

New algorithms shown to accelerate biopharmaceutical process

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Credit: Rensselaer Polytechnic Institute

Biopharmaceuticals are necessary, life-saving tools. But the process for



making them is time-consuming and costly, particularly when it comes to the process of purification—the removal of unwanted elements like proteins, viruses, and DNA.

A team of chemical and biological engineers at Rensselaer Polytechnic Institute has developed a method that significantly cuts down the amount of time needed for the purification process, laying the groundwork for faster, less expensive drug production. The findings were recently published in *Biotechnology and Bioengineering*, where the article was named an editor's choice.

"We're developing new technologies, new processes, and new modeling techniques for more efficient engineering processes for making drugs," said Steven Cramer, an endowed chair professor of chemical and biological engineering at Rensselaer who led this research.

Choosing which purification process to use on a biopharmaceutical is known as the process development phase. In this paper, Cramer and his team show that they can cut down the amount of time that process takes from months to days by employing screening methods and computer algorithms the Rensselaer researchers have developed.

Those <u>screening methods</u>, which help the team know exactly which unwanted molecules need to be removed, were developed after the researchers collected a large amount of data and built a database. The database enabled the team to develop <u>computer algorithms</u> to rank which sequences of chromatographic processes—methods of separating molecules—will work best.

"Using this new way, we were able to develop an overall process in one week, in contrast to typical time frames of four to six months," Cramer said. "We bring all the data together in our in silico modeling tool and the computer gives us a ranking of which processes to evaluate in the



lab."

When Cramer's team took the recommendations made by their algorithms into the lab, they found that the processes the <u>computer</u> suggested, worked. The successful results suggest, Cramer said, a new platform-like approach for rapidly designing <u>purification</u> processes.

More information: Nicholas Vecchiarello et al. A combined screening and in silico strategy for the rapid design of integrated downstream processes for process and product-related impurity removal, *Biotechnology and Bioengineering* (2019). DOI: 10.1002/bit.27018

Provided by Rensselaer Polytechnic Institute

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