

Influence of Ventilation design on the prevalence of anti-microbial resistant bacteria in homes

Stage 1: Microbiology

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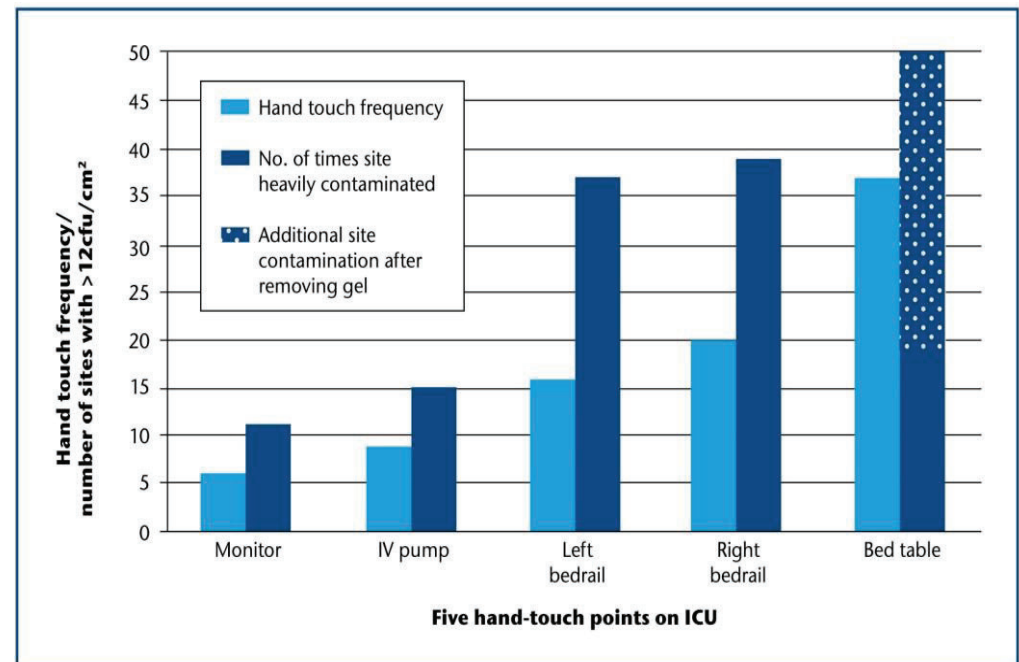


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Sites chosen for microbiological sampling

1. Bathroom (inside) Door Handle
2. Telephone
3. Kettle Handle
4. Bedside Table
5. Top of Door
6. TV Remote
7. Toilet Handle
8. Bedroom Window Sill

Figure 1: Hand touch frequency and gross microbial soil for five near patient sites on ICU



Dipslides

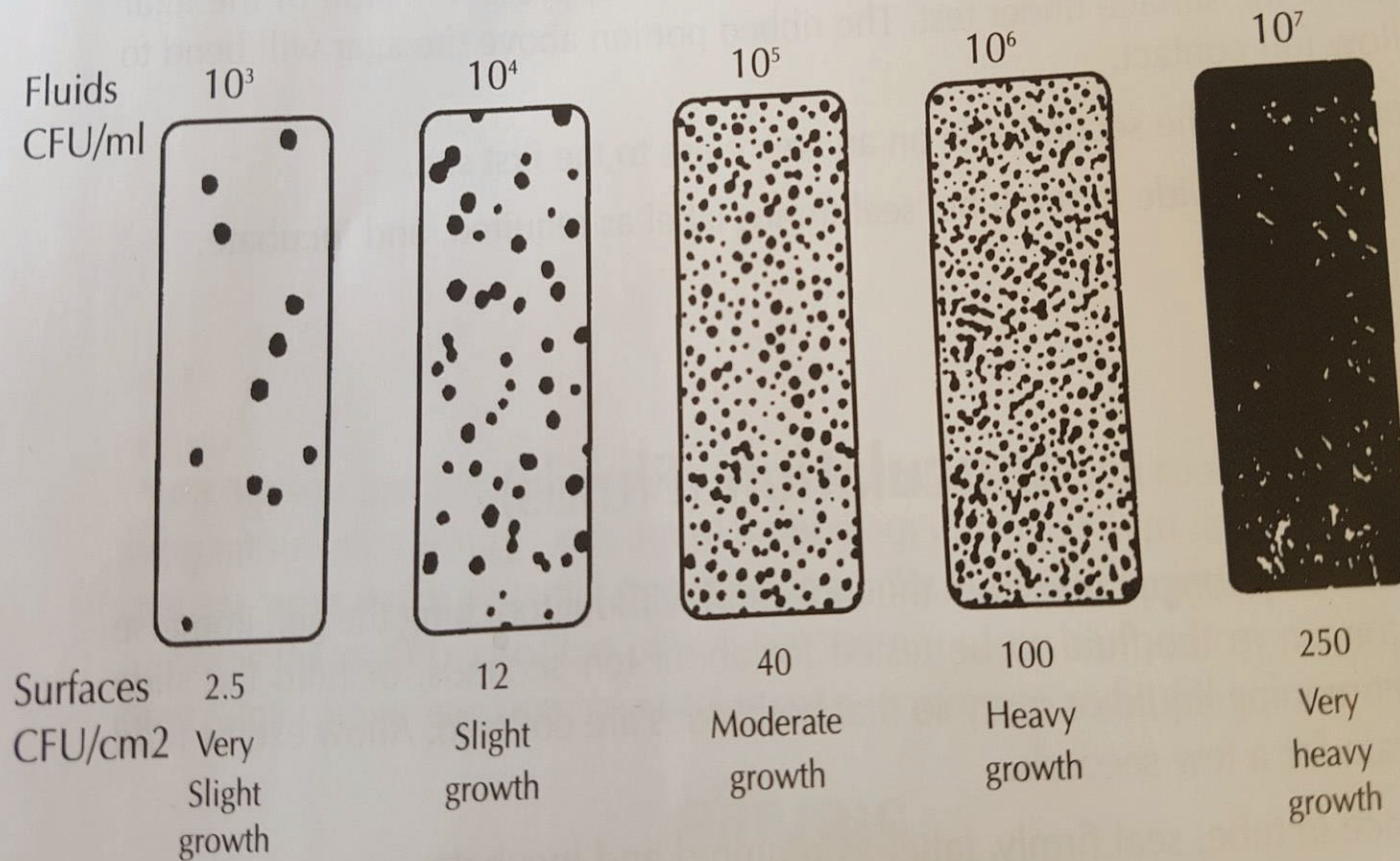


5 cfu/cm²



45 cfu/cm²

Comparison Charts Bacteria/Yeasts







Most common finding from all sites except top of the door



Site 5: Top of the Door



Site 8: Window sill



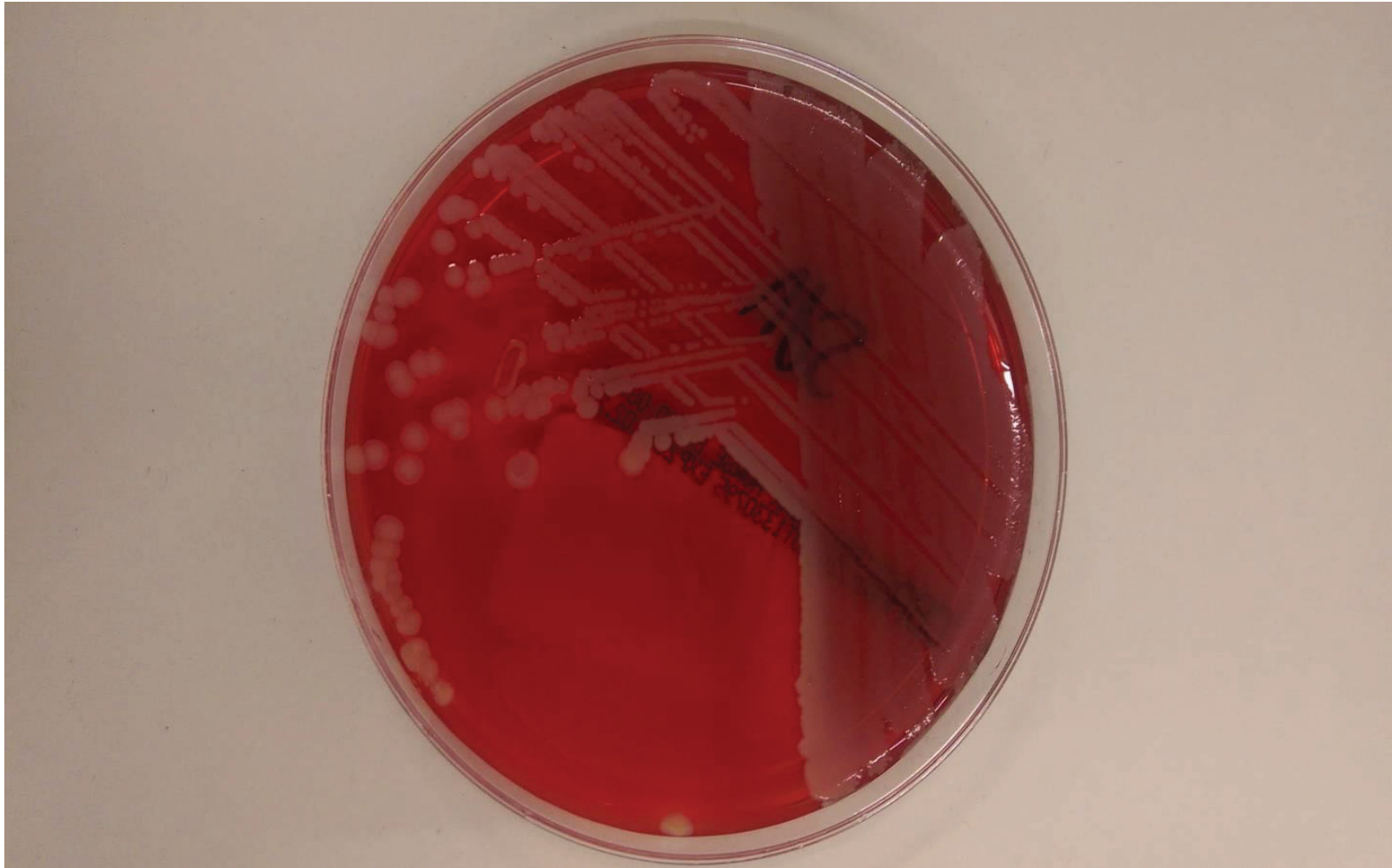
Site 7: Toilet handle



Site 1: Bathroom door handle (inside)



What's so special about *S.aureus*?



Baird Parker selective agar showing black colonies of *Staphylococcus* spp.



Site 3: Kettle handle



Sites 2 and 6: Telephone and TV Remote



Main microbiological characteristics of each site

Site 1 (**Bathroom door handle**): Staph, Micrococci, Gram neg, **Candida**

Site 2 (**Telephone**): Staph, Micrococci, Gram neg, ***S.aureus***

Site 3 (**Kettle handle**): Staph, Micrococci, Gram neg, ***S.aureus***

Site 4 (**Bedside table**): Staph, Micrococci, **Bacillus**, **Fungi**

Site 5 (**Door top**): **Bacillus**, Staph, Micrococci, **Fungi**

Site 6 (**TV remote**): Staph, Micrococci, ***S.aureus***, Gram neg

Site 7 (**Toilet handle**): Staph, **Gram neg** (20% *no growth!*)

Site 8 (**Bedroom window sill**): Staph, Micrococci, **Bacillus**, **Fungi**

***Staph aureus*; Gram-negative bacteria; and fungi**

Staph aureus: one or more sites positive in 35*/100 homes

Found predominantly on **kettle handle; TV remote; and telephone;**

No MRSA detected

Gram negative bacteria (mostly environmental): one or more sites positive in 70/100 homes; found predominantly on **kettle handle; TV remote; telephone; and bathroom door handle**

Fungi (filamentous) usually with *Bacillus* spp: one or more sites in 32/100 homes

Found predominantly on **door top; window sill; and bedside table**

Candida predominantly favoured bathroom door handle (site 1)

Most heavily contaminated site was the **top of the bedroom door;** the site most likely to yield 'no growth' was the **toilet handle!!**

**S.aureus is carried by approx. one third of humans*

Identification of cultivable Gram-negative bacteria

Most isolates identified as *Pantoea* spp. or *Pantoea agglomerans*

Also found:

Acinetobacter baumannii (standard resistance patterns)

Sphingomonas paucimobilis

Lerclercia adecarboxylata

Paracoccus yeei

Klebsiella pneumoniae (one isolate resistant to amoxicillin)

Enterobacter cloacae

Roseomonas gilardii

Acinetobacter lwoffii

Moraxella spp.

Methylobacteria

NB. Most awarded 'poor discrimination' by VITEK

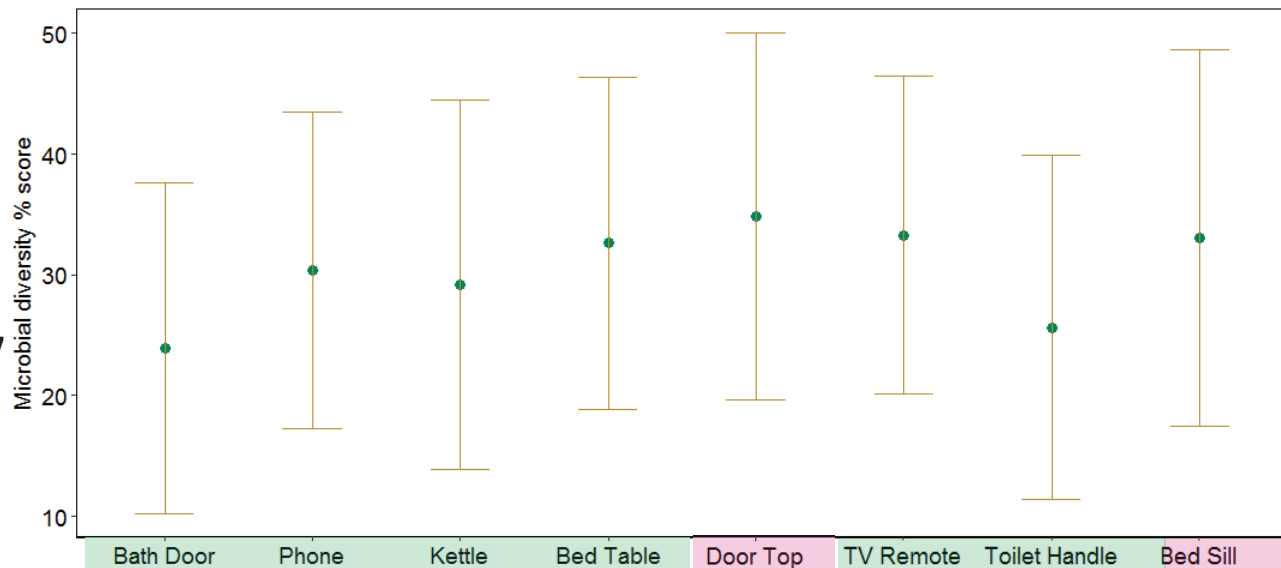
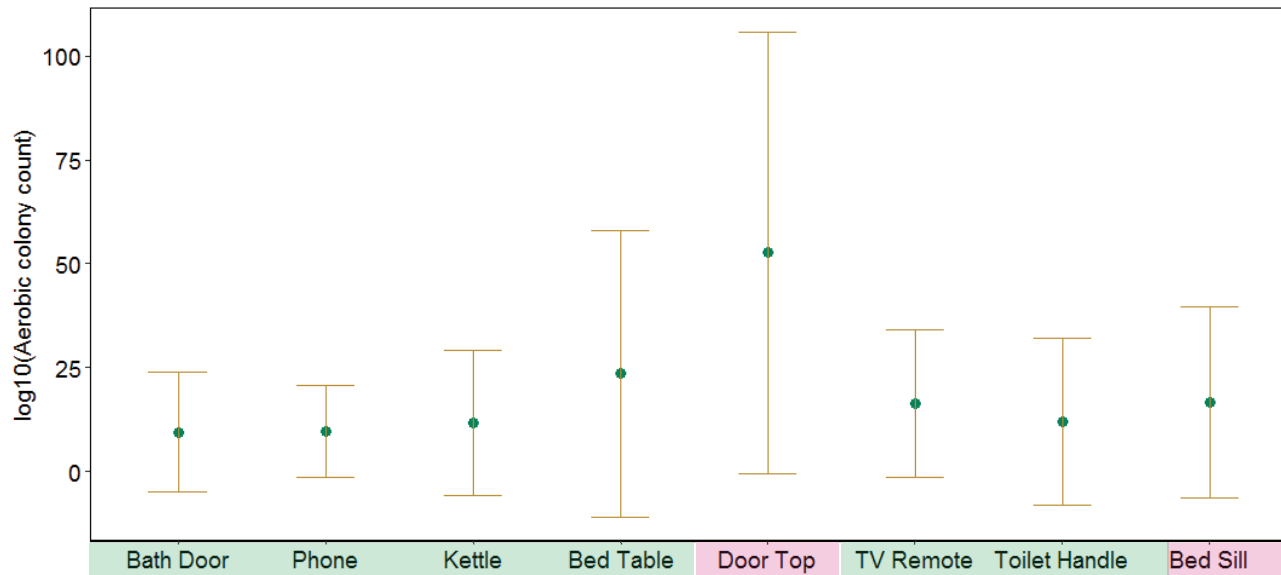
None were multiply resistant to antibiotics

Conclusions so far

1. Each site demonstrated common microbiological features;
2. Each home has its own unique microbiome;
3. Kettle handle, TV remote and telephone were most likely to host *S.aureus* and Gram-negative bacteria;
4. Fungi and Bacillus are most commonly found on bedroom door top, window sill and bedside table;
5. Yeasts were mostly recovered from bathroom door handle;
6. The site most likely to yield 'no growth' was the toilet handle!
7. While some bacterial pathogens were identified, none were multiply resistant to antibiotics

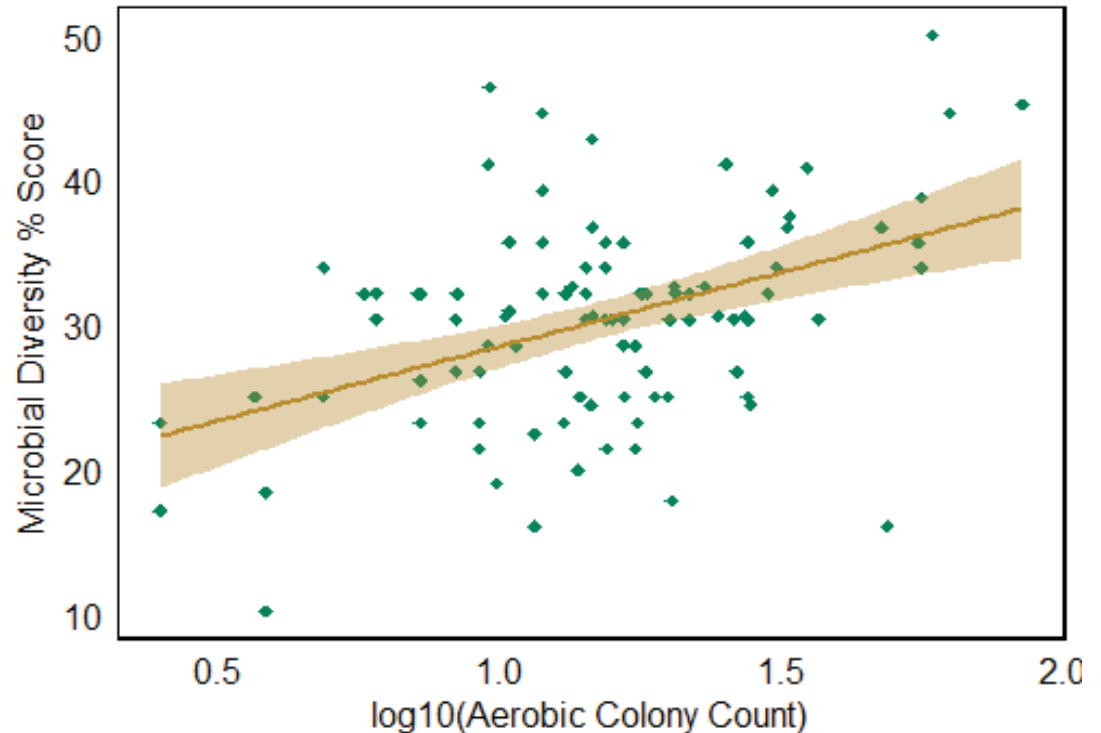
Microbial quantity and diversity

- Significant difference in microbial count between sites ($p < 0.001$) as well as **touch sites** vs **non-touch** sites (e.g. Door top vs Kettle)
- For each of the following: *Staphylococcus aureus*, *Micrococcus*, *Bacillus*, Fungi, Gram-positive cocci, Gram-negative rods and Gram-negative cocci a mark was given if they were found in a particular sample and the overall diversity score for that sample was calculated as a %.
- Bathroom surfaces had the lowest diversity ($p < 0.0001$) – mainly Gram positives
- Houses with young children showed higher ACC and higher diversity



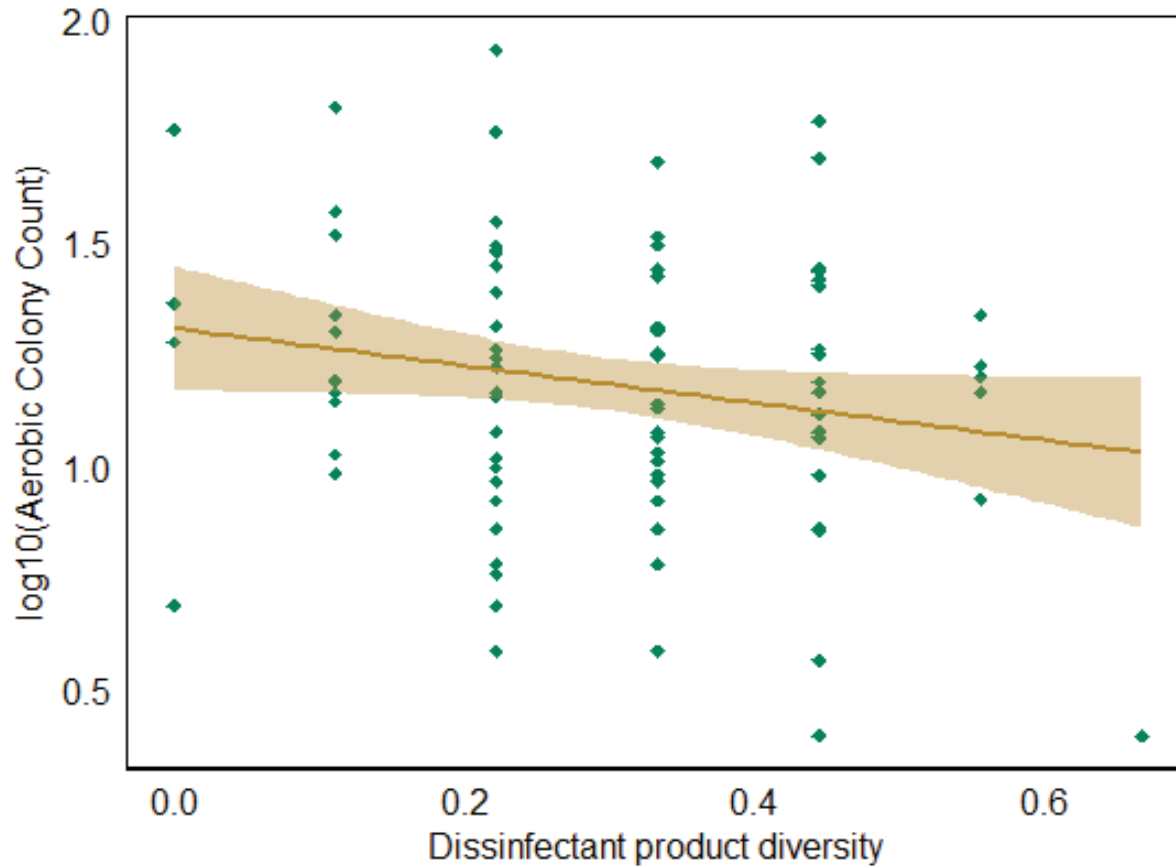
Microbial count and diversity

- As the microbial count increases at a site so does the range of species that are likely to be present ($p < 0.001$).
- Fungi presence correlated with increasing ACC count ($p < 0.05$) but presence of Gram-negatives did not correlate with increasing diversity.



Microbial diversity % score plotted against \log_{10} ACC

Disinfectant Range



Houses that use wider varieties of disinfectant products have a slightly lower ACC ($p < 0.05$), but there is no difference for bleach vs no bleach users.

Wider ranges of products also reduces chances of finding Gram-negatives. No effect on fungi

There were no statistically significant relationships between reported window opening, trickle vent usage or ventilation type and either the total ACC or with ACC from three specific sites in the bedroom: bedside table, window sill and door top.

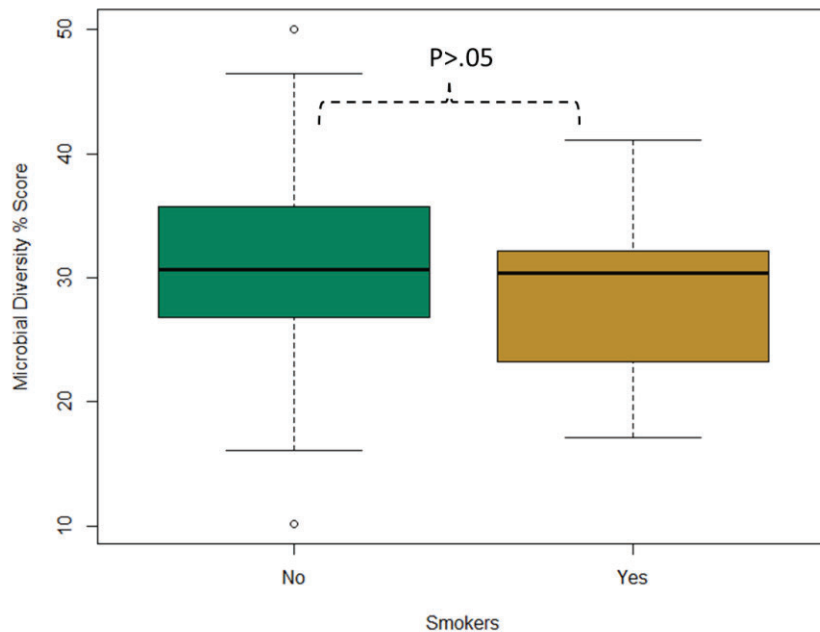
These sites were selected for analysis as they were considered to be the sites that might be most influenced by deposition of microorganisms from the air.

Heavy use of disinfectant may skew the findings for window opening, i.e. the tendency to have a contaminated home from never opening windows could be alleviated by lots of disinfectant use

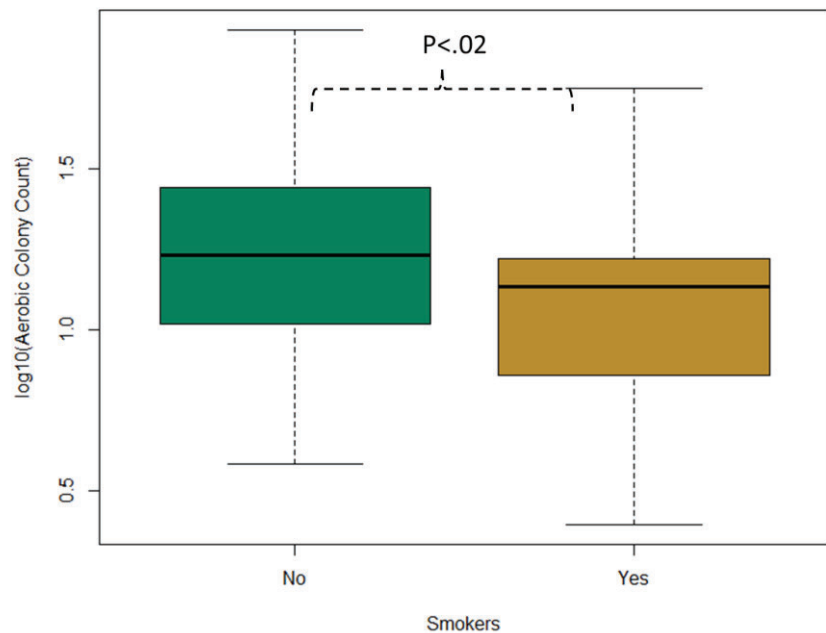


Smoking

- There is a reduction in overall microbial quantity in smoking households ($p < 0.02$) which seems to get stronger as the number of smokers increases ($p < 0.05$). But diversity cannot be distinguished nor is there a change in likelihood of finding Gram –ves (odds ratio = 1.7 , 95%CI 0.4-3.6)



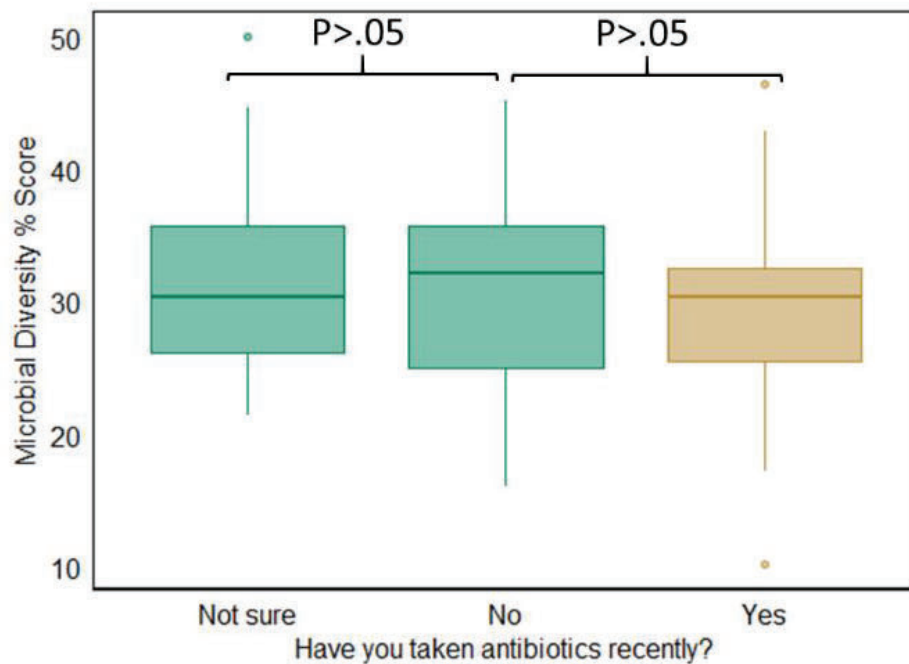
a) Microbial diversity score



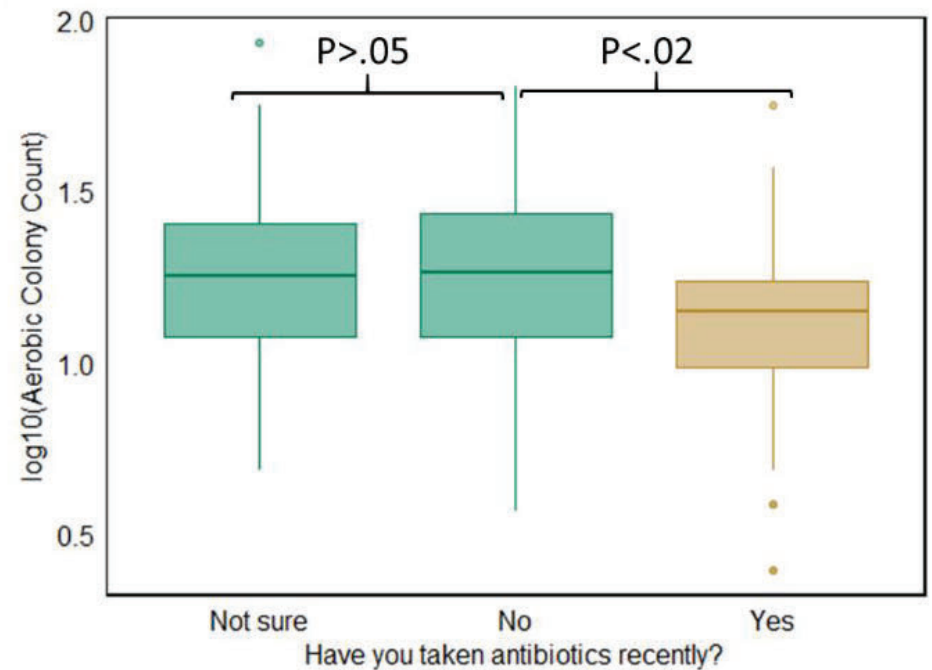
b) \log_{10} ACC

Effect of antibiotics

Those with recent antibiotic use show a small reduction in colony counts. Whether this is a direct effect of the drugs or whether people are more vigilant with cleaning when they have an infection is hard to say.



a) Microbial diversity score

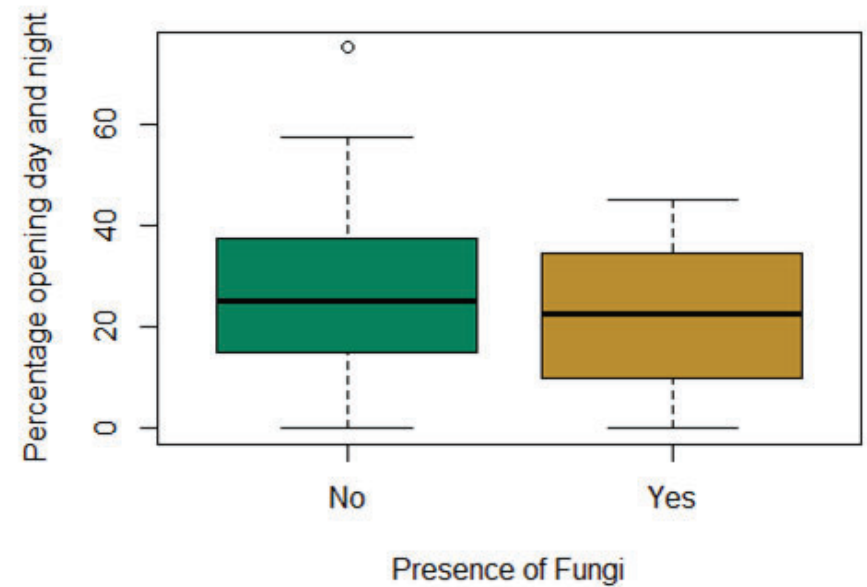
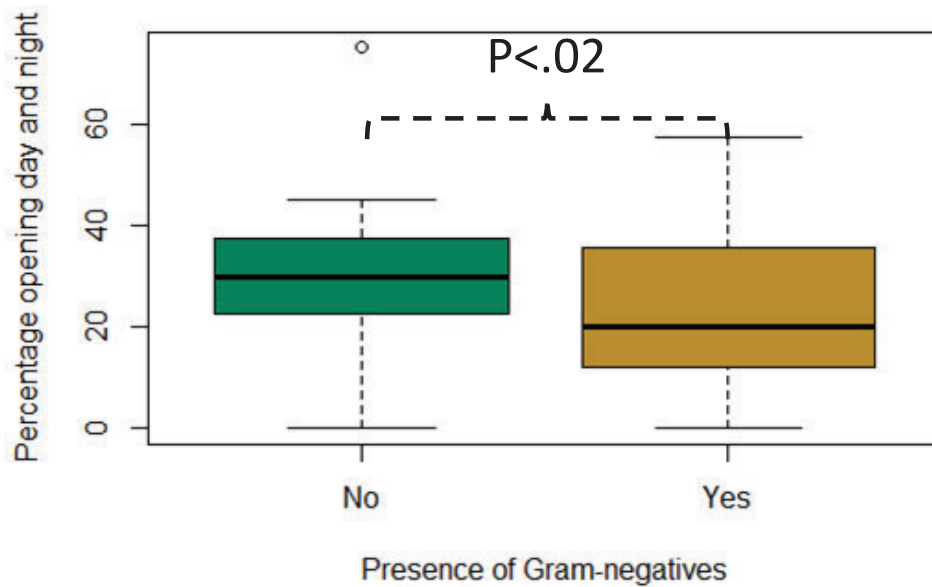


b) log10 ACC

Effect of Window Opening %

Window opening affects presence of Gram negative bacteria ($p < 0.02$). Also, more window opening = less Gram negatives (odds ratio 0.97).

Weak association with fungi presence.



‘Dirty water scandal at Glasgow NHS super hospital’



https://www.parliament.scot/S5_HealthandSportCommittee/Inquiries/HS-S5-19-HHHE-A2.pdf

Summary

Greater ACC correlated with higher diversity

Window opening frequency reduced Gram-negatives

Disinfectant range also reduced chances of finding Gram-negatives (but is this because these people simply cleaned more often?)

Smoking households had lower ACC and *tendency* to lower diversity

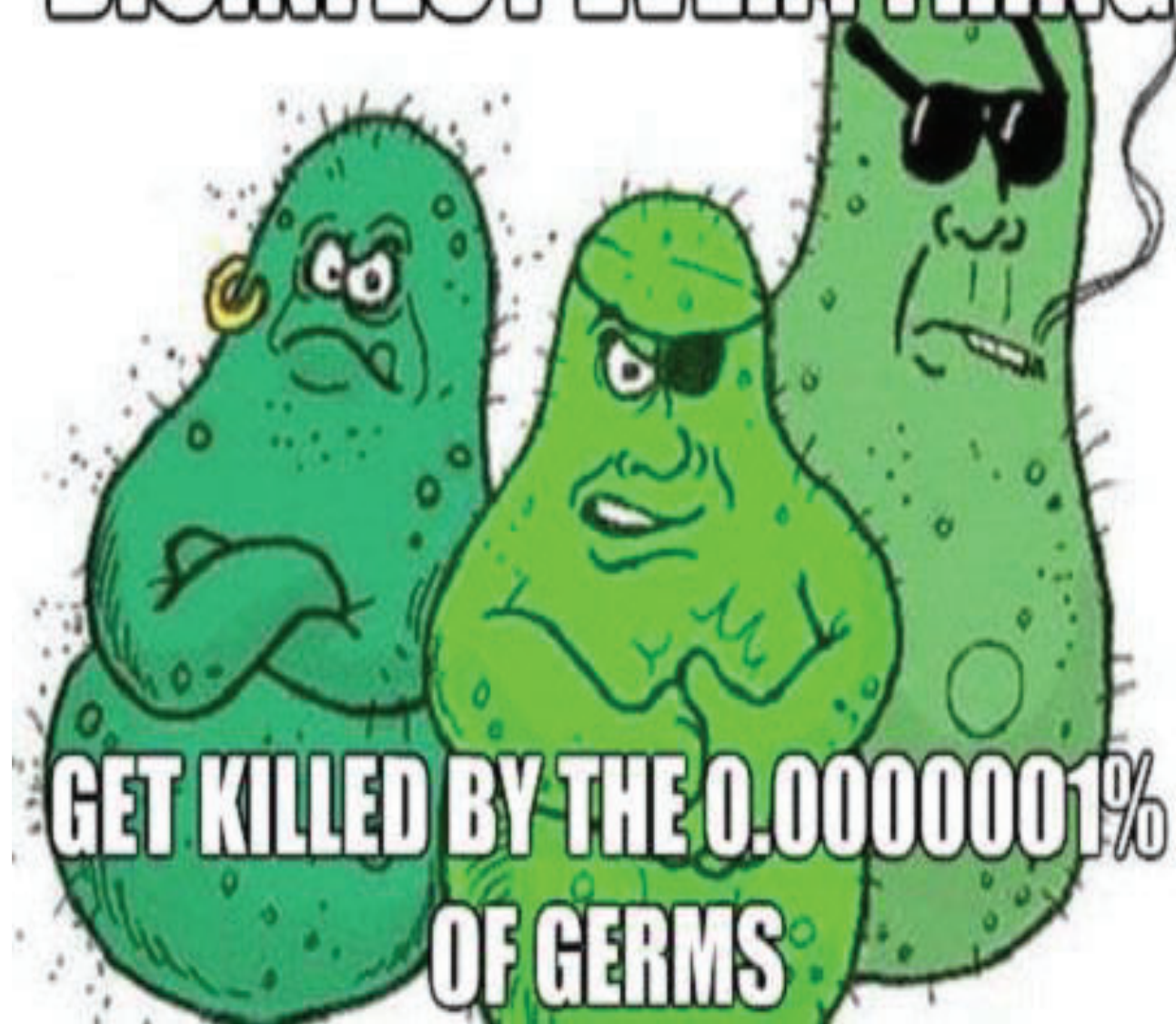
Antibiotic use appeared to reduce ACC and a *tendency* to reduce diversity

Acknowledgements

**All the staff in the Hairmyres Microbiology laboratory,
especially Janice McLaren and Louise McNally**



DISINFECT EVERYTHING



**GET KILLED BY THE 0.0000001%
OF GERMS**

Influence of Ventilation design on the prevalence of anti-microbial resistant bacteria in homes

Stage 2: Environmental Monitoring

Dr. Gráinne McGill, MEARU, Glasgow School of Art



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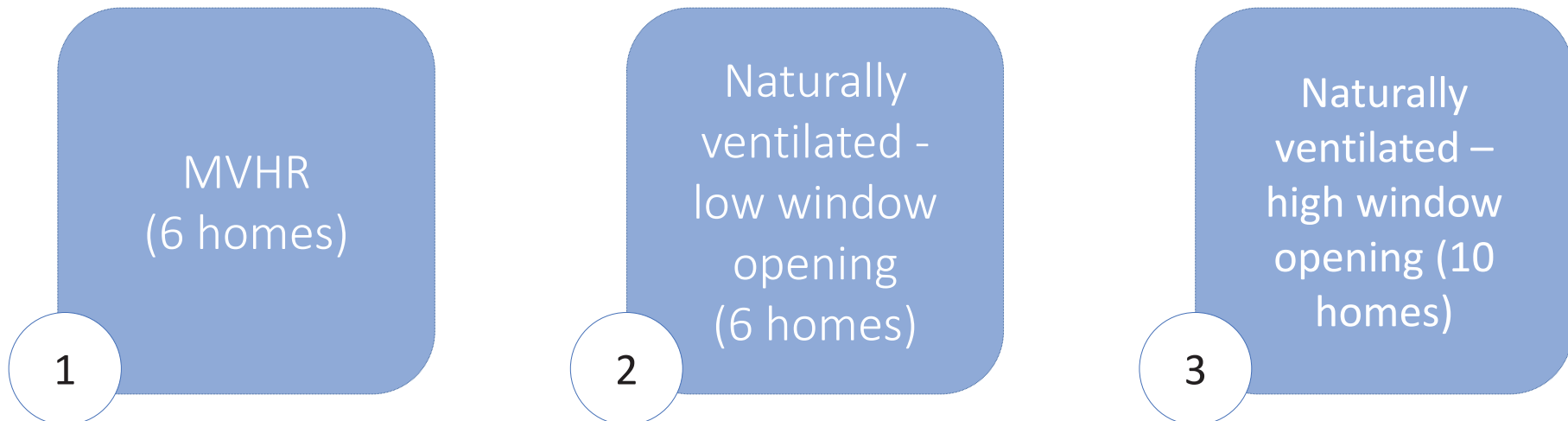
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Selection of homes

- Household survey data used to select homes for Stage 2
- 96 Households interested in participating (contact numbers available for 37)
- Delay due to fire may have affected recruitment
- Selection of homes based on:
 - Willingness to participate
 - Ventilation provision (homes selected with different ventilation strategies)
 - Ventilation behavior (homes selected based on reported window opening
 - Dwelling age (homes constructed since 2009)

Three main categories:



Selection of homes

| Cat. | Code | Site | Build year | Typology | No. people | Occupant age | Pets? | Bed window night | Total window opening freq |
|---|----------|--------------|------------|---------------|------------|--------------|---------|------------------|---------------------------|
| MVHR | House 19 | Lunestone | 2017 | Terraced | 3 | 21,22,52 | No | Never | 25% |
| | House 10 | Lunestone | 2017 | Flat | 2 | 62,65 | No | Never | 24% |
| | House 04 | Lunestone | 2017 | Flat | 1 | 66 | No | Never | 0% |
| | House 07 | Lunestone | 2017 | Flat | 2 | 48,77 | No | Never | 18% |
| | House 11 | Lunestone | 2017 | Flat | 2 | 70,76 | No | Never | 23% |
| | House 09 | Lunestone | 2017 | Terraced | 4 | 21,24,55,56 | Yes | Daily | 48% |
| Nat – Low window opening | House 16 | Doonholm | 2016 | Flat | 1 | 63 | No | Never | 0% |
| | House 18 | Doonholm | 2016 | Flat | 1 | 56 | 1 dog | Never | 0% |
| | House 01 | Murray | 2011 | Semi-detached | 2 | 68, 71 | No | Never | 3% |
| | House 22 | Mosspark | 2016 | Flat | 1 | 64 | Hamster | Never | 0% |
| | House 14 | Kirktonholme | 2009 | Flat | 1 | 75 | Cat | Never | 15% |
| | House 20 | Lincoln | 2017 | Flat | 1 | 60 | No | Never | 15% |
| Nat vent - high window opening | House 13 | Chantinghall | 2013 | Cottage | 2 | 78, 80 | Dog | Daily | 43% |
| | House 03 | Waterford | 2016 | Flat | 2 | 61, 64 | No | Daily | 45% |
| | House 15 | Fenwick | 2013 | Flat | 1 | 60, 63 | Dog | Never | 38% |
| | House 08 | Kirktonholme | 2009 | Flat | 2 | 69, 71 | No | Weekly | 43% |
| | House 17 | Lincoln | 2017 | Flat | 2 | 58, 60 | No | Weekly | 40% |
| | House 12 | Lincoln | 2017 | Flat | 1 | 63 | No | Daily | 45% |
| | House 21 | Lincoln | 2017 | Flat | 2 | 43, 59 | Dog | Daily | 38% |
| | House 05 | Slamannan | 2010 | Flat | 1 | 64 | Dog | Never | 38% |
| | House 06 | Slamannan | 2010 | Flat | 1 | 64 | Dog | Never | 38% |
| | House 02 | Slamannan | 2010 | Flat | 1 | 64 | No | Never | 38% |

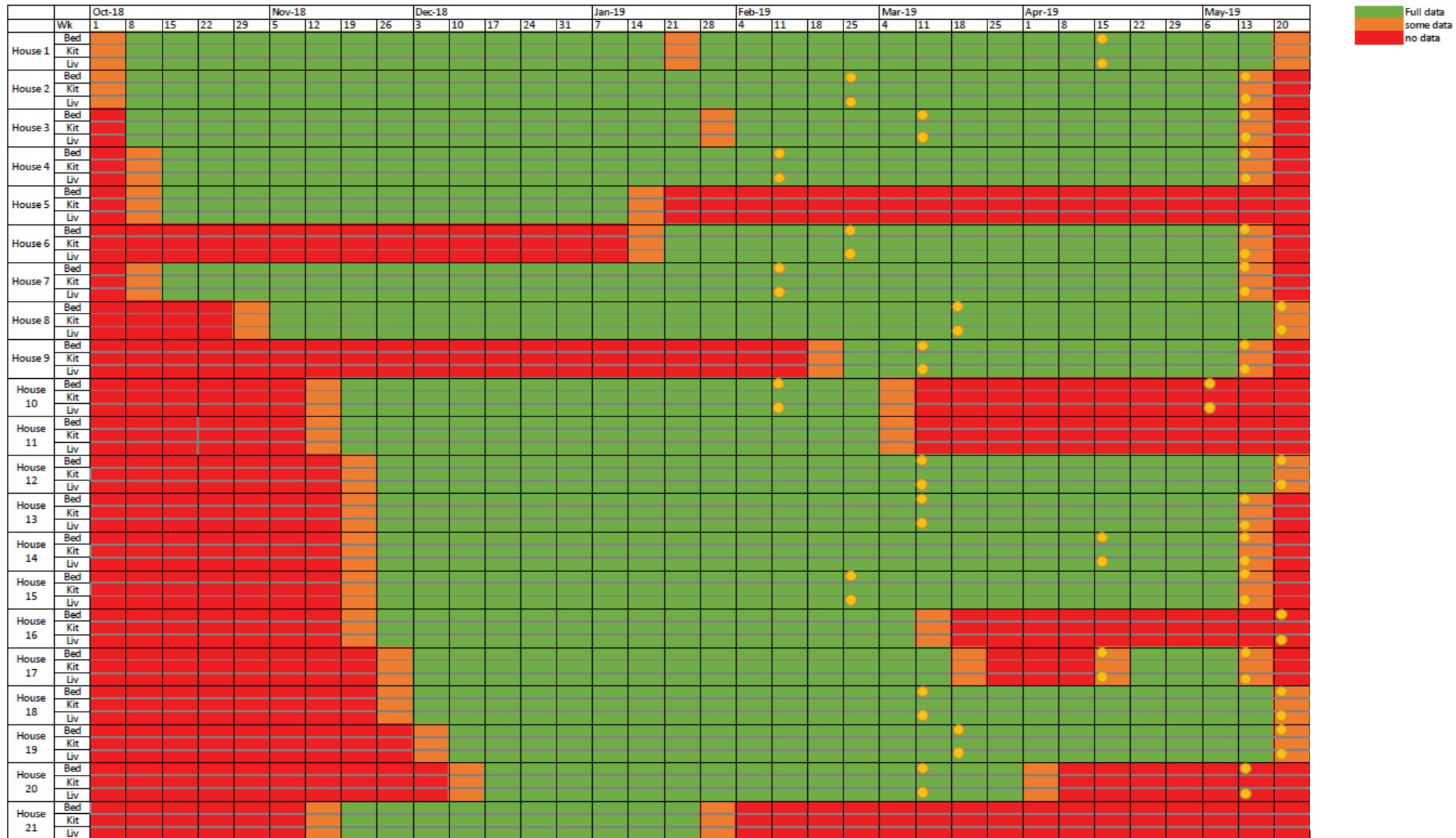
Environmental monitoring

- Information sheet & consent form
- Sensors installed in main bedroom, living room, kitchen & outside
- Temperature, RH & CO₂ logged at 10 minute intervals
- Low cost foobot IAQ sensors (temp, RH, TVOCs, PM2.5, CO₂)
- Mifi device – wifi connection
- First time using these devices – identified some technical issues (poor signal, no. sensors needed replaced, needed to change wifi passwords, lack of notification when sensors not logging)



Data availability

Temperature and RH data availability (microbial sampling in yellow)



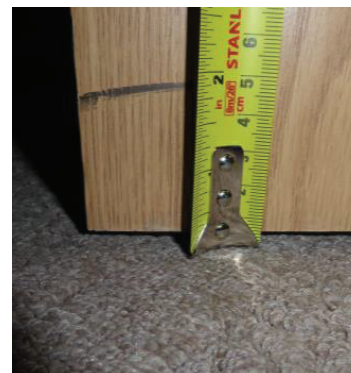
Data availability

Carbon dioxide data availability (microbial sampling in yellow)



Ventilation audit

- Airflow rate measurements – mechanical vent systems
- Trickle vent location, size & position
- Make and model of mechanical ventilation systems
- Operating condition of mechanical ventilation systems
- Filter condition
- Door undercuts
- Airflow pathways
- Occlusions to ventilation grilles / openings



Building survey

- Photo survey
- Building walk-through
 - Floor plan layouts, window locations etc.
 - Floor level
 - Type of heating source
 - Wall surfaces
 - Floor surfaces
 - Visible mould (if present)
 - Cleaning products
 - Meter readings



Occupant interview

- Semi-structured interviews undertaken with building occupants to gather data on:
 - Occupancy regimes and profiles
 - Cleaning regimes and products
 - Pet ownership (including use of antibiotics)
 - Bedroom night time conditions (door position, window position, time go to bed etc.)
 - Any problems with water damage / mould
 - Medication and health, particularly antibiotic use
 - Ventilation behaviour (knowledge and use of mechanical ventilation systems)
- Interviews recorded with Dictaphone

Microbial sampling

- Air samples: two seasons (Spring and Summer)
 - Bedroom (30 minutes)
 - Bathroom (15 minutes)
 - Living room (15 minutes)
 - Outside (15 minutes)
- Surface samples – 13-15 sites
- Finger tip samples (building occupants)
- Samples placed in cold storage box for transport
- Air samples posted to University of Leeds for analysis
- Surface and finger samples analysed by Hairmyres Microbiology Lab



Surface sampling locations

Bathroom door handle

Telephone/mobile

Kettle handle

Bedside table

Top door frame (bed)

Remote control

Toilet handle

Window ledge (bed)

Top door (bathroom)

Behind radiator (bed)

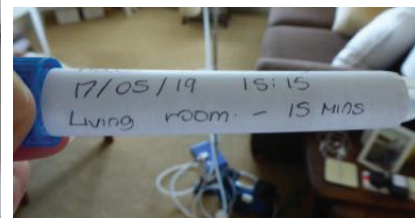
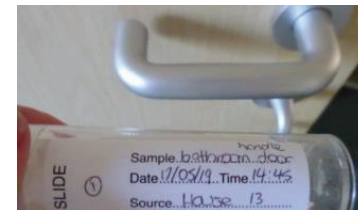
Extract (bathroom)

Supply (bed)

Top window frame (bed)

MVHR left filter

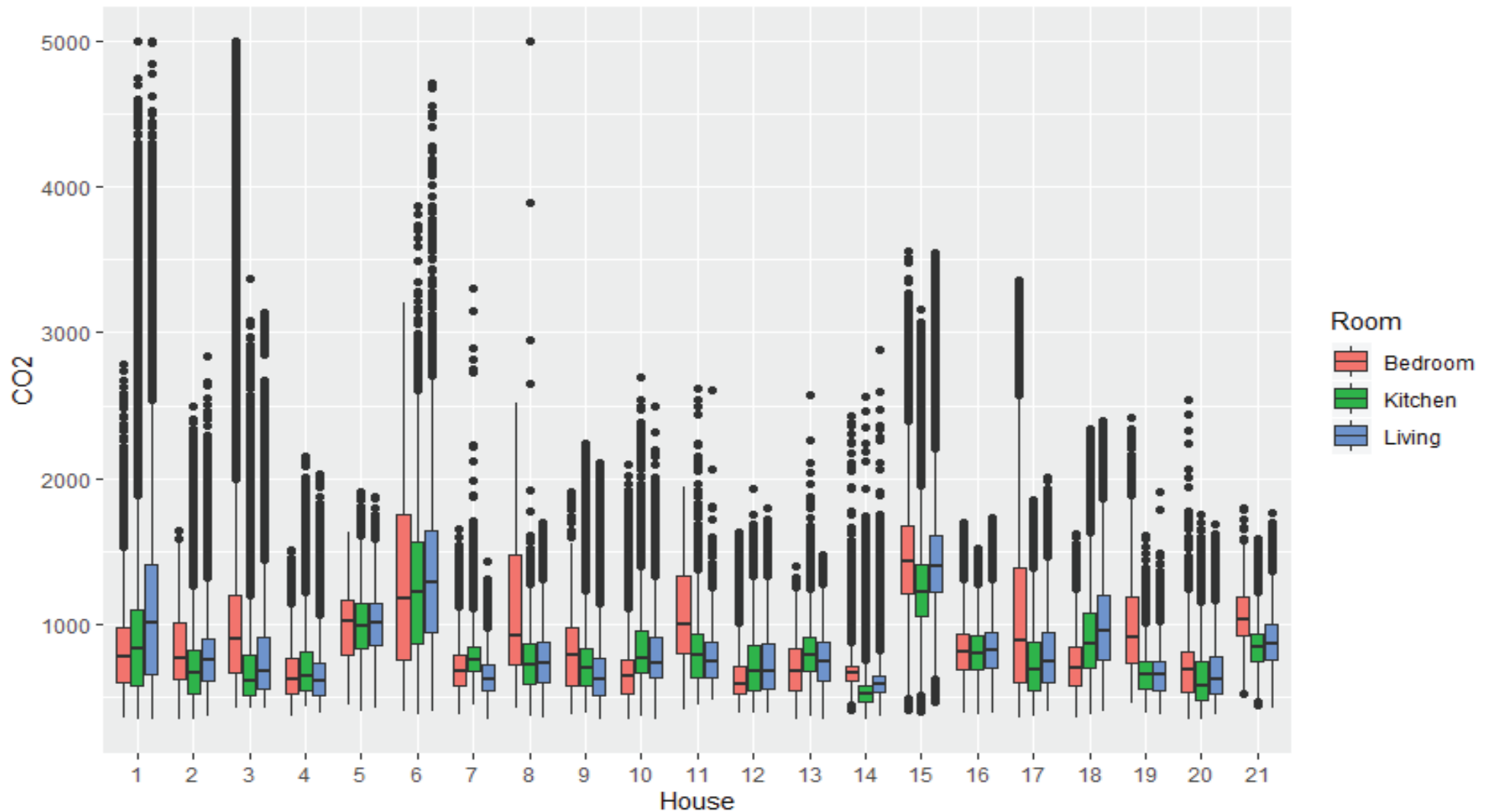
MVHR right filter



Initial results: Carbon dioxide

Overall CO₂ levels (whole monitoring period) by room, for each home

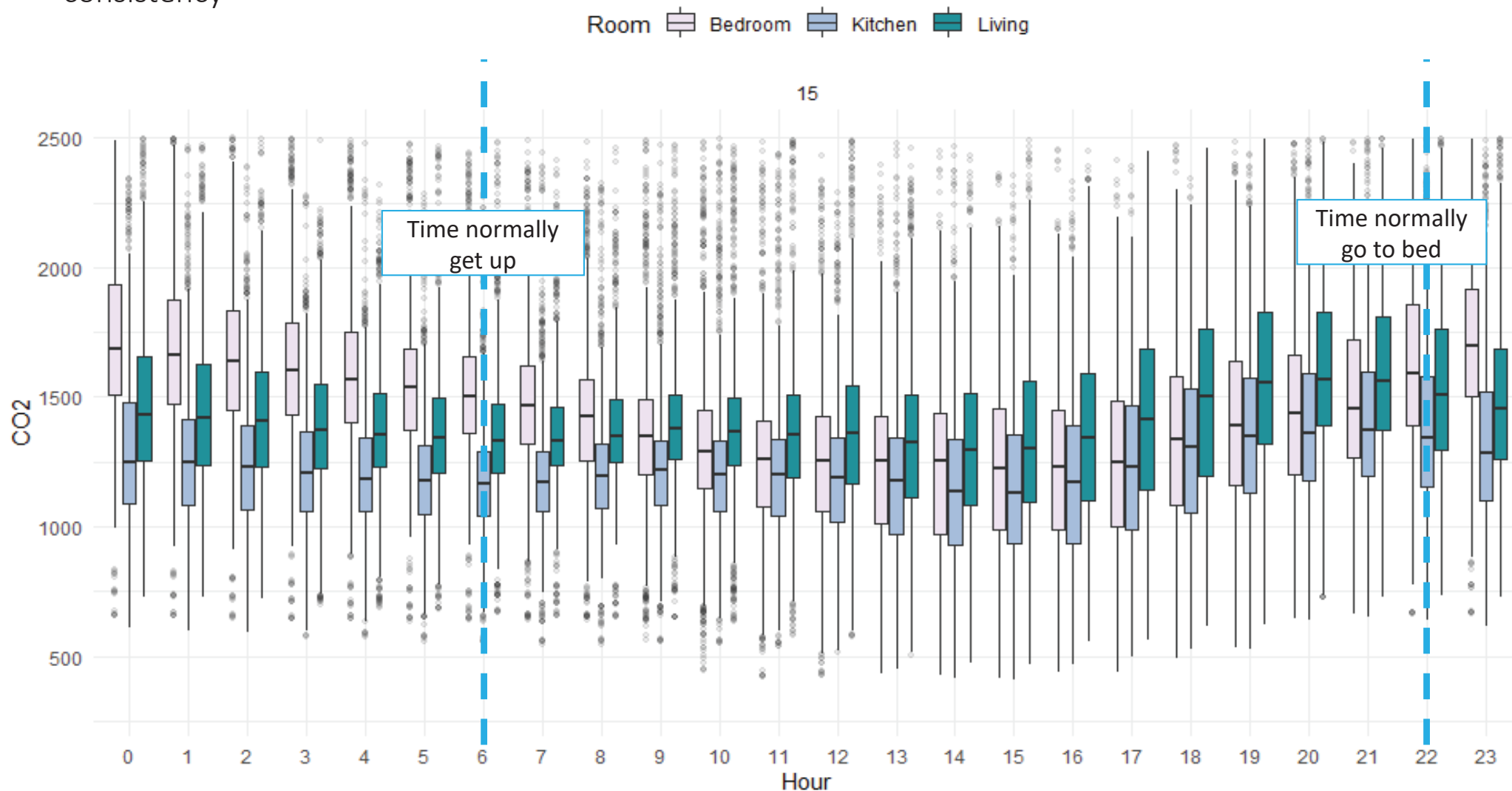
- CO₂ levels typically higher in bedroom
- CO₂ levels generally remained below 1,500ppm, however two homes (House 6 and 15) notably higher



Initial results: Carbon dioxide

Determining typical bedroom 'night time' occupied period

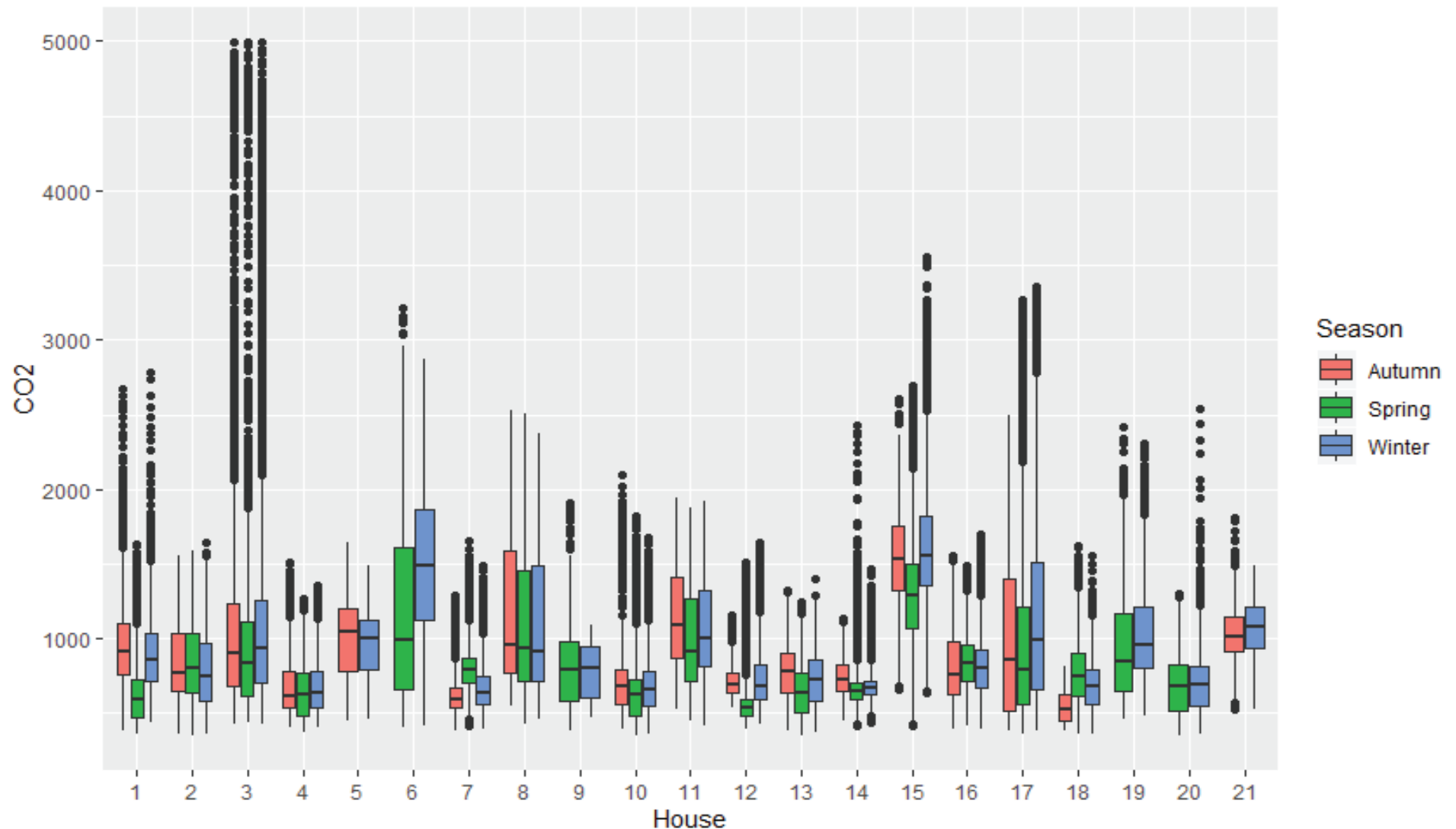
- Bedroom night time occupancy determined based on reported time occupants normally go to bed and get up each morning
- This information was compared with hourly bedroom CO2 data – found reasonable level of consistency



Initial results: Carbon dioxide

Bedroom carbon dioxide levels in each home, by season

- Bedroom carbon dioxide levels considerably higher during night time occupied periods
- Bedroom carbon dioxide levels did not vary considerably by season



Initial results: Carbon dioxide

Bedroom night time carbon dioxide levels

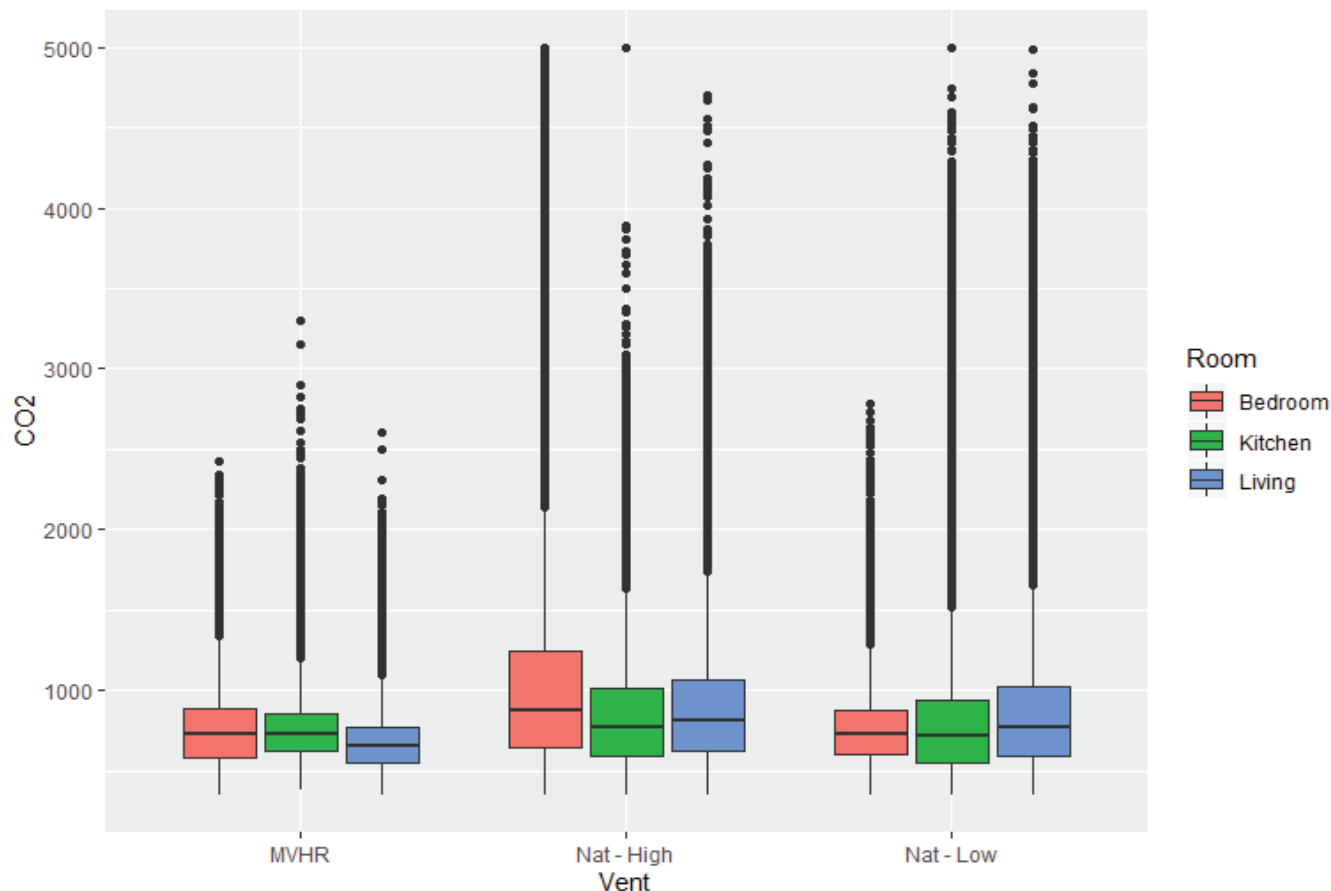
- Mean bedroom night time carbon dioxide levels ranged from 716ppm to 1678ppm
- All homes exceeded the recommended carbon dioxide guideline level of 1,000ppm at night.

| House No. | Vent cat | Window opening (%) | SD CO ₂ | Max CO ₂ | Mean CO ₂ | % time exceeding 1,000ppm |
|-----------|----------|--------------------|--------------------|---------------------|----------------------|---------------------------|
| 1 | N - Low | 3% | 290 | 1951 | 914 | 38% |
| 2 | N - High | 38% | 250 | 1565 | 950 | 51% |
| 3 | N - High | 45% | 773 | 4995 | 1465 | 78% |
| 4 | MVHR | 0% | 114 | 1509 | 760 | 1% |
| 5 | N - High | 38% | 253 | 1638 | 1067 | 77% |
| 6 | N - High | 38% | 558 | 2738 | 1520 | 76% |
| 7 | MVHR | 18% | 167 | 1522 | 720 | 6% |
| 8 | N - High | 43% | 422 | 2523 | 1566 | 89% |
| 9 | MVHR | 48% | 194 | 1906 | 988 | 45% |
| 10 | MVHR | 24% | 243 | 1912 | 821 | 17% |
| 11 | MVHR | 23% | 198 | 1878 | 1378 | 95% |
| 12 | N - High | 45% | 299 | 1611 | 739 | 31% |
| 13 | N - High | 43% | 159 | 1296 | 844 | 19% |
| 14 | N - Low | 15% | 93 | 1333 | 718 | 25% |
| 15 | N - High | 38% | 391 | 3555 | 1678 | 97% |
| 16 | N - Low | 0% | 182 | 1632 | 887 | 25% |
| 17 | N - High | 40% | 725 | 3311 | 1533 | 69% |
| 18 | N - Low | 0% | 213 | 1450 | 716 | 9% |
| 19 | MVHR | 25% | 433 | 2163 | 1059 | 63% |
| 20 | N - Low | 15% | 172 | 2535 | 755 | 6% |
| 21 | N - High | 38% | 385 | 1804 | 1017 | 71% |

Initial results: Carbon dioxide

Ventilation categories and measured CO₂ levels (by room)

- There was no clear association between ventilation categories and measured carbon dioxide levels
- Reported whole house window opening did not appear to correspond to measured carbon dioxide levels in any room
- May be explained by small sample size and wide range of factors affecting CO₂, such as mechanical ventilation systems, occupancy levels, floor area, door opening, ventilation pathways, cooking etc.



Initial results: Temperature

Percentage of time temperature thresholds were exceeded

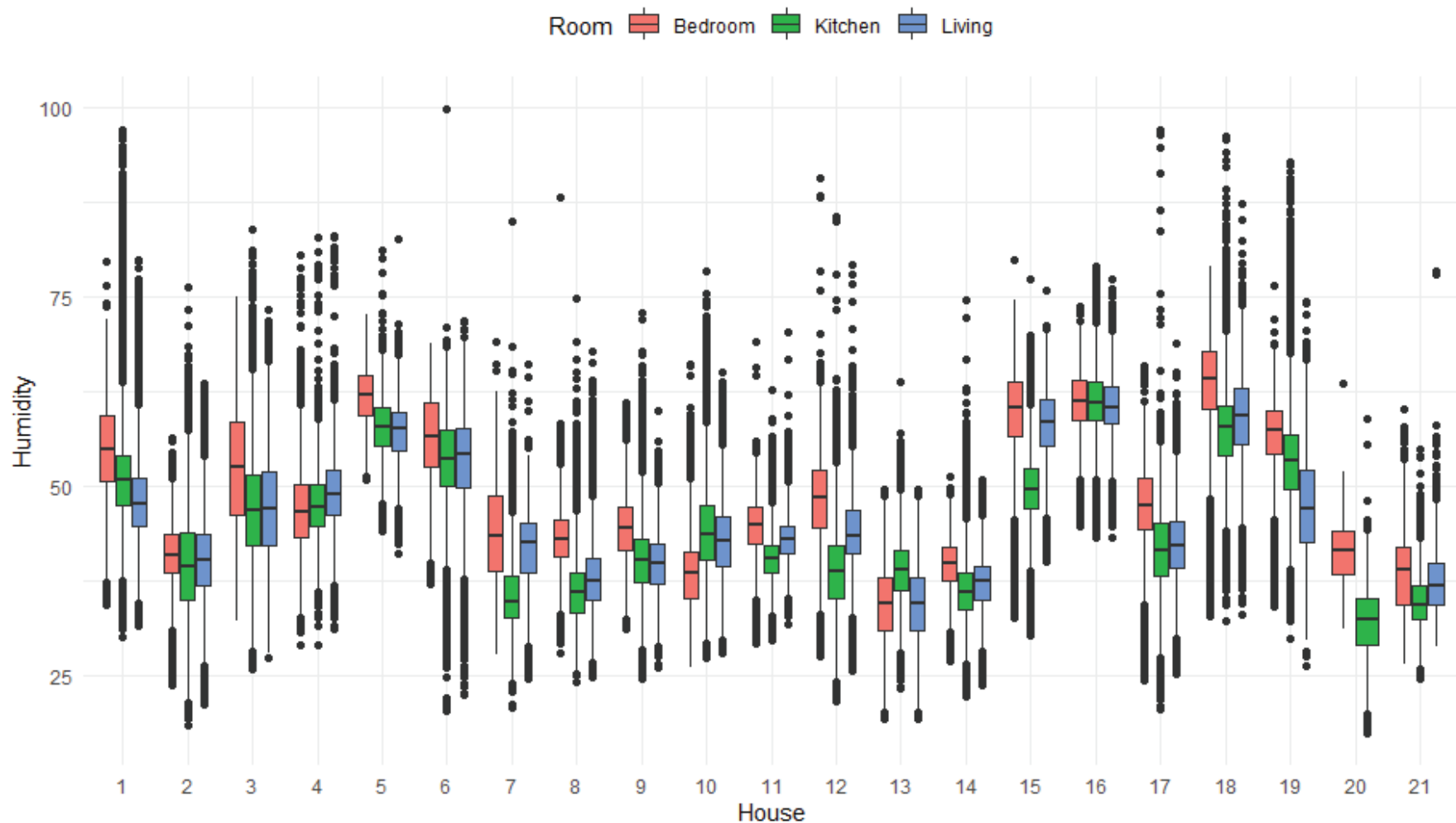
- While measurements were not undertaken during the summer season, high incidences of overheating were still observed.
- Overall, nine homes (43%) exceeded the CIBSE overheating threshold of 25 °C for more than 5% of the occupied time, in at least one of the monitored rooms.

| | Occupied hours (based on reported night time occupancy) | | | | | | | | | |
|-----------|---|---------------------|---------------------|---------------------|------------------------|------------------------|------------------------|-------------------------|-------------------------|-------------------------|
| House No. | % time bed temp <16 | % time bed temp >25 | % time bed temp >26 | % time bed temp >28 | % time living temp <16 | % time living temp >25 | % time living temp >28 | % time kitchen temp <16 | % time kitchen temp >25 | % time kitchen temp >28 |
| 1 | 13.0 | 0.0 | 0.0 | 0.0 | 0.0 | 11.8 | 0.1 | 0.0 | 5.3 | 0.0 |
| 2 | 4.8 | 0.7 | 0.1 | 0.0 | 5.1 | 11.0 | 0.0 | 4.9 | 29.8 | 1.0 |
| 3 | 10.8 | 0.0 | 0.0 | 0.0 | 0.7 | 0.0 | 0.0 | 1.4 | 0.0 | 0.0 |
| 4 | 0.1 | 1.0 | 0.4 | 0.0 | 4.3 | 0.0 | 0.0 | 1.3 | 0.0 | 0.0 |
| 5 | 41.8 | 0.0 | 0.0 | 0.0 | 23.4 | 0.0 | 0.0 | 23.6 | 0.0 | 0.0 |
| 6 | 25.9 | 0.0 | 0.0 | 0.0 | 6.4 | 0.0 | 0.0 | 4.1 | 0.1 | 0.0 |
| 7 | 0.0 | 10.2 | 2.0 | 0.0 | 0.0 | 13.4 | 2.3 | 0.0 | 59.4 | 0.7 |
| 8 | 0.8 | 0.1 | 0.0 | 0.0 | 0.4 | 26.6 | 0.2 | 0.3 | 62.9 | 10.7 |
| 9 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 2.2 | 0.0 | 0.0 | 1.0 | 0.0 |
| 10 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| 11 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 25.4 | 0.0 |
| 12 | 12.7 | 0.2 | 0.1 | 0.0 | 1.4 | 1.4 | 0.0 | 0.2 | 8.1 | 0.3 |
| 13 | 0.0 | 14.0 | 4.1 | 0.0 | 0.0 | 52.1 | 1.3 | 0.0 | 1.0 | 0.0 |
| 14 | 2.1 | 0.0 | 0.0 | 0.0 | 1.4 | 4.5 | 0.0 | 1.6 | 0.9 | 0.0 |
| 15 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.1 |
| 16 | 89.7 | 0.0 | 0.0 | 0.0 | 77.4 | 0.0 | 0.0 | 72.2 | 0.0 | 0.0 |
| 17 | 0.0 | 1.1 | 0.2 | 0.0 | 0.0 | 3.8 | 0.0 | 0.0 | 4.1 | 0.0 |
| 18 | 73.7 | 0.0 | 0.0 | 0.0 | 21.2 | 0.0 | 0.0 | 10.5 | 0.0 | 0.0 |
| 19 | 32.9 | 0.0 | 0.0 | 0.0 | 15.4 | 0.0 | 0.0 | 3.4 | 0.0 | 0.0 |
| 20 | 2.3 | 0.4 | 0.0 | 0.0 | n/a | n/a | n/a | 0.0 | 44.6 | 0.0 |
| 21 | 0.6 | 42.3 | 31.2 | 5.6 | 0.0 | 57.5 | 1.6 | 0.0 | 90.1 | 11.7 |

Initial results: Relative Humidity

Relative humidity levels (whole monitoring period) by room

- Homes with the lowest observed temperatures recorded highest levels of relative humidity, suggesting that high levels of RH in these homes were more likely a function of temperature
- An exception to this is House 15, where high levels of relative humidity were observed despite warm interior temperatures.



Preliminary conclusions

- Ventilation categories (based on reported window opening) do not appear to correlate with measured CO₂ levels
- Mean bedroom night time CO₂ levels exceeded 1,000ppm in 9/21 homes
- High incidences of overheating during heating season, with 9/21 homes exceeding CIBSE overheating threshold (25°C for >5% occupied time)
- Small no. of homes were noticeably colder, with temperatures rarely exceeding 18°C in one home (House 16)
- One home (House 15) measured considerably higher vapour pressure levels than other homes
- House 15 also recorded highest CO₂ levels, suggesting poor ventilation

Next steps

- Re-characterise the data based on the measured CO₂ and moisture levels
- Sequencing of bioaerosol samples – to determine breakdown of species
- Complete analysis of Stage 2 microbial sampling (air, surfaces, finger tips)
- Statistical analysis to compare differences between housing types, including environmental conditions, no. & type of microorganisms present and proportion resistant to antibiotics
- Undertake analysis of low-cost sensor data (and assess value of data as proxy indicators)
- Define protocols to allow understanding of levels of measurements needed for larger study
- Assess the implications for future design and legislation

Discussion

Questions and Answers