# Curriculum vitæ Vedran Miletić

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#### Core skills

In silico drug design and software development Scientific software development and research experience in **computational biochemistry**, **molecular modeling**, and its applications to *in silico* drug design

**GROMACS**: developed **conditional stopping** and **solvent acceleration** features, contributed 40+ patches (C/C++, MPI, CUDA/OpenCL, Python, CMake, Git) **RxDock**, a fork of **rDock**: project leader; approximately 100 commits in 2 years as scientific software architect at **RxTx Research** (C++, Eigen, {fmt}, JSON, Meson)

Network and system administration Planning, installation, configuration, and maintenance of **application-** and **network-layer services** on Linux and FreeBSD in **bare metal** and **virtualized** environments (including **containers**), installation and configuration **automation**; experience with network emulation and simulation (ns-3, SimGrid, CORE, Click)

Teaching and laboratory practice

Organizing and lecturing courses in **computer networks and system administration**, **parallel and distributed computing**, and **web development**; leading the writing of **publicly available teaching materials** (in Croatian, 320 000+ words) at **Group for apps and services on exascale research infrastructure (GASERI)** 

## Auxiliary skills

**Open source software porting, patching, packaging, and development**: Porting software to and packaging for Linux (RPM and deb) and FreeBSD (ports and pkg) using GCC and Clang compilers. Basic familiarity with porting to Windows, macOS, and illumos. Contributed patches to many open-source software projects: ns-3, Clang, LLVM, libclc, Mesa, CP2K, Sphinx, Meson Wrap DB, LaTeX Beamer class, Material for MkDocs, MoinMoin, Drogon, and Linux (kernel).

**Debugging, profiling, testing, and documentation**: Usage of GDB and Valgrind for debugging and profiling, including parallel and distributed applications. Test-driven development (GoogleTest and gMock). Broad experience with writing technical documentation for software development and system administration using various markup languages and tools: Markdown (MkDocs), reStructuredText (Sphinx), Doxygen, LaTeX, ConTeXt, and Pandoc; exported HTML and PDF post-processing.

**Scripting languages and the Web**: Experience with scripting using Python and PHP programming languages for automation of software development and system maintenance tasks as well as web application backend development. Knowledge of web application benchmarking with ApacheBench, Siege, and JMeter for scalability analysis and estimation of resource requirements for deployment.

### Languages

Croatian: Native, English: Fluent, Italian: Basic, German: Basic

## **Experience**

#### 2021 – today: Assistant Professor, FIDIT

**Institution**: Group for Applications and Services on Exascale Research Infrastructure (GASERI), Faculty of Informatics and Digital Technologies (FIDIT), University of Rijeka, Rijeka, Croatia

**Research:** Heterogeneous system architectures and parallel algorithms; biochemical/biophysical software applications for exascale supercomputers

**Teaching**: Computational Biochemistry and Biophysics, Big Data Infrastructure, Code Optimization (YUFE course), Web Programming, Computer System Administration, Computer Networks, Informatics for Pharmacists (Pharmacy study program)

Organization: Member of the University of Rijeka Supercomputing Resources Committee

**Promotion:** Regular public lectures on popular scientific topics, presentations at European Researchers Nights, participation in open laboratory days, organization of Faculty of Informatics and Digital Technologies anniversary day celebration

#### 2019 – 2021: Senior Lecturer, UniRi

**Institution**: Computer Networks, Parallelization, and Simulation Laboratory (CNPSLab), Department of Informatics, University of Rijeka, Rijeka, Croatia

**Research:** Distributed chemical compound database architectures; high-throughput virtual screening tools and their applications

**Teaching**: Computer Networks, Security of Information and Communication Systems, Code Optimization (YUFE course), Network and Mobile Operating Systems, Computer System Administration, Web Programming, Web Applications 2, Computer Networks 1, Computer Networks 2, Network Systems Management, Operating Systems 2, Distributed Systems, Parallel Programming on Heterogeneous Systems, Informatics (Biotechnology and Drug Research study program)

**Organization**: Member of the University of Rijeka Supercomputing Resources Committee, Representative for Teachers in the Department Council

**Promotion**: Regular public lectures on popular scientific topics, presentations at European Researchers Nights, participation in open laboratory days

#### ■ 2015 – 2018: Postdoctoral Researcher, HITS

**Institution**: *Molecular Biomechanics group* (MBM), Heidelberg Institute for Theoretical Studies (HITS), Heidelberg, Germany

**Research in computational biochemistry**: Used molecular dynamics to study mechanical indentation of biological membranes and water flow around peptides

**Scientific software development**: Developed conditional stopping and solvent acceleration features in GROMACS, patched Mesa, Clang, LLVM, and libclc to enable running GROMACS molecular dynamics simulator via OpenCL API on AMD Radeon GPUs "out of the box"

**Computer system maintenance**: Maintained group workstations, backup server, and computational biochemistry software collection on the institute and university supercomputers; installed, configured, and maintained group-owned heterogeneous CPU/GPU computer cluster

#### 2013 – 2014: Adjunct Teaching Assistant, RiTeh

**Institution**: *Department of Computer Engineering*, Faculty of Engineering (RiTeh), University of Rijeka, Rijeka, Croatia

Teaching: Auditory and laboratory exercises in the Computer Networks course

2009 – 2019: (Senior) Research and Teaching Assistant, UniRi

**Institution**: Computer Networks, Parallelization, and Simulation Laboratory (CNPSLab), Department of Informatics, University of Rijeka, Rijeka, Croatia

**Research**: Optical network reliability analysis, modeling, and simulation; applications of heterogeneous computing to network routing and e-learning recommender systems; computational biochemistry, molecular modeling, and *in silico* drug design

**Teaching**: Computer Networks 1, Computer Networks 2 (lead the development of the online laboratory exercises for the course), Network System Management (lectures entrusted in 2018), Operating Systems 1, Operating Systems 2, Distributed Systems, Parallel Programming on Heterogeneous Systems (obtained support from NVIDIA for the course under the CUDA Teaching Center / GPU Education Center program)

**System administration**: Set up and maintained University mirror for several projects including Eclipse, cygwin, GNU, TeX Users Group / CTAN, and LibreOffice / TDF

Organization: Represented Research and Teaching Assistants in Department Council

**Promotion**: Regular public lectures on popular scientific topics

### Education

#### ■ 2009 – 2015: Ph.D. in Computer Science

**Institution**: *Department of Telecommunications*, Faculty of Electrical Engineering and Computing (FER), University of Zagreb, Zagreb, Croatia

**Focus areas**: Optical telecommunication networks, communication network availability and reliability evaluation, computer network simulation and emulation

**Thesis**: *Method for optimizing availability of optical telecommunication network in presence of correlated failures*, supervised by Prof. Branko Mikac (retd.), defended 8th June 2015

#### 2004 – 2009: M.Ed. in Mathematics and Informatics

**Institution**: *Department of Mathematics*, Faculty of Arts and Sciences (FFRi), University of Rijeka, Rijeka, Croatia

**Focus areas**: Real and functional analysis, mathematical foundations of quantum physics, system and network administration, free and open source software movement

Thesis: Banach algebras, supervised by Prof. Cvjetan Jardas (decd.), defended 17th March 2009

End of regular CV; academic CV follows.

## Projects

#### ■ Present projects

2022 – 2024 DPU offload of force reduction calculations in molecular dynamics simulations, led by Dr. Vedran Miletić, funded by NVIDIA

	■ Past projects
2020 – 2022	EuroCC – National Competence Centres in the framework of EuroHPC, led by High-Performance Computing Center Stuttgart (HLRS), funded by Horizon 2020
2020 – 2022	Development of Code Optimization online course for YUFE virtual campus, led by Dr. Vedran Miletić, funded by the University of Rijeka
2019 – 2022	Modernization, modularization, and active maintenance of RxDock, a fast, versatile, and open-source program for docking ligands to proteins and nucleic acids, led by Dr. Vedran Miletić, funded by RxTx Research
2019 – 2021	Biochemistry on a supercomputer: development of new software, drug-design, and analysis of disease development on molecular level, led by Dr. Marta Žuvić, funded by the University of Rijeka, reference number uniri-prirod-18-132
2018-2021	Development of the International Education Program Veleri-OI IoT School, led by Dr. Alen Jakupović, funded by the European Social Fund
2014-2018	Mechano(bio)chemistry, led by Dr. Frauke Gräter, funded by various sources including the German Research Foundation, Heidelberg Institute for Theoretical Studies, and the University of Heidelberg
2014-2016	The development and commercialization of human DNA methyltransferase Dnmt1 inhibitor with a goal to reprogram functional organization of the genome of human cells, led by Dr. Željko Svedružić, funded by the University of Rijeka, reference number 13.11.1.2.04
2014-2016	Recommender system for computer-aided learning (ELARS), led by Dr. Nataša Hoić-Božić, funded by the University of Rijeka, reference number 13.13.1.3.05

2014 – 2016 RFID (Internet of Thing) based animal individual behavior intelligent identification technology and application in traceability (REMALLOY), led by Dr. Maja Matetić and Dr. Zetian Fu, funded by MZOS

**2012 – 2014** Establishment of CUDA Teaching Center at University of Rijeka, led by Vedran Miletić, funded by NVIDIA

**2012 – 2014** Development of Prototype WDM Network Simulator (PWNS), led by Vedran Miletić, no external funding

**2012** Development of e-learning materials for Computer Networks 2 course, led by Vedran Miletić, funded by the University of Rijeka

**2008 – 2011**Building the Future Optical Network in Europe: The e-Photon/ONe Network, led by BONE Consortium, funded by FP7

**2007 – 2013** Supporting e-Business by Distance Learning System Based on Dialogue, led by Dr. Božidar Kovačić, funded by MZOS

## Publications

### Book chapters

1. Nikolić, P., **Miletić, V.**, Odorčić, I. & Svedružić, Ž. M. In Silico Optimization of the First DNA-Independent Mechanism-Based Inhibitor of Mammalian DNA Methyltransferase DNMT1. Epi-Informatics 113–153 (2016). **doi:10.1016/B978-0-12-802808-7.00005-8** 

#### Research papers in journals

- Saftić Martinović, L., Birkić, N., Miletić, V., Antolović, R., Štanfel, D. & Wittine, K. Antioxidant Activity, Stability in Aqueous Medium and Molecular Docking/Dynamics Study of 6-Amino- and N-Methyl-6-amino-L-ascorbic Acid. Int. J. Mol. Sci. 24(2), 1410 (2023). doi:10.3390/ijms24021410 (WoS-SCIE, Q1 (2021), JIF: 6.208 (2021))
- 2. Svedružić, Ž. M, Vrbnjak, K., Martinović, M. & **Miletić, V.** Structural Analysis of the Simultaneous Activation and Inhibition of γ-Secretase Activity in the Development of Drugs for Alzheimer's Disease. Pharmaceutics 13(4), 514 (2021). **doi:10.3390/pharmaceutics13040514** (WoS-SCIE, Q1, JIF: 6.525; times cited: 3)
- 3. Herrera-Rodríguez, A., **Miletić, V.**, Aponte-Santamaría, C., & Gräter, F. Molecular dynamics simulations of molecules in uniform flow. Biophys. J. 116(6), 621–632 (2019). **doi:10.1016/j.bpj.2018.12.025** (WoS-SCIE, Q1, JIF: 3.854; times cited: 7)
- 4. Franz, F., Aponte-Santamaría, C., Daday, C., **Miletić, V.** & Gräter, F. Stability of Biological Membranes upon Mechanical Indentation. J. Phys. Chem. B 122(28), 7073–7079 (2018). doi:10.1021/acs.jpcb.8b01861 (WoS-SCIE, Q2, JIF: 2.923; times cited: 3)
- Miletić, V., Odorčić, I., Nikolić, P. & Svedružić, Ž. M. In silico design of the first DNA-independent mechanism-based inhibitor of mammalian DNA methyltransferase Dnmt1. PLOS ONE 12(4), e0174410 (2017). doi:10.1371/journal.pone.0174410 (WoS-SCIE, Q1, JIF: 2.766; times cited: 14)

#### Research papers in conference proceedings

- 1. **Miletić, V.**, Nikolić, P. & Kinkela, D. Structure-based Molecular Docking in the Identification of Novel Inhibitors Targeting SARS-CoV-2 Main Protease. in 2021 44th International Convention on Information, Communication, and Electronic Technology (MIPRO), 435–440 (2021). doi:10.23919/MIPRO52101.2021.9596660
- 2. **Miletić, V.**, Ašenbrener Katić, M. & Svedružić, Ž. High-throughput Virtual Screening Web Service Development for SARS-CoV-2 Drug Design. in 2020 43rd International Convention on Information, Communication, and Electronic Technology (MIPRO) 371–376 (2020). doi:10.23919/MIPRO48935.2020.9245082
- 3. **Miletić, V.**, Šubić, T. & Mikac, B. Optimizing maximum shared risk link group disjoint path algorithm using NVIDIA CUDA heterogeneous parallel programming platform. in Proceedings on the 2014 X International Symposium on Telecommunications (BIHTEL) (ed. Mrdović, S.; University of Sarajevo, Sarajevo, Bosnia and Herzegovina), 1–6 (IEEE, 2014). **doi:10.1109/BIHTEL.2014.6987645**

- Miletić, V., Holenko Dlab, M. & Hoić-Božić, N. Optimizing ELARS Algorithms Using NVIDIA CUDA Heterogeneous Parallel Programming Platform. in ICT Innovations 2014, Advances in Intelligent Systems and Computing (eds. Bogdanova, A. M. & Gjorgjevikj, D.; University of Skopje, Berlin, Heidelberg) 311, 135–144 (Springer International Publishing, 2015). doi:10.1007/978-3-319-09879-1 14
- Miletić, V., Maniadakis, D., Mikac, B. & Varoutas, D. On the influence of the underlying network topology on optical telecommunication network availability under shared risk link group failures. in Proceedings of the 2014 10th International Conference on the Design of Reliable Communication Networks (DRCN) (ed. Van Daele, P.; University of Ghent, Ghent, Belgium), 1–8 (IEEE, 2014). doi:10.1109/DRCN.2014.6816135
- 6. **Miletić, V.**, Mikac, B. & Džanko, M. Impact evaluation of physical length of shared risk link groups on optical network availability using Monte Carlo simulation. in Proceedings of the 2013 18th European Conference on Networks and Optical Communications (NOC) and 8th Conference on Optical Cabling & Infrastructure (OC&I) (ed. Leitgeb, E.; Technical University Graz, Graz, Austria), 249–256 (IEEE, 2013). doi:10.1109/NOC-OCI.2013.6582897
- 7. Džanko, M., Mikac, B., **Miletić, V.**, Gonzalez, N. A., Zervas, G. S. & Simeonidou, D. Analytical and simulation availability models of ROADM architectures. in Proceedings of the 12th International Conference on Telecommunications (ConTEL) (eds. Pripužić, K. & Banek, M.; University of Zagreb, Zagreb, Croatia), 39–45 (IEEE, 2013).
- 8. **Miletić, V.**, Mikac, B. & Džanko, M. Modelling optical network components: A network simulator-based approach. in Proceedings on the 2012 IX International Symposium on Telecommunications (BIHTEL) (ed. Mrdović, S.; University of Sarajevo, Sarajevo, Bosnia and Herzegovina), 1–6 (IEEE, 2012). doi:10.1109/BIHTEL.2012.6412064
- 9. Džanko, M., Mikac, B. & **Miletić, V.** Availability of all-optical switching fabrics used in optical cross-connects. in Proceedings on the 35th Convention International MIPRO 2012 (ed. Golubić, S.; MIPRO, Opatija, Croatia), 613–617 (IEEE, 2012).

#### Abstracts in conference proceedings

- Koren, R., Martinović, M., Nikolić, P., Odorčić, I., Ostojić, L., Visentin, D., Vrbnjak, K., Miletić, V. & Svedružić, Ž. M. Supercomputers as microscopes for the 21st century: substrate channeling, epigenetic regulation, and molecular basis of Alzheimer's disease. in 27HSKIKI Book of Abstracts, Zagreb, Croatia (Croatian Chemical Society, 2021).
- 2. **Miletić, V.**, Páll, S. & Gräter, F. LLVM AMDGPU for High Performance Computing: are we competitive yet? in 2017 European LLVM Developers' Meeting, Saarbrücken, Germany (2017).
- 3. **Miletić, V.**, Páll, S. & Gräter, F. Towards fully open source GPU accelerated molecular dynamics simulation. in 2016 European LLVM Developers' Meeting, Barcelona, Spain (2016).
- 4. Nikolić, P., Miletić, V. & Svedružić, Ž. M. DNA Methyltransferase Dnmt1: Regulation of Substrate Selectivity. in 6th OEGMBT Annual Meeting 2014 Abstract Book (eds. Khassidov, A., Glaser, W. & Klimek, C.; Austrian Association of Molecular Life Sciences; Biotechnology; Servicebetrieb ÖH-Uni Graz GmbH, Vienna, Austria), 129 (2014).

#### Mentions in research paper acknowledgments

Turalija, M., Petrović, M. & Kovačić, B. Towards General-Purpose Long-Timescale Molecular Dynamics Simulation on Exascale Supercomputers with Data Processing Units. in 2022 45th Jubilee International Convention on Information, Communication, and Electronic Technology (MIPRO), 300–306 (2022). doi:10.23919/MIPRO55190.2022.9803537

Rennekamp, B., Kutzki, F., Obarska-Kosinska, A., Zapp, C. & Gräter, F. Hybrid Kinetic Monte Carlo/Molecular Dynamics Simulations of Bond Scissions in Proteins. *J. Chem. Theory Comput.* 16(1), 553-563 (2020). doi:10.1021/acs.jctc.9b00786 (WoS-SCIE, Q2, JIF: 6.006; times cited: 6)

### ■ References

Prof. Dr	r. Frauke
Gräter	

Group Leader and former Scientific Director (2021 – 2022; deputy 2019 – 2020), Molecular Biomechanics group (MBM), Heidelberg Institute for Theoretical Studies (HITS); Professor for Molecular Biomechanics, Heidelberg University, Heidelberg,

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