

Type and abundance of mouth bacteria linked to lung cancer risk in non-smokers

December 14 2020



Credit: CC0 Public Domain

The type and abundance of bacteria found in the mouth may be linked to lung cancer risk in non-smokers, finds the first study of its kind,

published online in the journal *Thorax*.

Fewer species and high numbers of particular types of bacteria seem to be linked to heightened risk, the findings indicate.

Around one in four cases of [lung cancer](#) occurs in non-smokers and known [risk factors](#), such as second hand tobacco smoke, background radon exposure, air pollution, and family history of lung [cancer](#) don't fully explain these figures, say the researchers.

The type and volume of bacteria (microbiome), found in the mouth has been associated with a heightened risk of various cancers including those of the gullet, head and neck, and pancreas.

And the researchers wanted to find out if this association might also hold for lung cancer, given that the mouth is the entry point for bacteria to the lungs.

They drew on participants in The Shanghai Women's Health Study and the Shanghai Men's Health Study, all of whom were lifelong non-smokers, and whose health was monitored every 2-3 years after entry to the study between 1996 and 2006.

At enrolment, participants rinsed out their mouths to provide a profile of the resident bacteria, and information was obtained on lifestyle, diet, medical history and other environmental and workplace factors that might influence their disease risk.

In all, 90 of the women and 24 of the men developed lung cancer within around 7 years, on average.

These cases were matched with 114 non-smokers of the same age and sex, who also provided a mouth rinse sample. This comparison group

didn't have lung cancer but they had similar levels of education and family histories of lung cancer.

Comparison of both sets of rinse samples showed that the microbiome differed between the two groups. A wider range of bacterial species was associated with a [lower risk](#) of developing lung cancer. And a larger volume of particular types of species was also associated with lung cancer risk.

A larger volume of Bacteroidetes and Spirochaetes species was associated with lower risk while a larger volume of Firmicutes species was associated with heightened risk.

Specifically, within the Spirochaetes species, a greater abundance of Spirochaetia was associated with lower risk; and within the Firmicutes [species](#), a larger volume of organisms from the *Lactobacillales* order of microbes was associated with a heightened risk.

The associations remained when the analysis was restricted to those participants who had not taken any antibiotics in the 7 days before sample collection and after excluding those diagnosed with lung cancer within 2 years of sample provision.

This is an observational study, and therefore can't establish cause. And the researchers acknowledge several limitations. "While our study provides evidence that variation in the [oral microbiome](#) plays a role in lung cancer risk, the interpretation of our study must be done while considering the caveat that our findings are from a single time point in a single geographical location," they write.

In a linked editorial, Dr. David Christiani, of Harvard University, suggests that mouth bacteria may provoke chronic inflammation, boost cell proliferation and inhibit cell death, prompt DNA changes, and

switch on cancer genes and their blood supply, which would help to explain the findings.

The study findings raise several questions, he says. "First, how stable is the human oral microbiome over time? Second, if the human oral microbiome varies over time, what determines that variability? Third, how does the ambient environment such as exposure to air pollutants, affect the oral (and lung) microbiome?"

He adds: "It remains unclear whether the oral microbiome as measured in this (and other) epidemiological studies represents a causative agent or only a marker of disease or immune activity. If it is the former, then it will be important to understand whether the oral microbiome actually seeds the [lung microbiome](#) and thus acts locally."

More information: Variation in oral microbiome is associated with future risk of lung cancer among never-smokers, *Thorax* (2020). [DOI: 10.1136/thoraxjnl-2020-215542](https://doi.org/10.1136/thoraxjnl-2020-215542)

Editorial: The oral microbiome and lung cancer risk, *Thorax* (2020). [DOI: 10.1136/thoraxjnl-2020-216385](https://doi.org/10.1136/thoraxjnl-2020-216385)

Provided by British Medical Journal

Citation: Type and abundance of mouth bacteria linked to lung cancer risk in non-smokers (2020, December 14) retrieved 9 March 2023 from <https://medicalxpress.com/news/2020-12-abundance-mouth-bacteria-linked-lung.html>

<p>This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.</p>
--