
BIOGRAPHICAL SKETCH

NAME: Janovjak, Harald

POSITION TITLE: EMBL Australia Group Leader

EDUCATION/TRAINING:

INSTITUTION AND LOCATION	DEGREE (if applicable)	Completion Date MM/YYYY	FIELD OF STUDY
Biozentrum, University of Basel, Switzerland	Dipl./M.S.	11/2002	Molecular Biology
University of Technology Dresden, Germany	Dr.	12/2005	Biophysics, Biology
University of California Berkeley, USA	Post-doctoral	06/2010	Optogenetics, Neuroscience
Ludwig Maximilian University, Munich, Germany	Post-doctoral	02/2011	Optogenetics, Neuroscience

A. Personal Statement

My research has always revolved around understanding cell signaling and membrane receptors with new and innovative approaches. I followed this common thread throughout my career that brought me to the fields of protein folding/instrumentation development, optogenetics/neuroscience and cell signaling/physiology. My work resulted in well-cited publications at all stages of my career (May 1, 2018: 46 total publications: 29 peer-reviewed research articles, 8 review articles/perspectives and 9 book chapters/protocol articles; H-index 20; >2300 citations). More information about my current research can be found here: <http://www.janovjak-lab.com>

B. Positions and Honors

Positions:

- 2011 - 2017 Assistant Professor in Synthetic Physiology
Institute of Science and Technology Austria (IST Austria)
- 2018 - Group Leader, EMBL Australia Partnership Laboratory (EMBL Australia)
Monash University, Faculty of Medicine, Nursing and Health Sciences
- 2018 - Group Leader, Australian Regenerative Medicine Institute
Monash University, Faculty of Medicine, Nursing and Health Sciences

Honors:

- 2002 Graduate research with highest honors
- 2005 Doctorate with highest honors (*summa cum laude*)
- 2007 - 2009 Long-term Fellow of the European Molecular Biology Organization
- 2010 Returning Fellows Grant, Ministry of Science and Education, Düsseldorf
- 2011 Human Frontier Science Program Young Investigators Grant
- 2011 European Union Seventh Framework Programme Career Integration Grant
- 2018 JDRF/Macquarie Group Foundation Future Research Leader

Fellowships:

- 2007 Post-doctoral Fellowship, German Research Foundation
- 2007 Young Investigator Fellowship, Swiss National Science Foundation
- 2007 - 2009 Long-term Fellowship, European Molecular Biology Organization

Memberships:

- 2005 Biophysical Society (Germany)
- 2007 Biophysical Society (US)
- 2011 Competence Center Cancer Research (CCC, Austria)
- 2011 Biophysical Society (Austria)
- 2012 Austrian Association of Molecular Life Sciences and Biotechnology (OEGMBT)

2012	Society for Neuroscience (SFN, US)
2013	American Society for Cell Biology (ASCB)
2014	European Association for the Study of Diabetes (EASD)

Reviewer for funding agencies:

Alexander von Humboldt-Foundation (AvH), Boehringer Ingelheim Foundation (BIF), European Research Council (ERC), French National Alliance for Life and Health Sciences, French National Cancer Institute, German-Israeli Foundation for Scientific Research and Development, German Research Foundation (DFG), Israel Science Foundation (ISF), L'Agence nationale de la recherche (ANR), Ministry of Education and Science of the Russian Federation, Ministry of Health Italy, Netherlands Organisation for Scientific Research (NWO)

C. Contributions to Science

1. How do membrane proteins fold and unfold?

Protein folding is not yet understood. The discovery of parallel folding and unfolding pathways highlights that single-molecule techniques are required to address the folding problem. As a graduate student at TU Dresden, I applied forces generated by atomic force microscopy (AFM) to denature single membrane proteins. I was able to measure the energy stabilizing α -helices in the light-driven proton pump bacteriorhodopsin and provided the first experimental measure of roughness in a membrane protein's energy landscape. I also conducted the first combined thermal and mechanical unfolding experiment and, through instrumentation development, probed protein visco-elastics and extended the dynamic range of the technique. We also followed single proteins as they proceeded through the folding process revealing detailed insights into otherwise inaccessible kinetics and pathways. To complement the experiments, I started a collaboration with Marek Cieplak that yielded the first theoretical study of mechanical membrane protein unfolding.

a. "Transmembrane helices have rough energy surfaces"

H. Janovjak, H. Knaus & D.J. Müller

Journal of the American Chemical Society (2007) 129: 246-247.

b. "Observing folding pathways and kinetics of a single sodium-proton antiporter from *E. coli*"

A. Kedrov, **H. Janovjak**, C. Ziegler, W. Kühlbrandt & D.J. Müller

Journal of Molecular Biology (2006) 355: 2-8.

c. "Hydrodynamic effects in fast AFM single molecule force measurements"

H. Janovjak, J. Struckmeier & D.J. Müller

European Biophysics Journal (2005) 34: 91-96.

d. "Probing the energy landscape of the membrane protein bacteriorhodopsin"

H. Janovjak, J. Struckmeier, M. Hubain, M. Kessler, A. Kedrov & D.J. Müller

Structure (2004) 12: 871-879 (with front cover).

e. "Unfolding pathways of native bacteriorhodopsin depend on temperature"

H. Janovjak, M. Kessler, D. Oesterhelt, H.E. Gaub & D.J. Müller

EMBO Journal (2003) 22: 5220-5229.

2. Can we control nerve cell activity using light?

Genetically-targeted, light-gated ion channels can drive activity in selected nerve cells and thereby probe neural circuits. As a post-doctoral fellow at UC Berkeley, I created novel ion channels that are controlled by photochromic tethered ligands. I used protein engineering to redesign a glutamate receptor into a light-activated, K^+ -selective ion channel called HyLighter. This novel functionality was achieved by inserting a functional domain of a bacterial protein into its mammalian homologue. When expressed in neuronal cultures, brain slices and zebrafish and activated by a brief light pulse, HyLighter stably but reversibly inhibits action potential firing and behavior. The low light intensity required for HyLighter activation and its bi-stability proves advantageous for the dissection of neural circuitry *in vivo* where activity is first silenced and behavior is then analyzed in free animals. Motivated by the functional compatibility of a K^+ channel pore with a glutamate receptor demonstrated in HyLighter, I discovered a new family of glutamate receptors in lower invertebrates that carry a K^+ selectivity filter in their pore domains.

a. "A modern ionotropic glutamate receptor with a potassium-selectivity signature sequence"

H. Janovjak, G. Sandoz & E.Y. Isacoff

Nature Communications (2011) 2: 232.

b. "A light-gated, potassium-selective glutamate receptor for the optical inhibition of neuronal firing"

3. Optical control of cellular signals and cell behavior

The impact of light control on cell and developmental biology has been limited. In my independent laboratory at IST Austria, we established optogenetic tools that control cell signaling pathways with spatial and temporal precision. We pioneered light-activated receptor tyrosine kinases and light-activated serine/threonine kinases (called 'Opto-RTKs' and 'Opto-RSTKs', e.g. 'Opto-EGFR', our light-activated EGF receptor). We and others utilize these receptors to non-invasively control key signaling cascades (e.g. MAPK or PI3K) and cell behaviors (e.g. cell proliferation and differentiation) with unprecedented precision in different areas of cell and developmental biology. To advance the field as a whole, we share published (and often also unpublished) materials 'with-no-strings-attached', and groups at ETH, CNRS, Yale, CalTech, UCSF, Columbia, McGill, Princeton, UCL, to just name a few institutions, requested our constructs. We further pioneered the application of optogenetics to proteins whose function is currently not known. A large number human G-protein coupled receptors (GPCRs) are classified as 'orphans' as their ligands, downstream pathways and physiological roles remain elusive. To identify downstream signals of these proteins and to establish tools with which they can be studied in the relevant cellular context, we reengineered almost all human orphan GPCRs to be activated by light. Our results show that, in contrast to recent models, many orphan GPCRs are signaling-competent proteins that activate distinct signaling pathways.

a. "Green light-induced inactivation of receptor signaling using cobalamin-binding domains"

S. Kainrath, M. Stadler, E. Reichhart, M. Distel & **H. Janovjak**
Angewandte Chemie Int. Ed. (2017) 56: 4608-4611.

b. "Optogenetic control of nodal signaling reveals a temporal pattern of nodal signaling regulating cell fate specification during gastrulation"

K. Sako, S.J. Pradhan, V. Barone, A. Ingles-Prieto, K.W. Rogers, P. Müller, V. Ruprecht, D. Capek, S. Galande, **H. Janovjak** & C.P. Heisenberg
Cell Reports (2016) 16: 866-877.

c. "A phytochrome sensory domain permits receptor activation by red light"

E. Reichhart, A. Ingles-Prieto, A.M. Tichy, C. McKenzie & **H. Janovjak**
Angewandte Chemie Int. Ed. (2016) 55: 6339-6342.

d. "Spatio-temporally precise activation of engineered receptor tyrosine kinases by light"

M. Grusch, K. Schelch, R. Riedler, E. Reichhart, C. Differ, W. Berger, A. Ingles-Prieto & **H. Janovjak**
EMBO Journal (2014) 33: 1713-1726.

e. "Optical functionalization of human Class A orphan G-protein coupled receptors"

M. Morri, I. Sanchez-Romero, A.M. Tichy, S. Kainrath, E.J. Gerrard, P. Hirschfeld, J. Schwarz & **H. Janovjak**
Nature Communications (2018) 9: 1950.

f. We filed three patent applications connected to this work.

4. Light permits a new paradigm in drug screening

We and others proposed that the non-invasive nature of optogenetics may be exploited to create new experimental paradigms in small molecule screening. In my independent laboratory at IST Austria, we recently demonstrated the first optogenetics-assisted small molecule screen. In our 'all-optical' system, light acted both as activator and read-out to obviate the use of assay chemicals (e.g. peptide ligands or reagents for read-out), to limit the number of required operational steps (e.g. solution exchange or other invasive actions) and to improve specificity. Using the all-optical platform, we identified AV-951 (Tivozanib) as a potent and selective inhibitor of ROS1, an orphan kinase that we for the first time activated with light in the absence of the unknown ligand.

a. "Light-assisted small molecule screening against protein kinases"

A. Ingles-Prieto, E. Reichhart, M.K. Muellner, M. Nowak, S.M. Nijman, M. Grusch & **H. Janovjak**
Nature Chemical Biology (2015) 11: 952-954.

b. "Optogenetic methods in drug screening: Technologies and applications"

V. Agus & **H. Janovjak**
Current Opinion in Biotechnology (2017) 5: 8-14.

c. We filed one patent application connected to this work.