RYTIS STAKAITIS

RYTIS.STAKAITIS@LSMU.LT, ORCID, (+370) 641-1107 (LT)

WORK EXPERIENCE	
1 Sep 2018 – Present	Junior researcher
	LSMU Neuroscience institute Laboratory of Molecular Neurooncology (LT)
	 Gene expression analysis from qPCR and NGS experiments
	Data analysis from genomic databases
1 Sep 2014 – 1 Sep 2018	Technician
	Lithuanian University of Health Sciences (LT)
	 CpG methylation detection applying bisulfite modification and MS-PCR
	 qPCR analysis of total and small RNAs
1 Sep 2018 – 1 Sep 2019	Medical biologist
	Invitro Diagnostika Ltd. (LT)
	Blood smear microscopy
	 Interpretation and consulting on diagnostic tests
EDUCATION AND TRAINING	
1 Aug 2021 – 1 Aug 2022	Traineeship: "Computational analysis of sequencing data"
	Oregon Health & Science University, Don Conrad lab (USA)
	Single-cell sequencing data analysis
	 Sequencing data analysis on GenomeDK and Exacloud clusters
	Small RNA-seq data analysis
16 Sep 2019 – 30 May 2020	Traineeship: "NGS performance and analysis"
	Copenhagen University Hospital, Rigshospitalet, Kristian Almstrup group (DK)
	Small RNA sequencing on Illumina MiSeq
	 Mitochondrial DNA analysis using ONT MinION Cas9 approach
2 Jan 2019 – 1 Mar 2019	Certificate in data analysis using Python
	Kaunas Coding School Ltd. (LT)
	Data management in MySQL
	 Data analysis using Python Pandas and Scikit-learn
1 Sep 2018 – 30 Aug 2023	PhD (Biology)
	Lithuanian University of Health Sciences (LT)
	Thesis: ' <u>MiRNA studies of tumor tissue and blood extracellular vesicles for diagnosis and prognosis</u> of gliomas"
1 Sep 2016 – 1 Sep 2018	Master's degree in Medical Biology
	Lithuanian University of Health Sciences (LT)
	Thesis: "Semaphorin Sema3C molecular mechanisms involved in regulation of gliomagenesis"
1 Sep 2012 – 1 Mar 2016	Bachelor's degree in Molecular Genetics
	Lithuanian University of Health Sciences (LT)
	Thesis: "RAB40B and NPTX2 genes expression and promoter methylation analysis in different malignancies of astrocytic glioma"

Command line skills Conda, Singularity Slurm queue system Bash, Python3 Grants & Achievements 2 PhD student grants from Lithuanian University of Health Sciences Research foundation (each of 2,900 Eur) 2 Erasmus+ traineeship grants (each of 2,400 Eur) and 1 SMPE traineeship grant (3,000 Eur) 1 BAFE traineeship grant (30,000 USD) 3 incentive scholarships from Research Council of Lithuania (each 2,500 Eur) 9 Teer - reviewed publications 1. "Identification and comparison of m6A modifications in glioblastoma non-coding RNAs with MeRIP-seq and Nanopore dRNA-seq", Epigenetics – Taylor & Francis 2. "Antisense IncRNA CHROMR's Linked to glioma patient survival", Frontiers in Molecular, Bisciences – Frontiers 3. "Circulating levels and the bioactivity of mR-30b increase during pubertal progression in boys", Frontiers in Endocrinology – Frontiers 4. "Diverse Monogenic Subforms of Human Spermatogenic Failure", Nature Communications – Nature Portfolio 5. "Transcriptome-wide analysis of glioma stem cell specific m6A modifications in long-non-coding, RNAs", Spentlic Reports – Nature Portfolio 7. "Variant PNL/DC1, Defective piRNA Processing, and Azoospermia", NELM – NEJM Group 8. "Transcriptome-wide analysis of glioma stem cell specific m6A modifications in long-non-coding, RNAs", Spentlic Reports – Nature Portfolio 7. "Variant PNL/DC1, Defective piRNA Processing, and Azoospermia", NELM – NEJM Group 8. "Transcriptome-wide analysis of glioma stem cell sp	ADDITIONAL INFORMATION	
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 MeRIP-seq and Nanopore dRNA-seq", Epigenetics – Taylor & Francis "Antisense IncRNA <i>CHROMR</i> is linked to glioma patient survival", Frontiers in Molecular Biosciences – Frontiers "Circulating levels and the bioactivity of miR-30b increase during pubertal progression in boys", Frontiers in Endocrinology – Frontiers "Diverse Monogenic Subforms of Human Spermatogenic Failure", Nature Communications – Nature Portfolio "The piRNA-pathway factor FKBP6 is essential for spermatogenesis but dispensable for control of meiotic LINE-1 expression in humans", AJHG – Cell Press "Transcriptome-wide analysis of glioma stem cell specific m6A modifications in long-non-coding. RNAs", Scientific Reports – Nature Portfolio "Variant <i>PNLDC1</i>, Defective piRNA Processing, and Azoospermia", NEJM – NEJM Group "Small RNAs in Seminal Plasma as Novel Biomarkers for Germ Cell Tumors", Cancers – MDPI Group "Pasima-Derived miRNA-222 as a Candidate Marker for Papillary Thyroid Cancer", International Journal of Molecular Sciences – MDPI Group "Papillary Thyroid Carcinoma Tissue miR-146b, -21, -221, -222, -181b Expression in Relation with Clinicopathological Features", Diagnostics – MDPI Group "The Role of <i>CASC2 and miR-21</i> Interplay in Glioma Malignancy and Patient Outcome", International Journal of Molecular miR-181b/d Biomarkers and Health Related Quality of Life. Score in the Predictive Glioma Molecular miR-181b/d Biomarkers and Health Related Quality of Life. Score in the Predictive Glioma Molecular miR-181b/d Biomarkers and Health Related Quality of Life. Score in the Predictive Glioma Molecular miR-181b/d Biomarkers and Health Related Quality of Life. Score in the Predictive Glioma Molecular miR-181b/d Biomarkers and Health Related Quality of Life. 	Grants & Achievements	 2,900 Eur) 2 <u>Erasmus+</u> traineeship grants (each of 2,400 Eur) and 1 <u>SMPF</u> traineeship grant (3,000 Eur) 1 <u>BAFF</u> traineeship grant (30,000 USD)
 <u>Study</u>", <u>Cancers</u> – MDPI Group 14. "<u>High CHI3L1 expression is associated with glioma patient survival</u>", <u>BMC</u> – Springer Nature Group 	Peer - reviewed publications	 MeRIP-seq and Nanopore dRNA-seq", Epigenetics – Taylor & Francis "Antisense IncRNA <i>CHROMR</i> is linked to glioma patient survival", Frontiers in Molecular Biosciences – Frontiers "Circulating levels and the bioactivity of miR-30b increase during pubertal progression in boys", Frontiers in Endocrinology – Frontiers "Diverse Monogenic Subforms of Human Spermatogenic Failure", Nature Communications – Nature Portfolio "The piRNA-pathway factor FKBP6 is essential for spermatogenesis but dispensable for control of meiotic LINE-1 expression in humans", AJHG – Cell Press "Transcriptome-wide analysis of glioma stem cell specific m6A modifications in long-non-coding. RNAs", Scientific Reports – Nature Portfolio "Variant <i>PNLDC1</i>, Defective piRNA Processing, and Azoospermia", NEJM – NEJM Group "Small RNAs in Seminal Plasma as Novel Biomarkers for Germ Cell Tumors", Cancers – MDPI Group "Plasma-Derived miRNA-222 as a Candidate Marker for Papillary Thyroid Cancer", International Journal of Molecular Sciences – MDPI Group "Papillary Thyroid Carcinoma Tissue miR-146b, -21, -221, -222, -181b Expression in Relation with Clinicopathological Features", Diagnostics – MDPI Group "The Role of <i>CASC2</i> and <i>miR-21</i> Interplay in Glioma Malignancy and Patient Outcome", International Journal of Molecular Sciences – MDPI Group "Lingue Interplay between Molecular miR-181b/d Biomarkers and Health Related Quality of Life Score in the Predictive Glioma Models", International Journal of Molecular Sciences – MDPI Group "Association of miR-34a Expression with Quality of Life of Glioblastoma Patients: A Prospective Study", Cancers – MDPI Group "High CHI3L1 expression is associated with glioma patient survival", BMC – Springer Nature