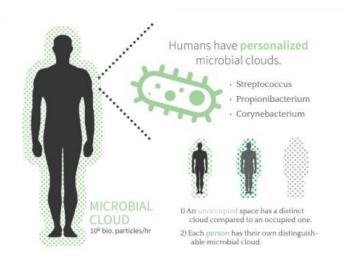


New research finds that people emit their own personal microbial cloud

September 22 2015





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METHODS

SETTLING DISHES Petri dishes collect surface biological particles.



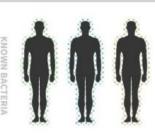
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AIR FILTERS Subjects are surrounded by cups with filters to collect air particles

Airborne particles were more **identifiable** than settled ones.



SUBJECT 1 Dolosigranulum SUBJECT 2 Streptococcus SUBJECT 3 Lactobacillus



Most of the 11 subjects had a **unique** microbial cloud.

PeerJ

Meadow et al. (2015). Humans differ in their personal microbial cloud. Peerl 3:e1258: DOI 10.7717/peerj.1258 https://peerl.com/articles/1258/



We each give off millions of bacteria from our human microbiome to the air around us every day, and that cloud of bacteria can be traced back to an individual. New research focused on the personal microbial cloud -- the airborne microbes we emit into the air -- examined the microbial connection we have with the air around us. The findings demonstrate the extent to which humans possess a unique 'microbial cloud signature.' Credit: Viputheshwar Sitaraman, of Draw Science.

We each give off millions of bacteria from our human microbiome to the air around us every day, and that cloud of bacteria can be traced back to an individual. New research focused on the personal microbial cloud—the airborne microbes we emit into the air—examined the microbial connection we have with the air around us. The findings demonstrate the extent to which humans possess a unique 'microbial cloud signature'.

To test the individualized nature of the personal microbial cloud, University of Oregon researchers sequenced microbes from the air surrounding 11 different people in a sanitized experimental chamber. The study found that most of the occupants sitting alone in the chamber could be identified within 4 hours just by the unique combinations of <u>bacteria</u> in the surrounding air. The findings appear in the September 22 issue of the open-access, peer-reviewed journal *PeerJ*.

The striking results were driven by several groups of bacteria that are ubiquitous on and in humans, such as Streptococcus, which is commonly found in the mouth, and Propionibacterium and Corynebacterium, both common skin residents. While these common human-associated microbes were detected in the air around all people in the study, the authors found that the different combinations of those bacteria were the



key to distinguishing among individual people.

The analyses, utilizing analysis of suspended particulate matter and shortread 16S sequencing, focused on categorizing whole <u>microbial</u> <u>communities</u> rather than identifying pathogens. The findings emerged from two different studies and more than 14 million sequences representing thousands of different types of bacteria found in the 312 samples from air and dust from the experimental chamber.

"We expected that we would be able to detect the human microbiome in the <u>air</u> around a person, but we were surprised to find that we could identify most of the occupants just by sampling their microbial cloud," said lead author James F. Meadow, a postdoctoral researcher formerly from the Biology and the Built Environment Center at the University of Oregon.

"Our results confirm that an occupied space is microbially distinct from an unoccupied one, and and demonstrate for the first time that individuals release their own personalized microbial cloud," the authors concluded.

The research sheds light on the extent to which we release our <u>human</u> <u>microbiome</u> to our surrounding environment, and might help understand the mechanisms involved in the spread of infectious diseases in buildings. The results also suggest potential forensic applications, for example to identify or determine where a person has been, though it is unclear whether individual occupants can be detected in a crowd of other people.

More information: Meadow JF, Altrichter AE, Bateman AC, Stenson J, Brown G, Green JL, Bohannan BJM. (2015) Humans differ in their personal microbial cloud. *PeerJ* 3:e1258 <u>dx.doi.org/10.7717/peerj.1258</u>



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