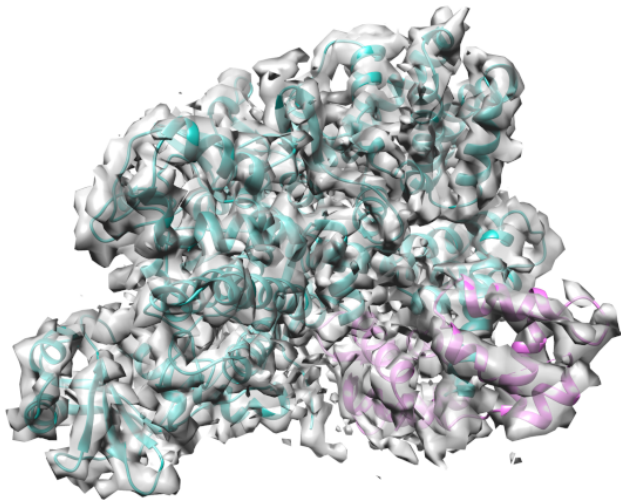


Researchers identify toxin responsible for Legionella growth

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A team of scientists led by EMBL group leader Sagar Bhogaraju and Ivan Dikic of Goethe University, Frankfurt, discovered that the toxin SidJ in Legionella bacteria enforces a unique modification on human proteins and helps legionella grow inside human cells. SidJ hijacks human protein Calmodulin to its own advantage in one of the classic examples of pathogenic bacteria exploiting the human molecular machinery and turning it against us. This makes SidJ an ideal target to curb Legionella infection.

Legionella – a complex bacterium

Pneumonia resulting from exposure to Legionella – although uncommon and affecting only 1 in 100,000 in Europe – has a higher than 10% fatality rate. The pathogenic bacterium Legionella pneumophila has more than 300 toxins that it uses to infect humans. Once the aerosols containing the bacteria are inhaled, Legionella enters the lungs where it starts infecting human cells, causing pneumonia.

Legionella toxins especially target the innate immune pathways facilitating the survival of the bacteria inside human cells and allowing the replication of the bacteria. Due to the large number of toxins it is difficult to see the effects of deleting one or multiple of these toxins on the Legionella infection capacities. This is further complicated by the fact that several toxins with similar functions exist inside the bacteria. This makes Legionella hard to target with specific drugs.

Focus on the SidJ toxin

Researchers from EMBL Grenoble and the Goethe University in Frankfurt have now studied the toxin SidJ in detail. It is an important toxic protein of *Legionella* that gets injected into the human cytoplasm and enables the successful infection and replication of the bacteria. In contrast to the other toxins in *Legionella*, the deletion of SidJ alone leads to a considerable growth defect of the bacteria in human cells. This makes SidJ one of the most important toxins of *Legionella* and an attractive target to curb *Legionella* infection.

While SidJ has been studied in the field for already more than a decade, the precise function of it remained unknown until today. "SidJ has no sequence similarity to any of the proteins with a known function. We had to resort to standard biochemical methods and mass spectrometry to determine its function", explains Bhogaraju. "While working out its mechanism proved to be challenging, it was also very exciting!"

In particular, the missing detailed molecular study of the toxin hindered the development of drugs that can target SidJ. The work by multidisciplinary scientists of Bhogaraju and Dikic groups now describes the molecular function of this protein in detail, elucidates its importance for *Legionella* infection and provides the identity of the human proteins that are targeted by SidJ.

Toxin at work

The group showed that SidJ possesses protein glutamylation activity: it attaches the amino acid glutamate to a target protein as post-translational modification. "This kind of activity is a first for bacterial proteins", says Ivan Dikic, Director of the Institute of Biochemistry II at Goethe University. SidJ glutamylates many human proteins that are involved in tackling microbial infections and innate immunity. In order to do this, SidJ interacts with the human protein Calmodulin – a highly conserved multifunctional intermediate calcium-binding messenger protein.