



## Projekta Izp-2020/2-0373 rezultāti

### No kukaiņu zarnu trakta mikrofloras izolētu DNS bakteriofāgu raksturošana

*Oriģināli zinātniskie raksti, kas publicēti zinātniskos žurnālos, rakstu krājumos vai konferenču rakstu krājumos, kuri ir indeksēti datu bāzēs Web of Science Core Collection, SCOPUS vai ERIH PLUS*

1. Zrelovs, N.; Dislers, A.; Kazaks, A. Genome characterization of nocturne116, novel lactococcus lactis-infecting phage isolated from moth. - Microorg., 2021, 9 (7), <https://doi.org/10.3390/microorganisms9071540>
2. Zrelovs, N.; Jansons, J.; Dislers, A.; Kazaks, A. Morganella Phage Mecenats66 Utilizes an Evolutionarily Distinct Subtype of Headful Genome Packaging with a Preferred Packaging Initiation Site. - Microorg., 2022, 10 (9), <https://doi.org/10.3390/microorganisms10091799>
3. Zrelovs, N.; Jansons, J.; Kazaka, T.; Kazaks, A.; Dislers, A. Three Phages One Host: Isolation and Characterization of Pantoea agglomerans Phages from a Grasshopper Specimen. - Int. J. Mol. Sci., 2023, 24 (3), <https://doi.org/10.3390/ijms24031820>

*Zinātniskās datubāzes un datu kopas, kas izstrādātas projekta ietvaros:*

1. Andris Dislers, Nikita Zrelovs, Andris Kazaks. 10x "Complete genome of phage X", where "X" is Pocis76; Nocturne116; Nifs112; Nufs112; Nafs113; Pocitis76; Mecenats66; Virsaitis27; Sarma103; Gungnir39. "The International Nucleotide Sequence Database Collaboration (INSDC)" through GenBank. Public biological sequence repository accession numbers of ten annotated complete genome sequences of insect-associated phages characterized during this project: MW689258; MW791312; OK570184; OK570185; OK570186; OL512804; OL614104; OL770309; OL856139; OL944604. [Home - Nucleotide - NCBI \(nih.gov\)](https://www.ncbi.nlm.nih.gov)

