Online supplementary figure S1

**Impact of Contrasting Poultry Exposures on Human, Poultry, and Wastewater Antibiotic Resistomes in Bangladesh.**

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**Supplementary Figure S1. Stacked bar charts showing the most abundant phyla and genera (relative abundances shown).** a) Phylum level breakdown of read compositions grouped by sample origin. The ‘other’ category includes reads belonging to bacterial phyla, archaea, eukaryote, and viral domains which sum to <1% in 38 of the total 40 samples. b) Genus level breakdown of read compositions grouped by sample origin. The ‘other’ category includes reads belonging to bacteria genera, archaea, eukaryote and viral domains which sum to <2% in 36 of the total 40 samples. Sample ID is displayed on the axis; (DL = urban wet market, TR= rural household, FL = low exposure human faecal, FH = high exposure human faecal, CL = poultry cloacal and WW = wastewater).