

Publication list Prof. Dr. Michael Bott

2021

219. **Ramp, P., Lehnert, A., Matamouros, S., Wirtz, A., Baumgart, M. and Bott, M. (2021)**
Metabolic engineering of *Corynebacterium glutamicum* for production of scyllo-inositol, a drug candidate against Alzheimer's disease
Metab. Eng. 67: 173-185 (<https://doi.org/10.1016/j.ymben.2021.06.011>)
218. **Wolf, S., Becker J., Tsuge, Y., Kawaguchi, H., Kondo, A., Marienhagen, J., Bott, M., Wendisch, V.F., and Wittmann, C. (2021)**
Advances in metabolic engineering of *Corynebacterium glutamicum* to produce high-value active ingredients for food, feed, human health, and well-being
Essays Biochem. 65: 197-212 (<https://doi.org/10.1042/EBC20200134>)
217. **Schweikert, S., Kranz, A., Yakushi, T., Filipchyk, A., Polen, T., Etterich, H., Bringer, S. and Bott, M. (2021)**
The FNR-type regulator GoxR of the obligatory aerobic acetic acid bacterium *Gluconobacter oxydans* affects expression of genes involved in respiration and redox metabolism
Appl. Environm. Microbiol. 87: e00195-21 (<https://doi.org/10.1128/AEM.00195-21>)
216. **Fricke, P.M., Klemm, A., Bott, M., and Polen, T. (2021)**
On the way toward regulatable expression systems in acetic acid bacteria: target gene expression and use cases
Appl. Microbiol. Biotechnol. 105: 3423–3456 (<https://doi.org/10.1007/s00253-021-11269-z>)
215. **Zelle, E., Pfelzer, N., Oldiges, M., Koch-Koerfges, A., Bott, M., Nöh, K., and Wiechert, W. (2021)**
An energetic profile of *Corynebacterium glutamicum* underpinned by measured biomass yield on ATP
Metab. Eng. 65: 66–78 (<https://doi.org/10.1016/j.ymben.2021.03.006>)
214. **Maeda, T., Koch-Koerfges, A. and Bott, M. (2021)**
Relevance of NADH dehydrogenase and alternative two-enzyme systems for growth of *Corynebacterium glutamicum* with glucose, lactate, and acetate
Front. Biotechnol. Bioeng. 8: 621213 (<https://doi.org/10.3389/fbioe.2020.621213>)

2020

213. **Zhu, L., Mack, C., Wirtz, A., Kranz, A., Polen, T., Baumgart, M. and Bott, M. (2020)**
Regulation of γ-aminobutyrate (GABA) utilization in *Corynebacterium glutamicum* by the PucR-type transcriptional regulator GabR and by alternative nitrogen and carbon sources
Front. Microbiol. 11: 544045 (<https://doi.org/10.3389/fmicb.2020.544045>)
212. **Bakkes, P.J., Ramp, P., Bida, A., Dohmen-Olma, A., Bott, M. and Freudl, R. (2020)**
Improved pEKEx2-derived expression vectors for tightly controlled production of recombinant proteins in *Corynebacterium glutamicum*
Plasmid 112: 102540 (<https://doi.org/10.1016/j.plasmid.2020.102540>)
211. **Fricke, P.M., Link, T., Gätgens, J., Sonntag, C., Otto, M., Bott, M. and Polen, T. (2020)**
A tunable L-arabinose-inducible expression plasmid for the acetic acid bacterium *Gluconobacter oxydans*
Appl. Microbiol. Biotechnol. 104: 9267-9282 (<https://doi.org/10.1007/s00253-020-10905-4>)
210. **Keppel, M., Hünnefeld, M., Filipchyk, A., Viets, U., Davoudi, C.-F., Krüger, A., Mack, C., Pfeifer, E., Polen, T., Baumgart, M., Bott, M. and Frunzke, J. (2020)**

- HrrSA orchestrates a systemic response to heme and determines prioritization of terminal cytochrome oxidase expression
Nucleic Acids Res. 48: 6547-6562 (<https://dx.doi.org/10.1093/nar/gkaa415>)
209. **Kircher, M., Bott, M., and Marienhagen, J. (2020)** The importance of biotechnology for the bioeconomy. In: Bioeconomy for beginners. J. Pietzsch (ed.), Springer-Verlag, Heidelberg, pp. 105-128 (http://dx.doi.org/10.1007/978-3-662-60390-1_5)
208. **Bott, M. (2020)**
Nutzung mikrobieller und pflanzlicher Diversität in der Biotechnologie
In: Spree, D., Kandarr, J. (Eds.), Biodiversität im Meer und an Land. Vom Wert biologischer Vielfalt. Potsdam: Deutsches Geoforschungszentrum GFZ, 50 - 53
(<https://dx.doi.org/10.2312/eskp.2020.1.2.3>)
207. **Küberl, A., Mengus-Kaya, A., Polen, T. and Bott, M. (2020)**
The iron deficiency response of *Corynebacterium glutamicum* and a link to thiamine biosynthesis.
Appl. Environm. Microbiol. 86: e00065-20 (<https://dx.doi.org/10.1128/AEM.00065-20>)
206. **Battling, S., Wohlers, K., Igwe, C., Kranz, A., Pesch, M., Wirtz, A., Baumgart, M., Büchs, J. and Bott, M. (2020)**
Novel plasmid-free *Gluconobacter oxydans* strains for production of the natural sweetener 5-ketofructose
Microb. Cell Fact. 19: 54 (<https://dx.doi.org/10.1186/s12934-020-01310-7>)
205. **Wolf, N., Bussmann, M., Koch-Koerfges, A., Katcharov, N., Schulte, J., Polen, T., Hartl, J., Vorholt, J.A., Baumgart, M. and Bott, M. (2019)**
Molecular basis of growth inhibition by acetate of an adenylate cyclase-deficient mutant of *Corynebacterium glutamicum*
Front. Microbiol. 11: 87 (<http://dx.doi.org/10.3389/fmicb.2020.00087>)
204. **Spielmann, A., Brack, Y., van Beek, H., Flachbart, L., Sundermeyer, L., Baumgart, M. and Bott, M. (2020)**
NADPH biosensor-based identification of an alcohol dehydrogenase variant with improved catalytic properties caused by a single charge reversal at the protein surface
AMB Expr 10: 14 (<http://dx.doi.org/10.1186/s13568-020-0946-7>)

2019

203. **Rohweder, B., Lehmann, G., Eichner, N., Polen, T., Rajendran, C., Linde, M., Treiber, T., Jung, O., Dettmer, K., Meister, G., Bott, M., Gronwald, W., and Sterner, R. (2019)**
Library selection with a randomized repertoire of $(\beta\alpha)_8$ -barrel enzymes results in unexpected induction of gene expression
Biochemistry 58: 4207-4217 (<http://dx.doi.org/10.1021/acs.biochem.9b00579>)
202. **Davoudi, C., Ramp, P., Baumgart, M., and Bott, M. (2019)**
Identification of Surf1 as an assembly factor of the cytochrome bc_1 -aa₃ supercomplex of *Actinobacteria*.
BBA Bioenergetics 1860: 148033 (<https://dx.doi.org/10.1016/j.bbabi.2019.06.005>)
201. **Kraxner, K.J., Polen, T., Baumgart, M., and Bott, M. (2019)**
The conserved actinobacterial transcriptional regulator FtsR controls expression of *ftsZ* and further target genes and influences growth and cell division in *Corynebacterium glutamicum*
BMC Microbiology 19: 179 (<http://dx.doi.org/10.1186/s12866-019-1553-0>)
200. **Kortmann, M., Baumgart, M. and Bott, M. (2019)**
Pyruvate carboxylase from *Corynebacterium glutamicum*: purification and characterization
Appl. Microbiol. Biotechnol. 103: 6571-6580 (<http://dx.doi.org/10.1007/s00253-019-09982-x>)
199. **Milke, L., Ferreira, P., Kallscheuer, N., Braga, A., Vogt, M., Kappelmann, J., Oliveira, J., Silva, R., Rocha, I., Bott, M., Noack, S., Faria, N., and Marienhagen, J. (2019)**

- Modulation of the central carbon metabolism of *Corynebacterium glutamicum* improves malonyl-CoA availability and increases plant polyphenol synthesis
Biotechnol. Bioeng. 116:1380–1391 (<http://dx.doi.org/10.1002/bit.26939>)
198. **Kallscheuer, N., Menezes, R., Fito, A., da Silva, M.H., Braga, A., Decker, W., Sevillano, D.M., Rosado-Ramos, R., Jardim, C., Oliveira, J., Ferreira, P., Rocha, I., Silva, A.R., Sousa, M., Allwood, J., Bott, M., Faria, N., Stewart, D., Ottens, M., Naesby, M., Santos, C. and Marienhagen, J. (2019)**
Identification of a phenol against Huntington's disease in raspberries and its microbial production
Plant Physiol. 179: 969–985 (<http://dx.doi.org/10.1104/pp.18.01074>)
202. **Davoudi, C., Ramp, P., Baumgart, M., and Bott, M. (2019)** Identification of Surf1 as an assembly factor of the cytochrome bc_1 -aa₃ supercomplex of *Actinobacteria*. BBA Bioenergetics 1860: 148033 (<http://dx.doi.org/10.1016/j.bbabi.2019.06.005>)
201. **Kraxner, K.J., Polen, T., Baumgart, M., and Bott, M. (2019)** The conserved actinobacterial transcriptional regulator FtsR controls expression of *ftsZ* and further target genes and influences growth and cell division in *Corynebacterium glutamicum*. BMC Microbiology 19: 179 (<http://dx.doi.org/10.1186/s12866-019-1553-0>)
200. **Kortmann, M., Baumgart, M. and Bott, M. (2019)** Pyruvate carboxylase from *Corynebacterium glutamicum*: purification and characterization. Appl. Microbiol. Biotechnol. 108: 6571–6580 (<http://dx.doi.org/10.1007/s00253-019-09982-x>)
199. **Milke, L., Ferreira, P., Kallscheuer, N., Braga, A., Vogt, M., Kappelmann, J., Oliveira, J., Silva, R., Rocha, I., Bott, M., Noack, S., Faria, N., and Marienhagen, J. (2019)** Modulation of the central carbon metabolism of *Corynebacterium glutamicum* improves malonyl-CoA availability and increases plant polyphenol synthesis. Biotechnol. Bioeng. 116:1380–1391 (<http://dx.doi.org/10.1002/bit.26939>)
198. **Kallscheuer, N., Menezes, R., Fito, A., da Silva, M.H., Braga, A., Decker, W., Sevillano, D.M., Rosado-Ramos, R., Jardim, C., Oliveira, J., Ferreira, P., Rocha, I., Silva, A.R., Sousa, M., Allwood, J., Bott, M., Faria, N., Stewart, D., Ottens, M., Naesby, M., Santos, C. and Marienhagen, J. (2019)** Identification of a phenol against Huntington's disease in raspberries and its microbial production. Plant Physiol. 179: 969–985 (<http://dx.doi.org/10.1104/pp.18.01074>)
197. **Kortmann, M., Baumgart, M., and Bott, M. (2019)** Pyruvate carboxylase variants enabling improved lysine production identified by biosensor-based high-throughput FACS screening
ACS Synth. Biol. 2019, 8, 274–281 (<http://dx.doi.org/10.1021/acssynbio.8b00510>)

2018

196. **Spielmann, A., Baumgart, M. and Bott, M. (2018)** NADPH-related processes studied with a SoxR-based biosensor in *Escherichia coli*. MicrobiologyOpen e785(<http://dx.doi.org/10.1002/mbo3.785>)
195. **Cormann, K.U., Baumgart, M. and Bott, M. (2018)** Structure-based design of versatile biosensors for small molecules based on the PAS domain of a thermophilic histidine kinase
ACS Synth. Biol. 7: 2888–2897 (<http://dx.doi.org/10.1021/acssynbio.8b00348>)
194. **Kranz, A., Steinmann, A., Degner, U., Mengus-Kaya, A., Matamouros, S., Bott, M. and Polen, T. (2018)** Global mRNA decay and 23S rRNA fragmentation in *Gluconobacter oxydans* 621H. BMC Genomics 19:753 (<http://dx.doi.org/10.1186/s12864-018-5111-1>)
193. **Morosov, X., Davoudi, C., Baumgart, M., Brocker, M., and Bott, M. (2018)** The copper-deprivation stimulon of *Corynebacterium glutamicum* comprises proteins for biogenesis of the actinobacterial cytochrome bc_1 -aa₃ supercomplex. J. Biol. Chem. 293: 15628–15640 (<http://dx.doi.org/10.1074/jbc.RA118.004117>)

192. **Alderwick, L., Birch, H.L., Krumbach, K., Bott, M., Eggeling, L. and Besra, G.S. (2018)**
AftD functions as an a1? 5 arabinofuranosyltransferase involved in the biosynthesis of the mycobacterial cell wall core. *Cell Surface* 1: 2-14 (<http://dx.doi.org/10.1016/j.tcs.2017.10.001>)
191. **Kranz, A., Busche, T., Vogel, A., Usadel, B., Kalinowski, J., Bott, M. and Polen, T. (2018)**
RNAseq analysis of α -proteobacterium *Gluconobacter oxydans* 621H. *BMC Genomics* 19:24 (<http://dx.doi.org/10.1186/s12864-017-4415-x>)
190. **Dudnik, A., Almeida, A.F., Andrade, R., Avila, B., Bañados, P., Barbay, D., Bassard, J.-E., Benkoulouche, M., Bott, M. et al. (2018).** BachBerry: BACterial Hosts for production of Bioactive phenolics from bERRY fruits. *Phytochem. Rev.* 17: 291–326 (<http://dx.doi.org/10.1007/s11101-017-9532-2>)

2017

189. **Kleine, B., Chattopadhyay, A., Polen, T., Pinto, D., Mascher, T., Bott, M., Brocker, M., and Freudl, R. (2017)** The three-component system EsrISR regulates a cell envelope stress response in *Corynebacterium glutamicum*. *Mol. Microbiol.* 106: 719-741 (<http://dx.doi.org/10.1111/mmi.13839>)
188. **Bott, M., Jäger, K.-E., Pietruszka, J., and Wiechert, W. (2017)** 40 Years of Biotechnology Research at Forschungszentrum Jülich. *J. Biotechnol.* 258: 1 (<http://dx.doi.org/10.1016/j.biote.2017.08.014>)
187. **Baumgart, M., Unthan, S., Kloß, R., Radek, A., Polen, T., Tenhaef, N., Kübel, A., Siebert, D., Brühl, N., Marin, K., Hans, S., Krämer, R., Bott, M., Kalinowski, J., Wiechert, W., Seibold, G., Frunzke, J., Rückert, C., Wendisch, V.F. and Noack, S. (2017)** *Corynebacterium glutamicum* chassis C1*: Building and testing a novel platform host for synthetic biology and industrial biotechnology. *ACS Synth. Biol.* 7:132-144 (<http://dx.doi.org/10.1021/acssynbio.7b00261>)
186. **Schulte, J., Baumgart, M., and Bott, M. (2017)** Development of a single-cell GlxR-based cAMP biosensor for *Corynebacterium glutamicum*. *J. Biotechnol.* 258: 33-40 (<http://dx.doi.org/10.1016/j.biote.2017.07.004>)
185. **Kallscheuer, N., Polen, T., Bott, M. and Marienhagen, J. (2017)** Reversal of β -oxidative pathways for the microbial production of chemicals and polymer building blocks. *Metab. Eng.* 42:33-42 (<http://dx.doi.org/10.1016/j.ymben.2017.05.004>)
184. **Kircher, M., Bott, M., and Marienhagen, J. (2017)** Die Bedeutung der Biotechnologie für die Bioökonomie. In: Bioökonomie für Einsteiger. J. Pietzsch (ed.), Springer Spektrum, Heidelberg, pp. 105-128 (http://dx.doi.org/10.1007/978-3-662-53763-3_5)
183. **Kiefler, I., Bringer, S. and Bott, M. (2017)** Metabolic engineering of *Gluconobacter oxydans* 621H for improved biomass yield. *Appl. Microbiol. Biotechnol.* 101: 5453-5467 (<http://dx.doi.org/10.1007/s00253-017-8308-3>)
182. **Kranz, A., Vogel, A., Degner, U., Kiefler, I., Bott, M., Usadel, B. and Polen, T. (2017)** High precision genome sequencing of engineered *Gluconobacter oxydans* 621H by combining long nanopore and short accurate Illumina reads. *J. Biotechnol.* 258:197-205 (<http://dx.doi.org/10.1016/j.biote.2017.04.016>)
181. **Kallscheuer, N., Vogt, M., Bott, M., and Marienhagen, J. (2017)** Functional expression of plant-derived O-methyltransferase, flavanone 3-hydroxylase, and flavonol synthase in *Corynebacterium glutamicum* for production of pterostilbene, kaempferol, and quercetin. *J. Biotechnol.* 258:190-196 (<http://dx.doi.org/10.1016/j.biote.2017.01.006>)
180. **Kallscheuer, N., Gätgens, J., Lübecke, M., Pietruszka, J., Bott, M., and Polen, T. (2017)** Improved production of adipic acid with *Escherichia coli* by reversal of β -oxidation. *Appl. Microbiol. Biotechnol.* 101: 2371-2382. (<http://dx.doi.org/10.1007/s00253-016-8033-3>)

179. **Bott, M. and Eggeling, L. (2017)** Chapter 13: Novel technologies for optimal strain breeding. In Amino Acid Fermentation. A. Yokota and M. Ikeda (eds.), Advances in Biochemical Engineering/Biotechnology 159: 227-254 (http://dx.doi.org/10.1007/10_2016_33)
178. **Schulte, J., Baumgart, M. and Bott, M. (2017)** Identification of the cAMP phosphodiesterase CpdA as novel key player in cAMP-dependent regulation in *Corynebacterium glutamicum*. Mol. Microbiol. 103: 534-552 (<http://dx.doi.org/10.1111/mmi.13574>)
177. **Hochheim, J., Kranz, A., Krumbach, K., Sokolowsky, S., Eggeling, L., Noack, S., Bocola, M., Bott, M. and Marienhagen, J. (2017)** Mutations in MurE, the essential UDP-N-acetylmuramoylalanyl-D-glutamate 2,6-diaminopimelate ligase of *Corynebacterium glutamicum*: Effect on L-lysine formation and analysis of systemic consequences. Biotechnol. Lett. 39: 283-288 (<http://dx.doi.org/10.1007/s10529-016-2243-8>)
176. **Limberg, M.H., Aryani, T., Schulte, J., Mahr, R., Baumgart, M., Bott, M., Wiechert, W. and Oldiges, M. (2017)** Metabolic profile of 1,5-diaminopentane producing *Corynebacterium glutamicum* under scale-down conditions: Blueprint for robustness to bioreactor inhomogeneities. Biotechnol. Bioeng. 114: 560-575 (<http://dx.doi.org/10.1002/bit.26184>)

----- 2016 -----

175. **Vogt, M., Brüsseler, C., van Ooyen, J., Bott, M., and Marienhagen, J. (2016)** Production of 2-methyl-1-butanol and 3-methyl-1-butanol in engineered *Corynebacterium glutamicum*. Metab. Eng. 38:436-445 (<http://dx.doi.org/10.1016/j.ymben.2016.10.007>)
174. **Graf, S., Fedotovskaya, O., Kao, W., Hunte, C., Ädelroth, P., Bott, M., von Ballmoos, C. and Brzezinski, P. (2016)** Rapid electron transfer within the III-IV supercomplex in *Corynebacterium glutamicum*. Sci. Rep. 6:34098 (<http://dx.doi.org/10.1038/srep34098>)
173. **Kao, W.-C., Kleinschroth, T., Nitschke, W., Baymann, F., Neehaul, Y., Hellwig, P., Richers, S., Vonck, J., Bott, M. and Hunte, C. (2015)** The obligate respiratory supercomplex from Actinobacteria. BBA Bioenergetics 1857: 1705-1714 (<http://dx.doi.org/10.1016/j.bbabi.2016.07.009>)
172. **Pahlke, J., Dostálová, H., Holátko, J., Degner, U., Bott, M., Pátek, M., and Polen, T. (2016)** The small 6C RNA of *Corynebacterium glutamicum* is involved in the SOS response. RNA Biology 13: 848-860 (<http://dx.doi.org/10.1080/15476286.2016.1205776>)
171. **Bringer, S. and Bott, M. (2016)** Central carbon metabolism and respiration in *Gluconobacter oxydans*. In: Acetic acid bacteria: ecology and physiology. K. Matsushita, H. Toyama, N. Tonouchi, and A. Kainuma (eds.). Chapter 11, pp. 235-253. Springer-Verlag, Berlin, Heidelberg, New York (http://dx.doi.org/10.1007/978-4-431-55933-7_11)
170. **Kallscheuer, N., Vogt, M., Stenzel, A., Gätgens, J., Bott, M. and Marienhagen, J. (2016)** Construction of a *Corynebacterium glutamicum* platform strain for the production of stilbenes and (2S)-flavanones. Metab. Eng. 38: 47-55 (<http://dx.doi.org/10.1016/j.ymben.2016.06.003>)
169. **Nguyen, G.T.T., Erlenkamp, G., Jäck, O., Küberl, A., Bott, M., Fiorani, F., Gohlke, H., and Groth, G. (2016)** Chalcone-based selective inhibitors of a C₄ plant key enzyme as novel potential herbicides. Sci. Rep. 6, 27333 (<http://dx.doi.org/10.1038/srep27333>)
168. **Küberl, A., Polen, T. and Bott, M. (2016)** The pupylation machinery is involved in iron homeostasis by targeting the iron storage protein ferritin. Proc. Natl. Acad. Sci. USA 113: 4806-4811 (<http://dx.doi.org/10.1073/pnas.1514529113>)
167. **Kallscheuer, N., Vogt, M., Kappelmann, J., Noack, S., Bott, M. and Marienhagen, J. (2016)** Identification of the *phd* gene cluster involved in phenylpropanoid utilization in *Corynebacterium*

glutamicum. Appl. Microbiol. Biotechnol. 100: 1871-1881 (<http://dx.doi.org/10.1007/s00253-015-7165-1>)

----- 2015 -----

166. **Kim, E., Um, Y., Bott, M. and Woo, H. M. (2015)** Engineering of *Corynebacterium glutamicum* for growth and succinate production from levoglucosan as a pyrolytic sugar substrate. FEMS Microbiol. Lett. 362: fnv161
165. **Kiefler, I., Bringer, S. and Bott, M. (2015)** SdhE-dependent formation of a functional *Acetobacter pasteurianus* succinate dehydrogenase in *Gluconobacter oxydans* – a first step toward a complete tricarboxylic acid cycle. Appl. Microbiol. Biotechnol. 99: 9147-9160
164. **Kallscheuer, N., Bott, M., van Ooyen, J. and Polen, T. (2015)** Single-domain FkpA from *Corynebacterium glutamicum* exhibits PPIase as well as chaperone activity and improves growth at increased temperature. Appl. Environ. Microbiol. 81: 7839-7850
163. **Michel, A., Koch-Koerfges, A., Krumbach, K., Brocker, M. and Bott, M. (2015)** Anaerobic growth of *Corynebacterium glutamicum* via mixed-acid fermentation. Appl. Environ. Microbiol. 81:7496 –7508
162. **Ostermann, S., Richhardt, J., Bringer, S., Bott, M., Wiechert, W., and Oldiges, M. (2015)** ¹³C tracers for glucose degrading pathway discrimination in *Gluconobacter oxydans* 621H. Metabolites 5, 455-474
161. **Eggeling, L. and Bott, M. (2015)** The Genus *Corynebacterium*. In: "Practical Handbook of Microbiology, Second Edition", (Goldman, E. and Green, L.H., eds.), CRC Press, Taylor & Francis Group, Boca Raton, Chapter 29, pp. 487-503
160. **Otten, A., Brocker, M. and Bott, M. (2015)** Metabolic engineering of *Corynebacterium glutamicum* for the production of itaconate. Metab. Eng. 30: 156-165
159. **Eggeling, L. and Bott, M. (2015)** A giant market and a powerful metabolism: L-lysine provided by *Corynebacterium glutamicum*. Appl. Microbiol. Biotechnol. 99: 3387-3394
158. **Küberl, A., Polen, T., and Bott, M. (2015)** Pupylierung – ein bakterielles Pendant zur Ubiquitinylierung. BIOSpektrum 02-2015,158-160
157. **Mustafi, N., Bott, M., and Frunzke, J. (2015)** Development and application of genetically-encoded biosensors for strain development and single cell analysis of *Corynebacterium glutamicum*. In: *Corynebacterium glutamicum*: systems biology, biotechnological applications and control. A. Burkovski (ed.), Caister Academic Press, Norfolk, UK, chapter 12, pp. 179-196
156. **Eikmanns, B.J. and Bott, M. (2015)** Engineering *Corynebacterium glutamicum* for production of organic acids and alcohols. In: *Corynebacterium glutamicum*: systems biology, biotechnological applications and control. A. Burkovski (ed.), Caister Academic Press, Norfolk, UK, chapter 8, pp. 111-137
155. **Witthoff, S., Schmitz, K., Niedenführ, S., Nöh, K., Noack, S., Bott, M. and Marienhagen, J. (2015)** Metabolic engineering of *Corynebacterium glutamicum* for the metabolization of methanol. Appl. Environm. Microbiol. 81: 2215-2225
154. **Vogt, M., Haas, S., Polen, T., van Ooyen, J. and Bott, M. (2015)** Production of 2-ketoisocaproate with *Corynebacterium glutamicum* strains devoid of plasmids and heterologous genes. Microb. Biotechnol. 8: 351-360
153. **Kortmann, M., Kuhl, V., Klaffl, S. and Bott, M. (2015)** A chromosomally encoded T7 RNA polymerase-dependent gene expression system for *Corynebacterium glutamicum*: construction and comparative evaluation at the single-cell level. Microb. Biotechnol. 8: 253-265

152. **Unthan, S., Baumgart, M., Radek, A., Herbst, M., Siebert, D., Brühl, N., Bott, M., Wiechert, W., Marin, K., Hans, S., Krämer, R., Seibold, G., Frunzke, J., Kalinowski, J., Rückert, C., Wendisch, V.F. and Noack, S. (2015)** Top-down towards a chassis organism – Identification and deletion of irrelevant gene clusters from *Corynebacterium glutamicum*. Biotechnol. J. 10: 290-301
151. **Vogt, M., Krumbach, K., Bang, W.-G., van Ooyen, J., Noack, S., Klein, B., Bott, M. and Eggeling, L. (2015)** The contest for precursors: Channeling L-isoleucine synthesis in *Corynebacterium glutamicum* without byproduct formation. Appl. Microbiol. Biotechnol. 99:791-800
150. **Bott, M. (2015)** Need for speed – finding productive mutations using transcription factor-based biosensors, FACS, and recombineering. Microb. Biotechnol. 8: 8-10
149. **Eggeling, L., Bott, M., and Marienhagen, J. (2015)** Novel screening methods – Biosensors. Curr. Opin. Biotechnol. 35: 30-36

2014

148. **Townsend, P.D., Jungwirth, B., Pojer, F., Bußmann, M., Money, V.A., Cole, S.T., Pühler, A., Tauch, A., Bott, M., Cann, M.J. and Pohl, E. (2014)** The crystal structures of apo and cAMP-bound GlxR from *Corynebacterium glutamicum* reveal structural and dynamic changes upon cAMP-binding in CRP/FNR family transcription factors. PLoS ONE 9(12): e113265
147. **Radek, A., Krumbach, K., Gärtgens, J., Wendisch, V.F., Wiechert, W., Bott, M., Noack, S., and Marienhagen, M. (2014)** Engineering of *Corynebacterium glutamicum* for minimized carbon loss during utilization of D-xylose containing substrates. J. Biotechnol. 192: 156-160
146. **Irzik, K., van Ooyen, J., Gärtgens, J., Krumbach, K., Bott, M., and Eggeling, L. (2014)** Acyl-CoA sensing by FasR to adjust fatty acid synthesis in *Corynebacterium glutamicum*. J. Biotechnol. 192: 96-101
145. **Lee, J., Sim, S. J., Bott, M., Um, Y., Oh, M.-K., and Woo, H. M. (2014)** Succinate production from CO₂-grown microalgal biomass as carbon source using engineered *Corynebacterium glutamicum* through consolidated bioprocessing. Scientific Rep. 4: 5819
144. **Raasch, K., Bocola, M., Labahn, J., Leitner,A., Eggeling, L and Bott, M. (2014)** Interaction of the 2-oxoglutarate dehydrogenase OdhA with Odhl of *Corynebacterium glutamicum*: Mutants and a model. J. Biotechnol. 191: 99-105
143. **Hentschel, E., Mack, C., Gärtgens, C., Bott, M., Brocker, M. and Frunzke, J. (2014)** Phosphatase activity of the histidine kinases ensures pathway specificity of the ChrSA and HrrSA two-component systems in *Corynebacterium glutamicum*. Mol. Microbiol. 92: 1326–1342
142. **Siedler, S., Bringer, S., Polen, T. and Bott, M. (2014)** NADPH-dependent reductive biotransformation with *Escherichia coli* and its *pfkA* deletion mutant: influence on global gene expression and role of oxygen supply. Biotechnol. Bioeng. 111: 2067-2075.
141. **Küberl, A., Fränzel, B., Eggeling, L., Polen, T., Wolters, D. A., and Bott, M. (2014)** Pupylated proteins in *Corynebacterium glutamicum* revealed by MudPIT analysis. Proteomics 14: 1531-1542
140. **Kang, M.-K., Lee, J., Um, Y., Lee T. S., Bott, M., Park, S. J. and Woo H. M. (2014)** Synthetic biology platform of CoryneBrick vectors for gene expression in *C. glutamicum* and its application to xylose utilization. Appl. Microbiol. Biotechnol. 98: 5991-6002
139. **Litsanov, B., Brocker, M., Oldiges, M. and Bott, M. (2014)** Chapter 16: Succinic acid. In ‘Bioprocessing of Renewable Resources to Commodity Bioproducts’, eds. Virendra S. Bisaria and Akihiko Kondo, Wiley & Sons, pp. 437-474

138. **Pelzer, A., Polen, T., Funken, H., Rosenau, F., Wilhelm, S., Bott, M., and Jaeger, K.-E. (2014)** Subtilase SprP exerts pleiotropic effects in *Pseudomonas aeruginosa*. *MicrobiologyOpen* 3: 89–103
137. **Vogt, M., Haas, S., Klaffl, S., Polen, T., Eggeling, L., van Ooyen, J., and Bott, M. (2014)** Pushing product formation to its limit: Metabolic engineering of *Corynebacterium glutamicum* for L-leucine overproduction. *Metab. Eng.* 22: 40-52
136. **Siedler, S., Schendzielorz, G., Binder, S., Eggeling, L., Bringer, S. and Bott, M. (2014)** SoxR as single-cell biosensor for NADPH-consuming enzymes in *Escherichia coli*. *ACS Synth. Biol.* 3: 41-47
135. **Schendzielorz, S., Dippong, M., Grünberger, A., Kohlheyer, D., Yoshida, A., Binder, S., Nishiyama, M., Nishiyama, C., Bott, M. and Eggeling, L. (2014)** Taking control over control: Use of product sensing in single cells to remove flux control at key enzymes in biosynthesis pathways. *ACS Synth. Biol.* 3: 21-29
134. **Platzen, L., Koch-Koerfges, A., Weil, B., Brocker, M. and Bott, M. (2014)** Role of flavohaemoprotein Hmp and nitrate reductase NarGHJI of *Corynebacterium glutamicum* for coping with nitrite and nitrosative stress. *FEMS Microbiol. Lett.* 350: 239-248

2013

133. **Witthoff, S., Mühlroth, A., Marienhagen, J., and Bott, M. (2013)** C1 metabolism in *Corynebacterium glutamicum*: an endogenous pathway for oxidation of methanol to carbon dioxide. *Appl. Environm. Microbiol.* 79: 6974-6983
132. **Hoffmann, K., Grünberger, A., Lausberg, F., Bott, M., and Eggeling, L. (2013)** Imbalances in sulfur assimilation and synthesis of sulfur-containing amino acids: Visualization at the single-cell level. *Appl. Environm. Microbiol.* 79: 6730-6736
131. **Baumgart, M., Unthan, S., Rückert, C., Sivalingam, J., Grünberger, A., Kalinowski, J., Bott, M., Noack, S. and Frunzke, J. (2013)** Construction of a prophage-free variant of *Corynebacterium glutamicum* ATCC 13032 - a novel platform strain for basic research and industrial biotechnology. *Appl. Environm. Microbiol.* 79: 6006–6015
130. **Klaffl, S., Brocker, M., Kalinowski, J., Eikmanns, B.J. and Bott, M. (2013)** Complex regulation of the PEP carboxykinase gene *pck* and characterization of its GntR-type regulator *lolR* as a repressor of *myo*-inositol utilization genes in *Corynebacterium glutamicum*. *J. Bacteriol.* 195: 4283-4296
129. **Richhardt, J., Luchterhand, B., Bringer, S., Büchs, J. and Bott, M. (2013)** Evidence for a key role of cytochrome *b*_o₃ oxidase in respiratory energy metabolism of *Gluconobacter oxydans*. *J. Bacteriol.* 195: 4210-4220
128. **van Ooyen, J., Noack, S., Bott, M. and Eggeling, L. (2013)** Proline addition increases the efficiency of L-lysine production by *Corynebacterium glutamicum*. *Eng. Life Sci.* 13: 393-398
127. **Polen, T., Spelberg, M. and Bott, M. (2013)** Toward biotechnological production of adipic acid and precursors from biorenewables. *J. Biotechnol.* 167: 75-84
126. **Binder, S., Siedler, S., Marienhagen, J., Bott, M. and Eggeling, L. (2013)** Recombineering in *Corynebacterium glutamicum* combined with optical nanosensors: A general strategy for fast producer strain generation. *Nucl. Acids Res.* 41: 6360-6369
125. **García-Nafría, J., Baumgart, M., Turkenburg, J.P., Wilkinson, A.J., Bott, M. and Wilson, K.S. (2013)** Crystal and solution studies reveal that the transcriptional regulator AcnR of *Corynebacterium glutamicum* is regulated by citrate:Mg²⁺ binding to a non-canonical pocket. *J. Biol. Chem.* 288: 15800-15812

124. **Richardt, J., Bringer, S. and Bott, M. (2013)** Role of the pentose phosphate pathway and the Entner-Doudoroff pathway in glucose metabolism of *Gluconobacter oxydans* 621H. *Appl. Microbiol. Biotechnol.* 97:4315–4323
123. **Koch-Koerfges, A., Pfelzer, N., Platzen, L., Oldiges, M. and Bott, M. (2013)** Conversion of *Corynebacterium glutamicum* from an aerobic respiration to an aerobic fermenting bacterium by inactivation of the respiratory chain. *BBA Bioenergetics* 1827: 699-708
122. **Hanke, T., Nöh, K., Noack, S., Polen, T., Bringer, S., Sahm, H., Wiechert, W. and Bott, M. (2013)** Combined fluxomics and transcriptomics analysis of glucose catabolism via a partially cyclic pentose phosphate pathway in *Gluconobacter oxydans* 621H. *Appl. Environm. Microbiol.* 79: 2336-2348
121. **Wieschalka, S., Blombach, B., Bott, M. and Eikmanns, B.J. (2013)** Bio-based production of organic acids with *Corynebacterium glutamicum*. *Microb. Biotechnol.* 6: 87-102
120. **Scheele, S., Oertel, D., Bongaerts, J., Evers, S., Hellmuth, H., Maurer, K.-H., Bott, M. and Freudl, R. (2013)** Secretory production of an FAD-cofactor containing cytosolic enzyme (sorbitol-xylitol oxidase from *Streptomyces coelicolor*) using the twinarginine translocation (Tat) pathway of *Corynebacterium glutamicum*. *Microb. Biotechnol.* 6: 202-206
119. **Litsanov, B., Brocker, M. and Bott, M. (2013)** Glycerol as a substrate for aerobic succinate production in minimal medium with *Corynebacterium glutamicum*. *Microb. Biotechnol.* 6: 189-195
118. **Marienhagen, J. and Bott, M. (2013)** Metabolic engineering of microorganisms for the synthesis of plant natural products. *J. Biotechnol.* 163: 166-178
117. **Siedler, S., Lindner, S. N., Bringer, S., Wendisch, V. F. and Bott, M. (2013)** Reductive whole-cell biotransformation with *Corynebacterium glutamicum*: improvement of NADPH generation from glucose by a cyclized pentose phosphate pathway using *pfkA* and *gapA* deletion mutants. *Appl. Microbiol. Biotechnol.* 97:143–152
116. **Bott, M. and Eikmanns, B. J. (2013)** TCA cycle and glyoxylate shunt of *Corynebacterium glutamicum*. In: "Biology and Biotechnology of *Corynebacterium glutamicum*" (Yukawa, H. and Inui, M., eds.), Microbiology Monographs 23 (A. Steinbüchel, ed.), Springer Verlag, Berlin, Heidelberg, pp. 281-313

2012

115. **Heyer, A., Gätgens, C., Kalinowski, J., Bott, M. and Frunzke, J. (2012)** The two-component system ChrSA is crucial for heme tolerance and interferes with HrrSA in heme-dependent gene regulation in *Corynebacterium glutamicum*. *Microbiology* 158: 3020-3031
114. **Funken, H., Bartels, K.-M., Wilhelm, S., Brocker, M., Bott, M., Bains, M., Hancock, R.E.W., Rosenau, F. and Jaeger, K.-E. (2012)** Specific association of lectin LecB with the surface of *Pseudomonas aeruginosa*: Role of outer membrane protein OprF. *PLoS ONE* 7(10):e46857
113. **Richardt, J., Bringer, S. and Bott, M. (2012)** Mutational analysis of the pentose phosphate pathway and the Entner-Doudoroff pathway in *Gluconobacter oxydans* reveals improved growth of an *edd-eda* deletion mutant on mannitol. *Appl. Environm. Microbiol.* 78: 6975-6986
112. **Witthoff, S., Eggeling, L., Bott, M. and Polen, T. (2012)** *Corynebacterium glutamicum* harbours a molybdenum cofactor-dependent formate dehydrogenase which alleviates growth inhibition in the presence of formate. *Microbiology* 158:2428-2439
111. **Heck, A., Tielker, D., Ernst, J. F., Freudl, R., Bott, M., Oldiges, M., Wiechert, W., Pietruszka, J., Wilhelm, S., Rosenau, F., Drepper, T., and Jaeger, K.-E. (2012)** Forschungsplattform ExpressO: Expressionsoptimierung in Mikroorganismen. *Biospektrum* 04.12, 449-451

110. **Koch-Koerfges, A., Kabus, A., Ochrombel, I., Marin, K. and Bott, M. (2012)** Investigation on ATP synthesis via oxidative phosphorylation in *C. glutamicum*: Measurement of oxygen consumption by an ATP synthase deletion mutant and the wild type strain with the SFR Shake Flask Reader. PreSens - Precision Sensing GmbH, Application note, pp.1-2
109. **Binder, S., Schendzielorz, G., Stäbler, N., Krumbach, K., Hoffmann, K., Bott, M., and Eggeling, L. (2012)** A high-throughput approach using metabolite sensors to identify single bacterial cells that produce high levels of small molecules. *Genome Biology* 13:R40
108. **Bott, M. and Brocker, M. (2012)** Two-component signal transduction in *Corynebacterium glutamicum* and other corynebacteria: on the way towards stimuli and targets. *Appl. Microbiol. Biotechnol.* 94: 1131–1150; Erratum *Appl. Microbiol. Biotechnol.* 96: 1105–1110
107. **van Ooyen, J., Noack, S., Bott, M., Reth, A., and Eggeling, L. (2012)** Improved L-lysine production with *Corynebacterium glutamicum* and systemic insight into citrate synthase flux and activity. *Biotechnol. Bioeng.* 109: 2070-2081
106. **Litsanov, B., Brocker, M. and Bott, M. (2012)** Towards homosuccinate fermentation: metabolic engineering of *Corynebacterium glutamicum* for anaerobic succinate production from glucose and formate. *Appl. Environm. Microbiol.* 78: 3325-3337
105. **Mustafi, N., Grünberger, A., Kohlheyer, D., Bott, M., and Frunzke, J. (2012)** The development and application of a single-cell biosensor for the detection of L-methionine and branched-chain amino acids. *Metab. Eng.* 14: 449-457
104. **Koch-Koerfges, A., Kabus, A., Ochrombel, I., Marin, K., and Bott, M. (2012)** Physiology and global gene expression of a *Corynebacterium glutamicum* ΔF_1F_0 -ATP synthase mutant devoid of oxidative phosphorylation. *BBA Bioenergetics* 1817: 370-380
103. **Lange, C. Mustafi, N., Frunzke, J., Kennerknecht, N., Wessel, M., Bott, M. and Wendisch, V. F. (2012)** Lrp of *Corynebacterium glutamicum* controls expression of the *bmFE* operon encoding the export system for L-methionine and branched-chain amino acids. *J. Biotechnol.* 158: 231-241
102. **Hanke, T., Richardt, J., Polen, T., Sahm, H., Bringer, S. and Bott, M. (2012)** Influence of oxygen limitation, absence of the cytochrome *bc*₁ complex and low pH on global gene expression in *Gluconobacter oxydans* 621H using DNA microarray technology. *J. Biotechnol.* 157: 359-372
101. **Siedler, S., Bringer, S., Blank, L.M., and Bott, M. (2012)** Engineering yield and rate of reductive biotransformation in *Escherichia coli* by partial cyclization of the pentose phosphate pathway and PTS-independent glucose transport. *Appl. Microbiol. Biotechnol.* 93:1459–1467
100. **Litsanov, B., Kabus, A., Brocker, M. and Bott, M. (2012)** Efficient aerobic succinate production from glucose in minimal medium with *Corynebacterium glutamicum*. *Microb. Biotechnol.* 5: 116-128

2011

99. **Baumgart, M., Mustafi, N., Krug, A. and Bott, M. (2011)** Deletion of the aconitase gene in *Corynebacterium glutamicum* causes a strong selection pressure for secondary mutations inactivating citrate synthase. *J. Bacteriol.* 193: 6864-6873
98. **Schelder, S., Zaade, D., Litsanov, B., Bott, M. and Brocker, M. (2011)** The two-component signal transduction system CopRS of *Corynebacterium glutamicum* is required for adaptation to copper-excess stress. *PLoS One* 6 (7): e22143.
97. **Siedler, S., Bringer-Meyer, S. and Bott, M. (2011)** Increased NADPH availability in *Escherichia coli*: Improvement of the product per glucose ratio in reductive whole-cell biotransformation. *Appl. Microbiol. Biotechnol.* 92: 929-937

96. **Baumgart, M. and Bott, M. (2011)** Biochemical characterisation of aconitase from *Corynebacterium glutamicum*. J. Biotechnol. 154: 163-170
95. **van Ooyen, J., Emer, D., Bussmann, M., Bott, M., Eikmanns, B.J. and Eggeling, L. (2011)** Citrate synthase in *Corynebacterium glutamicum* is encoded by two *gltA* transcripts which are controlled by RamA, RamB, and GlxR. J. Biotechnol. 154: 140-148
94. **Stäbler, N., Oikawa, T., Bott, M. and Eggeling, L. (2011)** *Corynebacterium glutamicum* as a host for the synthesis and export of D-amino acids. J. Bacteriol. 193: 1702-1709
93. **Broker, M., Mack, C. and Bott, M. (2011)** The response regulator MtrA of *Corynebacterium glutamicum*: target genes, consensus binding site and role of phosphorylation. J. Bacteriol. 193: 1237-1249
92. **Frunzke, J., Gätgens, C., Broker, M. and Bott, M. (2011)** Control of heme homeostasis in *Corynebacterium glutamicum* by the two-component system HrrSA. J. Bacteriol. 193: 1212-1221
91. **Bartels, K.-M., Funken, H., Knapp, A., Broker, M., Bott, M., Wilhelm, S., Jäger, K.-E., and Rosenau, F. (2011)** Glycosylation is required for outer membrane localization of the lectin LecB in *Pseudomonas aeruginosa*. J. Bacteriol. 193: 1107-1113

2010

90. **Woo, H. M., Noack, S., Seibold, G. M., Willbold, S., Eikmanns, B. J. and Bott, M. (2010)** A link between phosphate starvation and glycogen metabolism in *Corynebacterium glutamicum* revealed by metabolomics. Appl. Environm. Microbiol. 76: 6910-6919
89. **Garcia-Nafria, J., Baumgart, M., Bott, M., Wilkinson, A. J. and Wilson, K. S. (2010)** The *Corynebacterium glutamicum* aconitase repressor: Scratching around for crystals. Acta Crystallogr. F66, 1074-1077
88. **Bussmann, M., Baumgart, M. and Bott, M. (2010)** RosR (Cg1324), a hydrogen peroxide-sensitive MarR-type transcriptional regulator of *Corynebacterium glutamicum*. J. Biol. Chem. 285: 29305-29318
87. **Rehm, N., Georgi, T., Hiery, E., Degner, U., Schmiedl, Schmiedl, A., Burkovski, A., and Bott, M. (2010)** L-Glutamine as nitrogen source for *Corynebacterium glutamicum*: derepression of the AmtR regulon and implications for nitrogen sensing. Microbiology 156: 3180 - 3193.
86. **Boulahya, K., Guedon, E., Delaunay, S., Schultz, C., Boudrant, J., Bott, M. and Goergen, J.-L. (2010)** Odhl dephosphorylation kinetics during different glutamate production processes involving *Corynebacterium glutamicum*. Appl. Microbiol. Biotechnol. 87:1867-1874
85. **Zhang, Z., Buitenhuis, J., Cukkemane, A., Broker, M., Bott, M. and Dhont, J. (2010)** Charge reversal of rod-like colloidal fd virus through surface chemical modification. Langmuir 26:10593-10599.
84. **Krawczyk, S., Raasch, K., Schultz, C., Hoffelder, M., Eggeling, L. and Bott, M. (2010)** The FHA domain of Odhl interacts with the carboxyterminal 2-oxoglutarate dehydrogenase domain of OdhA in *Corynebacterium glutamicum*. FEBS Lett. 584: 1463–1468
83. **Bott, M., Bringer-Meyer, S., Broker, M., Eggeling, L., Freudl, R., Frunzke, J. and Polen, T. (2010)** Systemische Mikrobiologie – Etablierung bakterieller Produktionsplattformen für die Weiße Biotechnologie. Jahrbuch der Heinrich-Heine-Universität Düsseldorf 2008/2009, 227-242
82. **Bott, M. (2010)** Signal transduction by serine/threonine protein kinases in bacteria. In: "Bacterial signalling" (Krämer, R. and Jung, K., eds.), Wiley-VCH, Weinheim, Chapter 24, pp. 427-447

81. **Reher, M., Fuhrer, T., Bott, M. and Schönheit, P. (2010)** The non-phosphorylative Entner-Doudoroff pathway in the thermoacidophilic euryarchaeon *Picrophilus torridus* involves a novel 2-keto-3-deoxygluconate specific aldolase (KDGA). *J. Bacteriol.* 192: 964-974

----- 2009 -----

80. **Schultz, C., Niebisch, A., Schwaiger, A., Viets, U., Metzger, S., Bramkamp, M. and Bott, M. (2009)** Genetic and biochemical analysis of the serine/threonine protein kinases PknA, PknB, PknG and PknL of *Corynebacterium glutamicum*: evidence for non-essentiality and for phosphorylation of Odhl and FtsZ by multiple kinases. *Mol. Microbiol.* 74: 724–741
79. **Bussmann, M., Emer, D., Hasenbein, S., Degraf, S., Eikmanns, B.J. and Bott, M. (2009)** Transcriptional control of the succinate dehydrogenase operon *sdhCAB* of *Corynebacterium glutamicum* by the cAMP-dependent regulator GlxR and the LuxR-type regulator RamA. *J. Biotechnol.* 143: 173-182
78. **Brocke, M., Schaffer, S., Mack, C. and Bott, M. (2009)** Citrate utilization by *Corynebacterium glutamicum* is controlled by the CitAB two-component system through positive regulation of the citrate transport genes *citH* and *tctCBA*. *J. Bacteriol.* 191: 3869–3880
77. **Emer, D., Krug, A., Eikmanns, B. J. and Bott, M. (2009)** Complex expression control of the *Corynebacterium glutamicum* aconitase gene: identification of RamA as a third transcriptional regulator besides AcnR and RipA. *J. Biotechnol.* 140: 92-98
76. **Russo, S., Schweitzer, J.-E., Polen, T., Bott, M. and Pohl, E. (2009)** Crystal structure of the caseinolytic protease gene regulator, a transcriptional activator in actinomycetes. *J. Biol. Chem.* 284: 5208-5216
75. **Eggeling, L. and Bott, M. (2009)** The Genus *Corynebacterium*. In: "Practical Handbook of Microbiology, Second Edition", (Goldman, E. and Green, L.H., eds.), CRC Press, Taylor & Francis Group, Boca Raton, Chapter 26, pp. 355-373

----- 2008 -----

74. **Frunzke, J., Bramkamp, M., Schweitzer, J.-E. and Bott, M. (2008)** Population heterogeneity in *Corynebacterium glutamicum* ATCC 13032 caused by prophage CGP3. *J. Bacteriol.* 190: 5111-5119
73. **Frunzke, J. and Bott, M. (2008)** Regulation of iron homeostasis in *Corynebacterium glutamicum*. In "Corynebacteria: genomics and molecular biology" (Burkovski, A., ed.), pp. 241-266, Caister Academic Press, Norfolk, UK.
72. **Wendisch, V. F. and Bott, M. (2008)** Phosphorus metabolism and its regulation. In "Corynebacteria: genomics and molecular biology" (Burkovski, A., ed.), pp. 203-216, Caister Academic Press, Norfolk, UK.
71. **Sevvana, M., Vijayan, V, Zweckstetter, M., Reinelt, S., Madden, D.R., Herbst-Irmer, R., Sheldrick, G.M., Bott, M., Griesinger, C. and Becker, S. (2008)** A ligand-induced switch in the periplasmic domain of sensor histidine kinase CitA. *J. Mol. Biol.* 377: 512-523
70. **Frunzke, J., Engels, V., Hasenbein, S., Gätgens, C. and Bott, M. (2008)** Coordinated regulation of gluconate catabolism and glucose uptake in *Corynebacterium glutamicum* by two functionally equivalent transcriptional regulators, GntR1 and GntR2. *Mol. Microbiol.* 67: 305-322

----- 2007 -----

69. **Bott, M. (2007)** Offering surprises: TCA cycle regulation in *Corynebacterium*. *Trends Microbiol.* 15: 417-425

68. **Schaaf, S. and Bott, M. (2007)** Target genes and DNA-binding sites of the response regulator PhoR from *Corynebacterium glutamicum*. J. Bacteriol. 189: 5002-5011
67. **Polen, T., Schlüsener, D., Pötsch, A., Bott, M. and Wendisch, V.F. (2007)** Characterization of citrate utilization in *Corynebacterium glutamicum* by transcriptome and proteome analysis. FEMS Microbiology Lett. 273: 109-119
66. **Bott, M. (2007)** Corynebacteria: the good guys and the bad guys. Microbiology today, May 07, 74-77
65. **Schultz, C., Niebisch, A., Gebel, L. and Bott, M. (2007)** Glutamate production by *Corynebacterium glutamicum*: dependence on the oxoglutarate dehydrogenase inhibitor protein Odhl and protein kinase PknG. Appl. Microbiol. Biotechnol. 76: 691-700
64. **Kabus, A., Georgi, T., Wendisch, V.F. and Bott, M. (2007)** Expression of the *Escherichia coli* *pntAB* genes encoding a membrane-bound transhydrogenase in *Corynebacterium glutamicum* improves L-lysine formation. Appl. Microbiol. Biotechnol., 75: 47-53
63. **Kabus, A., Niebisch, A. and Bott, M. (2007)** Role of cytochrome *bd* oxidase from *Corynebacterium glutamicum* for growth and lysine production. Appl. Environm. Microbiol. 73: 861-868
62. **Cramer, A., Auchter, M., Frunzke, J., Bott, M. and Eikmanns, B. (2007)** RamB, the transcriptional regulator of acetate metabolism in *Corynebacterium glutamicum* is subject to regulation by RamA and RamB. J. Bacteriol. 189: 1145-1149

----- 2006 -----

61. **Broker, M. and Bott, M. (2006)** Evidence for activator and repressor functions of the response regulator MtrA from *Corynebacterium glutamicum*. FEMS Microbiol. Lett. 264: 205-212
60. **Förster-Fromme, K., Höschle, B., Mack, C., Bott, M., Armbruster, W. and Jendrossek, D. (2006)** Identification of genes and proteins involved in catabolism of acyclic terpenes and leucine/isovalerate in *Pseudomonas aeruginosa*. Appl. Environm. Microbiol. 72: 4819-4828
59. **Wendisch, V.F., Bott, M. and Eikmanns, B.J. (2006)** Metabolic engineering of *Escherichia coli* and *Corynebacterium glutamicum* for biotechnological production of organic acids and amino acids. Curr. Opin. Microbiol. 9: 268-274
58. **Wendisch, V.F., Bott, M., Kalinowski, J., Oldiges, M. and Wiechert, W. (2006)** Emerging *Corynebacterium glutamicum* systems biology. J. Biotechnol. 124: 74-92
57. **Reher, M., Bott, M. and Schönheit, P. (2006)** Characterization of glycerate kinase (2-phosphoglycerate forming), a key enzyme of the non-phosphorylative Entner-Doudoroff pathway, from the thermoacidophilic euryarchaeon *Picrophilus torridus*. FEMS Microbiol. Lett. 259: 113-119
56. **Niebisch, A., Kabus, A., Schultz, C., Weil, B. and Bott, M. (2006)** Corynebacterial protein kinase G controls 2-oxoglutarate dehydrogenase activity via the phosphorylation status of the Odhl protein. J. Biol. Chem. 281: 12300-12307
55. **Wennerhold, J. and Bott, M. (2006)** The DtxR regulon of *Corynebacterium glutamicum*. J. Bacteriol. 188: 2907-2918
54. **Cramer, A., Gerstmeir, R., Schaffer, S., Bott, M. and Eikmanns, B.J. (2006)** Identification of RamA, a novel LuxR-type transcriptional regulator of genes involved in acetate metabolism of *Corynebacterium glutamicum*. J. Bacteriol. 188: 2554-2567
53. **Kocan, M., Schaffer, S., Ishige, T., Sorger-Hermann, U., Wendisch, V.F. and Bott, M. (2006)**

Two-component systems of *Corynebacterium glutamicum*: deletion analysis and involvement of the PhoS-PhoR system in the phosphate starvation response. J. Bacteriol. 188: 724-732

----- 2005 -----

52. **Wennerhold, J., Krug, A. and Bott, M. (2005)** The AraC-type regulator RipA represses aconitase and other iron proteins from *Corynebacterium* under iron limitation and is itself repressed by DtxR. J. Biol. Chem. 280: 40500-40508
51. **Engels, S., Ludwig, C., Schweitzer, J.-E., Mack, C., Bott, M., and Schaffer, S. (2005)** The transcriptional activator ClgR controls transcription of genes involved in proteolysis and DNA repair in *Corynebacterium glutamicum*. Mol. Microbiol. 57: 576-591
50. **Wendisch, V.F. and Bott, M. (2005)** Phosphorus metabolism. In "Handbook of *Corynebacterium glutamicum*" (Eggeling, L. and Bott, M., eds), pp. 377- 396, CRC Press, Boca Raton, Florida.
49. **Bott, M. and Niebisch, A. (2005)** Respiratory energy metabolism. In "Handbook of *Corynebacterium glutamicum*" (Eggeling, L. and Bott, M., eds), pp. 305- 332, CRC Press, Boca Raton, Florida.
48. **Eggeling, L. and Bott, M. (2005)** Handbook of *Corynebacterium glutamicum*. CRC Press, Taylor & Francis Group, Boca Raton, Florida
47. **Krug, A., Wendisch, V.F. and Bott, M. (2005)** Identification of AcnR, a TetR-type repressor of the aconitase gene acn of *Corynebacterium glutamicum*. J. Biol. Chem. 280: 585-595

----- 2004 -----

46. **Möker, N., Brocker, M., Schaffer, S., Krämer, R., Morbach, S. and Bott, M. (2004)** Deletion of the genes encoding the MtrA-MtrB two-component system of *Corynebacterium glutamicum* has a strong influence on cell morphology, antibiotics susceptibility and expression of genes involved in osmoprotection. Mol. Microbiol. 54: 420-438
45. **Engels, S., Schweitzer, J., Ludwig, C., Bott, M. and Schaffer, S. (2004)** *c/pC* and *c/pP1P2* gene expression in *Corynebacterium glutamicum* is controlled by a regulatory network involving the transcriptional regulators ClgR and HspR as well as the ECF sigma factor σ^H . Mol. Microbiol. 52: 285-302
44. **Heinz, D., Bott, M., Kalinowski, J., Niefind, K., Takors, R. and Wendisch, V.F. (2004)** Die gläserne Zelle. In: Biotechnologie 2020 – von der gläsernen Zelle zum maßgeschneiderten Prozess, pp. 6-13, DECHEMA e.V.

----- 2003 -----

43. **Takors, R., Zelić, B., Gerharz, T. and Bott, M. (2003)** Pyruvat-Produktion aus Glucose mit rekombinanten *Escherichia coli*-Stämmen. In: Transkript, Sonderband Nachhaltige Biokatalyse, 96-99.
42. **Reineit, S., Hofmann, E., Gerharz, T., Bott, M. and Madden, D.R. (2003)** The structure of the periplasmic ligand-binding domain of the sensor kinase CitA reveals the first extracellular PAS domain. J. Biol. Chem. 40: 39189-39196
41. **Bott, M. and Niebisch, A. (2003)** The respiratory chain of *Corynebacterium glutamicum*. J. Biotechnol. 104: 129-153.
40. **Kalinowski, J., Bathe, B., Bischoff, N., Bott, M., Burkovski, A., Dusch, N., Eggeling, L., Eikmanns, B., Gaigalat, L., Goesmann, A., Hartmann, M., Huthmacher, K., Krämer , R., Linke, B., McHardy, A.C., Meyer, F., Möckel, M., Pfefferle, W., Pühler, A., Rey, D., Rückert, C., Sahm, H., Wendisch, V.F., Wiegräbe, I., Tauch, A. (2003)** The complete *Corynebacterium*

- glutamicum* ATCC 13032 genome sequence and its impact on the production of L-aspartate-derived amino acids and vitamins. J. Biotechnol. 104: 5-25.
39. **Bendt, A.K., Schaffer, S., Hermann, T., Bott, M., Farwick, M., and Burkovski, A. (2003)** Towards a phosphoproteome map of *Corynebacterium glutamicum*. Proteomics 3: 1637-1646.
 38. **Zelić, B., Gerharz, T., Bott, M., Vasić-Rački, D., Wandrey, C. and Takors, R. (2003)** Fed-batch process for pyruvate production by recombinant *Escherichia coli* YYC202 strain. Eng. Life Sci. 3: 299-305.
 37. **Ishige, T., Krause, M., Bott, M., Wendisch, V.F. and Sahm, H. (2003)** The phosphate starvation stimulon of *Corynebacterium glutamicum* as determined by DNA microarray analyses. J. Bacteriol. 185: 4519-4529.
 36. **Gerharz, T., Reinelt, S., Kaspar, S., Scapozza, L. and Bott, M. (2003)** Identification of basic amino acid residues important for citrate binding by the periplasmic receptor domain of the sensor kinase CitA. Biochemistry 42: 5917-5924.
 35. **Lange, C., Rittmann, D., Wendisch, V.F., Bott, M. and Sahm, H. (2003)** Global expression profiling and physiological characterization of *Corynebacterium glutamicum* grown in the presence of L-valine. Appl. Environm. Microbiol. 69: 2521-2532
 34. **Bott, M. (2003)** Muster des Lebens – die Analyse des Proteoms. In: Forschen in Jülich, Nr. 1/2003, pp. 12-13, Forschungszentrum Jülich GmbH
 33. **Niebisch, A. and Bott, M. (2003)** Purification of a cytochrome *bc₁-aa₃* supercomplex with quinol oxidase activity from *Corynebacterium glutamicum*. Identification of a fourth subunit of cytochrome aa₃ oxidase and mutational analysis of diheme cytochrome c₁. J. Biol. Chem. 278: 4339-4346.

2002

32. **Schneider, K., Kästner, C.N., Meyer, M., Wessel, M., Dimroth, P., and Bott, M. (2002)** Identification of a gene cluster in *Klebsiella pneumoniae* which includes *citX*, a gene required for the biosynthesis of the citrate lyase prosthetic group. J. Bacteriol. 184: 2439-2446.
31. **Kaspar, S. and Bott, M. (2002)** The sensor kinase CitA (DpiB) of *Escherichia coli* functions as a high affinity citrate receptor. Arch. Microbiol. 177: 313-321.

2001

30. **Schaffer, S., Weil, B., Nguyen, V.D., Dongmann, G., Günther, K., Nickolaus, M., Hermann, T., and Bott, M. (2001)** A high resolution reference map for cytoplasmic and membrane-associated proteins of *Corynebacterium glutamicum*. Electrophoresis 22: 4404-4422.
29. **Gerharz, T., Zelić, B., Takors, R., and Bott, M. (2001)** Produktion von Pyruvat aus Glucose mit *Escherichia coli*. In: *Biokatalyse*. Heiden, S. and Erb, R. (eds). Spektrum Akademischer Verlag (Heidelberg), pp. 29-33.
28. **Meyer, M., Dimroth, P. and Bott, M. (2001)** Catabolite repression of the citrate fermentation genes in *Klebsiella pneumoniae*: Evidence for the involvement of the cAMP receptor protein CRP. J. Bacteriol. 183: 5248-5256.
27. **Hermann, T., Pfefferle, W., Baumann, C., Busker, E., Schaffer, S., Bott, M., Sahm, H., Dusch, N., Kalinowski, J., Pühler, A., Bendt, A. K., Krämer, R., and Burkovski, A. (2001)** Proteome analysis of *Corynebacterium glutamicum*. Electrophoresis 22: 1712-1723.
26. **Niebisch, A. and Bott, M. (2001)** Molecular analysis of the cytochrome *bc₁-aa₃* branch of the *Corynebacterium glutamicum* respiratory chain containing an unusual di-heme cytochrome c₁. Arch. Microbiol. 175: 282-294

----- 1996 - 2000 -----

25. Schneider, K., Dimroth, P. and Bott, M. (2000) Identification of triphosphoribosyl-dephospho-CoA as precursor of the citrate lyase prosthetic group. FEBS Letters **483**: 165-168.
24. Schneider, K., Dimroth, P. and Bott, M. (2000) Biosynthesis of the prosthetic group of citrate lyase. Biochemistry **39**: 9438-9450
23. Kaspar, S., Perozzo, R., Reinelt, S., Meyer, M., Pfister, K., Scapozza, L., and Bott, M. (1999) The periplasmic domain of the histidine autokinase CitA functions as a highly specific citrate receptor. Mol. Microbiol. **33**: 858-872
22. Steuber, J., Krebs, W., Bott, M. and Dimroth, P. (1999) A membrane-bound NAD(P)⁺-reducing hydrogenase provides reduced pyridine nucleotides during citrate fermentation by *Klebsiella pneumoniae*. J. Bacteriol. **181**: 241-245
21. Pos, K.M., Dimroth, P. and Bott, M. (1998) The *Escherichia coli* citrate carrier CitT: a member of a novel eubacterial transporter family related to the 2-oxoglutarate/malate translocator from spinach chloroplasts. J. Bacteriol. **180**: 4160-4165.
20. Bott, M., Pfister, K., Burda, P., Kalbermatter, O., Woehlke, G. and Dimroth, P. (1997) Methylmalonyl-CoA decarboxylase from *Propionigenium modestum*: Cloning and sequencing of the structural genes and purification of the enzyme complex. Eur. J. Biochem. **250**: 590-599
19. Meyer, M., Dimroth, P. and Bott, M. (1997) *In vitro* binding of the response regulator CitB and of its carboxyterminal domain to A + T-rich DNA target sequences in the control region of the divergent *citC* and *citS* operons of *Klebsiella pneumoniae*. J. Mol. Biol. **269**: 719-731
18. Bott, M. (1997) Anaerobic citrate metabolism and its regulation in enterobacteria. Arch. Microbiol. **167**: 78-88

----- 1991 - 1995 -----

17. Bott, M., Meyer, M. and Dimroth, P. (1995) Regulation of anaerobic citrate metabolism in *Klebsiella pneumoniae*. Mol. Microbiol. **18**: 533-546
16. Bott, M., Thöny-Meyer, L., Loferer, H., Rossbach, S., Tully, R.E., Keister, D., Appleby, C.A. and Hennecke, H. (1995) *Bradyrhizobium japonicum* cytochrome c_{550} is required for nitrate respiration but not for symbiotic nitrogen fixation. J. Bacteriol. **177**: 2214-2217
15. Bott, M. and Dimroth, P. (1994) *Klebsiella pneumoniae* genes for citrate lyase and citrate lyase ligase: localization, sequencing, and expression. Mol. Microbiol. **14**: 347-356
14. Pos, K.M., Bott, M. and Dimroth, P. (1994) Purification of two active fusion proteins of the Na⁺-dependent citrate carrier of *Klebsiella pneumoniae*. FEBS Lett. **347**: 37-41
13. Loferer, H., Bott, M. and Hennecke, H. (1993) *Bradyrhizobium japonicum* TlpA, a novel membrane-anchored thioredoxin-like protein involved in the biogenesis of cytochrome aa₃ and development of symbiosis. EMBO J. **12**: 3373-3383
12. Hennecke, H., Anthamatten, D., Babst, M., Bott, M., Fischer, H.M., Kaspar, T., Kullik, I., Loferer, H., Preisig, O., Ritz, D. and Weidenhaupt, M. (1993) Genetic and physiologic requirements for optimal bacteroid function in the *Bradyrhizobium japonicum*-soybean symbiosis. In: *Advances in Molecular Genetics of Plant-Microbe Interactions*. Nester, E.W. and Verma, D.P.S. (eds.). Kluwer Academic Publishers (Dordrecht), pp. 199-207

11. **Ritz, D., Bott, M. and Hennecke, H. (1993)** Formation of several bacterial c-type cytochromes requires a novel membrane-anchored protein that faces the periplasm. *Mol. Microbiol.* **9**: 729-740
10. **Bott, M., Preisig, O. and Hennecke, H. (1992)** Genes for a second terminal oxidase in *Bradyrhizobium japonicum*. *Arch. Microbiol.* **158**: 335-343
9. **Bott, M., Ritz, D. and Hennecke, H. (1991)** The *Bradyrhizobium japonicum cycM* gene encodes a membrane-anchored homolog of mitochondrial cytochrome c. *J. Bacteriol.* **173**: 6766-6772

----- 1985 - 1990 -----

8. **Hennecke, H., Bott, M., Ramseier, T., Thöny-Meyer, L., Fischer, H.-M., Anthamatten, D., Kullik, I. and Thöny, B. (1990)** A genetic approach to analyze the critical role of oxygen in bacteroid metabolism. In: *Nitrogen Fixation: Achievements and Objectives*. Gresshoff, P.M., Roth L.E., Stacey, G. und Newton, W.E. (eds.). Chapman and Hall (New York, London), pp. 293-300
7. **Bott, M., Bolliger, M. and Hennecke, H. (1990)** Genetic analysis of the cytochrome c-aa₃ branch of the *Bradyrhizobium japonicum* respiratory chain. *Mol. Microbiol.* **4**: 2147-2157
6. **Karrasch, M., Bott, M. and Thauer, R.K. (1989)** Carbonic anhydrase activity in acetate-grown *Methanosarcina barkeri*. *Arch. Microbiol.* **151**: 137-142
5. **Bott, M. and Thauer, R.K. (1989)** The active species of "CO₂" formed by carbon monoxide dehydrogenase from *Peptostreptococcus productus*. *Z. Naturforsch.* **44c**: 392-396
4. **Bott, M. and Thauer, R.K. (1989)** Proton translocation coupled to the oxidation of carbon monoxide to CO₂ and H₂ in *Methanosarcina barkeri*. *Eur. J. Biochem.* **179**: 469-472
3. **Bott, M. and Thauer, R.K (1987)** Proton-motive-force-driven formation of CO from CO₂ and H₂ in methanogenic bacteria. *Eur. J. Biochem.* **168**: 407-412
2. **Bott, M., Eikmanns, B. and Thauer, R.K. (1986)** Coupling of carbon monoxide oxidation to CO₂ and H₂ with the phosphorylation of ADP in acetate-grown *Methanosarcina barkeri*. *Eur. J. Biochem.* **159**: 393-398
1. **Bott, M., Eikmanns, B. and Thauer, R.K. (1985)** Defective formation and/or utilization of carbon monoxide by H₂/CO₂ fermenting methanogens dependent on acetate as carbon source. *Arch. Microbiol.* **143**: 266-269