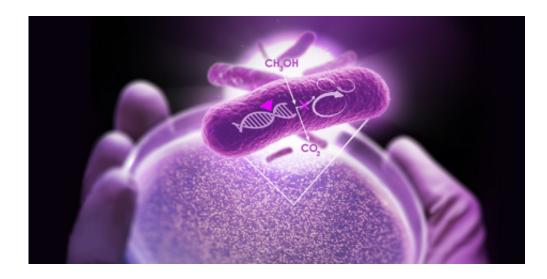


How a bacterium can live on methanol

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ETH Zurich researchers study bacteria that can use methanol (CH3OH) as a carbon source. Credit: ETH Zurich

ETH Zurich researchers have identified all the genes required by a bacterium to use methanol as a food source. The results will help scientists advance the use of this resource in the field of biotechnology.

Many chemists are currently researching how small carbon molecules, such as methane and <u>methanol</u>, can be used to generate larger molecules. The earth is naturally rich in methane, and artificial processes like the fermentation of biomass in biogas plants also produce it in abundance. Methanol can be generated from methane. Both are simple molecules containing only a single carbon atom. However, using them to produce larger molecules with several carbon atoms is complex.



While challenging for chemists, bacteria learned long ago to build large molecules out of small ones: Some bacteria use methanol as a carbon source in order to create energy carriers and cellular building material. They live primarily on <u>plant leaves</u> and occur in large numbers on every leaf. The bacterium most extensively researched is called Methylobacterium extorquens. A team led by Julia Vorholt, Professor of Microbiology, has now identified all the genes required by this bacterium to live on methanol.

Bacteria utilize a plant waste product

"Plant leaves are a natural source of methanol, which is produced as a <u>waste product</u> during plant cell wall biosynthesis," explains Vorholt. Much like a hybrid car can be fuelled by either petrol or electricity, Methylobacterium extorquens can use either larger carbon molecules (such as carboxylic acids) or methanol from plants as a nutrient, depending on availability. "When using methanol, the bacteria construct all the complex chemical compounds they need using this small molecule. This is an extraordinary accomplishment," says Vorholt.

As the scientists have now been able to demonstrate, Methylobacterium extorquens has nearly 150 genes that it needs specifically to grow on methanol. Of these, 95 had not been known before.

In order to find these genes, Vorholt's group worked together with the group of Beat Christen, Professor of Experimental Systems Biology, to create about a million bacterial mutants. These were placed in a culture medium with methanol on one hand, and in a normal nutrient medium containing succinic acid on the other hand. The researchers were looking for mutants that grew normally in the usual medium, but poorly or not at all when fed with medium containing only methanol. This allowed the scientists to identify when specific genes required for using methanol had been damaged.



"The method allowed us to determine whether any given gene in the bacterium was needed for methanol-based growth or not. This ultimately allowed us to identify all of the required genes," explains Andrea Ochsner, doctoral student in Vorholt's group and the first author of the study published in the journal *Current Biology*.

Signal for switch to methanol mode

The researchers were particularly surprised by one of the genes detected using this method. The gene was previously known from plants and bacteria that metabolize CO2 from the air, but Methylobacterium extorquens does not do this. The gene provides the instructions for an enzyme that produces a sugar crucial for CO2 use.

The researchers could now demonstrate that Methylobacterium extorquens is also able to produce this sugar when it encounters methanol. However, In contrast to <u>plants</u>, the bacterium does not use this sugar as a cellular building material. Instead, the scientists believe that it is used as a signal in order to switch to methanol consumption.

After uncovering the significance of this particular gene, the scientists now want to further investigate the other newly identified genes.

New abilities of microorganisms

Studying the genetics of Methylobacterium extorquens is also valuable for biotechnology. The bacterium is already used in research laboratories to produce complex molecules. The new results could help scientists manipulate the <u>bacterium</u> in order to create desired <u>molecules</u> in greater quantities.

Since all the genes relevant for methanol-based growth have now been



identified, it may also be possible to introduce these <u>genes</u> into other microorganisms, allowing them to use methanol – and thus enable various biotechnical applications for this resource in the future.

More information: Andrea M. Ochsner et al. Transposon Sequencing Uncovers an Essential Regulatory Function of Phosphoribulokinase for Methylotrophy, *Current Biology* (2017). <u>DOI:</u> <u>10.1016/j.cub.2017.07.025</u>

Provided by ETH Zurich

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