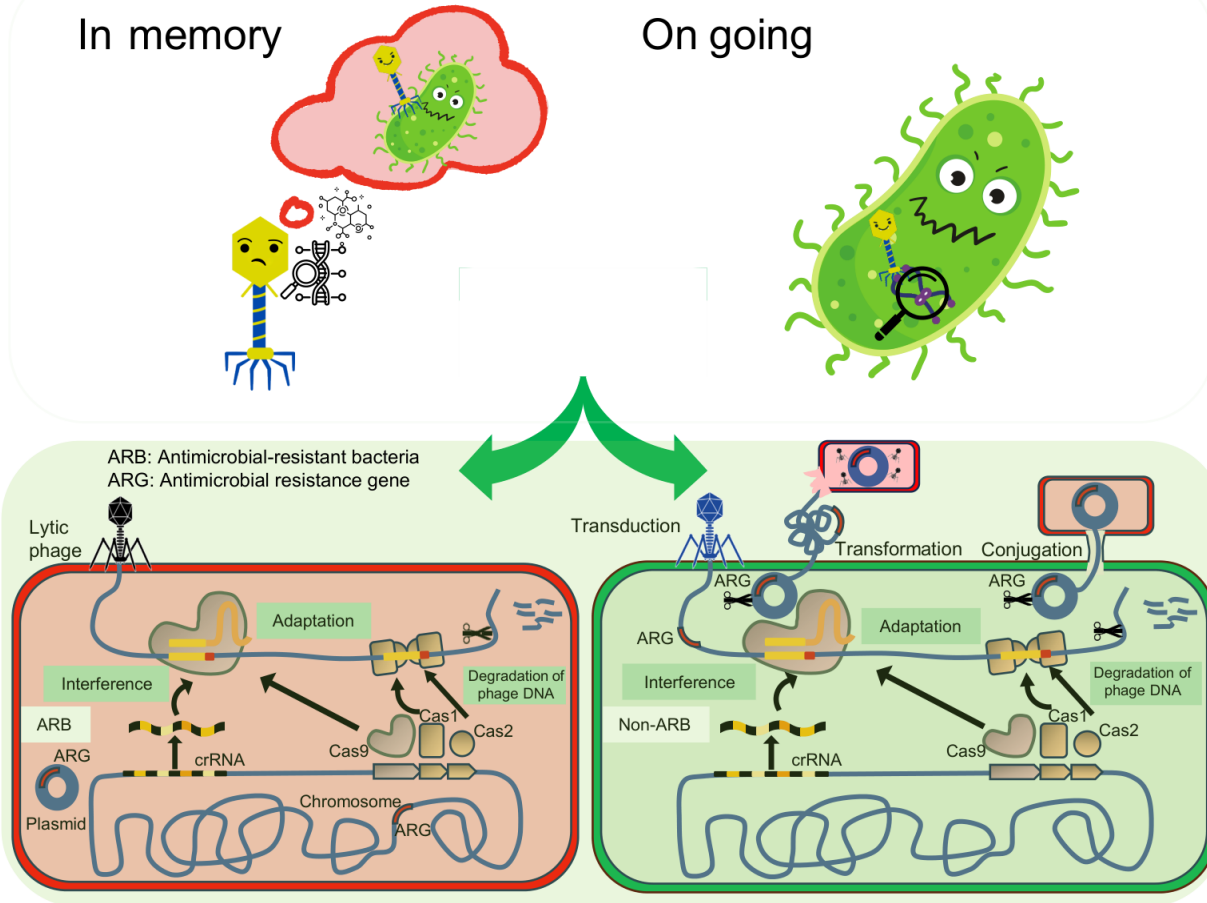
After the emergence of phage immunity, compared to the wild-type pairs, **the frequency of acquiring the RP4 plasmid via conjugation significantly decreased by 98.7%–99.8%** (dropping from 9.05×10^{-6} to 1.91×10^{-8} , $p < 0.05$).



Take-home messages

In memory

On going



- By integrating short/long-read metagenomics, metatranscriptomics, and *in situ* Hi-C interactions, combined with *in vitro* conjugation validation, **we systematically resolved the impact of the phage-host arms race on the fate of ARGs in anaerobic digestion.**
- We discovered that the proportion of phages carrying ARGs is extremely low, and transduction events are rare; **phages act more in reducing ARGs rather than as their transmission vectors.**
- **This study revealed the dual ecological role of ADS:** on one hand, protecting ARB from phage lysis and promoting their long-term persistence; on the other hand, inhibiting plasmid-mediated horizontal transfer of ARGs.

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