PERSONAL INFORMATION

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EDUCATION

09/2015	Dr. rer. nat. in Biology (summa cum laude)
	Dresden International PhD Program / TU Dresden (Germany)

01/2011 **Diploma in Biochemistry** (excellent) University of Tübingen (Germany)

STAGES OF ACADEMIC CAREER & RESEARCH EXPERIENCE

Since 06/23	W1 TT W2 Juniorprofessor at Freie Universität Berlin
	Department of Biology, Chemistry, Pharmacy
	Institute for Chemistry and Biochemistry

- 03/22-05/23 **Postdoctoral researcher at University Medical Center Göttingen** Molecular Biology department Group of Prof. Dr. Markus Bohnsack
- 07/16-12/21 **Postdoctoral associate at Massachusetts Institute of Technology (MIT)** Biology department Group of Prof. Gene-Wei Li, PhD
- 10/15-06/16 **Postdoctoral associate at Yale University** Molecular Biophysics and Biochemistry department Group of Prof. Karla M. Neugebauer, PhD
- 06/11-09/15 Graduate study at the Max Planck Institute of Molecular Cell Biology and Genetics (MPI-CBG) and Yale University Supervisor: Prof. Karla M. Neugebauer, PhD Thesis title: "Co-transcriptional splicing in two yeasts"

05/10-10/10 **Diploma thesis at Harvard Medical School** Supervisor: Prof. Charles J. Weitz, MD, PhD Thesis title: "Biochemistry of circadian clocks - 24 hour high-resolution analysis of clock gene expression and isolation of cytoplasmic PERIOD 2 complexes"

2009 Student assistant at the Center for Plant Molecular Biology (ZMBP) of the University of Tübingen 6 months Supervisor: Prof. Dr. Rita Gross-Hardt

RECOGNITION

2018	Infinite Kilometer Award to recognize the outstanding contributions made by postdoctoral researchers and research staff at MIT
2017	Personal grant by the Helen Hay Whitney Foundation (3-year postdoctoral fellowship)
2017	Personal grant by the Deutsche Forschungsgemeinschaft (DFG, 2-year post- doctoral fellowship, declined)
2015	Best poster award at the Annual Yale Department of Molecular Biophysics & Biochemistry Retreat
2013	Travel award Dresden International Graduate School for Biomedicine and Bioengineering
2012	Travel grant for the EMBO Conference "Gene Transcription in Yeast"
2012	Travel grant for the COST (Cooperation in Science and Technology) Training School on Next Generation Sequencing
2011	Max Planck Society PhD stipend
2010	5 month stipend and travel grant of the German Academic Exchange Service to work on the diploma thesis at the Harvard Medical School (Boston, USA)

TEACHING ACTIVITIES

SS 24	Advanced Biochemistry I: lecture + seminar "Complex, but precise bacterial gene expression" for master students in Biochemistry, FU Berlin
WS 23/24	Biochemistry II (metabolism & regulation): 4 lectures + seminars for bache- lor students in Biochemistry, FU Berlin
WS 23/24	Enzyme models lecture for master students in Molecular Medicine, Charité Berlin
WS 23/24	Lecture & hands-on practical for high school students in the program "NATürlich Studium", NATlab FU Berlin
2018	Guest lecture "Kinetics of RNA splicing from sequencing nascent RNA" at MIT undergraduate course 7.09 "Quantitative & Computational Biology" taught by Prof. Chris Burge and Prof. Gene-Wei Li
2018	Completion of the Kaufman Teaching Certificate Program at MIT
2006 - 2008	Teaching Assistant "Chemistry and Biochemistry" for medical students at the University of Tübingen (recitation, practical course, premedical exam preparation)

PUBLICATIONS

2024	The spliceosome impacts morphogenesis in the human fungal pathogen <i>Can- dida albicans</i> . Lash E, Maufrais C, Janbon G, Robbins N, Herzel L *, Cowen LE*. mBio. 2024 Jul 9, *co-corresponding authors PMID:38980041
2024	Co-transcriptional gene regulation: common ground across eukaryotes and prokaryotes. Shine M, Gordon J, Schärfen L, Zigackova D, Herzel L* , and Neugebauer KM*. Nat Rev Mol Cell Biol. 2024 Mar 20, *co-corresponding authors PMID:38509203
2023	N2-methylguanosine modifications on human tRNAs and snRNA U6 are important for cell proliferation, protein translation and pre-mRNA splicing. Wang C, Ulryck N, Herzel L , Pythoud N, Kleiber N, Guérineau V, Jactel V, Moritz C, Bohnsack MT, Carapito C, Touboul D, Bohnsack KE, Graille M Nucleic Acids Res. 2023 Jun 7 PMID:37283053
2022	Ubiquitous mRNA decay fragments in <i>E. coli</i> redefine the functional tran- scriptome. Herzel L , Stanley JA, Yao CC, Li GW Nucleic Acids Research. 2022 May 7 PMID:35524564
2020	Widespread Transcriptional Readthrough Caused by Nab2 Depletion Leads to Chimeric Transcripts with Retained Introns. Alpert T, Straube K, Carrillo Oesterreich F, Herzel L , Neugebauer KM Cell Reports. 2020 Oct 27 PMID:33113357
2018	Long-read sequencing of <i>S. pombe</i> nascent RNA reveals coupling among RNA processing events. Herzel L , Straube K, Neugebauer KM Genome Research. 2018 Jun 14 PMID:29903723
2018	Evolutionary Convergence of Pathway-Specific Enzyme Expression Stoichio- metry. Lalanne JB, Taggart JC, Guo MS, Herzel L , Schieler A, Li GW Cell. 2018 Mar 23 PMID:29606352
2017	Splicing and transcription touch base: co-transcriptional spliceosome as- sembly and function. Herzel L*, Ottoz DSM*, Alpert T, Neugebauer KM Nat Rev Mol Cell Biol. 2017 Oct 18, * co-first authors PMID:28792005
2017	Dynamic RNA-protein interactions underlie the zebrafish maternal-to-zygotic transition. Despic V, Dejung M, Gu M, Krishnan J, Zhang J, Herzel L , Straube K, Gerstein MB, Butter F, Neugebauer KM

 2016 Perfect timing: splicing and transcription rates in living cells. Alpert T, Herzel L and Neugebauer KM WIREs RNA. 2016 Nov 21 PMID:27873472 2016 Splicing of nascent RNA coincides with intron exit from RNA polymerase II. Carrillo Oesterreich F*, Herzel L*, Straube K, Hujer K, Howard J, and Neugebauer KM Cell. 2016 Mar 24, * co-first authors, listed alphabetically PMID:27020755 2015 Quantification of co-transcriptional splicing from RNA-Seq data. Herzel L, Neugebauer KM Methods. 2015 Apr 27 PMID:25929182 2013 Counting on co-transcriptional splicing. Brugiolo M*, Herzel L*, Neugebauer KM F1000Prime Rep. 2013 Apr 2, * co-first authors, listed alphabetically PMID:23638305 		Genome Research. 2017 Apr 5 PMID:28381614
 Carrillo Oesterreich F*, Herzel L*, Straube K, Hujer K, Howard J, and Neugebauer KM Cell. 2016 Mar 24, * co-first authors, listed alphabetically PMID:27020755 Quantification of co-transcriptional splicing from RNA-Seq data. Herzel L, Neugebauer KM Methods. 2015 Apr 27 PMID:25929182 Counting on co-transcriptional splicing. Brugiolo M*, Herzel L*, Neugebauer KM F1000Prime Rep. 2013 Apr 2, * co-first authors, listed alphabetically 	2016	Alpert T, Herzel L and Neugebauer KM WIREs RNA. 2016 Nov 21
 Herzel L, Neugebauer KM Methods. 2015 Apr 27 PMID:25929182 2013 Counting on co-transcriptional splicing. Brugiolo M*, Herzel L*, Neugebauer KM F1000Prime Rep. 2013 Apr 2, * co-first authors, listed alphabetically 	2016	Carrillo Oesterreich F*, Herzel L *, Straube K, Hujer K, Howard J, and Neuge- bauer KM Cell. 2016 Mar 24, * co-first authors, listed alphabetically
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	2013	Brugiolo M*, Herzel L *, Neugebauer KM F1000Prime Rep. 2013 Apr 2, * co-first authors, listed alphabetically

Berlin, July 25, 2024