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- Humans in urban environments spend near 90% of their time inside buildings.
- Usage of antimicrobial chemicals is widespread.
- resistance genes (ARG) in pathogenic bacteria, as seen in culture.
- (MGE) in the dust microbial community remains largely unknown.
- Research questions: Is there evidence of transfer of ARG via MGE in dust? Does the rate of
- transfer increase in the presence of antimicrobials?

#### RESULTS Abundance Antibiotic class Top 20 ARG (RPKM) Integron multidrug 2.29 CRP AAC(3)-VIIIa 2.15 aminoglycosides AAC(3)-VIIa aminoglycosides 1.71 1.54 erm(33) macrolides 1.37 multidrug mexT 1.26 tetQ tetracycline 1.16 multidrug emrR Plasmid aminoglycosides 1.13 acrB U 0.95 tetracycline tetW lincosamides InuA 0.81 0.88 multidrug срхА 0.87 tetO tetracycline multidrug 0.76 qacA penicillin SRT-1 0.82 Transposon -(betalactamase) 0.81 macrolides ermB others 0.78 dfrC (trimethoprim) 0.68 acrD aminoglycosides 10 0.62 vanRO glycopeptides macrolides ErmO 0.61 0.61 tetracylin tet41



## **OUTCOMES AND PERSPECTIVES**

- Based on correlations with  $\alpha$ -diversity, mobile ARG are likely carried by specific bacterial clades.
- Triclosan may foster ARG dissemination in dust.

# The built environment resistome: potential of indoor dust microbial community for dissemination of antibiotic resistance genes

It is expected that 66% of the world population will be living in urban areas by 2050.

• Antimicrobials are commonly found in dust and can trigger transfer of antibiotic

The effects of antimicrobials on the transfer of ARG through mobile genetic elements





• Sample size: 167 samples from 63 distinct buildings Collected measurements: total microbial DNA and antimicrobial concentration (triclosan, triclocarban and parabens).

### **Data processing:**

 Metagenomic sequences quality-controlled with Kneaddata. • Antibiotic resistance profiles in each sample were determined with ShortBRED. MGE and associated mobile ARG were defined from assembled and annotated metagenomic contigs with Kiki and RAST respectively. Pearson correlations were calculated with R.

Assays on dust bacterial isolates for triclosan resistance and triclosanrelated mobile ARG transfers (conjugation with plasmids carrying a gidb gene) are being performed to further confirm our results.

