

Poster presentations – „Mechanism of Gene Regulation 2014“

Author(s)	Institution	Title	Poster No.
<u>Pinto D.</u> , Huang X., Fritz G., Mascher T.	LMU München	Actinobacterial Intelligence: a comprehensive survey of signal transduction in this phylum	P01
<u>Tobias Busche</u> , Christian Rückert, Jörn Kalinowski	Center for Biotechnology, Bielefeld University	ROSE: an <i>in vitro</i> RNA-seq based approach to dissect hierarchies and interactions in sigma factor regulons	P02
Jennifer Pahlke, Michael Bott, Miroslav Patek, Julia Frunzke and <u>Tino Polen</u>	Forschungszentrum Jülich	Small 6C RNA from <i>Corynebacterium glutamicum</i>	P03
<u>Dharmender Kumar</u> , Sheila MacIntyre	University of Reading	Unraveling the expression control of <i>Yersenia pestis</i> F1 antigen: a novel vaccine target against plague	P04
<u>S. Wörner</u> , P. Steinmetz, M. Zeltner and G. Uden	Johannes Gutenberg-Universität Mainz	Sensing of C4-Dicarboxylates in the DcuB/DcuS and the DctA/DcuS Sensor complexes	P05
<u>A. Kretzschmar</u> , S. Nilkens and G. Uden	Johannes Gutenberg-Universität Mainz	The DxxxQ phosphatase motif of the sensor kinase NreB of <i>Staphylococcus carnosus</i>	P06
<u>Marc Keppel</u> , Eva Hentschel, Thomas Gensch and Julia Frunzke	Forschungszentrum Jülich	Coordinated control of heme-homeostasis by the ChrSA and HrrSA two-component systems in <i>Corynebacterium glutamicum</i>	P07
<u>Oliver Goldbeck</u> , Anna M. Bartsch, Ines Ochrombel, Kay Marin, Reinhard Krämer, Gerd M. Seibold	Universität zu Köln	The second messenger cyclic di-AMP controls potassium uptake in <i>Corynebacterium glutamicum</i> by binding to the RCK domain of the channel protein CgIK	P08
<u>Stephanie Trauth</u> , Ilka B. Bischofs	Universität Heidelberg	Fluorescent promoter reporters exhibiting minimal noise for quantitative single cell studies in <i>B. subtilis</i>	P09
<u>Steffi Heyber</u> , Jenny Jacobs, Elisabeth Härtig, Richard Münch, Dieter Jahn	Technische Universität Braunschweig	Light-dependent regulation of bacteriochlorophyll biosynthesis in the aerobic anoxygenic phototrophic bacteria <i>Dinoroseobacter shibae</i>	P10
<u>Yvonne Engel</u> , Safia Hemati and Helge B. Bode	Goethe Universität Frankfurt	Heterogeneity in the Secondary Metabolism of <i>Photorhabdus luminescens</i> and <i>Xenorhabdus nematophila</i>	P11

<u>Eugen Pfeifer</u> , Stefan Helfrich, Christina Krämer, Dietrich Kohlheyer, Katharina Nöh, and Julia Frunzke	<i>Forschungszentrum Jülich</i>	Spatiotemporal analysis of SOS and prophage dynamics in <i>Corynebacterium glutamicum</i> populations	P12
<u>Stefan Helfrich</u> , Christina Krämer, Eugen Pfeifer, Wolfgang Wiechert, Julia Frunzke, Dietrich Kohlheyer, and Katharina Nöh	<i>Forschungszentrum Jülich</i>	Microfluidics as platform for the spatiotemporal analysis of bacterial populations at single-cell level	P13
<u>Lydia Segler</u> and Ute Lechner	<i>Martin-Luther Universität Halle-Wittenberg</i>	Transcriptional Regulation of Reductive Dehalogenase-Encoding Genes in <i>Dehalococcoides mccartyi</i> by a MarR-Type Regulator	P14
Nora Kuhlmann, Dimitar P. Petrov, Alexander W. Henrich, <u>Gerd M. Seibold</u>	<i>Universität zu Köln</i>	PTS controls maltose metabolism in <i>Corynebacterium glutamicum</i> : Identification of <i>malP</i> -transcription as target of PTS-dependent regulation	P15
Anja Heinz, Nicole Pasian, Marie-Luise Hoffmann, <u>Falk Kalamorz</u>	<i>Martin-Luther Universität Halle-Wittenberg</i>	Revisiting the Expression of the <i>atp</i> Operon in <i>Escherichia coli</i>	P16
<u>Hannes Breddermann</u> and Karin Schnetz	<i>Universität zu Köln</i>	Feedback control of <i>leuO</i> encoding a pleiotropic regulator and H-NS antagonist in <i>Escherichia coli</i>	P17
<u>N. Brühl</u> , A. Uhde, C. Matano, V. F. Wendisch, R. Krämer, G.M. Seibold	<i>Universität zu Köln</i>	The transcriptional repressor NanR controls sialic acid catabolism in <i>Corynebacterium glutamicum</i>	P18
<u>Meike Baumgart</u> , Kerstin Luder, Shipra Grover, Gurdyal S Besra, Julia Frunzke	<i>Forschungszentrum Jülich</i>	IpsA, a novel LacI-type regulator required for cell wall biosynthesis in Corynebacteria and Mycobacteria	P19
<u>Joana Simen</u> , Michael Löffler, Alexander Broicher, Andreas Freund and Ralf Takors	<i>Universität Stuttgart</i>	Modeling Regulation of Carbon and Nitrogen Metabolism in Large-Scale Processes with <i>Escherichia coli</i>	P20

<u>Timo Fettweiss</u> , Kathrin Emmi Scholz, Martin Diener, Karl-Erich Jaeger and Ulrich Krauss	<i>Heinrich-Heine-Universität Düsseldorf</i>	Light, Oxygen, Voltage (LOV) photoreceptor signalling probed by the design of artificial light-gated histidine kinases	P21
<u>Raphael Freiherr von Boeselager</u> , Regina Mahr, Stefan Helfrich, Katharina Nöh, Dietrich Kohlheyer, Julia Frunzke	<i>Forschungszentrum Jülich</i>	High-throughput screening of a fluorescent reporter library for novel metabolite biosensors	P22
<u>Christina Krämer</u> , Eugen Pfeifer, Andrea Rocker, Stefan Helfrich, Katharina Nöh, Anton Meinhart, Wolfgang Wiechert, Julia Frunzke, Dietrich Kohlheyer	<i>Forschungszentrum Jülich</i>	Bacterial Single-Cell Stress Studies in Controlled Microfluidic Environments: Enabling a Phenotypical Insight on Bacteria Tackling for Survival	P23