Stefan Badelt

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Educational Background

2011 – 2016	Graduate studies in Molecular Biology, University of Vienna
	PhD thesis with Prof. Ivo Hofacker:
	Control of RNA function by conformational design
	Graduation with distinction, PhD
2004 – 2011	Undergraduate studies in Molecular Biology, University of Vienna Master's thesis with Prof. Ivo Hofacker:
	RNA folding kinetics including pseudoknots
	Graduation with distinction, Mag. rer. nat.

Professional Experience

2020/06 – present	Postdoctoral Scholar
	with Ivo L. Hofacker – Theoretical Biochemistry
	Institute for Theoretical Chemistry, Vienna, Austria
2016/04 - 2020/05	Postdoctoral Scholar - Compilers for DNA strand displacement systems
	with Erik Winfree – Division of Biology and Biological Engineering
	California Institute of Technology, Pasadena, USA
2011/10 - 2016/03	Ph.D. thesis – Control of RNA function by conformational design
	with Ivo L. Hofacker – Theoretical Biochemistry
	Institute for Theoretical Chemistry, Vienna, Austria
WS2013 - SS2015	Teaching – Exercises for Foundations of Bioinformatics
	University of Vienna, Austria
2009/05 - 2011/09	Master's thesis – RNA folding kinetics including pseudoknots
	with Ivo L. Hofacker – Theoretical Biochemistry
	Institute for Theoretical Chemistry, Vienna, Austria
2008/07 - 2008/09	Internship – Chromosome degradation in apoptotic cells
	with Reinhard Ullmann – Molecular Cytogenetics
	Max Planck Institute for Molecular Genetics, Berlin, Germany
2008/03 - 2008/04	Internship – Interaction of Stat1-GRDBD-Stat1 and GRE
	with Pavel Kovarik – Infection Biology
	Max F. Perutz Laboratories, Vienna, Austria
2006/07 - 2009/03	Technician – Plasmid library administration, genotyping
	Max F. Perutz Laboratories, Vienna, Austria
	Group Kovarik – Infection Biology

Funding Awards

2016/05	Caltech Biology and Biological Engineering Division Fellowship
2011/05	Max F. Perutz Laboratories PhD program selection - Track RNA biology

Skills

- Languages: German (native), fluent English
- Computer Skills: Python, Perl, Bash, LaTeX, julia, R, C, C++, ...
- Lab-Techniques: PCR, Real Time PCR, Tissue Culture work (including Nucleofection), Immunfluorescence, Immunoprecipitation, Nuclear Extract, Western Blot Analysis, Electrophoretic Mobility Shift Assay, DNA/RNA Extraction, DNA/RNA Gel Electrophoresis, Reverse Transcription, Array CGH, Oligoarray, BAC Array, ChIP on Chip.
- Snowboard and Windsurfing instructor

Selected Publications

- [2020] S. Badelt, C. Grun, K. Sarma, B. Wolfe, S. W. Shin, and E. Winfree, "A domain-level DNA strand displacement reaction enumerator allowing arbitrary non-pseudoknotted secondary structures," *Journal of the Royal Society Interface*, vol. 17, p. 20190866, 2020.
- [2018] J. Berleant, C. Berlind, S. Badelt, F. Dannenberg, J. Schaeffer, and E. Winfree, "Automated sequence-level analysis of kinetics and thermodynamics for domain-level DNA stranddisplacement systems," *Journal of the Royal Society Interface*, vol. 15, p. 20180107, 2018.
- [2017] S. Badelt, S. W. Shin, R. F. Johnson, Q. Dong, C. Thachuk, and E. Winfree, "A generalpurpose CRN-to-DSD compiler with formal verification, optimization, and simulation capabilities," in *International Conference on DNA-Based Computers*, pp. 232–248, Springer, 2017.
- [2016] S. Badelt, Control of RNA function by conformational design. PhD thesis, University of Vienna, 2016.
- [2016] M. Tajaddod, A. Tanzer, K. Licht, M. T. Wolfinger, S. Badelt, F. Huber, O. Pusch, S. Schopoff, M. Janisiw, I. Hofacker, and M. F. Jantsch, "Transcriptome-wide effects of inverted SINEs on gene expression and their impact on RNA Polymerase II activity," *Genome Biology*, vol. 17, p. 220, 2016.
- [2016] S. Badelt, C. Flamm, and I. L. Hofacker, "Computational design of a circular RNA with prionlike behavior," Artificial Life, vol. 22, pp. 1–13, 2016.
- [2015] S. Petkovic, S. Badelt, S. Block, C. Flamm, M. Delcea, I. L. Hofacker, and S. Müller, "Sequence-controlled RNA self-processing: computational design, biochemical analysis and visualization by AFM," *RNA*, vol. 21, pp. 1249–1260, 2015.
- [2015] S. Badelt, S. Hammer, C. Flamm, and I. L. Hofacker, "Thermodynamic and kinetic folding of riboswitches," in *Methods in Enzymology*, vol. 553, pp. 193–213, Elsevier, 2015.
- [2014] M. Marz, A. R. Gruber, C. Höner zu Siederdissen, F. Amman, S. Badelt, S. Bartschat, S. H. Bernhart, W. Beyer, S. Kehr, R. Lorenz, A. Tanzer, D. Yusuf, H. Tafer, I. L. Hofacker, and P. F. Stadler, "Animal snoRNAs and scaRNAs with exceptional structures," *RNA biology*, vol. 8, pp. 938–946, 2011.

Selected Conferences, Talks & Posters

- FNANO19 Conference in Snowbird, Utah, April 15 18, 2019 Talk: Enumeration, condensation and simulation of pseudoknot-free domain-level DNA strand displacement systems
- Gordon Research Conference on RNA nanotechnology in Ventura, USA, Jan 13 18, 2019 Poster: Compilation and verification of nucleic acid reaction networks
- DNA23 Conference in Austin, Texas, Sept 24 28, 2017 Talk: A General-Purpose CRN-to-DSD Compiler with Formal Verification, Optimization, and Simulation Capabilities
- DNA22 Conference in Munich, Germany, Sept 4 8, 2016 Talk: Energy landscapes and folding kinetics of pairwise interacting RNAs Poster: Nuskell: A verifying and optimizing CRN-to-DSD compiler
- Gordon Research Conference on RNA nanotechnology in Ventura, USA, Feb 1 6, 2015 Poster: Design of XOR riboswitches
- Artificial Life Conference in New York, USA, Jul 30 Aug 2, 2014 Talk: Design of a circular RNA with prion-like behavior
- International Synthetic and Systems Biology Summer School in Taormina, Italy, Jun 15 19, 2014
 Poster: Sequence-controlled RNA self-processing: computational design, biochemical analysis and visualization by AFM
- *Herbstseminar Bioinformatik* in Decin, Czech Republic, Oct 2 7, 2013 Talk: Circularization and multimerization of synthetic ribozymes
- *TBI Winterseminar* in Bled, Slovenia, Feb 13 20, 2011 Talk: Energy barriers in pseudoknot conformation space
- *TBI Winterseminar* in Bled, Slovenia, Feb 14 21, 2010 Talk: Design of artificial RNA-switches