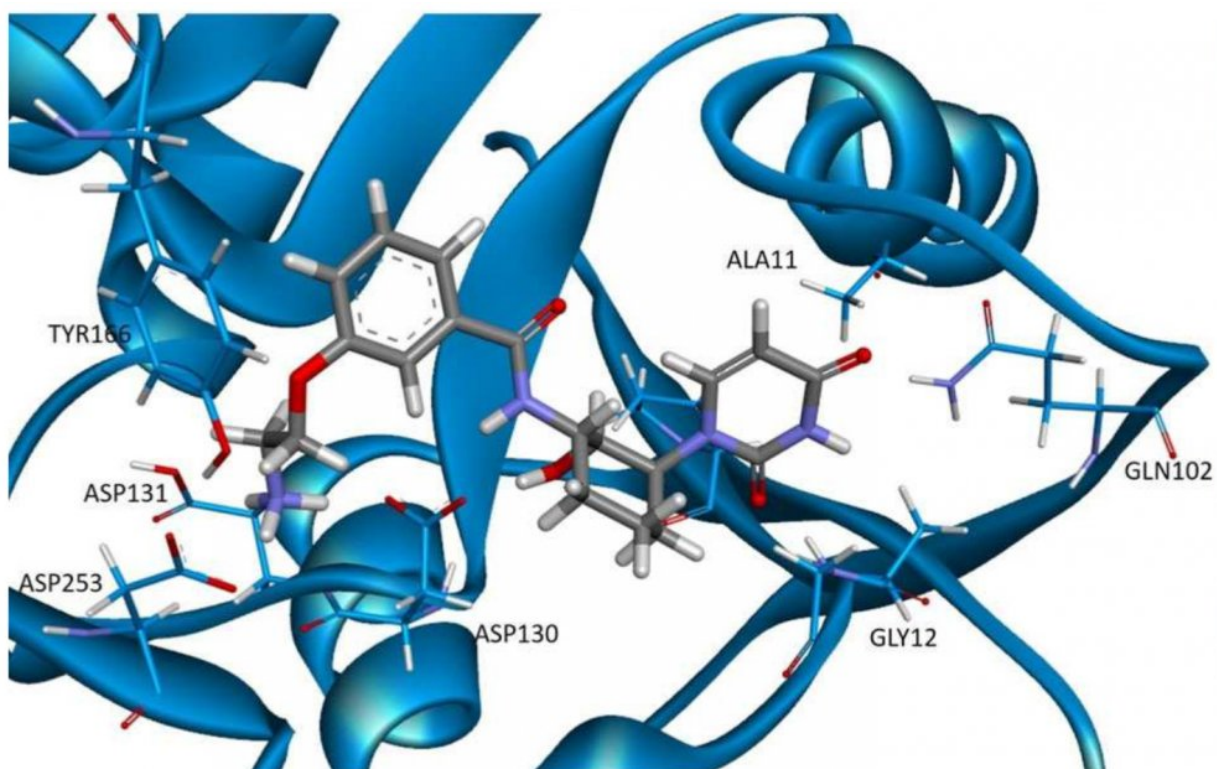


# Computers discover compounds that could reduce *Listeria's* virulence

March 12 2018, by Tracey Peake



3-D docking pose of potential GalU inhibitor. Credit: North Carolina State University

In a proof-of-concept study, researchers from North Carolina State University have pinpointed new compounds that may be effective in containing the virulence—or ability to produce disease - of *Listeria*, a

well-known bacterium that can cause severe food poisoning and even death.

Listeria are bacteria most commonly found in soil. Humans come into contact with Listeria via contaminated meat or milk products and can contract listeriosis, which can lead to severe illness or death—particularly in very young, elderly and/or immunocompromised populations.

Denis Fourches, assistant professor of computational chemistry, postdoctoral researcher Melaine Kuenemann and Paul Orndorff, professor emeritus of microbiology, knew that inhibiting a particular enzyme of Listeria—known as glucose-1-phosphate uridylyltransferase (GalU) - led to dramatic modifications of the bacterial cell surface. These chemical modifications in turn rendered the Listeria much less virulent - in other words, less able to cause illness.

The researchers turned their attention to identifying potential compounds that could inhibit the function of GalU. Using computers and cheminformatics methods, they characterized, analyzed and virtually screened more than 88,000 compounds with the potential to inhibit GalU. Computer models found 37 compounds promising enough to be tested in vitro. Of the 37, three were deemed effective enough to warrant further study, although many of the other, less active compounds yielded key information about how their chemical structures relate to their activity in inhibiting the enzyme's function.

"We can derive several predictive structure-activity relationships based on those 37 compounds and these relationships will help us design even more effective GalU inhibiting [compounds](#)," Fourches says. "We plan to use our computers to virtually generate thousands of new analogues, virtually screen them, and select another batch of up to 50 molecules to be tested experimentally in the future. This is true research at the

interface of disciplines."

Interestingly, inhibiting GalU also served to make the *Listeria* more vulnerable to cefotaxime, an antibiotic to which the bacteria are naturally resistant.

"While our ultimate objective is to get away from antibiotics altogether, in the near term the antibiotic susceptibility opens up the possibility of combinatorial therapies that could include a GalU inhibitor and a known antibiotic such as cefotaxime," Orndorff says. "Ultimately, we believe if the GalU inhibitor is effective enough, the host (human or animal) should be able to eliminate the listerial population without [antibiotics](#). For farmers working toward antibiotic-free farms, this could be a wonderful solution."

"This proof-of-concept study shows that small molecules can actually be developed to shut down the activity of one specific bacterial enzyme, leading to the suppression of virulence," Fourches says. "This is clearly a new avenue for fighting drug-resistant bacteria."

The research appears in *Molecular Informatics*.

**More information:** Melaine A. Kuenemann et al, In silico Predicted Glucose-1-phosphate Uridyltransferase (GalU) Inhibitors Block a Key Pathway Required for *Listeria* Virulence, *Molecular Informatics* (2018). [DOI: 10.1002/minf.201800004](https://doi.org/10.1002/minf.201800004)

Provided by North Carolina State University

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