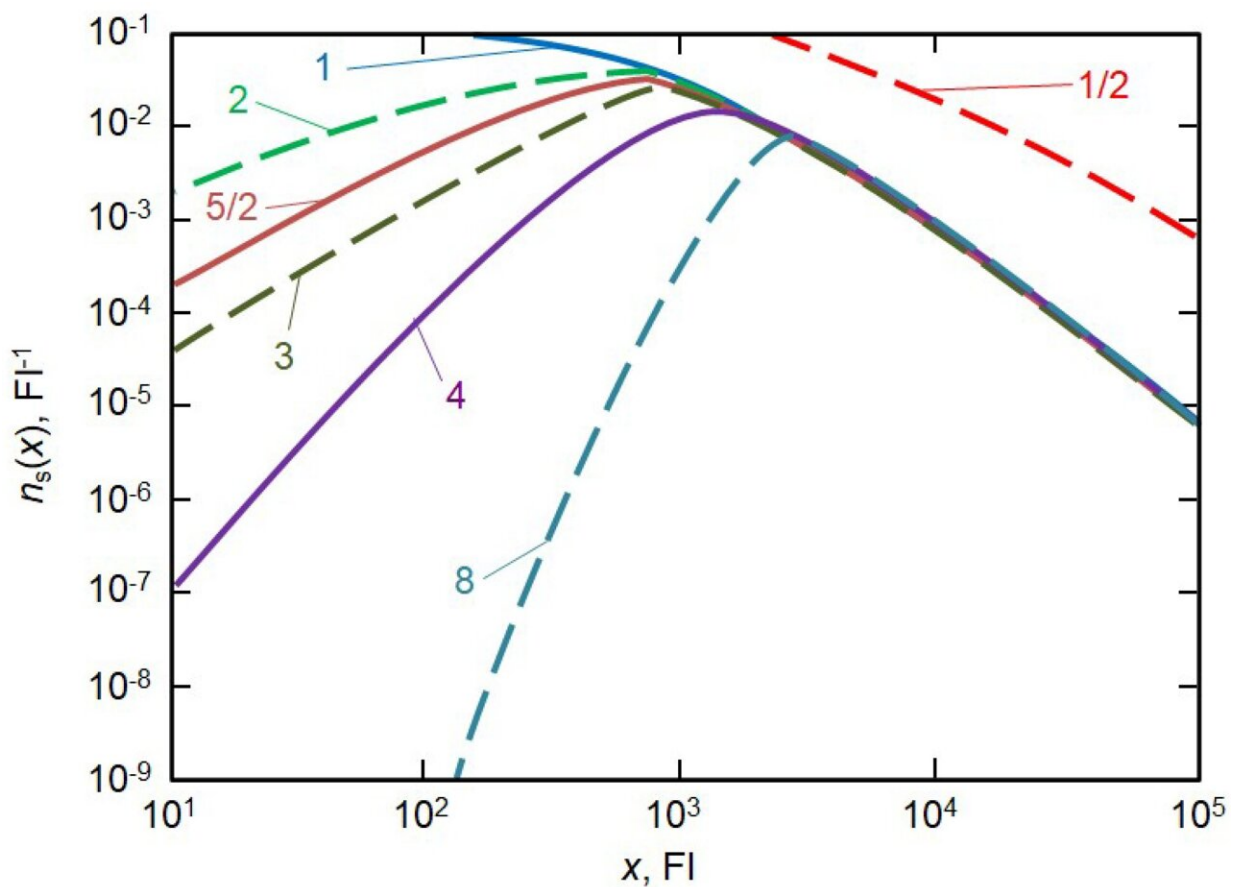


Scientists have created a mathematical model for the dynamics of nanoparticles and viruses in cells

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The steady-state density distribution functions $n_s(x)$ at different $I(x)$ and v (numbers at the curves). Credit: *Crystals* (2022). DOI: 10.3390/cryst12081159

Physicists and mathematicians from the Ural Federal University (UrFU) have created a complex mathematical model that calculates the distribution of nanoparticles (in particular, viruses) in living cells. The mathematical model helps with finding how the nanoparticles cluster (merge into a single particle) inside cells, namely in cellular endosomes, which are responsible for sorting and transporting proteins and lipids.

These calculations will be useful for medical purposes because they show the behavior of [viruses](#) when they enter cells and seek to replicate. The model also allows for the accurate calculation of the amount of medication needed for therapy to ensure that the treatment is as effective as possible and with minimal side effects. The model description and the results of calculations were published in the journal *Crystals*.

"The processes in cells are extremely complex, but in simple words, the viruses use different variants to reproduce. Some of them deliver the genetic material directly to the cytoplasm. Others use the endocytosis pathway: they deliver the viral genome by releasing it from the endosomes. If viruses linger in the endosomes, the acidity increases and they die in the lysosomes," says Dmitri Alexandrov, Head of the Laboratory of Multiscale Mathematical Modeling at UrFU.

"So, our model has allowed to find out, first, when and which viruses 'escape' from endosomes in order to survive. For example, some [influenza viruses](#) are low pH-dependent viruses; they fuse with the [endosome](#) membrane and release their genome into the cytoplasm. Secondly, we found that it is easier for viruses to survive in endosomes during clustering, when two particles merge and tend to form a single particle."

As the scientists explain, the [mathematical model](#) will also be useful in tumor targeting therapy: many cancer therapies depend on when and how [nanoparticles](#) of a drug saturate cancer cells. And the model will

help to calculate this parameter.

In addition, understanding the behavior of viruses in cells is important for the development of vaccines and drugs, as well as for [gene therapy](#), which treats diseases that conventional medicine cannot cope with. For example, various adenovirus-based vectors and lipid particles are used as a platform for gene delivery to treat the disease. But their limited ability to "slip out" of the endosomes also limits their use as deliverers.

"Nanoparticles smaller than 100 nanometers are becoming increasingly important tools in modern medicine. Its applications range from nanodiagnostics to radiation therapy for cancer. For example, pH-sensitive nanoparticles mimicking viruses are used for targeted delivery of antitumor drugs. This is how drugs are delivered from whole organs to [individual cells](#)," says Head of the Laboratory of Stochastic Transport of Nanoparticles in Living Systems (UrFU) Eugeny Makoveeva.

More information: Eugeny V. Makoveeva et al, Analysis of Smoluchowski's Coagulation Equation with Injection, *Crystals* (2022). [DOI: 10.3390/cryst12081159](https://doi.org/10.3390/cryst12081159)

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