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Key Publications

Metabolic Engineering

- S. Ostergaard; L. Olsson; M. Johnston; J. Nielsen (2000) Increasing galactose consumption by Saccharomyces
 cerevisiae through metabolic engineering of the GAL gene regulatory network. Nature Biotechnol. 18:1283-1286
- K.-K. Hong; W. Vongsangnak; G.N. Vemuri; J. Nielsen (2011) Unravelling evolutionary strategies of yeast for improving galactose utilization through integrated systems level analysis. Proc. Nat. Acad. Sci. USA 108:12179-12184
- L. Caspeta; J. Nielsen (2013) Economic and environmental impacts of microbial biodiesel. Nature Biotechnol. 31:789-793
- J.C. Qin; Y.J. Zhou; A. Krivoruchko; M. Huang; L. Liu; S. Khoomrung; V. Siewers; B. Jiang; J. Nielsen (2015) Modular pathway rewiring of *Saccharomyces cerevisiae* enables high-level production of L-ornitine. *Nature Com.* 6:8224
- M. Huang; Y. Bai; S.L. Sjostrom; B.M. Hallström; Z. Liu; D. Petranovic; M. Uhlen; H.N. Joensson; H. Andersson-Svahn; J. Nielsen (2015) Microfluidic screening and whole genome sequencing identifies mutations associated with improved protein secretion by yeast. *Proc. Nat. Acad. Sci. USA* 112:E4689-96
- 6. Y. Zhou; N. A. Buijs; Z. Zhu; J. Qin; V. Siewers; **J. Nielsen** (2016) Production of fatty acid derived oleochemicals and biofuels by synthetic yeast cell factories. *Nature Com.* **7**:11709
- Y. Zhou; N.A. Buijs; Z. Zhu; D.O. Gomez; A. Boonsombuti; V. Siewers; J. Nielsen (2016) Harnessing peroxisomes for production of fatty acid-derived biofuels and chemicals in yeast. J. Am. Chem. Soc. 138:15368-15377
- 8. Z. Zhu; Y.J. Zhou; A. Krivoruchko; M. Grininger; Z.K. Zhao; J. Nielsen (2017) Expanding the product portfolio of fungal type I fatty acid synthases. *Nature Chem. Biol.* 13:360-362
- 9. M. Huang; J. Bao; B.M. Hallström; D. Petranovic; J. Nielsen (2017) Efficient protein production by yeast requires global tuning of metabolism. *Nature Com.* **8**:1131
- R. Ferreira; P.G. Teixeira; V. Siewers; J. Nielsen (2018) Redirection of lipid flux towards phospholipids in yeast increases fatty acid turnover and secretion. Proc. Nat. Acad. Sci. USA 115:1262-1267
- Z. Dai; M. Huang; Y. Chen; V. Siewers; J. Nielsen (2018) Global rewiring of cellular metabolism renders Saccharomyces cerevisiae Crabtree-negative. Nature Com. 9:3059
- 12. T. Yu; Y. Zhou; M. Huang; Q. Liu; R. Pereira; F. David; J. Nielsen (2018) Reprogramming yeast metabolism from alcoholic fermentation to lipogenesis. *Cell* 174:1-10
- M. Huang; G. Wang; J. Qin; D. Petranovic; J. Nielsen (2018) Engineering the protein secretory pathway of Saccharomyces cerevisiae enables improved protein production. Proc. Nat. Acad. Sci. USA 115:E11025-E11032
- 14. Y. Liu; Q. Liu; A. Krivoruchko; S. Khoomrung; **J. Nielsen** (2020) Engineering yeast phospholipid metabolism for *de novo* oleoylethanolamide production. *Nature Chem. Biol.* **16**:197-205
- 15. Z. Zhu; Y. Hu; P.G. Teixeira; R. Pereira; Y. Chen; V. Siewers; J. Nielsen (2020) Multidimensional engineering of *Saccharomyces cerevisiae* for efficient synthesis of medium-chain fatty acids. *Nature Cat.* **3**:64-74
- J. Qin, A. Krivoruchko; B. Ji; Y. Chen; M. Kristensen, E. Özdemir; J. D. Keasling; M.K. Jensen; J. Nielsen (2021) Engineering yeast metabolism for the discovery and production of polyamines and polyamine analogues. *Nature Cat.* 4:498-509
- 17. X. Li; Y. Wang; G. Lo; Q. Liu; R. Pereira; Y. Chen; J. Nielsen (2021) Metabolic network remodeling enhances yeast's fitness on xylose using aerobic glycolysis. *Nature Cat.* **4**:783-796
- 18. Q. Liu; Y. Liu; G. Li; O. Savolaiinen; Y. Chen; J. Nielsen (2021) *De novo* biosynthesis of bioactive isoflavonoids by engineered yeast cell factories. *Nature Com.* **12**:6085
- 19. N. Qin; L. Li; X. Ji; R. Pereira; Y. Chen; S. Yin; C. Li; X. Wan; D. Qiu; J. Jiang; H. Lou; Y. Zhang; G. Dong; Y. Zhang; S. Shi; H.J. Jessen; J. Xia; Y. Chen; C. Larsson; T. Tan; Z. Liu; J. Nielsen (2023) Flux regulation through glycolysis and respiration is balanced by inositol pyrophosphates. *Cell* 186:748-76

Systems Biology

- J. Förster; I. Famili; P. Fu; B. Ø. Palsson; J. Nielsen (2003) Genome-scale reconstruction of the Saccharomyces cerevisiae metabolic network. Genome Res. 13:244-253
- 2. I. Borodina; P. Krabben; **J. Nielsen** (2005) Genome-scale analysis of *Streptomyces coelicolor* A3(2) metabolism. *Genome Res.* 15:820-829
- 3. K. R. Patil, **J. Nielsen** (2005) Uncovering transcriptional regulation of metabolism by using metabolic network topology. *Proc. Nat. Acad. Sci.* 102:2685-2689
- M. R. Andersen; M. L. Nielsen; J. Nielsen (2008) Metabolic model integration of the bibliome, genome, metabolome and reactome of Aspergillus niger. Mol. Systems Biol. 4:178
- 5. M. R. Andersen; W. Vongsangnak; G. Panagiotou; M. P. Salazar; L. Lehmann; **J. Nielsen** (2008) A trispecies *Aspergillus* microarray: Comparative transcriptomics of three *Aspergillus* species. *Proc. Nat. Acad. Sci.* 105:4387-4392
- R. Agren; L. Liu; S. Shoaie; W. Vongsangnak; I. Nookaew; J. Nielsen (2013) The RAVEN toolbox and its use for generating a genome-scale metabolic model for *Penicillium chrysogenum*. *PLoS Comp. Biol.* 9:e1002980
- L. Caspeta; Y. Chen; P. Ghiaci; A. Feizi; S. Buskov; B.M. Hallström; D. Petranovic; J. Nielsen (2014) Altered sterol
 composition renders yeast thermotolerant. Science 346:75-78
- 8. J.C. Nielsen; S. Grijseels; S. Prigent; B. Ji; J. Dainat; K.F. Nielsen; J.C. Frisvad; M. Workman; **J. Nielsen** (2017) Global analysis of biosynthetic gene clusters reveals vast potential of secondary metabolite production in *Penicillium* species. *Nature Microbiol.* **2**:17044
- 9. P.-J. Lahtvee; B.J. Sanchez; A. Smialowska; S. Kasvandik; I. Elsemman; F. Gatto; **J. Nielsen** (2017) Absolute quantification of protein and mRNA abundances demonstrate variability in gene-specific translation efficiency in yeast. *Cell Systems* **4**:495-504

- 10. B.J. Sanchez; C. Zhang; A- Nilsson; P.-J. Lahtvee; E. Kerkhoven; J. Nielsen (2017) Improving the phenotype predictions of a yeast genome-scale metabolic model by incorporating enzymatic constraints. *Mol. Systems Biol.* 13:935
- 11. Y. Chen; J. Nielsen (2019) Energy metabolism controls phenotypes by protein efficiency and allocation. *Proc. Nat. Acad. Sci. USA* 116:17592-17597
- 12. H. Lu; F. Li; B.J. Sanchez; Z. Zhu; G. Li; I. Domenzain; S. Marcisauskas; P.M. Anton; D. Lappa; C. Lieven; M.E. Beber; N. Sonnenschein; E.J. Kerkhoven; **J. Nielsen** (2019) A consensus *S. cerevisiae* metabolic model Yeast8 and its ecosystem for comprehensively probing cellular metabolism. *Nature Com.* **10**:3586
- 13. R. Yu; K. Campbell; R. Pereira; J. Björkeroth; Q. Qi; E. Vorontsov; C. Sihlbom; J. Nielsen (2020) Nitrogen limitation reveals large reserves in metabolic and translational capacities of yeast. *Nature Com.* 11:1881
- 14. K. Campbell; J. Westholm; S. Kasvendik; F. Di Bartolomei; M. Mormino; J. Nielsen (2020) Building blocks are synthesized on demand during the yeast cell cycle. *Proc. Nat. Acad. Sci. USA* 117:7575-7583
- F. Di Bartolomeo; C. Malina; K. Campbell; M. Mormini; J. Fuchs; E. Vorontsov; C.M. Gustafsson; J. Nielsen (2020)
 Absolute yeast mitochondrial proteome quantification reveals trade-off between biosynthesis and energy generation
 during diauxic shift. *Proc. Nat. Acad. Sci. USA* 117:7524-7535
- 16. J. Björkeroth; K. Campbell; C. Malina; R. Yu; F. Di Bartolomeo; J. Nielsen (2020) Proteome re-allocation from amino acid biosynthesis to ribosomes enables yeast to grow faster in rich media. *Proc. Nat. Acad. Sci. USA* 117:21804-21812
- 17. Y. Chen; F. Li; J. Mao; Y. Chen; J. Nielsen (2021) Yeast optimizes metal utilization based on metabolic network and enzyme kinetics. *Proc. Nat. Acad. Sci. USA* 118:e2020154118
- 18. Y. Chen; **J. Nielsen** (2021) In vitro turnover numbers do not reflect in vivo activities of yeast enzymes. *Proc. Nat. Acad. Sci. USA* **118**:e2108391118
- 19. G. Li; Y. Hu; J. Zrimec; H. Lou; H. Wang; A. Zelezniak; B. Ji; J. Nielsen (2021) Bayesian genome scale modeling identifies termal determinants of yeast metabolism. *Nature Com.* 12:190
- 20. Y. Chen; F. Li; J. Mao; Y. Chen; J. Nielsen (2021) Yeast optimizes metal utilization based on metabolic network and enzyme kinetics. *Proc. Nat. Acad. Sci. USA* 118:e2020154118
- 21. R. Yu; E. Vorontsov; C. Sihlbom; **J. Nielsen** (2021) Quantifying absolute gene expression profiles reveals distinct regulation of central carbon metabolism genes in yeast. *eLife* **10**:e65722
- 22. H. Lu; F. Li; L. Yuan; I. Domenzain; R. Yu; H. Wang; G. Li; Y. Chen; B. Ji; E.J. Kerkhoven; **J. Nielsen** (2021) Yeast metabolic innovations emerged via expanded metabolic network and gene positive selection. *Mol. Systems Biol.* 17:e10427
- F. Li; L. Yuan; H. Lu; G. Li; Y. Chen; M.K.M. Engqvist; E.J. Kerkhoven; J. Nielsen (2022) Deep learning based k_{cat} prediction enables improved enzyme constrained model reconstruction. Nature Cat. 5:662-672
- 24. F. Li; Y. Chen; Q. Qi; Y. Wang; L. Yuan; M. Huang; I.E. Elsemman; A. Feizi; E.J. Kerkhoven; J. Nielsen (2022) Improving recombinant protein production by yeast through genome-scale modelling using proteome constraints. *Nature Com.* 13:2969
- J. Xia; B.J. Sanchez; Y. Chen; K. Campbell; S. Kasvandik; J. Nielsen (2022) Proteome allocation change linearly with specific growth rate of Saccharomyces cerevisiae under glucose-limitation. Nature Com. 13:2819
- I. Domenzain; B. Sanchez; M. Anton; E.J. Kerkhoven; A. Millan-Oropeza; C. Henry; V. Siewers; J.P. Morrissey; N. Sonnenschein; J. Nielsen (2022) Reconstruction of a catalogie of genome-scale metabolic models with enzymatic constraints using GECKO2.0. Nature Com. 13:3766

Human Metabolism

- 1. F. H. Karlsson; F. Fåk; I. Nookaew; V. Tremaroli; B. Fagerberg; D. Petranovic; F. Bäckhed*; J. Nielsen* (2012) Symptomatic atherosclerosis is associated with an altered gut metagenome. *Nature Com.* 3:1245
- F. Karlsson; V. Tremaroli; I. Nookaew; G. Bergström; C.J. Behre; B. Fagerberg; J. Nielsen*; F. Bäckhed* (2013) Gut metagenome in European women with normal, impaired and diabetic glucose control. Nature 498:99-103
- A. Mardinoglu; R. Agren; C. Kampf; A. Asplund; I. Nookaew; P. Jacobsen; A.J. Walley; P. Froguel; L.M. Carlsson; M. Uhlen; J. Nielsen (2013) Integration of clinical data with a genome-scale metabolic model of the human adipocyte.
 Mol. Systems Biol. 9:649
- F. Gatto; I. Nookaew; J. Nielsen (2014) Chromosome 3p loss of heterozygosity is associated with a unique metabolic network in clear cell renal carcinoma. *Proc. Nat. Acad. Sci.* 111:E866-E875
- A. Mardinoglu; R. Agren; K. Kampf; A. Asplund; M. Uhlen; J. Nielsen (2014) Genome-scale metabolic modeling of hepatocytes reveals serine deficiency in patients with non-alcoholic fatty liver disease. *Nature Com.* 5:3083
- 6. R. Agren; A. Mardinoglu; C. Kampf; A. Asplund; M. Uhlen; J. Nielsen (2014) Identification of anticancer drugs for hepatocellular carcinoma through personalized genome-scale metabolic modeling. *Mol. Systems Biol.* 10:721
- 7. M. Uhlén, L. Fagerberg, B.M. Hallström, C. Lindskog, P. Oksvold, A. Mardinoglu, Å. Sivertsson, C. Kampf, E. Sjöstedt, A. Asplund, I. Olsson, K. Edlund, E. Lundberg, S. Navani, C.A.-K. Szigyarto, J. Odeberg, D. Djureinovic, J.O. Takanen, S. Hober, T. Alm, H. Berling, H. Tegel, J. Mulder, J. Rockberg, P. Nilsson, J.M. Schwenk, M. Hamsten, K. von Feilitzen, M. Forsberg, L. Persson, F. Johansson, M. Zwahlen, G. von Heijne, J. Nielsen; F. Ponten (2015) Tissue based map of the humane proteome. *Science* 347:1260419,1-9S
- 8. Shoaie; P. Ghaffari; P. Kovatcheva-Datchary; A. Mardinoglu; P. Sen; E. Pujos-Guillot; T. de Wouters; C. Juste; S. Rizkalla; J. Chilloux; L. Hoyles; J.K. Nicholson; ANR MicroObese Consortium; J. Dore; M.E. Dumas; K. Clement; F. Bäckhed; J. Nielsen (2015) Quantifying diet-induced metabolic changes of the human gut microbiome. *Cell Metabolism* 22:320-331
- 9. A. Mardinoglu; S. Shoaie; M. Bergentall; P. Ghaffari; C. Zhang; E. Larsson; F. Bäckhed; J. Nielsen (2015) The gut microbiome modulates host amino acid and glutathione metabolism in mice. *Mol. Systems Biol.* 11:834

- F. Gatto; I. Nookaew; H. Nilsson; M. Maruzzo; A. Roma; M. E. Johansson; U. Steiner; S. Lundstam; N. Volpi; U. Basso; J. Nielsen (2016) Measurements of glycosaminoglycans in plasma and urine for diagnosis of clear cell renal cell carcinoma. *Cell Rep.* 15:1-15
- 11. P. Babaei; S. Shoaei; B. Ji; **J. Nielsen** (2018) Challenges in modeling the human gut microbiome. *Nature Biotechnol.* **16**:682-686
- 12. J.L. Robinson; A. Feizi; M. Uhlen; J. Nielsen (2019) A systematic investigation of the malignant functions and diagnostic potential of the cancer secretome. *Cell Rep.* 10:2622-2635
- 13. A. Nilsson, E. Björnson; M. Flockhart; F.J. Larsen; J. Nielsen (2019) Complex I is bypassed during high intensity exercise. *Nature Com.* 10:5072
- J.L. Robinson; P. Kocabas; H. Wang; P.-E. Cholly; D. Cook; A. Nilsson; M. Anton; R. Ferreira; I. Domenzain; V. Billa;
 A. Limeta; A. Hedin; J. Gustafsson; E.J. Kerkhoven; L.T. Svensson; B.O. Palsson; A. Mardinoglu; L. Hansson; M. Uhlen; J. Nielsen (2020) An Atlas of Human Metabolism. Science Signal. 13:eaaz1482
- A. Nilsson; J.R. Haanstra; M. Engqvist; A. Gerding; B.M. Bakker; U. Klingmüller; B. Teusink; J. Nielsen (2020)
 Quantitative analysis of amino acid metabolism in liver cancer links glutamate excretion to nucleotide synthesis. *Proc. Nat. Acad. Sci. USA* 117:10294-10304
- 16. J. Geng; B. Ji: G. Li; F. Lopez-Isunza; J. Nielsen (2021) CODY enables quantitatively spatiotemporal predictions on in vivo gut microbial variability induced by diet-intervention. *Proc. Nat. Acad. Sci. USA* 118:e2019336118
- H. Wang; J.L. Robinson; P. Kocabas; J. Gustafsson; M. Anton; P.-E. Cholley; S. Huang; J. Gobom; T. Svensson; M. Uhlen; H. Zetterberg; J. Nielsen (2021) Genome-scale metabolic network reconstruction of model animals as a platform for translational research. *Proc. Nat. Acad. Sci. USA* 118:e2102344118

Reviews

- S. Ostergaard; L. Olsson; J. Nielsen (2000) Metabolic engineering of Saccharomyces cerevisiae. Microb. Mol. Biol. Rev. 64:34-50
- 2. J. Nielsen (2001) Metabolic engineering. Appl. Microbiol. Biotechnol. 55:263-283
- 3. K.-K. Hong; **J. Nielsen** (2012) Metabolic engineering of *Saccharomyces cerevisiae*: A key cell factory platform for future biorefineries. *Cell. Mol. Life Sci.* **16**:2671-2690
- 4. M. Uhlen; B.M. Hallström; C. Lindskog; A. Mardinoglu; F. Ponten; J. Nielsen (2016) Transcriptomics resources of human tissues and organs. *Mol. Sys. Biol.* 12:862
- J. Nielsen; J. Keasling (2016) Engineering Cellular Metabolism. Cell 164:1185-1197
- J. Nielsen (2017) Systems Biology of Metabolism: A Driver for Developing Personalized and Precision Medicine. Cell Met. 25:572-579
- 7. **J. Nielsen** (2017) Systems Biology of Metabolism. *Ann. Rev. Biochem.* **86**:245-275
- 8. K. Campbell; J. Xia; **J. Nielsen** (2017) The impact of systems biology on bioprocessing. *Trends Biotechnol.* **35**:1156-1168
- 9. A. Mardinoglu; J. Boren; U. Smith; M. Uhlen; **J. Nielsen** (2018) Systems biology in hepatology: Approaches and applications *Nature Rev. Gastro. Hep.* **15**:365-377
- Y. Zhou; E. Kerkhoven; J. Nielsen (2018) Barriers and opportunities in bio-based production of hydrocarbons. *Nature Energy* 3:925-935
- M. Kumar; B. Ji; K. Zengler; J. Nielsen (2019) Modeling approaches for studying the gut microbiota. *Nature Microbiol.* 4:1253-1267
- 12. Z. Liu; K. Wang; Y. Chen; T. Tan; J. Nielsen (2020) Third-generation biorefineries as a mean to produce fuels and chemicals from CO₂. *Nature Cat.* **3**:274-288
- H. Lu; E.J. Kerkhoven; J. Nielsen (2022) Multiscale models quantifying yeast physiology: towards a whole-cell model. Trends Biotechnol. 40:291-305
- 14. X. Tan; J. Nielsen (2022) The integration of bio-catalysis and electrocatalysis to produce fuels and chemicals from carbon dioxide. *Chem. Soc. Rev.* **51**:4753
- 15. **J. Nielsen**; C.B. Tillegreen; D. Petranovic (2022) Innovation trends in industrial biotechnology. *Trends Biotechnol.*, in press

Commentaries and Perspectives

- 1. J. Nielsen (2007) Principles of optimal metabolic network operation. Mol. Sys. Biol. 3:126
- 2. J. Nielsen (2011) Transcriptional control of metabolic fluxes. Mol. Systems Biol. 7:478
- 3. J. Nielsen (2011) Chimeric Synthetic Pathways. Nature Chem. Biol. 7:195-196
- 4. **J. Nielsen**; J. Keasling (2011) Synergies between synthetic biology and metabolic engineering. *Nature Biotechnol.* **29**:693-695
- 5. **J. Nielsen**; M. Fussenegger; J. Keasling; S.Y. Lee; J.C. Liao; K. Prather; B. Palsson (2014) Engineering synergy in biotechnology. *Nature Chem. Biol.* **10**:319-322
- 6. J. Nielsen (2014) Synthetic Biology for Engineering Acetyl Coenzyme A Metabolism in Yeast. mBio 5:e02153-14
- 7. J. Pronk; S.Y. Lee; J. Lievense; J. Pierce; B. Palsson; M. Uhlen; J. Nielsen (2015) How to set up collaborations between academia and industrial biotech companies. *Nature Biotechnol.* 33:237-240
- 8. J. Nielsen (2015) Yeast cell factories on the horizon. Science 349:1050-1051
- 9. **J. Nielsen** (2017) Built on stable catalysts. *Nature Microbiol.* **2**:17085
- 10. J. Nielsen (2019) Designer Microbes Serving Society. Cell Met. 29:50
- 11. J. Nielsen (2019) Cell factory engineering for improved production of natural products. Nat. Prod. Rep. 36;1233

- 12. J. Nielsen (2019) Antibiotic lethality is impacted by nutrient availabilities: New insights from machine learning. Cell **177**:1373-1374
- 13. J. Nielsen (2019) A stress-coping strategy for yeast cells. *Nature* 572:184-185
 14. J. Nielsen (2022) Bioactive metabolites: The double-edged sword in your food. *Cell* 185:4469-4471

Complete List of Publications

Research papers in international journals with peer review

1988

1. K. Nikolajsen; J. Nielsen; J. Villadsen (1988) In-line flow injection analysis for monitoring lactic acid fermentations. Anal. Chim. Acta 214:137-145

1989

- J. Nielsen; K. Nikolajsen; J. Villadsen (1989) FIA for on-line monitoring of important lactic acid fermentation variables. Biotechnol. Bioeng. 33:1127-1134
- 3. **J. Nielsen**; C. Emborg; K. Halberg; J. Villadsen (1989) Compartment model concept used in the design of fermentation with recombinant microorganisms *Biotechnol. Bioeng.* 34:478-486

1990

- J. Nielsen; K. Nikolajsen; S. Benthin; J. Villadsen (1990) Application of flow-injection analysis in the on-line monitoring of sugars, lactic acid, protein, and biomass during lactic acid fermentations. *Anal. Chim. Acta* 237:165-175
- G. Wehnert; K.-D. Anders; C. Bittner; R. Kammeyer; U. Hübner; J. Nielsen; T. Scheper (1990) Ein kombinierter Fluoreszenz-/Streulichtsensor und dessen Einsatz zur Prozessbeobachtung in der Biotechnologie. *Chem.-Ing.-Tech.* 62:211-212

1991

- S. Benthin; J. Nielsen; J. Villadsen (1991) A simple and reliable method for the determination of cellular RNA content. Biotechnol. Techn., 5:39-42
- 7. **J. Nielsen**; K. Nikolajsen; J. Villadsen (1991) Structured modelling of a microbial system 1. A theoretical study of the lactic acid fermentation. *Biotechnol. Bioeng.* 38:1-10
- 8. **J. Nielsen**; K. Nikolajsen; J. Villadsen (1991) Structured modelling of a microbial system 2. Experimental verification of a structured lactic acid fermentation model. *Biotechnol. Bioeng.* 38:11-23
- 9. K. Nikolajsen; J. Nielsen; J. Villadsen (1991) Structured modelling of a microbial system 3. Growth on mixed substrates. *Biotechnol. Bioeng.* 38:24-29
- J. Nielsen; A. G. Pedersen; K. Strudsholm; J. Villadsen (1991) Modelling fermentations with recombinant microorganisms: Formulation of a structured model. *Biotechnol. Bioeng.* 37:802-808
- 11. S. Benthin; **J. Nielsen**; J. Villadsen (1991) Characterisation and application of precise and robust flow injection analyzers for on-line measurement during fermentations. *Anal. Chim. Acta* 247:45-50
- 12. L. H. Christensen; J. Nielsen; J. Villadsen (1991) Monitoring of substrates and products during fed-batch penicillin fermentations on complex media. *Anal. Chim. Acta* 249:123-136
- 13. L. H. Christensen; **J. Nielsen**; J. Villadsen (1991) Delay and dispersion in an in-situ membrane probe for bioreactors. *Chem. Eng. Sci.* 46:3304-3307

1992

- 14. S. Benthin; J. Nielsen; J. Villadsen (1992) Flow Injection Analysis of micromolar concentrations of glucose and lactate in fermentation media. *Anal. Chim. Acta* 261:145-153
- S. Benthin; J. Nielsen; J. Villadsen (1992) Anomeric specificity of glucose uptake systems in *Lactococcus cremoris*, *Escherichia coli* and *Saccharomyces cerevisiae*: Mechanisms, kinetics and implications. *Biotechnol. Bioeng.* 40:137-146
- 16. J. Nielsen (1992) On-line monitoring of microbial processes by flow injection analysis. Proc. Control Qual. 2:371-384
- 17. C. L. Johansen; L. H. Christensen; J. Villadsen; J. Nielsen (1992) Monitoring and control of fed-batch penicillin fermentation. *Comp. Chem. Eng.* 16:S297-S304
- 18. K. Strudsholm; J. Nielsen; C. Emborg (1992) Product formation during batch fermentation with recombinant *Escherichia coli* containing a runaway plasmid. *Bioprocess Eng.* 8:173-181

1993

- 19. G. Pedersen; M. Bundgaard; O. Hassager; J. Nielsen; J. Villadsen (1993) Rheological characterization of media containing *Penicillium chrysogenum*. *Biotechnol. Bioeng*. 41:162-164
- 20. **J. Nielsen** (1993) A simple morphologically structured model describing the growth of filamentous microorganisms. *Biotechnol. Bioeng.* 41:715-727
- 21. M. Carlsen; L. H. Christensen; J. Nielsen (1993) Flow-injection analysis for measurement of penicillin V in fermentation samples. *Anal. Chim. Acta* 274:117-123
- 22. M. Carlsen; H. Meier; F. Lantreibecq; C. Johansen; R. W. Min; **J. Nielsen** (1993) On-line monitoring of penicillin V during penicillin fermentations: A comparison of two different methods based on FIA. *Anal. Chim. Acta* 279:51-58
- 23. S. Benthin; **J. Nielsen**; J. Villadsen (1993) Transport of sugars via two anomer-specific sites on mannose-phosphotransferase system in *Lactococcus cremoris*: In vivo study of mechanism, kinetics and adaption. *Biotechnol. Bioeng.* 42:440-448
- 24. J. Nielsen (1993) Simulation of bioreactions. Comp. Chem. Eng. 18:S615-S620
- S. Benthin; J. Nielsen; J. Villadsen (1993) Two uptake systems for fructose in *Lactococcus lactis* subsp. *cremoris* FD1 produce glucolytic and gluconeogenic fructose phosphates and induce oscillations of growth and lactic acid formation. *Appl. Environ. Microbiol.* 59:3206-3211

- S. Benthin; J. Nielsen; J. Villadsen (1994) Galactose expulsion during lactose metabolism in *Lactococcus lactis* subsp. cremoris FD1 due to dephosphorylation of intracellular galactose-6-phosphate. Appl. Environ. Microbiol. 60:1254-1259
- 27. A. G. Pedersen; M. Bundgaard-Nielsen; J. Nielsen; J. Villadsen (1994) Characterization of mixing in stirred tank bioreactors equipped with Rushton turbines. *Biotechnol. Bioeng.* 44:1013-1017
- 28. A. G. Pedersen; H. Andersen; J. Nielsen; J. Villadsen (1994) A novel technique based on ⁸⁵Kr for quantification of gas-liquid mass transfer in bioreactors. *Chem.Eng. Sci.* 49:803-810
- S. Benthin; U. Schulze; J. Nielsen; J. Villadsen (1994) Growth energetics of Lactococcus cremoris FD1 during energy-, carbon- and nitrogen limitation in steady state and transient cultures. Chem. Eng. Sci. 49:589-610
- 30. L. H. Christensen; G. Mandrup; **J. Nielsen**; J. Villadsen (1994) A robust LC method for studying the penicillin fermentation. *Anal. Chim. Acta* 296:51-62
- 31. L. H. Christensen; J. Nielsen; J. Villadsen (1994) Degradation of penicillin V in fermentation media. *Biotechnol. Bioeng.* 44:165-169
- 32. M. A. Hjortso; **J. Nielsen** (1994) A conceptual model of autonomous oscillations in microbial cultures. *Chem. Eng. Sci.* 49:1083-1095
- 33. J. Nielsen; C. L. Johansen; J. Villadsen (1994) Culture fluorescence measurements during batch and fed-batch cultivations with *Penicillium chrysogenum*. J. Biotechnol 38:51-62
- 34. M. Carlsen; J. Marcher; J. Nielsen (1994) An improved FIA-system for measuring α-amylase in cultivation *media*, *Biotechnol. Tech.* 8:479-482

- 35. H. Jørgensen; J. Nielsen; J. Villadsen; H. Mølgaard (1995) Analysis of the penicillin V biosynthesis during fed-batch cultivations with a high yielding strain of *Penicillium chrysogenum*, *Appl. Microbiol. Biotechnol.* 43:123-130
- 36. **J. Nielsen**; P. Krabben (1995) Hyphal growth and fragmentation of *P. chrysogenum* in submerged cultures. *Biotechnol. Bioeng.* 46:588-598
- 37. **J. Nielsen**; C. L. Johansen; M. Jacobsen; P. Krabben; J. Villadsen (1995) Pellet formation and fragmentation in submerged cultures of *Penicillium chrysogenum* and its relation to penicillin production, *Biotechnol. Prog.* 11:93-98
- 38. H. Jørgensen; J. Nielsen; J. Villadsen; H. Mølgaard (1995) Metabolic flux distributions in *Penicillium chrysogenum* during fed-batch cultivations. *Biotechnol. Bioeng.* 46:117-131
- 39. R. Mørkeberg; M. Carlsen; **J. Nielsen** (1995) Induction and repression of α-amylase production in recombinant and wild-type strains of *Aspergillus oryzae*, *Microbiol*. 141:2449-2454
- 40. **J. Nielsen**; H. S. Jørgensen (1995) Metabolic control analysis of the penicillin biosynthetic pathway in a *high yielding strain of Penicillium chrysogenum*, *Biotechnol. Prog.* 11:299-305
- 41. R. Lejeune; J. Nielsen; G. Baron (1995) Morphology of *Trichoderma reesei* QM 9414 in submerged cultures, *Biotechnol. Bioeng.* 47:609-615
- 42. L. H. Christensen; U. Schulze; J. Nielsen; J. Villadsen (1995) Acoustic gas analysis for fast and precise monitoring of bioreactors, *Chem. Eng. Sci.* 50:2101-2110
- 43. R. Lejeune; J. Nielsen; G. Baron (1995) Influence of pH on the morphology of *Trichoderma reesei* QM 9414 in submerged cultures. *Biotechnol. Lett.* 17:341-344
- 44. Rong Wei Min; **J. Nielsen**; J. Villadsen (1995) Simultaneous monitoring of glucose, lactic acid and penicillin by Sequential Injection Analysis. *Anal. Chim. Acta* 312:149-156
- 45. L. H. Christensen; C. M. Henriksen; J. Nielsen; J. Villadsen; M. Egel-Mitani (1995) Continuous cultivation of *P. chrysogenum*. Growth on glucose and penicillin production. *J. Biotechnol.* 42:95-107
- 46. M. A. Hjortso; **J. Nielsen** (1995) Population balance models of autonomous microbial oscillations. *J. Biotechnol.* 42:255-269
- 47. Rong Wei Min; M. Carlsen; **J. Nielsen**; J. Villadsen (1995) Measurements of α-amylase activity by Sequential Injection Analysis. *Biotechnol. Techn.* 9:765-768

- 48. M. Carlsen; J. Nielsen; J. Villadsen (1996) Kinetic studies of acid-inactivation of α-amylase from *Aspergillus oryzae*. *Chem. Eng. Sci.* 51:37-43
- 49. M. Carlsen; A. B. Spohr; J. Nielsen; J. Villadsen (1996) Morphology and physiology of an α-amylase producing strain of *Aspergillus oryzae* during batch cultivations, *Biotechnol. Bioeng.* 49:266-276
- C. M. Henriksen; L. H. Christensen; J. Nielsen; J. Villadsen (1996) Growth energetics and metabolic fluxes in continuous cultures of *Penicillium chrysogenum*, J. Biotechnol. 45:149-164
- 51. L. H. Christensen; J. Marcher; U. Schulze; M. Carlsen; R. W. Min; J. Nielsen; J. Villadsen (1996) Semi-on-line analysis for fast and precise monitoring of bioreaction processes. *Biotechnol. Bioeng*. 52:237-247
- 52. C. M. Henriksen; S. S. Holm; H. S. Jørgensen; J. Nielsen; J. Villadsen (1997) Kinetic studies of the carboxylation of 6-amino-penicillanic acid to 8-hydroxy-penillic acid, *Proc. Biochem.* 32:85-91
- 53. M. Carlsen; J. Nielsen; J. Villadsen (1996) Growth and α-amylase production of *Aspergillus oryzae* during continuous cultivations. *J. Biotechnol.* 45:81-93
- Rong Wei Min; J. Nielsen; J. Villadsen (1996) On-line monitoring of glucose and penicillin by Sequential Injection Analysis. Anal. Chim. Acta 320:199-205
- 55. P. de N. Pissarra; J. Nielsen; M. J. Bazin (1996) Pathway kinetics and metabolic control analysis of a high-yielding strain of *Penicillium chrysogenum* during fed-batch cultivations. *Biotechnol. Bioeng.* 51:168-176
- U. Schulze; G. Liden; J. Nielsen; J. Villadsen (1996) Physiological effects of nitrogen starvation in an anaerobic batch culture of Saccharomyces cerevisiae. Microbiology 142:2299-2310

- 57. **J. Nielsen**; H. S. Jørgensen (1996) Kinetic model for the penicillin biosynthetic pathway in *Penicillium chrysogenum*. *Control Eng. Prac.*4:765-771
- 58. C. J. L. Klein, L. Olsson, B. Rønnow, J. D. Mikkelsen, **J. Nielsen** (1996) Alleviation of glucose repression on maltose metabolism by *MIG1* disruption in *Saccharomyces cerevisiae*. *Appl. Environ. Microbiol*. 62:4441-4449

- 59. **J. Nielsen** (1997) Metabolic control analysis of biochemical pathways based on a thermokinetic description of reaction rates. *Biochem. J.* 321:133-138
- 60. T. L. Nissen; U. Schulze; J. Nielsen; J. Villadsen (1997) Flux distributions in anaerobic, glucose limited continuous cultures of *Saccharomyces cerevisiae*. *Microbiology* 143:203-218
- 61. K. Schmidt; M. Carlsen; J. Nielsen; J. Villadsen (1997) Modelling isotopmer distributions in biochemical networks Using isotopomer mapping matrices. *Biotechnol. Bioeng.* 55:831-840
- 62. T. C. Zangirolami; C. L. Johansen; J. Nielsen; S. B. Jørgensen (1997) Simulation of penicillin production in fed-batch cultivations using a morphologically structured model. *Biotechnol. Bioeng.* 56:593-604
- 63. P. de N. Pissarra; **J. Nielsen** (1997) Thermodynamics of metabolic pathways for penicillin production: Analysis of thermodynamic feasibility and free energy changes during fed-batch cultivations. *Biotechnol. Prog.* 13:156-165
- 64. M. Carlsen; K. V. Jocumsen; C. Emborg; J. Nielsen (1997) Modeling the growth and proteinase A production in continuous cultures of recombinant *Saccharomyces cerevisiae*. *Biotechnol*. *Bioeng*. 55:447-454
- 65. P. Krabben; J. Nielsen; M. L. Michelsen (1997) Analysis of single hyphal growth and fragmentation in submerged cultures using a population model. *Chem. Eng. Sci.* 52:2641-2652
- 66. A. B. Spohr; M. Carlsen; **J. Nielsen**; J. Villadsen (1997) Morphological characterization of recombinant strains of *Aspergillus oryzae* producing α-amylase during batch cultivations. *Biotechnol. Letters* 19:257-261
- 67. L. Olsson; M. E. Larsen; B. Rønnow; J. D. Mikkelsen; **J. Nielsen** (1997) Silencing *MIG1* in *Saccharomyces cerevisiae*: Effects of Antisense *MIG1* Expression and *MIG1* Gene Disruption. *Appl. Environ. Microbiol.* 63:2366-2371
- 68. B. Aa. Theilgaard; C. M. Henriksen; K. Kristiansen; J. Nielsen (1997) Purification and characterization of δ-(L-α-aminoadipyl)-L-cysteinyl-D-valine synthetase (ACVS) from Penicillium chrysogenum. Biochem. J. 327:185-191

1998

- 69. **J. Nielsen** (1998) Metabolic Engineering: Techniques for analysis of targets for genetic manipulations. *Biotechnol. Bioeng.* 58:125-132
- 70. A. Spohr; M. Carlsen; **J. Nielsen**; J. Villadsen (1998) α-Amylase production in recombinant *Aspergillus oryzae* during fed-batch and continuous cultivations. *J. Ferment. Bioeng.* 86:49-56
- 71. K Schmidt; A. Marx; A. A. de Graaf; W. Wiechert; H. Sahm; J. Nielsen; J. Villadsen (1998) ¹³C Tracer experiments and metabolite balancing for metabolic flux analysis: Comparing two approaches. *Biotechnol. Bioeng.* 58:254-257
- 72. T. Agger; A. B. Spohr; M. Carlsen; **J. Nielsen** (1998) Growth and product formation of *Aspergillus oryzae* during submerged cultivations: Verification of a morphologically structured model using fluorescent probes. *Biotechnol. Bioeng.* 57:321-329
- 73. M. Henriksen; J. Nielsen; J. Villadsen (1997) Influence of dissolved oxygen concentration on the penicillin biosynthetic pathway in steady state cultures of *Penicillium chrysogenum*. *Biotechnol. Prog.* 13:776-782
- A. B. Spohr; C. Dam Mikkelsen; M. Carlsen; J. Nielsen; J. Villadsen (1998) On-line study of fungal morphology during submerged growth in a small flow-through cell. *Biotechnol. Bioeng.* 58:541-553
- 75. S. Ostergaard; H. B. Aa. Theilgaard; J. Nielsen (1998) Identification and purification of O-acetyl-L-serine sulfhydrylase in *Penicillium chrysogenum*. Appl. Microbiol. Biotechnol. 50:663-668
- K. Klein; L. Olsson; B. Rønnow; J. D. Mikkelsen; J. Nielsen (1998) Glucose and maltose metabolism in MIG1disrupted and MAL-constitutive strains of Saccharomyces cerevisiae. Food Technol. Biotechnol. 35:287-292
- H. P. Smits; A. Cohen; T. Buttler; J. Nielsen; L. Olsson (1998) Clean-up and analysis of sugar phosphates in biological extracts by using solid phase extraction and anion-exchange chromatography with pulsed amperometric detection. *Anal. Biochem.* 261:36-42
- J. Holmalahti; O. Raatikainen; A. von Wright; H. Laatsch; A. Spohr; O. K. Lyngberg; J. Nielsen (1998) Production of dihydroabikoviromycin by *Streptomyces anulatus*. Production parameters and chemical characterization of genotoxicity. *J. Appl. Microbiol*. 85:61-68
- C. J. L. Klein; L. Olsson; J. Nielsen (1998) Nitrogen-limited continuous cultivations as a tool to quantify glucose control in Saccharomyces cerevisiae. Enz. Microbial. Technol. 23:91-100
- 80. C. M. Henriksen; **J. Nielsen**; J. Villadsen (1998) High exogeneous concentrations of phenoxyacetic acid are crucial for a high penicillin V productivity in *Penicillium chrysogenum*. *Microbiol*. 144:2001-2006
- 81. C. M. Henriksen; J. Nielsen; J. Villadsen (1998) Modelling of the protonophoric uncoupling by phenoxyacetic acid of the plasma membrane of *Penicillium chrysogenum*. *Biotechnol. Bioeng*. 60:761-767
- 82. J. Dynesen; H. P. Smits; L. Olsson; **J. Nielsen** (1998) Carbon catabolite repression of invertase during batch cultivations of *Saccharomyces cerevisiae*: The role of glucose, fructose, and mannose. *Appl. Microbiol. Biotechnol.* 50:579-582

- 83. K. Schmidt; **J. Nielsen**; J. Villadsen (1999) Quantitative analysis of metabolic fluxes in *E. coli*, using 2 dimensional NMR spectroscopy and complete isotopomer models. *J. Biotechnol*. 71:175-190
- 84. A. Aleksenko; W. Liu; Z. Gojkovic; **J. Nielsen**; J. Piskur (1999) Structural and transcriptional analysis of the *pyrABCN*, *pyrC*, and *PyrF* genes in *Aspergillus nidulans* and the evolutionary origin of fungal dihydroorotases. *Mol. Microbiol.* 33:599-611

- 85. H. Pedersen; M. Carlsen; J. Nielsen (1999) Identification of enzymes and quantification of metabolic fluxes in the wild type and in a recombinant strain of *Aspergillus oryzae* strain. *Appl. Environ. Microbiol.* 65:11-19
- 86. A. L. Santerre Henriksen; S. Even; C. Müller; P. J. Punt; C. A. M. J. J. van den Hondel; **J. Nielsen** (1999) Study of the glucoamylase promoter in *Aspergillus niger* using green fluorescence protein. *Microbiol*. 145:729-734
- 87. T. Christiansen; A. Spohr; **J. Nielsen** (1999) On-line study of growth kinetics of single hyphae of *Aspergillus oryzae* in a flow-through cell. *Biotechnol. Bioeng*. 63:147-153
- 88. A. Heydorn; T. Suhr-Jessen; **J. Nielsen** (1999) Growth and production kinetics of a teicoplanin producing strain of *Actinoplanes teicomyceticus*. *J. Antibio*. 52:40-44
- 89. H. Aae. Theilgaard; **J. Nielsen** (1999) Metabolic control analysis of the penicillin biosynthetic pathway: The influence of the LLD-ACV:bisACV ratio on the flux control. *Antonie van Leeuvenhoek* 75:145-154
- 90. C. J. L. Klein; J. J. Rasmussen; B. Rønnow; L. Olsson; **J. Nielsen** (1999) Investigation of the impact of *MIG1* and *MIG2* on the phyiology of *Saccharomyces cerevisiae*. *J. Biotechnol*. 68:197-212
- 91. M. Anderlund; T. L. Nissen; **J. Nielsen**; J. Villadsen; J. Rydström; B. Hahn-Hägerdal; M. C. Kielland-Brandt (1999) Expression of the *E. coli pntA* and *pntB* genes encoding nicotinamide nucleotide transhydrogenase in *Saccharomyces cerevisiae* and its effect on product formation during anaerobic glucose fermentation. *Appl. Environ. Microbiol.* 65:2333-2340
- K. Schmidt; L. C. Nørregaard; B. Pedersen; A. Meissner; J. Ø. Duus; J. Nielsen; J. Villadsen (1999) Quantification of intracellular metabolic fluxes from fractional enrichment and ¹³C-¹³C coupling constraints on the isotopomer distribution in labelled biomass components. *Metabol. Eng.* 1:166-179
- 93. B. Christensen; J. Nielsen (1999) Isotopomer analysis using GC-MS. Metabol. Eng. 1:282-290
- A. Santerre Henriksen; M. Carlsen; H. de Bang; J. Nielsen (1999) Kinetics of α-amylase secretion in Aspergillus oryzae. Biotechnol. Bioeng. 65:76-82
- 95. B. Rønnow; L. Olsson; **J. Nielsen**; J. D. Mikkelsen (1999) Derepression of galactose metabolism in melibiase producing baker's and distillers' yeast *J. Biotechnol.* 72:213-228
- T. Agger; J. Nielsen (1999) Genetically structured modelling of protein production in filamentous fungi. Biotechnol. Bioeng. 66:164-170

- 97. H. Pedersen; **J. Nielsen** (2000) The influence of nitrogen sources on the α-amylase productivity of *Aspergillus oryzae* in continuous culture. *Appl. Microb. Biotechnol.* 53:278-281
- 98. C. Müller; A. Spohr; **J. Nielsen** (2000) Role of substrate concentration on mitosis and hyphal extension of *Aspergillus*. *Biotechnol. Bioeng.* 67:390-397
- 99. H. Pedersen; B. Christensen; C. Hjort; J. Nielsen (2000) Construction and characterization of an oxalic acid non-producing strain of *Aspergillus niger*. *Metabol. Eng.* 2:34-41
- 100. H. Pedersen; M. Beyer; **J. Nielsen** (2000) Glucoamylase production in batch, chemostat and fed-batch cultivations by an industrial strain of *Aspergillus niger*. *Appl. Microbiol. Biotechnol.* 53:272-277
- 101. S. Ostergaard; C. Roca; B. Rønnow; J. Nielsen; L. Olsson (2000) Physiological studies in aerobic batch cultivations of *Saccharomyces cerevisiae* strains harbouring the *MEL1* gene. *Biotechnol. Bioeng.* 68:252-259
- 102. H. Pedersen; C. Hjort; J. Nielsen (2000) Cloning and characterization of *oah*, the gene encoding oxaloacetate hydrolase from *Aspergillus niger*. *Mol. Gen. Genetics*. 263:281-286
- 103. L. Olsson; **J. Nielsen** (2000) The role of metabolic engineering in the improvement of *Saccharomyces cerevisiae*: Utilization of industrial media. *Enz. Microbiol. Technol.* 26:785-792
- 104. T. L. Nissen; C. W. Hamann; M. C. Kielland-Brandt; J. Nielsen; J. Villadsen (2000) Anaerobic and batch cultivations of *Saccharomyces cerevisiae* mutants impaired in glycerol synthesis. *Yeast* 16:463-474
- 105. T. L. Nissen; M. C. Kielland-Brandt; J. Nielsen; J. Villadsen (2000) Optimisation of ethanol production in Sacchaoromyces cerevisiae by metabolic engineering of the ammonia assimilation. Metabol. Eng. 2:69-77
- 106. A. Heydorn, B. O. Pedersen, J. Ø. Duus, S. Bergmann, T. Suhr-Jessen, J. Nielsen (2000) Biosynthetic studies of the glycopeptide teicoplanin by 1H and 13C NMR. J. Biol. Chem. 275:6201-6206
- 107. J. P. van Dijken et al. (2000) An inter-laboratory comparison of physiological and genetic properties of four Saccharomyces cerevisiae strains. Enz. Microb. Technol. 26:706-714
- 108. T. C. Zangirolami; M. Carlsen; J. Nielsen; S. B. Jørgensen (2000) Selection and characterization of a high α-amylase-producing variant in glucose-limited continuous cultures of *Aspergillus oryzae*. *Myc. Res.* 104:1241-1249
- 109. B. Christensen; J. Nielsen (2000) Metabolic network analysis on *Penicillium chrysogenum* using 13C-labelled glucose. *Biotechnol. Bioeng.* 68:652-659
- B. Christensen; J. Thykær; J. Nielsen (2000) Metabolic characterization of high and low yielding strains of *Penicillium chrysogenum*. Appl. Microbiol. Biotechnol. 54:212-217
- 111. **J. Nielsen** (2000) The role of metabolic engineering in the improvement of industrial processes. *Thai J. Biotechnol.* 2:14-25
- 112. H. P. Smits; J. Hauf; S. Müller; T. J. Hobley; F. K. Zimmermann; B. Hahn-Hägerdal; **J. Nielsen**; L. Olsson (2000) Simultaneous over-expression of enzymes of the lower part of glycolysis can enhance the fermentative capacity of *Saccharomyces cerevisiae*. *Yeast* 16:1325-1334
- 113. L. F. Bautista; M. Hentzer; A. Santerre Henriksen; A. Aleksenko; **J. Nielsen** (2000) Antisense silencing of the *creA* gene in *Aspergillus nidulans*. *Appl. Environ. Microbiol*. 66:4579-4581
- 114. S. Ostergaard; L. Olsson; M. Johnston; J. Nielsen (2000) Increasing galactose consumption by *Saccharomyces cerevisiae* through metabolic engineering of the *GAL* gene regulatory network. *Nature Biotechnol.* 18:1283-1286

- 115. T. Agger; A. Spohr; **J. Nielsen** (2001) α-Amylase production in high cell density submerged cultivations of *Aspergillus oryzae* and *Aspergillus nidulans. Appl. Microbiol. Biotechnol.* 55:81-84
- 116. H. Aae Theilgaard; M. van den Berg; C. Mulder; R. A. L. Bovenberg; **J. Nielsen** (2001) Quantitative analysis of *Penicillium chrysogenum* Wis54-1255 transformants over-expressed in the penicillin biosynthetic genes. *Biotechnol. Bioeng.* 72:379-388
- 117. E. Jonsbu; T. Elingsen; J. Nielsen (2001) Effects of nitrogen sources on cell growth and production of nystatin by Streptomyces noursei J. Antibiot. 53:1354-1362
- 118. M. McIntyre; J. Dynesen; J. Nielsen (2001) Morphological characaterization of *Aspergillus nidulans*: Growth, septation and fragmentation. *Microbiol*. 147:239-246
- 119. A. K. Gombert; M. M. dos Santos; B. Christensen; J. Nielsen (2001) Network identification and flux quantification in the central metabolism of *Saccharomyces cerevisiae* at different conditions of glucose repression. *J. Bacteriol*. 183:1441-1451
- 120. S. Ostergaard; L. Olsson; **J. Nielsen** (2001) *In vivo* dynamics of galactose metabolism in *Saccharomyces cerevisiae*: Metabolic fluxes and metabolite levels. *Biotechnol. Bioeng.* 73:412-425
- 121. S. Ostergaard; K. O. Walløe; C. S. G. Gomes; L. Olsson; **J. Nielsen** (2001) The impact of *GAL6*, *GAL80*, and *MIG1* on glucose control of the *GAL* system in *Saccharomyces cerevisiae*. *FEMS Yeast Research*. 1:47-55
- 122. T. L. Nissen; M. Anderlund; M. C. Kielland-Brandt; J. Nielsen; J. Villadsen (2001) Expression of a cytoplasmic transhydrogenase in *Saccharomyces cerevisiae* results in formation of 2-oxoglutarate due to depletion of the NADPH pool. *Yeast* 18:19-32
- 123. K. Møller; L. D. Tidemand; J. R. Winther; L. Olsson; J. Piskur; J. Nielsen (2001) Production of a heterologous proteinase A by Saccharomyces kluyveri. Appl. Microbiol. Biotechnol. 57:216-219
- 124. B. Christensen; T. Christiansen; A. K. Gombert; J. Thykær; **J. Nielsen** (2001) Simple and robust method for estimation of the split between the oxidative pentose phosphate pathway and the Embden-Meyerhof Parnas pathway in microorganisms. *Biotechnol. Bioeng.* 74:517-523
- 125. E. Jonsbu; B. Christensen; **J. Nielsen** (2001) Changes in *in vivo* fluxes through central metabolic pathways during production of nystatin by *Streptomyces noursei* in batch culture. *Appl. Microbiol. Biotechnol.* 56:93-100
- 126. M. Carlsen; J. Nielsen (2001) Influence of carbon source on α-amylase production by Aspergillus oryzae. Appl. Microbiol. Biotechnol. 57:346-349
- 127. J. Robin; M. Jacobsen; M. Beyer; H. Noorman; **J. Nielsen** (2001) Physiological characterisation of *Penicillium chrysogenum* strains expressing the expandase gene from *Streptomyces clavuligerus* during batch cultivations. Growth and adipoyl-7-ADCA production. *Appl. Microbiol. Biotechnol.* 57:357-362
- 128. T. C. Zangirolami; M. Carlsen; J. Nielsen; S. B. Jørgensen (2002) Growth and enzyme production during continuous cultures of a high amylase-producing variant of *Aspergillus oryzae*. *Brazilian J. Chem. Eng.* 19:55-68

- 129. T. Agger; J. B. Petersen; S. M. O'Connor; R. L. Murphy; J. M. Kelly; **J. Nielsen** (2002) Physiological characterisation of recombinant *Aspergillus nidulans* strains with different *creA* genotypes expressing *A. oryzae* α-amylase. *J. Biotechnol.* 92:279-285
- 130. K. Møller; B. Christensen; J. Förster; J. Piskur; J. Nielsen; L. Olsson (2002) Aerobic glucose metabolism of Saccharomyces kluyveri: Growth, metabolite production and quantification of metabolic fluxes. Biotechnol. Bioeng. 77:186-193
- 131. T. Christiansen; B. Christensen; **J. Nielsen** (2002) Metabolic network analysis of *Bacillus clausii* on minimal and semirich medium using ¹³C-labeled glucose. *Metabolic Eng.* 4:159-169
- 132. J. Thykaer; B. Christensen; J. Nielsen (2002) Metabolic network analysis of an adipoyl-7-ADCA producing strain of *Penicillium chrysogenum*: Elucidation of adipate degradation. *Metabolic Eng.* 4:151-158
- 133. M. McIntyre; J. Breum; J. Arnau; J. Nielsen (2002) Growth physiology and dimorphism of *Mucor circinelloides* (syn. *racemosus*) during submerged batch cultivation. *Appl. Microbiol. Biotechnol.* 58:495-502
- 134. T. Christiansen; J. Nielsen (2002) Growth energetics of an alkaline serine protease producing strain of *Bacillus clausii* during continuous cultivation. *Bioproc. Biosystems Eng.* 24:329-339
- 135. B. Christensen; **J. Nielsen** (2002) Reciprocal ¹³C-labeling: A method for investigating the catabolism of co-substrates. *Biotechnol. Prog.* 18:163-166
- 136. C. Müller; M. McIntyre; K. Hansen; **J. Nielsen** (2002) Metabolic engineering of the morphology of *Aspergillus oryzae* by altering chitin synthesis. *Appl. Environ. Microbiol.* 68:1827-1836
- 137. A. G. Vara; A. Hochkoeppler; **J. Nielsen**; J. Villadsen (2002) Production of teicoplanin by *Actinoplanes teichomyceticus* in continuous fermentation. *Biotechnol. Bioeng.* 77:589-598
- 138. **J. Nielsen**; L. Olsson (2002) An expanded role for microbial physiology in metabolic engineering and functional genomics: Moving towards systems biology. *FEMS Yeast Res.* 2:175-181
- 139. N. B. S. Jensen; B. Christensen; **J. Nielsen**; J. Villadsen (2002) The simultaneous biosynthesis and uptake of amino acids by *Lactococcus lactis* studied by ¹³C labelling experiments. *Biotechnol. Bioeng.* 78:11-16
- 140. K. Møller; C. Bro; J. Piskur; J. Nielsen; L. Olsson (2002) Steady-state and transient-state analysis of aerobic fermentation in *Saccharomyces kluyveri*. *FEMS Yeast Res.* 2:233-244
- 141. T. Christiansen; J. Nielsen (2002) Production of extracellular protease and glucose uptake in *Bacillus clausii* in steady state and transient continuous cultures. *J. Biotechnol*. 97:265-273
- 142. J. Förster; A. K. Gombert; J. Nielsen (2002) A functional genomics approach using metabolomics and *in silico* pathway analysis. *Biotechnol. Bioengi* 79:703-712
- 143. J. Zaldivar; A. Borges; B. Johansson; H. P. Smits; S. G. Villas-Boas; J. Nielsen; L. Olsson (2002) Fermentation performance and intracellular metabolite patterns in laboratory and industrial xylose fermenting *Saccharomyces cerevisiae*. Appl. Microbiol. Biotechnol. 59:436-442

- 144. C. Müller; C. M. Hjort; K. Hansen; **J. Nielsen** (2002) Altering the expression of two chitin synthase genes differentially affects the growth and morphology of *Aspergillus oryzae*. *Microbiology* 148:4025-4033
- 145. B. Christensen; A. K. Gombert; **J. Nielsen** (2002) Analysis of flux estimates based on ¹³C-labeling experiments. *Eur. J. Biochem.* 269:2795-2800
- 146. E. Jonsbu; M. McIntyre; **J. Nielsen** (2002) The influence of carbon sources and morphology on nystatin production by *Streptomyces noursei*. *J. Biotechnol*. 95:133-144
- 147. M. D. W. Piper; P. Daran-Lapujade; C. Bro; B. Regenberg; S. Knudsen; J. Nielsen; J. T. Pronk (2002) Reproducibility of oligonucleotide microarray transcriptome analysis: an interlaboratory comparison using chemostat cultures of Saccharomyces cerevisiae. J. Biol. Chem. 277:37001-37008

- 148. C. Müller; K. Hansen; P. Szabo; **J. Nielsen** (2003) Effect of deletion of chitin synthesis on mycelial morphology and culture viscosity in *Aspergillus oryzae*. *Biotechnol. Bioeng.* 81:525-534
- 149. T. Christiansen; S. Michaelsen; M. Wümpelmann: J. Nielsen (2003) Production of Savinase and population viability of *Bacillus clausii* during high cell density fed-batch cultivations. *Biotechnol. Bioeng.* 83:344-352
- 150. J. Robin; G. Lettier; M. McIntyre; H. Noorman; **J. Nielsen** (2003) Continuous cultivations of *Penicillium chrysogenum* strain expressing the expandase gene from *Streptomyces clavuligerus*: Growth yields and morphological characterisation. *Biotechnol. Bioeng.* 83:353-360
- 151. J. Robin; P. Bruheim; M. L. Nielsen; H. Noorman; J. Nielsen (2003) Continuous cultivations of *Penicillium chrysogenum* strain expressing the expandase gene from *Streptomyces clavuligerus*: Kinetics of adipoyl-7-aminodeacetoxycephalosporanic acid and by-product formation. *Biotechnol. Bioeng.* 83:361-368
- 152. J. Förster; I. Famili; P. Fu; B. Ø. Palsson; J. Nielsen (2003) Genome-scale reconstruction of the *Saccharomyces cerevisiae* metabolic network. *Genome Res.* 13:244-253
- 153. T. L. Lübbehüsen; J. Nielsen; M. McIntyre (2003) Characterization of the *Mucor circinelloides* life cycle by on-line image analysis. *J. Appl. Microbiol.* 95:1152-1160
- 154. J. Robin; S. Bonneau; D. Schipper; H. Noorman; **J. Nielsen** (2003) Influence of the adipate and dissolved oxygen concentrations on the β-lactams production during continuous cultivations of a *Penicillium chrysogenum* strain expressing the expandase gene from *Streptomyces clavuligerus*. *Metabol*. *Eng*. 5:42-48
- 155. N. Gunnarsson; P. Bruheim; J. Nielsen (2003) Production of the glycopeptide antibiotic A40926 by *Nonomuria* sp. ATCC 39727: influence of medium composition in batch fermentation. *J. Ind. Microbiol. Biotechnol.* 30:150-156
- 156. J. Dynesen; **J. Nielsen** (2003) Surface hydrophobicity of *Aspergillus nidulans* conidiophores and its role in pellet formation. *Biotechnol. Prog.* 19:1049-1052
- 157. M. M. dos Santos; A. K. Gombert; B. Christensen; L. Olsson; **J. Nielsen** (2003) Identification of *in vivo* enzyme activities in the co-metabolism of glucose and acetate by *Saccharomyces cerevisiae* using ¹³C-labeled substrates. *Eukar. Cell.* 2:599-608
- 158. J. Förster; I. Famili; B. Ø. Palsson; J. Nielsen (2003) Large-scale Evaluation of *in silico* Gene Deletions in *Saccharomyces cerevisiae*. *Omics J. Integrative Biol*. 7:193-202
- 159. T. L. Lübbehüsen; J. Nielsen; M. McIntyre (2003) Morphology and physiology of the dimorphic fungus *Mucor circinelloides* (syn. *M. racemosus*) during anaerobic growth. *Mycol. Res.* 107:223-230
- 160. C. Roca; **J. Nielsen**; L. Olsson (2003) Metabolic engineering of ammonium assimilation in xylose fermenting *Saccharomyces cerevisiae* improves ethanol production. *Appl. Env. Microbiol.* 69:4732-4736
- 161. J. Nielsen (2003) It is all about metabolic fluxes. J. Bacteriol., 185:7031-7035
- 162. C. Bro; B. Regenberg; G. Lagniel; J. Labarre; M. Montero-Lomeli; J. Nielsen (2003) Transcriptional, proteomics and metabolic response to lithium in galactose-grown yeast cells. J. Biol. Chem. 278:32141-32149
- 163. W. Prathumpai; J. B. Gabelgaard; P. Wanchanthuek; P. J. I. van de Vondervoort; M. J. L. de Groot; M. McIntyre; J. Nielsen (2003) Metabolic control analysis of xylose catabolism in *Aspergillus*. *Biotechnol*. *Prog*. 19:1136-1141
- 164. H. David; M. Åkesson; **J. Nielsen** (2003) Reconstruction of the central carbon metabolism of *Aspergillus niger*. *Eur. J. Biochem.* 270: 4243-4253
- 165. M. M. dos Santos; G. Thygesen; P. Kötter; L. Olsson; J. Nielsen (2003) Aerobic physiology of redox-engineered Saccharomyces cerevisiae strains modified in the ammonium assimilation for increased NADPH availability. FEMS Yeast Res. 4:59-68
- 166. I. Famili; J. Förster; **J. Nielsen**; B. Palsson (2003) *Saccharomyces cerevisiae* phenotypes can be predicted using constraint based analysis of a genome-scale reconscructed metabolic network. *Proc. Nat. Acad, Sci.* 100:13134-13139
- 167. J. Dynesen; J. Nielsen (2003) Branching is coordinated with mitosis in growing hyphae of *Aspergillus nidulans*. Fungal Gen. Biol. 40:15-24
- 168. S. G. Villas-Boas; D. G. Delicado; M. Åkesson; **J. Nielsen** (2003) Simultaneous analysis of amino and nonamino organic acids as methyl chloroformate derivatives using gas chromatography-mass spectrometry. *Anal. Biochem.* 322:134-138

- 169. N. Gunnarson; P. Bruheim; J. Nielsen (2004) Glucose metabolism in the antibiotic producing Actinomycete Nonomuiaea sp. Biotechnol. Bioeng. 88: 652-663
- 170. T. L. Lübbehüsen, **J. Nielsen**, M. McIntyre (2004) Aerobic and anaerobic ethanol production by *Mucor circinelloides* during submerged growth. *Appl. Microbiol. Biotechnol.* 63:543-548
- 171. T. L. Lübbehüsen, V. G. Polo, S. Rossi, **J. Nielsen**, S. Moreno, M. McIntyre, J. Arnau (2004) Protein kinase A is involved in the control of morphology and branching during aerobic growth in *Mucor circinelloides*. *Microbiol*. 150:143-150
- 172. C. Bro; B. Regenberg; **J. Nielsen** (2004) Genome-wide transcriptional response of a *Saccharomyces cerevisiae* strain with an altered redox metabolism. *Biotechnol. Bioeng.* 85:269-276

- 173. W. Prathumpai; M. McIntyre; **J. Nielsen** (2004) The effect of CreA in glucose and xylose catabolism in *Aspergillus nidulans*. *Appl. Microbiol. Biotechnol.* 63:748-753
- 174. C. Bro; J. Nielsen (2004) Impact of 'Ome' analysis on inverse metabolic engineering. Met. Eng. 6:204-211
- 175. K. Møller; C. Bro; R. B. Langkjær; J. Piskur; J. Nielsen; L. Olsson (2004) Pyruvate decarboxylases from the petitenegative yeast Saccharomyces kluyveri. Mol. Genet. Genomics 270:558-568
- 176. T. Grotkjær; M. Åkesson; B. Christensen; A. K. Gombert; **J. Nielsen** (2004) Impact of transamination reactions and protein turnover on labeling dynamics in ¹³C-labelling experiments. *Biotechnol. Bioeng.* 86:209-216
- 177. M. Åkesson; J. Förster; **J. Nielsen** (2004) Integration of gene expression data into genome-scale metabolic models. *Met. Eng*, 6:285-293
- 178. G. G. Larsen; K. F. Appel; A.-M. Wolff, J. Nielsen; J. Arnau (2004) Characterization of the *Mucor circinelloides* regulated promoter gpd1P and construction of a constitutive derivative. *Curr. Genet.* 45:225-234
- 179. N. Gunnarson; U. H. Mortensen; M. Sosio; **J. Nielsen** (2004) Identification of the Entner-Doudoroff pathway in an antibiotic producing actinomycete species. *Mol Microbiol* 52: 895-902
- 180. N. Eckert-Boulet; P. S. Nielsen; C. Friis; M. M. dos Santos; **J. Nielsen**; M. C. Kielland-Brandt; B. Regenberg (2004) Transcriptional profiling of extracellular amino acid sensing in *Saccharomyces cerevisiae* and the role of Stp1p and Stp2p. *Yeast* 21: 635-648
- 181. S. L. Westergaard; C. Bro; L. Olsson; **J. Nielsen** (2004) Elucidation of the role of Grr1p on glucose sensing by *Saccharomyces cerevisiae* through genome-wide transcription analysis. *FEMS Yeast Res.* 5:193-204
- 182. V. Raghevendran; A. K. Gombert; B. Christensen; P. Kötter; **J. Nielsen** (2004) Phenotypic characterization of glucose repression mutants of *Saccharomyces cerevisiae* using experiments with ¹³C-labelled glucose. *Yeast* 21:769-779
- 183. T. Grotkjær; J. Nielsen (2004) Enhancing yeast transcription analysis through integration of heterogeneous data. *Cur. Genomics* 5:673-686
- 184. M. M. dos Santos; V. Raghevendran; P. Kötter; L. Olsson; **J. Nielsen** (2004) Manipulation of the malic enzyme in *Saccharomyces cerevisiae* for increasing NADPH production capacity aerobically in different cellular compartments. *Met. Eng.* 6:352-363
- 185. B. Regenberg; U. Krühne; M. Beyer; L. H. Pedersen; M. Simón; O. R.T. Thomas; J. Nielsen; T. Ahl (2004) Use of laminar flow patterning for miniaturised biochemical assays. *Lab. Chip* 4:654-657
- 186. W. Prathumpai; S. J. Flitter; M. McIntyre; **J. Nielsen** (2004) Lipase production by recombinant strains of *Aspergillus niger* expressing a lipase-encoding gene from *Thermomyces lanuginosus*. *Appl. Microbiol. Biotechnol.* 65:714-719
- 187. C. Cannizzaro; B. Christensen; **J. Nielsen**; U. von Stockar (2004) Metabolic network analysis on *Phaffia rhodozyma* yeast using ¹³C-labeled glucose and gas chromatography-mass spectrometry. *Met. Eng.* 6:340-351

- 188. S. G. Villas-Bôas; J. H. Pedersen; M. Åkesson; J. Smedsgaard; J. Nielsen (2005) Global metabolite analysis of yeast: evaluation of sample preparation methods. *Yeast* 22:1155-1169
- 189. G. Panagiotou, S. G. Villas-Boas, P. Christakopoulos, **J. Nielsen**, L. Olsson (2005) Intracellular metabolite profiling of *Fusarium oxysporum* converting glucose to ethanol. *J. Biotechnol*. 115:425-434
- 190. J. Smedsgaard, **J. Nielsen** (2005) Metabolite profiling of fungi and yeast: from phenotype to metabolome by MS and informatics. *J. Exp. Bot.* 56:273-286
- K. R. Patil, J. Nielsen (2005) Uncovering transcriptional regulation of metabolism by using metabolic network topology. Proc. Nat. Acad. Sci. 102:2685-2689
- 192. I. Borodina, C. Schöller, A. Eliasson, **J. Nielsen** (2005) Metabolic network analysis of *Streptomyces tenebrarius*, a *Streptomyces* with Entner-Doudoroff pathway. *Appl. Environ. Microbiol.* 71:2294-2302
- 193. S. G. Villas-Bôas; M. Åkesson; **J. Nielsen** (2005) Biosynthesis of glyoxylate from glycine in *Saccharomyces cerevisiae*. *FEMS Yeast Res.* 5:703-709
- 194. S. G. Villas-Bôas; J. F. Moxley; M. Åkesson; G. Stephanopoulos; J. Nielsen (2005) High-throughput metabolic state analysis: The missing link in integrated functional genomics of yeasts. *Biochem. J.* 388:669-677
- 195. N. Eckert- Boulet; B. Regenberg; **J. Nielsen** (2005) Grr1p is required for transcriptional indusction of amino acid permease genes and proper transcriptional regulation of genes in carbon metabolism of *Saccharomyces cerevisiae*. *Curr. Genet.* **47**:139-149
- 196. T. Seker; K. Møller; **J. Nielsen** (2005) Analysis of acyl CoA ester intermediates of the mevalonate pathway in *Saccharomyces cerevisiae*. *Appl. Microbiol. Biotechnol.* 67:199-124
- 197. I. Borodina; P. Krabben; **J. Nielsen** (2005) Genome-scale analysis of *Streptomyces coelicolor* A3(2) metabolism. *Genome Res.* 15:820-829
- 198. A. P. Oliveira; **J. Nielsen**; J. Förster (2005) Modeling *Lactococcus lactis* using a genome-scale flux model. *BMC Microbiol*, 5:39
- 199. C. Bro; S. Knudsen; B. Regenberg; L. Olsson; **J. Nielsen** (2005) Improvement of galactose uptake in *Saccharomyces cerevisiae* through overexpression of phosphoglucomutase: Example of transcript analysis as a tool in inverse metabolic engineering. *Appl. Environ. Microbiol.*, 71:6465-6472
- 200. H. David; A. M. Krogh; C. Roca; M. Åkesson; **J. Nielsen** (2005) CreA influences the metabolic fluxes of *Aspergillus nidulans* during growth on glucose and xylose. *Microbiol.* 151:2209-2221
- 201. V. Raghevendran; J. Nielsen; L. Olsson (2005) Teaching microbial physiology using glucose repression phenomenon in Baker's yeast as an example. *Biochem. Mol. Biol. Edu.* 33:404-410
- 202. M. C. Jewett; A. P. Oliveira; K. R. Patil; J. Nielsen (2005) The role of high-throughput transcriptome analysis in metabolic engineering. *Biotechnol. Bioproc. Eng.* 10:385-399
- 203. T. Grotkjær; P. Christakopoulos; J. Nielsen; L. Olsson (2005) Comparative metabolic network analysis of two xylose fermenting recombinant Saccharomyces cerevisiae strains. Met. Eng. 7:437-444

204. K. R. Patil; I. Rocha; J. Förster; **J. Nielsen** (2005) Evolutionary programming as a platform for *in silico* metabolic engineering. *BMC Bioinformatics* 6:308

2006

- 205. T. Cakir; K. R. Patil; Z. I. Önsan; K. Ö. Ülgen; B. Kirdar; J. Nielsen (2006) Integration of metabolome data with metabolic networks reveals reporter reactions. *Mol. Systems Biol.* 2:50
- 206. T. Grotkjær; O. Winther; B. Regenberg; J. Nielsen; L. K. Hansen (2006) Robust multi-scale clustering of large DNA microarray datasets with the consensus algorithm. *Bioinformatics* 22:58-67
- 207. N. Eckert-Boulet; K. Larsson; B. Wu; P. Poulsen; B. Regenberg; **J. Nielsen**; M. Kielland-Brandt (2006) Deletion of *RTS1*, encoding a regulatory subunit of protein phosphatase 2A, results in constitutive amino acid signaling via increased Stp1p processing. *Eukaryotic Cell* 5:174-179
- 208. C. Bro; B. Regenberg; J. Förster; **J. Nielsen** (2006) *In silico* aided metabolic engineering of *Saccharomyces cerevisiae* for improved bioethanol production. *Met. Eng.* 8:102-111
- 209. R. Mustachi; S. Hohmann; J. Nielsen (2006) Yeast systems biology to unravel the network of life. Yeast 23:227-238
- 210. J. Mogensen; H. B. Nielsen; G. Hofmann; J. Nielsen (2006) Transcription analysis using high-density micro-arrays of *Aspergillus nidulans* wild-type and D*creA* mutant during growth on glucose and ethanol. *Fung. Gen. Bio.* 43:593-603
- 211. V. Raghevendran; K. R. Patil; L. Olsson; J. Nielsen (2006) Hap4 is not essential for activation of respiration at low specific growth rates in Saccharomyces cerevisiae. J. Biol. Chem. 281:12308-12314
- 212. R. Usaite; K. R. Patil; T. Grotkjær; J. Nielsen; B. Regenberg (2006) Globl transcriptional and physiological responses of Saccharomyces cerevisiae to ammoniu, L-alanine or L-glutamine limitation. Appl. Environ. Microbiol. 72:6194-6203
- 213. A. Diano; S. Bekker-Jensen; J. Dynesen; J. Nielsen (2006) Polyol synthesis in *Aspergillus niger*: Influence of oxygen availability, carbon and nitrogen sources on the metabolism. *Biotechnol. Bioeng.* 94:899-908
- 214. B. Regenberg; T. Grotkjær; O. Winther; A. Fausbøll; M. Åkesson; C. Bro; L. K. Hansen; S. Brunak; J. Nielsen (2006) Growth-rate regulated genes have profound impact on interpretation of transcriptome profiling in *Saccharomyces cerevisiae*. Genome Biol. 7:R107
- 215. H. David; G. Hofmann; A. P. Oliveira; H. Jarmer; J. Nielsen (2006) Metabolic network driven analysis of genome-wide transcription data from *Aspergillus nidulans*. *Genome Biol*. 7:R108

2007

- 216. M. J. van der Werf; R. Takors; J. Smedsgaard; J. Nielsen; T. Ferenci; J. C. Portais; C. Wittmann; M. Hooks; A. Tomassini; M. Oldiges; J. Fostel; U. Sauer (2007) Standard reporting requirements for biological samples in metabolomics experiments: Microbial and in vitro biology experiments. *Metabolomics* 3:189-194
- 217. S. L. Westergaard; A. P. Oliveira; C. Bro; L. Olsson; J. Nielsen (2007) A systems biology approach to study glucose repression in the yeast *Saccharomyces cerevisiae*. *Biotechnol*. *Bioeng*, 96:134-145
- 218. G. N. Vemuri; M. A. Eiteman; J. E. McEwen; L. Olsson; J. Nielsen (2007) Increasing NADH oxidation reduces overflow metabolism in *Saccharomyces cerevisiae*. *Proc. Nat. Acad. Sci.* 104:2402-2407
- 219. H. J. Pel, J.H. de Winde, D.B. Archer, P.S. Dyer, G. Hofmann, P.J. Schaap, G. Turner, R.P. de Vries, R. Albang, K. Albermann, M.R. Andersen, J.D. Bendtsen, J.A.E. Benen, M. van den Berg, S. Breestraat, M.X. Caddick, R. Contreras, M. Cornell, P.M. Coutinho, E.G.J. Danchin, A.J.M. Debets, P. Dekker, P.W.M. van Dijck, A. van Dijk, L. Dijkhuizen, A.J.M. Driessen, C. d'Enfert, S. Geysens, C. Goosen, G.S.P. Groot, P.W.J. de Groot, T. Guillemette, B. Henrissat, M. Herweijer, J.P.T.W. van den Hombergh, C.A.M.J.J. van den Hondel, R.T.J.M. van der Heijden, R.M. van der Kaaij, F.M. Klis, H.J. Kools, C.P. Kubicek, P.A. van Kuyk, J. Lauber, X. Lu, M.J.E.C. van der Maarel, R. Meulenberg, H. Menke, M.A. Mortimer, J. Nielsen, S.G. Oliver, M. Olsthoorn, K. Pal, N.N.M.E. van Peij, A.F.J. Ram, U. Rinas, J.A. Roubos, C.M.J. Sagt, M. Schmoll, J. Sun, D. Ussery, J. Varga, W. Vervecken, P.J.J. van de Vondervoort, H. Wedler, H.A.B.Wösten, A.-P. Zeng, A.J.J. van Ooyen, J. Visser and H. Stam (2007) Genome sequencing and analysis of the versatile cell factory *Aspergillus niger* CBS513.88. *Nature Biotech*. 25:221-231
- 220. S. Mass; S. G. Villas-Boas; M. E. Hansen; M. Ålesson; J. Nielsen (2007) A comparison of direct infusion MS and GC-MS for metabolic footprinting of yeast mutants. *Biotechnol. Bioeng.* 96:1014-1022
- 221. S. Wattanachaisaereekul; A. Eliasson Lantz; M. L. Nielsen; O. S. Andresson; J. Nielsen (2007) Optimization of heterologous production of the polyketide 6-MSA in *Saccharomyces cerevisiae*. *Biotechnol. Bioeng*. 97:893-900
- 222. C. J. Baxter; H. Redestig; N. Schauer; D.Repsilber; K. R. Patil; J. Nielsen; J. Selbig; J. Liu; A. R. Fernie; L. J. Sweetlove (2007) The metabolic response of heterotrophic *Arabidopsis* cells to oxidative stress. *Plant Phys.* 143:312-325
- 223. K. Y. Arga; Z. I. Onsan; B. Kirdar; K. O. Ulgen; J. Nielsen (2007) Understanding signaling in yeast: Insights from network analysis. *Biotechnol. Bioeng*, 97:1246-1258
- 224. T. Cakir; B. Kirdar; Z. I. Onsan; K. O. Ulgen; J. Nielsen (2007) Effect of carbon source pertubations on transcriptional regulation of metabolic fluxes in *Saccharomyces cierevisiae*. *BMC Systems Biol*. 1:18
- 225. J. Thykaer; **J. Nielsen** (2007) Evidence, through C13-labelling analysis, of phosphoketolase activity in fungi. *Proc. Biochem.* 42:1050-1055
- 226. S. Meier; G. Panagiotou; L. Olsson; **J. Nielsen** (2007) Physiological characterization of xylose metabolism in *Aspergillus niger* under oxygen limited conditions. *Biotechnol. Bioeng.* 98:462-475
- 227. I. Nookaew; A. Meechai; C. Thammarongtham; K. Loateng; V. Ruanglek; S. Cheevadhanarak; **J. Nielsen**; S. Bhumiratana (2007) Identification of flux regulation coefficients from elementary flux modes: A systems biology tool for analysis of metabolic networks. *Biotechnol. Bioeng.* **97**:1535-1549
- 228. M. L. Nielsen; W. A. de Jongh; S. L. Meijer; J. Nielsen; U. Mortensen (2007) Transient marker system for iterative gene targeting of a prototrophic fungus. *Appl. Environ. Microbiol.* 73:7240-7245

- 229. J. Maury; M. A. Asadollahi; K. Møller; M. Schalk; A. Clark; L. R. Formenti; **J. Nielsen** (2008) Reconstruction of a bacterial isoprenoid biosynthetic pathway in *Saccharomyces cerevisiae*. *FEBS Let*. 582:4032-4038
- 230. R. Usaite; J. Nielsen; L. Olsson (2008) Physiological characterization of glucose repression in the strains with SNF1 and SNF4 genes deleted. J. Biotechnol. 133:73-81
- 231. R. Usaite; J. Wohlschlegel; J. D. Venable; S. K. Park; J. Nielsen; L. Olsson John R. Yates III (2008) Characterization of global yeast quantitative proteome data generated from the wild type and glucose repression *Saccharomyces cerevisiae* strains: the comparison of two quantitative algorithms. *J. Proteom Res.* 7:266-275
- 232. W. de Jongh; J. Nielsen (2007) Enhanced citrate production through gene insertion in *Aspergillus niger. Met. Eng.* 10:87-96
- 233. M. Asadollahi; J. Maury; K. Møller; K. F. Nielsen; M. Schalk; A. Clark; **J. Nielsen** (2008) Production of plant sesquiterpenes in *Saccharomyces cerevisiae*: Effect of *ERG9* repression on sesquiterpene biosynthesis. *Biotechnol. Bioeng.* 99:666-677
- 234. M. R. Andersen; M. L. Nielsen; J. Nielsen (2008) Metabolic model integration of the bibliome, genome, metabolome and reactome of *Aspergillus niger*. Mol. Systems Biol. 4:178
- 235. A. P. Oliveira; K. Patil; J. Nielsen (2008) Architecture of transcriptional regulatory circuits is knitted over the topology of bio-molecular interaction networks. BMC Systems Biol. 2:17
- 236. J. Thykaer; K. Rueksomtawin; H. Noorman; **J. Nielsen** (2008) NADPH-Dependent glutamate dehydrogenase in *Penicillium chrysogenum* is involved in regulation of b-lactam production. *Microbiol.* 154:1242-1250
- 237. H. David; I. Ozcelik; G. Hofmann; **J. Nielsen** (2008) Analysis of *Aspergillus nidulans* metabolism at the genome-scale. *BMC Genomics* 9:163
- 238. W. A. de Jongh; C. Bro; S. Ostergaard; B. Regenberg; L. Olsson; J. Nielsen (2008) The roles of galactitol, galactose-1-phosphate and phosphoglucomutase in galactose-induced toxicity in *Saccharomyces cerevisiae*. *Biotechnol. Bioeng*. 101:317-326
- 239. G. Panagiotou; M. R. Andersen; T. Grotkjær; T. B. Regueira; G. Hofmann; **J. Nielsen**; L. Olsson (2008) Systems analysis unfolds the relationship between the phosphoketolase pathway and growth in *Aspergillus nidulans*. *PLoS One* 3:e2847
- 240. M. R. Andersen; W. Vongsangnak; G. Panagiotou; M. P. Salazar; L. Lehmann; **J. Nielsen** (2008) A trispecies *Aspergillus* microarray: Comparative transcriptomics of three *Aspergillus* species. *Proc. Nat. Acad. Sci.* 105:4387-4392
- 241. W. Vongsangnak; P. Olsen; K. Hansen; S. Krogsgaard; J. Nielsen (2008) Improved annotation through genome-scale metabolic modeling of *Aspergillus oryzae*. *BMC Genomics* 9:245
- 242. A. Fazio; M. C. Jewett; P. Daran-Lapujade; R. Mustacchi; R. Usaite; J. Pronk; C. Workman; J. Nielsen (2008) Transcription factor control of growth rate dependent genes in *Saccharomyces cerevisiae*: A three factor design. *BMC Genomics* 9:341
- 243. S. Wattanachaisaereekul; A. E. Lantz; M. L. Nielsen; J. Nielsen (2008) Production of the polyketide 6-MSA in yeast engineered for increased malonyl-CoA supply. *Metabolic Eng.* 10:246-254
- 244. I. Nookaew; M. C. Jewett; A. Meecha; C. Thammarongtham; K. Laoteng; S. Cheevadhanarak; J. Nielsen; S. Bhumiratana (2008) The genome-scale metabolic model iIN800 of Saccharomyces cerevisiae and its validation: a scaffold to query lipid metabolism. *BMC Sys. Biol.* 2:71
- 245. F. Pizarro; M. C. Jewett; J. Nielsen; E. Agosin (2008) Growth temperature exerts a differential physiological and transcriptional response in laboratory and wine strains of Saccharomyces cerevisiae. Appl. Environ. Microbiol. 74:6358-6368
- 246. I. Borodina; J. Siebring; J. Zhang; C. P. Smith; G. van Keulen; L. Dijkhuizen; J. Nielsen (2008) Antibiotic overproduction in *Streptomyces coelicolor* A3(2) mediated by phosphofructokinase deletion. *J. Biol. Chem.* 283:25186-25199
- 247. J. Højer-Pedersen; J. Smedsgaard; J. Nielsen (2008) The yeast metabolome addressed by electrospray ionization mass spectrometry: Initiation of a mass spectral library and its application for metabolic footprinting by direct infusion mass spectrometry. *Metabolomics* 4:393-405
- 248. M. J. Herrgard; N. Swainston; P. Dobson; W.B. Dunn; K.Y. Arga; M. Arvas; N. Blüthgen; S. Borger; R. Costenoble; M. Heinemann; M. Hucka; N. Ke NovereM P. Li; W. Liebermeister; M.L. Mo; A.P. Oliveira; D. Petranovic; S. Pettifer; E. Simeonidis; K. Smallbone; I. Spasic; D. Weichart; R. Brent; D. S. Broomhead; H.V. Westerhoff; B. Kirdar; M. Penttilä; E. Klipp; B.O. Palsson; U. Sauer; S.G. Oliver; P. Mendes; J. Nielsen; D. B. Kell (2008) A consensus yeast metabolic network obtained from a community approach to systems biology. *Nature Biotechnol*. 26:1155-1160
- 249. M. Rocha; P. Maia; R. Mendes; J. P. Pinto; E. C. Ferreira; K. Patil; **J. Nielsen**; I. Rocha (2008) Natural computation meta-heuristics for the *in silico* optimization of microbial strains. *BMC Bioinformatics* 9:499

- 250. M. Salazar; W. Vongsangnak; G. Panagiotou; M. R. Andersen; J. Nielsen (2009) Uncovering transcriptional regulation of glycerol metabolism in Aspergilli through genome-wide gene expression data analysis. *Mol. Genet. Genom.* 282:571-586
- 251. A. Diano; J. Peeters; J. Dynesen; J. Nielsen (2009) Physiology of *Aspergillus niger* in oxygen-limited continuous cultures: Influence of aeration, carbon source concentration and dilution rate. *Biotechnol. Bioeng.* 103:956-965
- 252. J. R. Wortman; J.M. Gilsenan; V. Joardar; J. Deegan; J. Clutterbuck; M.R. Andersen; D. Archer; M. Bencina; G. Braus; P. Coutinho; H. von Döhren; A.J. Driessen E. Espeso; E. Fekete; M. Flipphi; C.G. Estrada; S. Geysens; G. Goldman; P.W. de Groot; K. Hansen; S.D. Harris; T. Heinekamp; K. Helmstaedt; B. Henrissat; G. Hofmann; T. Homan; T. Horio; H. oriuchi; S. James; M. Jones, L. Karaffa; Z. Karanyi; M. Kato; N. Keller; D.E. Kelly; J.A. Kiel; J.M. Kim; I.J. van der Klei; F.M. Klis; A. Kovalchuk; N. Krasevec; C.P. Kubicek; B. Liu; A. Maccabe; V. Meyer; P. Mirabito; M. Miskei; M. Mos; J. Mullins; D.R. Nelson; J. Nielsen; B.R. Oakley; S.A. Osmani; T. Pakula; A. Paszewsko; I. Paulsen; S. Pilsyk; I. Pocsi; P.J. Punt; A.F. Ram; Q. Ren; X. Robellet; G. Robinson; B. Seiboth; P. van Solingen; T. Specht; J. Sun; N.

- Taheri-Talesh; N. Takeshita; D. Ussery; P.A van Kuyk; H. Visser; P.J. van der Vondervoort; R.P. de Vries; J. Walton; X. Xiang; Y. Xiong; A.P. Zeng; B.W. Brandt; M.J. Cornell; C.A. van den Hondel; J. Visser; S.G. Oliver; G. Turner (2009) The 2008 update of the *Aspergillus nidulans* genome annotation: A community effort. *Fungal Gen. Biol.* 46:S2-S13
- 253. K. R. Kjeldsen; J. Nielsen (2008) In silico genome-scale reconstruction and validation of the Corynebacterium glutamicum metabolic network. Biotechnol. Bioeng. 102:583-597
- 254. P. M. Coutinho; M.R. AndersenM K. Kolenova; P.A. van Kuyk; I. Benott; B.S. Gruben; B. Trejo-Aguilar; P. van Solingen; T. Pakula; B. Seiboth; E. Battaglia; G. Aguilar-Osorio; J.F. de Jong; R.A. Ohm; M. Aguilar; B. Henrissat; J. Nielsen; H. Stålbrandt; R.P. de Vries (2009) Post-genomic insights into the plant polysaccharide degradation potential of Aspergillus nidulans and comparison to Asperpergillus niger and Aspergillus oryzae. Fungal Gen. Biol. 46:S161-S169
- 255. G. Hofmann; A. Diano; J. Nielsen (2009) Recombinant bacterial haemoglobin alters metabolism of Aspergillus niger. Met. Eng. 11:8-12
- 256. J. Thykaer; K. Rueksomtawin; H. Noorman; J. Nielsen (2009) Disruption of the NADPH-dependent Glutamate Dehydrogenase Affects the Morphology of Two Industrial Strains of *Penicillium chrysogenum*. J. Biotechnol. 139:280-282
- 257. T. S. Christensen; A. P. Olivaira; J. Nielsen (2009) Reconstruction and logical modeling of glucose repression signaling pathways in Saccharomyces cerevisiae. BMC Systems Biol. 3:7
- 258. S. Meijer; J. Otero; M. R. Andersen; L Olsson; **J. Nielsen** (2009) Overexpression of isocitrate lyase glyoxylate byplass influence on metabolism in *Aspergillus niger*. *Metabolic Eng.* 11:107-116
- 259. D. Cimini; K. R. Patil; C. Shiraldi; J. Nielsen (2009) Global transcriptional response of *Saccharomyces cerevisiae* to the deletion of SDH3. *BMC Systems Biol.* 3:17
- 260. J. F. Moxley; M. C. Jewett; M. R. Antoniewicz; S. G. Villas-Boas; H. Alper; R. T. Wheeler; L. Tong; A. G. Hinnebusch; T. Ideker; J. Nielsen; G. Stephanopoulos (2009) Linking high-resolution metabolic flux phenotypes and transcriptional regulation in yeast modulated by the global regulator Gcn4p. *Proc. Nat. Acad. Sci.* 106:6477-6482
- 261. G. Panagiotou; M. R. Andersen; T. Grotkjær; T. B. Regueira; J. Nielsen; L. Olsson (2009) Studies of the production of fungal polyketides in *Aspergillus nidulans* by using systems biology tools. *Appl. Environ, Microbiol.* 75:2212-2220
- 262. S. Meijer; M. L. Nielsen; L. Olsson; J. Nielsen (2009) Gene deletion of cytosolic ATP: Citrate lyase leads to altered organic acid production in *Aspergillus niger*. J. Ind. Microbiol. Biotechnol. 36:1275-1280
- 263. M. R. Andersen; L. Lehmann; J. Nielsen (2009) Systemic analysis of the response of *Aspergillus niger* to ambient pH. *Genome Biol.* 10:R47
- 264. M. A. Asadollahi; J. Maury; K. R. Patil; M. Schalk; A. Clark; J. Nielsen (2009) Enhancing sesquiterpene production in *Saccharomyces cerevisiae* through *in silico* driven metabolic engineering. *Met. Eng.* 11;328-334
- 265. D. Rossouw; R. Olivares-Hernandes; J. Nielsen; F.F. Bauer (2009) A comparative 'omics' approach to investigate differences in wine yeast physiology and metabolism during fermentation. Appl. Environ. Microbiol. 75:6600-6612
- 266. S.Meijer; W.A.de Jongh; L. Olsson; J. Nielsen (2009) Physiological characterization of acuB deletion in Aspergillus niger. Appl. Microbiol. Biotechnol. 84:157-167
- 267. D. B. Flagfeldt; V. Siewers; L. Huang; J. Nielsen (2009) Characterization of chromosomal integration sites for Khooheterologous gene expression in Saccharomyces cerevisiae. Yeast 26:545-551
- V. Siewers; X. Chen; L. Huang; J. Zhang; J. Nielsen (2009) Heterologous production of the non-ribosomal peptide LLD-ACV in Saccharomyces cerevisiae. Met. Eng. 11:391-397
- 269. R. Usaite; M. C. Jewett; A. P. Oliveira; J. R. Yates III; L. Olsson; J. Nielsen (2009) Reconstruction of the yeast Snfl kinase regulatory network reveals its role as a global energy regulator. *Mol. Sys. Biol.* 5:319
- 270. W. Vongsangnak; M. Salazar; K. Hansen; J. Nielsen (2009) Genome-wide analysis of maltose utilization and regulation in aspergilla. *Microbiol.* 155:3893-3902

- 271. J.M. Otero; W. Vongsangak; M.A. Asadollahi; R. Olivares-Hernandes; J.Maury; L.Farinelli; L.Barlocher; M.Østerås; M.Schalk; A. Clark; J. Nielsen (2010) Whole genome sequencing of *Saccharomyces cerevisiae*: from genotype to phenotype for improved metabolic engineering applications. *BMC Genomics* 11:723
- 272. M. A. Asadollahi; J. Maury; M. Schalk; A. Clark; J. Nielsen (2010) Enhancement of farnesyl diphosphate pool as direct precursor of sesquiterpenes through metabolic engineering of the mevalonate pathway in *Saccharomyces cerevisiae*. *Biotechnol. Bioeng.* 106:86-96
- 273. K. van Eunen; J. Bouwman; P. Daran-Lapujade; J. Postmus; A. B. Canelas; F. I. Mensonides; R. Orij; I. Tuzun; J. van den Brink; G. J. Smits; W. M. van Gulik; S. Brul; J. J. Heijnen; J. H. de Winde; M. J. Teixeira de Mattos; C. Kettner; J. Nielsen; H. V. Westerhoff; B. M. Bakker (2010) Measuring enzyme activities under standardized in-vivo like conditions for systems biology. FEBS J. 277:749-760
- 274. W. Vongsangnak; I. Nookaew; M. Salazar; **J. Nielsen** (2010) Analysis of genome-wide co-expression and co-evolution of *Aspergillus oryzae* and *Aspergillus niger*. *Omics J.* 14:165-175
- S. Bordel; J. Nielsen (2010) Identification of flux control in metabolic networks using non-equilibrium thermodynamics. *Met. Eng.* 12:369-377
- 276. V. Siewers; R. San-Bento; **J. Nielsen** (2010) Implementation of communication-mediating domains for non-ribosomal peptide production in *Saccharomyces cerevisiae*. *Biotechnol. Bioeng*. 106:841-844
- 277. I. Rocha; P. M. P. Evangelista; P. Vilaça; S. Soares; J. P Pinto; **J. Nielsen**; K. R Patil; E. C. Ferreira; M. Rocha (2010) OptFlux: an open-source software platform for in silico metabolic engineering. *BMC Systems Biol.* 4:45
- 278. J. Thykaer; **J. Nielsen**; W. Wohlleben; T. Weber; M. Gutknecht; A. E. Lantz; E. Stegmann (2010) Increased glycopeptides production after overexpression of shikimate pathway genes being part of the balhimycin biosynthetic gene cluster. *Met. Eng.* 12:455-461

- 279. M. Cvijovic; R. Olivares-Hernandez; R. Agren; W. Vongsangnak; I. Nookaew; K. R. Patil; **J. Nielsen** (2010) BioMet Toolbox: Genome-wide analysis of metabolism. *Nuc. Acid Res.* 38:W144-W149
- 280. S. K. Nandy; P. Jouhten; J. Nielsen (2010) Reconstruction of the yeast protein-protein interaction network involved in nutrient sensing and global metabolic regulation. *BMC Sys. Biol.* 4:68
- 281. J. Zhang; L. Olsson; **J. Nielsen** (2010) The β-subunits of the Snf1 kinase in *Saccharomyces cerevisiae*, Gal83 and Sip2, but not Sip1, are redundant in glucose derepression and regulation of sterol biosynthesis. *Mol. Microbiol.* 77:371-383
- 282. S. Bordel; R. Agren; **J. Nielsen** (2010) Sampling the solution space in genome-scale metabolic networks reveals transcriptional regulation in key enzymes. *PLoS Comp. Biol.* 6:e100859
- 283. I. Nookaew; B. G. Gabrielsson; A: Holmäng; A.-S. Sandberg; **J. Nielsen** (2010) Identifying molecular effects of diet through systems biology: Influence of herring diet on sterol metabolism and protein turnover in mice. *PLoS One* 5:e12361
- 284. M. Papini; I. Nookaew; G. Scalcinati; V. Siewers; J. Nielsen (2010) Phosphoglycerate mutase knock-out mutant *Saccharomyces cerevisiae*: Physiological investigation and transcriptome analysis. *Biotechnol. J.* 5:1016-1027
- 285. R. Olivares-Hernandez; R. Usaite; **J. Nielsen** (2010) Integrative analysis using proteome and transcriptome data from yeast to unravel regulatory patterns at post-transcriptional level. *Biotechnol. Bioeng.* 107:865-875
- 286. R. Olivares-Hernandez; H. Sunner; J. C. Frisvad; L. Olsson; **J. Nielsen**; G. Panagiotou (2010) Combining substrate specificity analysis with support vector classifiers reveals feruloyl esterase as a phylogenetically informative protein group. *PLoS One* 5:e12781
- 287. S. Partow; V. Siewers; S. Bjørn; J. Nielsen; J. Maury (2010) Characterization of different promoters for designing a new expression vector in *Saccharomyces cerevisiae*. *Yeast* 27:955-964
- 288. J.M. Otero; M.A. Papadakis; D.B.R.K. Gupta Udatha; **J. Nielsen**; G. Panagiotou (2010) Yeast biological networks unforld the interplay of antioxidants, genome and phenotype, and reveal a novel regulator of the oxidative stress response. *PLoS One* 5:e13606
- 289. A.Canelas; N.Harrison; A.Fazio; J.Zhang; J.-P. Pitkänen, J. van den Brink, B.M. Bakker, L. Bogner, J. Bouwman, J.I. Castrillo, A. Cankorur, P. Chumnanpuen, P. Daran-Lapujade, D. Dikicioglu, K. van Eunen, J.C. Ewald, J. J. Heijnen, B. Kirdar, I. Mattila, F.I.C. Mensonides, A. Niebel, M. Penttilä, J.T. Pronk, M. Reuss, L. Salusjärvi, U. Sauer, D. Sherman, M. Siemann-Herzberg, H. Westerhoff, J. de Winde, D. Petranovic, S.G. Oliver, C.T. Workman, N. Zamboni and J. Nielsen (2010) Integrated multilaboratory systems biology reveals differences in protein metabolism between two reference yeast strains. *Nature Comm.* 1:145

- 290. P. Chumnapuen; C. Brackmann; S.K. Nandy; S. Chatzipapdopoulos; J. Nielsen; A. Enejder (2011) Lipid biosynthesis monitored at single cell level in *Saccharomyces cerevisiae*. *Biotechnol. J.* 6:1-8
- 291. W. Vongsangnak; K. Hansen; **J. Nielsen** (2011) Integrated analysis of the global transcriptional response to α–amylase over-production in *Aspergillus oryzae*. *Biotechnol. Bioeng.* 108:1130-1139
- 292. L. Albertsen; Y. Chen; L.S. Bach; S. Rattleff; J. Maury; S. Brix; J. Nielsen; U. Mortensen (2011) Diversion of flux toward sesquiterpene production in *Saccharomyces cerevisiae* by fusion of host and heterologous enzymes. *Appl. Environ. Microbiol.* 77:1033-1040
- 293. R. Ruenwai; A. Neiss; K. Laoteng; W. Vongsangnak; A.B. Dalfard; S. Cheevadhanarak; D. Petranovic; J. Nielsen (2011) Heterologous production of polyunsaturated fatty acids in *Saccharomyces cerevisiae* causes a global transcriptional response resulting in reduced proteasomal activity and increased oxidative stress. *Biotechnol. J.* 6:343-356
- 294. R. Olivares-Hernandez; S. Bordel; **J. Nielsen** (2011) Codon usage variability determines the correlation between proteome and transcriptome fold changes. *BMC Systems Biol.* 5:33
- 295. S. Pabinger; R. Rader; R. Agren; J. Nielsen; Z. Trajanoski (2011) MEMOSys: Bioinformatics platform for genomescale metabolic models. *BMC Systems Biol.* 5:20
- 296. T.B. Regueira; K.R. Kildegaard; B.G. Hansen; U.H. Mortensen; C. Hertweck; J. Nielsen (2011) Molecular basis for mycophenolic acid biosynthesis of *Penicillium brevicompactum*. Appl. Environ. Microbiol. 77:3035-3043
- 297. B.G. Hansen; B. Salomonsen; M.T. Nielsen; N. B. Hansen; J.B. Nielsen; K.F. Nielsen; T.B. Regueira; **J. Nielsen**; K.R. Patil; U.H. Mortensen (2011) Versatile enzyme expression and characterization system for *Aspergillus nidulans* with the *Penicillium brevicompactum* polyketide synthase gene from the mycophenolic acid gene cluster as a test case. *Appl. Environ. Microbiol.* 77:3044-3051
- 298. S.E. van Mulders; C. Stassen; L. Daenen; B. Devreese; V. Siewers; R.G.E. van Eijsden; J. Nielsen; F.R. Delvaux; R. Willaert (2011) The influence of microgravity on invasive growth in *Saccharomyces cerevisiae*. *Astrobiol*. 11:45-55
- 299. F.H. Karlsson; D.W.Ussery; **J. Nielsen**; I. Nookaew (2011) A closer look at *Bacteroides*: Phylogenetic relationship and genomic implications of a life in the human gut. *Microb. Ecol.* 61:473-485
- 300. M.R. Andersen; M.P- Salazar; P.J. Schaap, P.J.I. van de Vondervoort, D. Culley, J. Thykaer, J.C. Frisvad, K.F. Nielsen, R. Albang, K. Albermann, R.M. Berka, G.H. Braus, S.A. Braus-Stromeyer, L.M. Corrochano, Z. Dai, P.W.M. van Dijck, G. Hofmann, L.L. Lasure, J.K. Magnuson, H. Menke, M. Meijer, S.L. Meijer, J.B. Nielsen, M.L. Nielsen, A.J.J. van Ooyen, H.J. Pel, L. Poulsen, R. A. Samson, H. Stam, A. Tsang, J.M. van den Brink, A. Atkins, A. Aerts, H. Shapiro, J. Pangilinan, A. Salamov, Y. Lou, E. Lindquist, S. Lucas, J. Grimwood, I.V. Grigoriev, C.P. Kubicek, D. Martinez, N.N.M.E. van Peij, J.A. Roubous, J. Nielsen; Scott Baker (2011) Comparative genomics of citric-acid producing Aspergillus niger ATCC 1015 versus enzyme-producing CBS 513.88. Genome Res. 21:885-897
- 301. K.M. Madsen; G.D.B.R.K. Udatha; S. Semba; J.M. Otero; P. Koetter; **J. Nielsen**; Y. Ebizuka; T. Kushiro; G. Panagiotou (2011) Linking genotype and phenotype of *Saccharomyces cerevisiae* strains reveals metabolic engineering targets and leads to triterpene hyper-producers. *PLoS One* 6:e14763

- 302. W.A. Rodriguez-Limas; K.E.J. Tyo; J. Nielsen; O.T. Ramirez; L.A. Palomares (2011) Molecular and process design for rotavirus-like particle production in Saccharomyces cerevisiae, Microbial, Cell Factory 10:33
- 303. M.P. Piddocke; A. Fazio; W. Vongsangnak; M.L.Wong; H.P. Heldt-Hansen; C. Workman; **J. Nielsen**; L. Olsson (2011) Revealing the beneficial effect of protease supplementation to high gravity beer fermentations using "-omics" techniques. *Microbial Cell Factory* 10:27
- 304. M. Lindqvist; I. Nookaew; I. Brinkenberg; E. Samuelson; K. Thörn; **J. Nielsen**; A.M.Harandi (2011) Unraveling molecular signatures of immunostimulatory adjuvants in the female genital tract through systems biology. *PLoS One* 6:e20448
- 305. K.-K. Hong; W. Vongsangnak; G.N. Vemuri; J. Nielsen (2011) Unravelling evolutionary strategies of yeast for improving galactose utilization through integrated systems level analysis. *Proc. Nat. Acad. Sci. USA* 108:12179-12184
- 306. J.S. Rokem; W. Vongsangnak; J. Nielsen (2011) Comparative metabolic capabilities for *Micrococcus luteus* NCTC 2665, the "Fleming" Strain, and Actinobacteria. *Biotechnol. Bioeng.* 108:2770-2775
- 307. J. Zhang; S. Vaga; P. Chumnapuen; R. Kumar; G.N. Vemuri; R. Aebershold; J. Nielsen (2011) Mapping the interaction of Snf1 with Tor1 in Saccharomyces cerevisiae. Mol. Systems Biol. 7:545

- 308. L. Pedersen; K. Hansen; J. Nielsen; A. Eliasson Lantz; J. Thykaer (2012) Industrial glucoamylase fed-batch benefits from oxygen limitation and high osmolarity. *Biotechnol. Bioeng.* 109:116-120
- 309. L. Pedersen; K. Hansen; J. Nielsen; A.E. Lantz; J. Thykaer (2012) Industrial glucoamyase fed-batch benefits from oxygen limitation and high osmolarity. *Biotechnol. Bioeng.* 109:116-124
- 310. E. Larsson; V. Tremaroli; Y.S. Lee; O. Koren; I. Nookaew; A. Fricker; **J. Nielsen**; R. E. Ley; F. Bäckhed (2012) Analysis of the gut microbial regulation of host gene expression along the length of the gut and regulation of gut microbial ecoloty through MyD88. *Gut* doi:10.1136/gutjnl-2011-301104
- 311. W. Vongsangnak; L.F. Figueiredo; J. Förster; T. Weber; J. Thykaer; E. Stegmann; W. Wohlleben; J. Nielsen (2012) Genome-scale metabolic representation of *Amycolatopsis balhamycina*. *Biotechnol*. *Bioeng*. 109:1798-1807
- 312. L. Zihe; K.E.J. Tyo; J.L. Martinez; D. Petranovic; J. Nielsen (2012) Different expression systems for production of recombinant proteins in *Saccharomyces cerevisiae*. *Biotechnol*. *Bioeng*. 109:1259-1268
- 313. K.E.J. Tyo; L. Zihe; D. Petranovic; **J. Nielsen** (2012) Imbalance of heterologous protein folding and disulfide bond formation rates yields runaway oxidative stress. *BMC Biol.* 10:16
- 314. J. Hou; K.E.J. Tyo; Z. Liu; D. Petranovic; J. Nielsen (2012) Engineering of vesicle trafficking improves heterologous protein secretion in *Saccharomyces cerevisiae*. *Met. Eng.* 14:120-127
- 315. G. Scalcinati; C. Knuf; S. Partow; Y. Chen; J. Maury; M. Schalk; L. Daviet; **J. Nielsen**; V. Siewers (2012) Dynamic control of gene expression in *Saccharomyces cerevisiae* engineered for the production of plant sesquiterpene α-santalene in a fed-batch mode. *Met. Eng.* 14:91-103
- 316. M. Papini; I. Nookaew; V. Siewers; **J. Nielsen** (2012) Physiological characterization of recombinant *Saccharomyces cerevisiae* expressing the *Aspergillus nidulans* phosphoketolase pathway: validation of activity through ¹³C-based metabolic flux analysis. *Appl. Microbiol. Biotechnol.* 95:1001-1010
- 317. S. Shi; J.O. Valle-Rodriguez; S. Khoomrung; V. Siewers; **J. Nielsen** (2012) Functional expression and characterization of five wax ester synthases in *Saccharomyces cerevisae* and their utility for biodiesel production. *Biotechnol. Biofuels* 5:7
- 318. L. Caspeta; S.; R. Agren; I. Nookaew; **J. Nielsen** (2012) Genome-scale metabolic reconstructions of *Pichia stiidis* and *Pichia pastoris* and *in silico* evaluation of their potentials. *BMC Systems Biol.* 6:24
- 319. J.F. Nijkamp; M. van den Broek; E. Datema, S. de Kok; L. Bosman; M.A. Luttik; P. Daran-Lapujade; W. Vongsangnak; J. Nielsen; W.H.M. Heijne; P. Klaassen; C.J. Paddon; D. Platt; P. Kötter; R.C. van Ham; M.J.T. Reinders; J.T. Pronk; D. de Ridder; J.-M. Daran (2012) De novo sequencing, assembly and analysis of the genome of the laboratory strain Saccharomyces cerevisiae CEN.PK113-7D, a model for modern industrial biotechnology. *Microb. Cell Factories* 11:36
- 320. G. Scalcinati; J. Otero; J. van Vleet; T. Jeffries; L. Olsson; **J. Nielsen** (2012) Evolutionary engineering of *Saccharomyces cerevisiae* for efficient xylose consumption. *FEMS Yeast Res.* 12:582-597
- 321. R. Agren; S. Bordel; A. Mardinoglu; N. Pornputtapong; I. Nookaew; J. Nielsen (2012) Reconstruction of genome-scale active metabolic networks for 69 human cell types and 16 cancer types using INIT. *PLoS Comp. Biol.* 8:e1002518
- 322. Y. Chen; S. Partow; G. Scalcinati; V. Siewers; J. Nielsen (2012) Enhancing the copy number of episomal plasmids in *Saccharomyces cerevisiae* for improved protein production. *FEMS Yeast Res.* 12:598-607
- 323. S. Khoomrung; P. Chumnapuen; S. Jansa-ard; I Nookaew; **J. Nielsen** (2012) Fast and accurate preparation fatty acid acid methyl esters by microwave-assisted derivatization in the yeast *Saccharomyces cerevisiae*. *Appl. Microbiol*. *Biotechnol*. 94:1637-1646
- 324. P. Chumnapuen; J. Zhang; I. Nookaew; **J. Nielsen** (2012) Integrated analysis of transcriptome and lipid profiling reveals the co-influences of inositol-choline and Snf1 in controlling lipid analysis in yeast. *Mol. Genetics Genom.* 287:541-554
- 325. K.-K. Hong; J. Hou; S. Shoaie; J. Nielsen; S. Bordel (2012) Dynamic 13C-labelling experiments prove important differences in protein turnover rate between two Saccharomyces cerevisiae strains. FEMS Yeast Res. 12:741-747
- 326. M. R. Andersen; M. G. Jacobsen; R. P. de Vries; **J. Nielsen** (2012) Mapping the polysaccharide degradation potential of *Aspergillus niger*. *BMC Genomics* 13:313
- 327. I. Nookaew; M. Papini; N. Pornputtpong; G. Scalcinati; L. Fagerberg; M. Uhlen; **J. Nielsen** (2012) A comprehensive comparisons of RNA-Seq based transcriptome analysis from reads to differential gene expression and cross comparison with microarrays: A case study in *Saccharomyces cerevisiae*. *Nucleic Acid Res.* 40:10084-10097
- 328. Y. Chen; V. Siewers; **J. Nielsen** (2012) Profiling of cytosolic and peroxisomal acetyl-CoA metabolism in *Saccharomyces cerevisiae*. *PLoS One* 7:e42475

- 329. K.-K. Hong; **J. Nielsen** (2012) Recovery of phenotypes obtained by adaptive evolution through inverse metabolic engineering. *Appl. Environ. Microbiol.* 78:7579-7586
- 330. G. Scalcinati; S. Partow; V. Siewers; M. Schalk; L. Daviet; **J. Nielsen** (2012) Combined metabolic engineering of precursor and co-factor supply to increase alpha-santalene production by *Saccharomyces cerevisiae*. *Microb. Cell Fac.* 11:117
- 331. M. Papini; I. Nookaew; M. Uhlen; J. Nielsen (2012) Scheffersomyces stipitis: a comparative systems biology study with the Crabtree positive yeast Saccharomyces cerevisiae. Microb. Cell Fac. 11:136
- 332. K. Kocharin; Y. Chen; V. Siewers; J. Nielsen (2012) Engineering of acetyl-CoA metabolism for the improved production of polyhydroxybutyrate in *Saccharomyces cerevisiae*. *AMB Express* 2:52
- 333. C. Geijer; I. Pirkov; W. Vongsangnak; A. Ericsson; **J. Nielsen**; M. Krantz; S. Hohmann (2012) Time course gene expression profiling of yeast spore germination reveals a network of transcription factors orchestrating the global response. *BMC Genomics* 13:554
- 334. F. H. Karlsson; F. Fåk; I. Nookaew; V. Tremaroli; B. Fagerberg; D. Petranovic; F. Bäckhed*; **J. Nielsen*** (2012) Symptomatic atherosclerosis is associated with an altered gut metagenome. *Nature Comm.* 3:1245
- 335. J. K. Michener; J. Nielsen; C. D. Smolke (2012) Identification and treatment of heme depletion attributable to overexpression of a lineage of evolved P450 monooxygenases. *Proc. Nat. Acad. Sci.* 109:19504-19509
- 336. S. Partow; V. Siewers; L. Daviet; M. Schalk; J. Nielsen (2012) Reconstruction and evaluation of the synthetic bacterial MEP pathway in *Saccharomyces cerevisiae*. *PLoS One* 7(12):e52498

- 337. J. Hou; T. Österlund; L. Zihe; D. Petranovic; J. Nielsen (2012) Heat shock response improves heterologous protein production in *Saccharomyces cerevisiae*. *Appl. Microbiol. Biotechnol.* **97**:3559-3568
- 338. I. Nookaew; P.-A. Svensson; P. Jacobsen; M. Jernås; M. Taube; I. Larsson; J.C. Andersson-Assarsson; L. Sjöström; P. Frougel; A. Walley; J. Nielsen*; L.M.S. Carlsson* (2013) Adipose tissue resting energy expenditure and expression of genes involved in mitochondrial function are higher in women than in men. *J. Clin. Endocrin. Met.* **98**:E370-E378
- 339. Y. Chen; L. Daviet; M. Schalk; V. Siewers; J. Nielsen (2013) Establishing a platform cell factory through engineering of yeast acetyl-CoA metabolism. *Met. Eng.* 15:48-54
- 340. J.M. Otero; D. Cimini; K.R. Patil; S.G. Poulsen; L. Olsson; J. Nielsen (2013) Industrial systems biology of Saccharcomyces cerevisiae enables novel succinic acid cell factory. PLoS One 8:e54144
- L.Zihe; J. Hou; J.L.Martinez; D.Petranovic; J. Nielsen (2013) Correlation of cell growth and heterologous protein production by Saccharomyces cerevisiae. Appl. Microbiol. Biotechnol. 97:3559-3568
- 342. K.-K.Hong; **J.Nielsen** (2013) Adaptively evolved yeast mutants on galactose show trade-offs in carbon utilization on glucose. *Metabolic Eng.* 16:78-86
- 343. K. Sanli; F. Karlsson; I. Nookaew; **J. Nielsen** (2013) FANTOM: Functional and taxonomic analysis of metagenomes. *BMC Bioinformatics* 14:38
- 344. L. Väremo; J. Nielsen; I. Nookaew (2013) Enriching the gene set analysis of genome-wide data by incorporating directionality of gene expression and combining statistical hypotheses and methods. *Nucleic Acid Res.* 41:4378-4391
- 345. I. Thiele; Neil Swainston; R.M.T. Fleming, A. Hoppe, S. Sahoo, M.K. Aurich, H. Haraldsdottir, M.L. Mo, O. Rolfsson, M.D. Stobbe, S.G. Thorleifsson, R. Agren, C. Bölling, S. Bordel, A.K. Chavali, P. Dobson, W.B. Dunn, L. Endler, D. Hala, M. Hucka, D. Hull, D. Jameson, N. Jamshidi, J.J. Jonsson, N. Juty, S. Keating, I. Nookaew, N. Le Novère, N. Malys, A. Mazein, J.A. Papin, N.D. Price, E. Selkov, Sr, M.I. Sigurdsson, E. Simeonidis, N. Sonnenschein, K. Smallbone, A. Sorokin, J.H.G.M. van Beek, D. Weichart, I. Goryanin, J. Nielsen; Hans V. Westerhoff; D. B. Kell; P. Mendes; B. Palsson (2013) A community-driven global reconstruction of human metabolism. *Nature Biotechnol*. 31:419-425
- 346. L. Caspeta; N.A.A. Buijs; **J. Nielsen** (2013) The role of biofuels in the future energy supply. *Energy Environ. Sci.* 6:1077-108
- 347. A. Mardinoglu; R. Agren; C. Kampf; A. Asplund; I. Nookaew; P. Jacobsen; A.J. Walley; P. Froguel; L.M. Carlsson; M. Uhlen; J. Nielsen (2013) Integration of clinical data with a genome-scale metabolic model of the human adipocyte. *Mol. Systems Biol.* 9:649
- 348. L. Zihe; T. Österlund; J. Hou; D. Petranovic; **J. Nielsen** (2013) Anaerobic α-amylase production and secretion with fumarate as the final electron acceptor in yeast. *Appl. Environ. Microbiol.* 79:2962-2967
- 349. S. E. F. Borgos; S. Bordel; H. Sletta; H. Ertesvåg; Ø. Jacobsen; P. Bruheim; T. E. Ellingsen; J. Nielsen; S. Valla (2013) Mapping global effects of the anti-sigma factor MucA in Pseudomonas fluorescens SBW25 through genome-scale metabolic modeling. *BMC Systems Biol.* 7:19
- 350. R. Agren; L. Liu; S. Shoaie; W. Vongsangnak; I. Nookaew; J. Nielsen (2013) The RAVEN toolbox and its use for generating a genome-scale metabolic model for *Penicillium chrysogenum*. *PLoS Comp. Biol.* 9:e1002980
- 351. K. Kocharin; J. Nielsen (2013) Specific growth rate and substrate dependent polyhydroxybutyrate production in *Saccharomyces cerevisiae*. *AMB Express* 3:18
- 352. K. Kocharin; V. Siewers; **J. Nielsen** (2013) Improved polyhydroxybutyrate production by *Saccharomyces cerevisiae* through the use of the phosphoketolase pathway. *Biotechnol. Bioeng.* 110:2216-2224
- 353. F. Karlsson; V. Tremaroli; I. Nookaew; G. Bergström; C.J. Behre; B. Fagerberg; J. Nielsen*; F. Bäckhed* (2013) Gut metagenome in European women with normal, impaired and diabetic glucose control. *Nature* 498:99-103
- 354. E. Petelenz-Kirdziel; C. Kuehn; B. Nordlander; D. Klein: K.-K. Hong; T. Jacobson; P. Dahl; J. Schaber: J. Nielsen; S. Hohmann; E. Klipp (2013) Quantiative analysis of glycerol accumulation, glycolysis and growth under hyper osmotic stress. *PLoS Comp. Biol.* 9:e1003084
- 355. T. Osterlund; I. Nookaew; S. Bordel; J. Nielsen (2013) Mapping condition dependent regulation of metabolism in yeast through genome-scale modeling. *BMC Systems Biol.* 7:36

- 356. P. Chumnapuen; I. Nookaew; **J. Nielsen** (2013) Integrated analysis, transcriptome-lipidome, reveals the effects of INO-level (INO2 and INO4) on lipid metabolism in yeast. *BMC Systems Biol.* 7 Suppl 2:S7
- 357. L. Caspeta; **J. Nielsen** (2013) Economic and environmental impacts of microbial biodiesel. *Nature Biotechnol.* 31:789-793
- 358. R.G.E. van Eijsden; C. Stassen; L. Daenen; S.E. van Mulders; P.M. Bapat; V. Siewers; K.V. Gossens; J. Nielsen; F.R. Delvaux; P. van Hummelen; B. Devreese; R.G. Willaert (2013) A universal fixation method based on quaternary ammonium salts (RNAlater) for omics-technologies: *Saccharomyces cerevisiae* as a case study. *Biotechnol. Lett.* 35:891-900
- 359. R. Agren; J. M. Otero; J. Nielsen (2013) Genome-scale metabolic modeling enables metabolic engineering of *Saccharomyces cerevisiae* for succinic acid production. *J. Ind. Microbiol. Biotechnol.* 40:735-747
- 360. A. Krivorucho; C. Serrano-Amatriain; Y. Chen; V. Siewers; J. Nielsen (2013) Improving biobutanol production in engineered Saccharomyces cerevisiae by manipulation of acetyl-CoA metabolism. J. Ind. Microbiol. Biotechnol. 40:1051-1056
- 361. C. Knuf; I. Nookaew; S. Brown; M. McCulloch; A. Berry; **J. Nielsen** (2013) Investigation of malic acid production in *Aspergillus oryzae* under nitrogen starvation conditions. *Appl. Environ. Microbiol.* 79:6050-6058
- 362. S. Shoaie; F. Karlsson; A. Mardinoglu; I. Nookaew; S. Bordel; J. Nielsen (2013) Elucidating the interactions between bacteria in the human gut through metabolic modeling. *Sci. Rep.* 3:2532
- 363. A. Feizi; T. Österlund; D. Petranovic; S. Bordel; J. Nielsen (2013) Genome-scale modeling of the protein secretory machinery in yeast. *PLOS One* 8:e63284
- 364. M.C. Jewett; C.T.Workman; I. Nookaew; F.A. Pizarro; E. Agosin; L.I. Hellgren; J. Nielsen (2013) Mapping condition dependent regulation of lipid metabolism in *Saccharomyces cerevisiae*. *G3* 3:1979-1995
- 365. J. Anfelt; B. Hallström; J. Nielsen; M. Uhlen; E.P. Hudson (2013) Using transcriptomics to improve butanol tolerance in cyanobacteria. *Appl. Environ. Microbiol.* 79:7419-7427
- 366. S.Khoomrung; P. Chumnanpuen; S. Jansa-Ard; M. Ståhlman; I. Nookaew; J. Boren; J. Nielsen (2013) Rapid quantification of yeast lipid using microwave-assisted total lipid extraction and HPLC-CAD. *Anal. Chem.* 85:4912-4919
- 367. A.Hussain; I. Nookaew; S.Khoomrung; L. Andersson; I. Larsson; L. Hulthén; N. Jansson; R. Jakubowicz; S. Nilsson; A.-S.Sandberg; J. Nielsen; A. Holmäng (2013) A maternal diet of fatty fish reduces body fat of offspring compared with a maternal diet of beef and post-weaning diet of fish improves insulin sensitivity and lipid profile in adult C57BL/6 male mice. Acta Physiol. 209:220-234
- 368. I. Nookaew; K. Thorell; K. Worah; E. Wang; M.L. Hibberd; H. Sjövall; S. Pettersson; **J. Nielsen**; S. B. Lundin (2013) Transcriptome signatures in *Helicobacter pylori* infected mucosa identifies acidic mammalian chitinase loss as a corpus atrophy marker. *BMC Med. Genomics* 6:41

- 369. N. Pornputtapong; K. Wanichthanarak; A. Nilsson; I. Nookaew; J. Nielsen (2014) A dedicated database system for handling multi-level data in systems biology. *Source Code Biol. Med.* 9:17
- 370. A. Mardinoglu; R. Agren; K. Kampf; A. Asplund; M. Uhlen; **J. Nielsen** (2014) Genome-scale metabolic modeling of hepatocytes reveals serine deficiency in patients with non-alcoholic fatty liver disease. *Nature Comm.* 5:3083
- 371. P. Chumnapuen; M.A. Edberg; Jörn Smedsgaard; J. Nielsen (2014) Dynamic metabolic footprinting reveals the key components of metabolic networks in yeast *Saccharomyces cerevisiae*. *Int. J. Genomics* 894296:1-14
- 372. J.O.Valle-Rodriguez; S. Shi; V. Siewers; **J. Nielsen** (2014) Metabolic engineering of *Saccharomyces cerevisiae* for production of fatty acid ethyl esters, an advanced biofuel, by eliminating non-essential fatty acid utilization pathways. *Appl. Energy* 115:226-232
- 373. L.Liu; J.L.Martinez; D. Petranovic; J. Nielsen (2014) Balanced globin protein expression and heme biosynthesis improves production of human hemoglobin in *Saccharomyces cerevisiae*. *Met. Eng.* 21:9-16
- 374. N. Jensen; T. Strucko; K. Kildegaard; D. Florian; J. Maury; U. Mortensen; J. Forster; **J. Nielsen**; I. Borodina (2013) EasyClone: method for iterative chromosomal integration of multiple genes in *Saccharomyces cerevisiae*. *FEMS Yeast Res.* 14:238-248
- 375. R. Ledesma-Amaro; E.J. Kerkhoven; J.L. Revuelta; J. Nielsen (2014) Genome-scale metabolic modeling of the riboflavin overproducer *Ashbya gossypii*. *Biotechnol. Bioeng*. 111:1191-1199
- 376. S.L. Sjostrom; Y. Bai; M. Huang; Z. Liu; J. Nielsen; H.N. Joensson; H.A. Svahn (2014) High-throughput screening for industrial enzyme production hosts by droplet microfluidics. *Lab. Chip.* 14:806-813
- 377. L. Fagerberg; B.M. Hallstrom; P. Oksvold; C. Kampf; D. Djureinovic; J. Odeberg; M. Habuka; S. Tahmasebpoor; A. Danielsson; K. Edlund; A. Asplund; E. Sjostedt; E. Lundberg; C. Al-Khalili Szigyarto; M. Skogs; J. O. Takanen; H. Berling; H. Tegel; J. Mulder; P. Nilsson; J. M. Schwenk; C. Lindskog; F. Danielsson; A. Mardinoglu; A. Sivertsson; K. von Felitzen; M. Forsberg; M. Zwahlen; I.M. Olsson; S. Navani; M. Huss; J. Nielsen; F. Ponten; M. Uhlen (2014) Analysis of the human tissue-specific expression by genome-wide integration of transcriptomics and antibody-based proteomics. *Mol. Cell. Proteomics* 13:397-406
- 378. Y. Chen; J. Bao; I.-K. Kim; V. Siewers; J. Nielsen (2014) Coupled incremental precursor and co-factor supply improves 3-hydroxypropionic acid production by *Saccharomyces cerevisiae*. *Met. Eng.* 22:104-109
- 379. F. Gatto; I. Nookaew; J. Nielsen (2014) Chromosome 3p loss of heterozygosity is associated with a unique metabolic network in clear cell renal carcinoma. *Proc. Nat. Acad. Sci.* 111:E866-E875
- 380. B.W. de Jong; S. Shi; V. Siewers; **J. Nielsen** (2014) Improved production of fatty acid ethyl esters in *Saccharomyces cerevisiae* through up-regulation of the ethanol degradation pathway and expression of the heterologous phosphoketolase pathway. *Microb. Cell Fac.* 13:39
- 381. R. Agren; A. Mardinoglu; C. Kampf; A. Asplund; M. Uhlen; J. Nielsen (2014) Identification of anticancer drugs for hepatocellular carcinoma through personalized genome-scale metabolic modeling. *Mol. Systems Biol.* 10:721

- 382. C. Kampf; A. Mardinoglu; L. Fagerberg; B.M. Hallström; K. Edlund; E. Lundberg; F. Ponten; **J. Nielsen**; M. Uhlen (2014) The human liver-specific proteome defined by transcriptomics and antibody-based profiling. *FASEB J.* 28:2901-2914
- 383. J. L. Martinez; S. Bordel; K.-K. Hong; **J. Nielsen** (2014) Gcn4p and the Crabtree effect of yeast: Drawing the causal effect of the Crebtree effect of *Saccharomyces cerevisiae* and explaining evolutionary trade-offs of adaptation to galactose through systems biology. *FEMS Yeast Res.* 14:654-662
- 384. S. Shi; J. O. Valle-Rodriguez; V. Siewers; **J. Nielsen** (2014) Engineering of chromosomal wax ester synthase integrated *Saccharomyces cerevisiae* mutants for improved biosynthesis of fatty acid ethyl esters. *Biotechnol. Bioeng.* 111:1740-1747
- 385. I.E. El-Semman; F.H. Karlsson; S. Shoaie; I. Nookaew; T.H. Soliman; **J. Nielsen** (2014) Genome-scale metabolic reconstructions of *Bifidobacterium adolescent* L2-32 and *Faecalibacterium prausnitzii* A2-165 and their interaction using CommunityML. *BMC Systems Biol.* 8:41
- 386. S. Shi; Y. Chen; V. Siewers; **J. Nielsen** (2014) Improving production of malonyl-CoA derived metabolites by abolishing Snf1 dependent regulation of Acc1. *mBio* 5:e01130-14
- 387. C. Knuf; I. Nookaew; I. Remmers; S. Khoomrung; S. Brown; A. Berry; J. Nielsen (2014) Physiological characterization of the high malic acid-producing *Aspergillus oryzae* strain 2103a-68. *Appl. Microbiol. Biotechnol.* 98:3517-3527
- 388. K.E.J. Tyo; Z. Liu; Y. Magnusson; D. Petranovic; J. Nielsen (2014) Impact of protein uptake and degradation on recombinant protein secretion in yeast. *Appl. Microbiol. Biotechnol.* 98:7149-7159
- 389. M. Garcia-Albornoz; S. Thankaswamy-Kosalia; A. Nilsson; L. Väremo; I. Nookaew; **J. Nielsen** (2014) BioMet Toolbox 2.0: Genome-wide analysis of metabolism and omics data. *Nuc. Acid Res.* 42:W175-W181
- 390. F.H. Karlsson; I. Nookaew; J. Nielsen (2014) Metagenomic data utilization and analysis and construction of a global gut microbial gene catalogue. *PLOS Comp. Biol.* 10:e1003706
- 391. J. Hou; H. Tang; L. Zihe; T. Österlund; J. Nielsen; D. Petranovic (2014) Management of the endoplasmic reticulum stress by activation of the heat shock response in yeast. FEMS Yeast Res. 14:481-494
- 392. L. Liu; A. Feizi; T. Österlund; C. Hjort; J. Nielsen (2014) Genome-scale analysis of the high-efficient protein secretion system of *Aspergillus oryzae*. *BMC Systems Biol*. 8:73
- 393. L. Zihe; L. Liu; T. Österlund; J. Hou; M. Huang; L. Fagerberg; D. Petranovic; M. Uhlen; J. Nielsen (2014) Improved production of a heterologous amylase by *Saccharomyces cerevisiae* by inverse metabolic engineering. *Appl. Environ. Microbiol.* 80:5542-5550
- 394. C. Kampf; A. Mardinoglu; L. Fagerberg; B.M. Hallström; A. Danielsson; **J. Nielsen**; F. Ponten; M. Uhlen (2014) Defining the gall bladder proteome by transcriptomics and affinity proteomics. *Proteomics* 14:2498-2507
- 395. P.-J.P. Aspuria; S.Y.Lunt; L. Väremo; L. Vergnes; M. Gozo; J.A. Beach; B.Salumbides; K. Reue; W.R. Wiedemeyer; J. Nielsen; B.Y. Karlan; S. Orsulic (2014) Succinate dehydrogenase inhibition leads to epithelial-mesenchymal transition and reprogrammed carbon metabolism. *Cancer Met.* 4:21
- 396. A. Mardinoglu; C. Kampf; A. Asplund; L. Fagerberg; B. Hallström; K. Edlund; M. Blüher; F. Ponten; M. Uhlen; J. Nielsen (2014) Defining the human adipose tissue proteome to reveal metabolic alterations in obesity. *J. Prot. Res.* 11:5106-5119
- 397. K. Kildegaard; B. M. Hallström; T.H.Blicher; N. Sonnenschein; S. Sherstyk; N.B. Jensen; S. Harrison; J. Maury; M. Herrgard; A. Juncker; J. Förster; J. Nielsen; I. Borodina (2014) Evolution reveals a cyclic glutathione-dependent mechanism of 3-hydroxypropionic acid detoxification. *Met. Eng.* 26:57-66
- 398. L. Väremo; F. Gatto; J. Nielsen (2014) Kiwi: a tool for integration and visualization of network topology and gene-set analysis. *BMC Bioinfor*. 15:408
- 399. L.Caspeta; Y. Chen; P. Ghiaci; A. Feizi; S. Buskov; B.M. Hallström; D. Petranovic; J. Nielsen (2014) Altered sterol composition renders yeast thermotolerant. *Science* 346:75-78
- 400. C. Navarrete; V. Siewers; **J. Nielsen** (2014) Enhanced ethanol production and reduced glycerol formation in *fps1* mutants of *Saccharomyces cerevisiae* engineered for improved redox balancing. *AMB Express* 4:86

- 401. M. Uhlén, L. Fagerberg, B.M. Hallström, C. Lindskog, P. Oksvold, A. Mardinoglu, Å. Sivertsson, C. Kampf, E. Sjöstedt, A. Asplund, I. Olsson, K. Edlund, E. Lundberg, S. Navani, C.A.-K. Szigyarto, J. Odeberg, D. Djureinovic, J.O. Takanen, S. Hober, T. Alm, H. Berling, H. Tegel, J. Mulder, J. Rockberg, P. Nilsson, J.M. Schwenk, M. Hamsten, K. von Feilitzen, M. Forsberg, L. Persson, F. Johansson, M. Zwahlen, G. von Heijne, J. Nielsen; F. Ponten (2015) Tissue based map of the humane proteome. Science 347:1260419,1-9
- 402. J. L. Martinez; L. Lifang, D. Petranovic; J. Nielsen (2015) Engineering the oxygen sensing regulation results in an enhanced recombinant human hemoglobin production by *Saccharomyces cerevisiae*. *Biotechnol. Bioeng.* 112:181-188
- 403. N.A. Buijs; Y.J. Zhou; V. Siewers; **J. Nielsen** (2015) Long-chain alkane production by the yeast *Saccharomyces cerevisiae*. *Biotechnol. Bioeng*. 112:1275-1279
- 404. P. Ghaffari; A. Mardinoglu; A. Asplund; S. Shoaie; C. Kampf; M. Uhlen; J. Nielsen (2015) Identifying anti-growth factors for human cancer cell lines through genome-scale metabolic modeling. *Sci. Rep.* 5:8183
- 405. I. Borodina; K. Kildegaard; N.B. Jensen; T.H. Blicher; J. Maury; S. Sherstyk; K. Schneider; P. Lamosa; M.J. Herrgård; I. Rosenstand; F. Öberg; J. Förster; J. Nielsen (2015) Establishing a synthetic pathway for high-level production of 3-hydroxypropionic acid in *Saccharomyces cerevisiae* via beta-alanine. *Met. Eng.* 27:57-64
- 406. C. Zhang; B. Ji; A. Mardinoglu; J. Nielsen; Q. Hua (2015) Logical transformation of genome scale metabolic models for genetic applications and analysis. *Bioinfor*. 31:2324-2331
- 407. B. de Jong; S. Shi; J.O. Valle-Rodriguez; V. Siewers; **J. Nielsen** (2015) Metabolic pathway engineering for fatty acid ethyl ester production in *Saccharomyces cerevisiae* using stable chromosomal integration. *J. Ind. Microbiol. Biotechnol.* 42:477-486

- 408. Y. Chen; Y. Zhang; V. Siewers; **J. Nielsen** (2015) Ach1 is involved in shutting mitochondrial acetyl units for cytosolic C2 provision in *Saccharomyces cerevisiae* lacking pyruvate carboxylase. *FEMS Yeast Res.* 15:fov015
- 409. R. Nicastro; F. Tripodi; C. Guzzi; V. Reghellin; S. Khoomrung; C. Airoldi; **J. Nielsen**; L. Alberghina; P. Cocetti (2015) Enhanced amino acid utilization sustains growth of cells lacking Snf1/AMPK. *BBA Mol. Cell Res.* 1853:1615-1625
- 410. K.V.Y. Gossens; F.S. Ielasi; I. Nookaew; I. Stals; L. Alonso-Sarduy; L. Daenen; S.E. Van Mulders; R.G.E. van Eijsden; C. Stassen; V. Siewers; F.R. Delvaux; S. Kasas; **J. Nielsen**; B. Devreese; R.G. Willaert (2015) Molecular mechanisms of flocculation self-recognition in yeast and its role in mating and survival. *mBio* 6:e00427-15
- 411. T. Österlund; S. Bordel; J. Nielsen (2015) Controllability analysis of transcriptional regulatory networks reveals circular control patterns among transcription factors. *Integr. Biol.* 7:560-568
- 412. S. Tymoshenko; R.D. Oppenheim; R. Agren; J. Nielsen; D. Soldati-Favre; V. Hatzimanikatis (2015) Metabolic needs and capabilities of *Toxoplasma gondii* through combined computational and experimental analysis. *PLOS Comp. Biol.* 11:e1004261
- 413. L. Väremo; C. Scheele; C. Broholm; A. Mardinoglu; C. Kampf; A. Asplund; I. Nookaew; M. Uhlen; B.K. Pedersen; J. Nielsen (2015) Transcriptome and proteome driven reconstruction of the human myocyte metabolic network and its use for identification of markers for diabetes. *Cell Rep.* 11:1-13
- 414. M. Engqvist; J. Nielsen (2015) ANT: Software for generating and evaluationg degenerate codons for natural and expandic genetic codes. ACS Synthetic Biol. 4:935-938
- 415. A. Feizi; A. Banaei-Esfahani; J. Nielsen (2015) HCSD: the human cancer secretome database. Database bav051:1-8
- 416. J. Lopez; K. Essus; I.-K. Kim; R. Pereira; J. Herzog; V. Siewers; J. Nielsen; E. Agosin (2015) Production of β-ionone by combined expression of carotenogenic and plant CCD1 genes in *Saccharomyces cerevisiae*. *Microb. Cell Fac.* 14:84
- 417. Y. Zhang; Z. Dai; A. Krivoruchko; Y. Chen; V. Siewers; J. Nielsen (2015) Functional pyruvate formate lyase pathway expressed in different electron donors in *Saccharomyces cerevisiae* at aerobic growth. *FEMS Yeast Res.* 15:fov024
- 418. F. Gatto; H. Miess; A. Schulze; J. Nielsen (2015) Flux balance analysis predicts essential genes in clear cell renal cell carcinoma metabolism. *Sci. Rep.* 5:10738
- 419. C. Lindskog; J. Linne; L. Fagerberg; B.M. Hallström; C.-J. Sundberg; M. Lindholm; M. Huus; C. Kampf; H. Choi; A. Liem; P. Ping; L. Väremo; A. Mardinoglu; J. Nielsen; E. Larsson; F. Ponten; M. Uhlen (2015) The human cardiac and skeletal muscle proteome defined by transcriptomics and antibody-based profiling. *BMC Genom.* 16:475
- 420. S. Shoaie; P. Ghaffari; P. Kovatcheva-Datchary; A. Mardinoglu; P. Sen; E. Pujos-Guillot; T. de Wouters; C. Juste; S. Rizkalla; J. Chilloux; L. Hoyles; J.K. Nicholson; ANR MicroObese Consortium; J. Dore; M.E. Dumas; K. Clement; F. Bäckhed; J. Nielsen (2015) Quantifying diet-induced metabolic changes of the human gut microbiome. *Cell Metabolism* 22:320-331
- 421. L. Caspeta; **J. Nielsen** (2015) Thermotolerant yeast strains adapted by laboratory evolution show trade-off at ancestral temperatures to other stresses. *mBio* 6:e00431-15
- 422. V. Tremaroli; F. Karlsson; M. Weling; M. Ståhlman; P. Kovatcheva-Datchary; T. Olbers; L. Fändriks; C. W. le Roux; J. Nielsen; F. Bäckhed (2015) Roux-en-Y Gastric Bypass and Vertical Banded Gastroplasty Induce Long-Term Changes on the Human Gut Microbiome Contributing to Fat Mass Regulation. *Cell Metabolism* 22:228-238
- 423. N. Pornputtaporn; I. Nookaew; **J. Nielsen** (2015) Human metabolic atlas: An online resource for human metabolism. *Database* bay068:1-9
- 424. M. Garcia-Albornoz; **J. Nielsen** (2015) Finding directionality and gene-disease predictions in disease associations. *BMC Systems Biol.* 9:35
- 425. M.H. Medema; R. Kottmann; P. Yilmaz; M. Cummings; J.B. Biggins; K. Blin: I. de Bruijn; Y.H. Chooi; J. Claesen; R.C. Coates; P. Cruz-Morales; S. Duddela; S. Düsterhus; D.J. Edwards; D.P. Fewer; N. Garg; C. Geiger; J.P. Gomez-Escribano; A. Greule; M. Hadjithomas; A.S. Haines; E.J.N. Helfrich; M.L. Hillwig; K. Ishida; A.C. Jones; C.S. Jones; K. Jungmann; C. Kegler; H.U. Kim; P. Kötter; D. Krug; J. Masschelein; A.V. Melnik; S.M. Mantovani; E.A. Monroe; M. Moore; N. Moss; H.-W. Nützmann; G. Pan; A. Pati; D. Petras; J. Reen; F. Rosconi; Z. Rui; Z. Tian; N.J. Tobias; Y. Tsunematsu; P. Wiemann; E. Wyckoff; X. Yan; G. Yim; F. Yu; Y. Xie; B. Aigle; A.K. Apel; C.J. Balibar; E.P. Balskus; F. Barona-Gómez; A. Bechthold; H.B. Bode; R. Borriss; S.F. Brady; A.A. Brakhage; P. Caffrey; Y.-Q. Cheng; J. Clardy; R.J. Cox; R. De Mot; S. Donadio; M.S. Donia; W.A. van der Donk; P.C. Dorrestein; S. Doyle; A.J.M. Driessen; M. Ehling-Schulz; K.-D. Entian; M.A. Fischbach; L. Gerwick; W.H. Gerwick; H. Gross; B. Gust; C. Hertweck; M. Höfte; S.E. Jensen; J. Ju; L. Katz; L. Kaysser; J.L. Klassen; N.P. Keller; J. Kormanec; O.P. Kuipers; T. Kuzuyama; N.C. Kyrpides; H.-J. Kwon; S. Lautru; R. Lavigne; C. Lee; B. Linquan; X. Liu; W. Liu; A. Luzhetskyy; T. Mahmud; Y. Mast; C. Méndez; M. Metsä-Ketelä; J. Micklefield; D.A. Mitchell; B.S. Moore; L.M. Moreira; R. Müller; B.A. Neilan; M. Nett; J. Nielsen; F. O'Gara; H. Oikawa; A. Osbourn; M.S. Osburne; B. Ostash; S.M. Payne; J.-L. Pernodet; M. Petricek; J. Piel; O. Ploux; J.M. Raaijmakers; J.A. Salas; E.K. Schmitt; B. Scott; R.F. Seipke; B. Shen; D.H. Sherman; K. Sivonen; M.J. Smanski; M. Sosio; R.D. Süssmuth; K. Tahlan; C.M. Thomas; Y. Tang; A.W. Truman; M. Viaud; J.D. Walton; C.T. Walsh; T. Weber; G.P. van Wezel; B. Wilkinson; J.M. Willey; W. Wohlleben; G.D. Wright; N. Ziemert; C. Zhang; S.B. Zotchev; R. Breitling; E. Takano; F.O. Glöckner (2015) The Minimum Information about a Biosynthetic Gene cluster (MIBiG) specification. Nature Chem. Biol. 11:625-631
- 426. L. Liu; Y. Zhang; Z. Liu; D. Petranovic; **J. Nielsen** (2015) Improving heterologous protein secretion at aerobic conditions by activating hypoxia induced genes in *Saccharomyces cerevisiae*. *FEMS Yeast Research* 15:fov070
- 427. Y. Zhang; G. Liu; M.K.M. Engqvist; A. Krivoruchko; B.M. Hallström; Y. Chen; V. Siewers; J. Nielsen (2015) Adaptive mutations in sugar metabolism restores growth on glucose in a pyruvate decarboxylase negative yeast strain. *Microb. Cell Fac.* 14:116
- 428. J.C. Qin; Y.J. Zhou; A. Krivoruchko; M. Huang; L. Liu; S. Khoomrung; V. Siewers; B. Jiang; **J. Nielsen** (2015) Modular pathway rewiring of *Saccharomyces cerevisiae* enables high-level production of L-ornitine. *Nature Com*. 6:8224

- 429. A. Rodriguez; K.R. Kildegaard; M. Li; I. Borodina; J. Nielsen (2015) Establishment of a yeast platform strain for production of p-coumaric acid through metabolic engineering of aromatic amino acid biosynthesis. *Met. Eng.* 31:181-188
- 430. M. Huang; Y. Bai; S.L. Sjostrom; B.M. Hallström; Z. Liu; D. Petranovic; M. Uhlen; H.N. Joensson; H. Andersson-Svahn; J. Nielsen (2015) Microfluidic screening and whole genome sequencing identifies mutations associated with improved protein secretion by yeast. *Proc. Nat. Acad. Sci. USA* 112:E4689-96
- 431. M. Li; K.R. Kildegaard; Y. Chen; A. Rodriguez; I. Borodina; J. Nielsen (2015) *De novo* production of resveratrol from glucose or ethanol by engineered *Saccharomyces cerevisiae*. *Met. Eng.* 32:1-11
- 432. E. Fletcher; A. Feizi; S. Kim; V. Siewers; J. Nielsen (2015) RNAseq analysis of *Pichia anomaia* reveals important mechansisms required for survival at low pH. *Mic. Cell Fac.* 14:143
- 433. A. Mardinoglu; J. Helker; D. Gärtner; E. Bjornson; M.R. Schön; G. Flehmig; N. Klöting; M. Fasshauer; M. Stumvoll; J. Nielsen; M. Blüher (2015) Extensive weight loss reveals distinct gene expression changes in human subcutaneous and visceral adipose tissue. Sci. Rep. 5:14841
- 434. A. Mardinoglu; S. Shoaie; M. Bergentall; P. Ghaffari; C. Zhang; E. Larsson; F. Bäckhed; J. Nielsen (2015) The gut microbiome modulates host amino acid and glutathione metabolism in mice. *Mol. Systems Biol.* 11:834
- 435. E. Björnson; B. Mukhopadhyay; A. Asplund; N. Pristovsek; R. Cinar; S. Romeo; M. Uhlen; G. Kunos; J. Nielsen; A. Mardinoglu (2015) Stratification of Hepatocellular Carcinoma Patients based on Acetate Utilization. Cell Rep. 13:2014-2026
- 436. S. Khoomrung; J.L. Martinez; S. Tippmann; S. Jansa-Ard; M. Buffing; R. Nicastro; J. Nielsen (2015) Expanded metabolite coverage of *Saccharomyces cerevisiae* extract through improved chemical derivatization. *Anal. Chem. Res.* 6:9-16
- 437. K.R. Kildegaard; Y. Chen; Z. Wang; J. Nielsen; I. Borodina (2015) Production of 3-hydroxypropionic acid from glucose and xylose by metabolically engineered *Saccharomyces cerevisiae*. *Met. Eng. Com.* 2:132-136
- 438. J. Anfelt; D. Kaczmarzyk; K. Shabestary; B. Renberg; J. Rockberg; J. Nielsen; M. Ühlen; E.P. Hudson (2015) Genetic and nutrient modulation of acetyl-CoA levels in *Synechocystis* for n-butanol production. *Mic. Cell Fac.* 14:167

- 439. Z.A. Irani; E. Kerkhoven; S.A. Shojaosadati; J. Nielsen (2015) Genome-scale metabolic model of *Pichia pastoris* with native and humanized glycosylation of recombinant proteins. *Biotechnol. Bioeng.* 113:961-969
- 440. E. Kerkhoven; K. Pmoraning; S. Baker; J. Nielsen (2015) Regulation of amino acid metabolism controls flux to lipid accumulation in *Yarrowia lipolytica*. npj Systems Biol. Appl. 2:16005
- 441. S. Shi; J. Haichuan; V. Siewers; J. Nielsen (2015) Improved production of fatty acids by *Saccharomyces cerevisiae* through screening a cDNA library from the oleaginous yeast *Yarrowia lipolytica*. FEMS Yeast Res. 16:fv108
- 442. B. de Jong; V. Siewers; J. Nielsen (2016) Physiological and transcriptional characterization of *Saccharomyces cerevisiae* engineered for production of fatty acid ethyl esters. *FEMS Yeast Res.* Fov105:1-9
- 443. S. Tippmann; G. Scalcinati; V. Siewers; **J. Nielsen** (2015) Production of farnesene and santalane by *Saccharomyces cerevisiae* using fed-batch cultivations with RQ-controlled feed. *Biotechnol. Bioeng.* 113:72-81
- 444. S. Tippmann; J. Nielsen; S. Khoomrung (2016) Improved quantification of farnesene during microbial production from *Saccharomyces cerevisiae* in two-liquid-phase fermentations. *Talanta* 146:100-106
- 445. F. David; J. Nielsen; V. Siewers (2016) Flux control at the malonyl-CoA node through hierarchical dynamic pathway regulation in *Saccharomyces cerevisiae*. ACS Syn. Biol. 5:224-233
- 446. L. Pedersen; M. Idorn; G.H. Olofsson; B. Lauenborg; I. Nookaew; R.H. Hansen; H.H. Johannesen; J.C. Becker; K.S. Pedersen; C. Dethlefsen; J. Nielsen; J. Gehl; B.K. Pedersen; P.T. Straten; P. Hojman (2016) Voluntary running suppresses tumor growth through epinephrine- and IL-6-dependent NK cell mobilization and redistribution. *Cell Met.* 23:1-9
- 447. J, Maury; S.M. Germann; S.A.B. Jacobsen; N.B. Jensen; K.R. Kildegaard; M.J. Heergård; K. Schneider; A. Koza; J. Förster; J. Nielsen; I. Borodina (2016) EasyCloneMulti: A set of vectors for simultaneous and multiple genomic integrations in *Saccharomyces cerevisiae*. *PLOS One* 0150394:1-22
- 448. F. Gatto; J. Nielsen (2016) Systematic analysis of overall survival and interactions between tumor metabolism and drug treatment. J. Hematology Oncology 9:15
- 449. A. Nilsson; J. Nielsen (2016) Metabolic trade-offs in yeast are caused by F1F0-ATP synthase. Sci. Rep. 6:22264
- 450. K. Kildegaard; N.B. Jensen; K. Schneider; E. Czarnotta; E. Özdemir; T. Klein; J. Maury; B.E. Ebert; H.B. Christensen; Y. Chen; I.-K. Kim; M.J. Herrgård; L.M. Blank; J. Forster; J. Nielsen; I. Borodina (2016) Engineering and systems-level analysis of Saccharomyces cerevisiae for production of 3-hydroxypropionic acid via malonyl-CoA reductase-dependent pathway. *Microb. Cell Fac.* 15:53
- 451. Y. Zhou; N. A. Buijs; Z. Zhu; J. Qin; V. Siewers; **J. Nielsen** (2016) Production of fatty acid derived oleochemicals and biofuels by synthetic yeast cell factories. *Nature Com.* 7:11709
- 452. I. E. Elsemman; A. Mardinoglu; S. Shoaie; T.H. Soliman; **J. Nielsen** (2016) Systems biology analysis of metabolism reveals the role of copy number increase in parts of chromosome 1q in hepatitis C virus associated hepatocellular carcinoma. *Mol. Biosystems* 12:1496-1506
- 453. G. Liu; D. Bergenholm; J. Nielsen (2016) Genome-wide mapping of binding sites reveals multiple biological functions of the transcription factor Cst6p in *Saccharomyces cerevisiae*. mBio 7:e00559-16
- 454. F. Gatto; N. Volpi; H. Nilsson; I. Nookaew; M. Maruzzo; A. Roma; M. E. Johansson; U. Steiner; S. Lundstam; U. Basso; J. Nielsen (2016) Glycosaminoglycan Profiling in Patients' Plasma and Urine Predicts the Occurrence of Metastatic Clear Cell Renal Cell Carcinoma. Cell Rep. 15:1-15
- 455. S.L. Svahn; L. Väremo; B. Gabrielsson; E. Peris; I. Nookaew; L. Grahnemo; A.-S. Sandberg; I.W. Asterholm; J.-O. Jansson; **J. Nielsen**; M.E. Johansson (2016) Six tissue transcriptomics reveals specific immune suppression in spleen by dietary polyunsaturated fatty acids. *PLOS One* 11:e0155099

- 456. R. Pereira; J. Nielsen; I. Rocha (2016) Improving the flux distributions simulated with genome-scale metabolic models of *Saccharomyces cerevisiae*, *Met. Eng. Com.* 3:153-163
- 457. S. Lee; M. Kilicarslan; C. Zhang; E. Björnson; B.M. Hallström; A.K. Groen; E. Ferrannini; M. Laakso; M. Blüher; M. Uhlen; J. Nielsen; U. Smith; J. Boren; M. Serlie; A. Mardinoglu (2016) Integrated network analysis reveals an association between increased plasma mannose levels and insulin resistance and secretion. *Cell Met.* 24:172-184
- 458. L. Caspeta; Y. Chen; J. Nielsen (2016) Thermotolerant yeasts selected by adaptive evolution express heat stress response at 30C. *Sci. Rep.* 6:27003
- 459. S. Lee; A. Marginoglu; C. Zhang; D. Lee; **J. Nielsen** (2016) Dysregulated signaling hubs of liver metabolism reveal hepatocellular carcinoma pathogenesis. *Nuc. Acid Res.* 44:5529-5539
- 460. S. Busti; V. Mapelli; F. Tripodi; R. Sanvito; F. Magni; M.C. Jewett; J. Nielsen; L. Alberghina; M. Vanoni (2016) Respiratory metabolism and calorie restriction relieve persistent endoplasmic reticulum stress induced by calcium shortage in yeast. *Sci. Rep.* 6:27942
- 461. F. Gatto; A. Schultze; J. Nielsen (2016) Systematic analysis reveals that cancer mutations converge on deregulated metabolism of arachinodate and xenobiotics. *Cell Rep.* 16:878-895
- 462. P.-J. Lahtvee; R. Kumar; B.M. Hallström; J. Nielsen (2016) Adaption to different types of stress converge on mitochondrial metabolism. *Mol. Biol. Cell* 27:2505-2514
- 463. A.S. Rajkumar; G. Liu; D. Bergenholm; D. Arsovska; M. Kristensen; J. Nielsen; M.K. Jensen; J.D. Keasling (2016) Engineering of synthetic, stress sensitive yeast promoters. *Nuc. Acid Res.* 44:e136
- 464. J.L. Martinez; E. Meza; D. Petranovic; J. Nielsen (2016) The impact of respiration and oxidative stress response on recombinant α-amylase production by *Saccharomyces cerevisiae*. *Metab*. *Eng. Com.* 3:205-210
- 465. S. Grijseels; J.C. Nielsen; M. Randelovic; **J. Nielsen**; K.F. Nielsen; M. Workman; J.C. Frisvad (2015) *Penicillium arizonense*, a new, genome sequenced fungal species, reveals a high chemical diversity in secreted metabolites. *Sci. Rep.* 6:35112
- 466. J. Casey; A. Mardinoglu; **J. Nielsen**; D. Kari (2016) Adaptive evolution of phosphorous metabolism in *Prochlorococcus. mSystems* 1:e00065-16
- 467. Y. Zhou; N.A.. Buijs; Z. Zhu; D.O. Gomez; A. Boonsombuti; V. Siewers; J. Nielsen (2016) Harnessing peroxisomes for production of fatty acid-derived biofuels and chemicals in yeast. J. Am. Chem. Soc. 138:15368-15377
- 468. M. Li; K.R. Kildegaard; Y. Chen; A. Rodriguez; I. Borodina; J. Nielsen (2016) Engineering yeast for high-level production of stilbenoid antioxidants. *Sci. Rep.* **6**:36827
- 469. E. Fletcher; A. Feizi; B. Hallström, M. Bisschops; S. Koomrung; V. Siewers; J. Nielsen (2016) Evolutionary engineering revewals divergent paths when yeast is adapted to different acidic environments. *Met. Eng.* 113:1164-1170
- 470. F. Gatto; M. Maruzzo; C. Magro; U. Basso; J. Nielsen (2016) Prognostic value of plasma and urine glycosaminoglycan scores in clear cell renal cell carcinoma. *Front. Oncol.* **6**:253
- 471. A. Bergman; V. Siewers; **J. Nielsen**; Y. Chen (2016) Functional expression and evaluation of heterologous phosphoketolases in *Saccharomyces cerevisiae*. *AMB Express* **6**:115

- 472. Z. Zhu; Y.J. Zhou; A. Krivoruchko; M. Grininger; Z.K. Zhao; J. Nielsen (2017) Expanding the product portfolio of fungal type I fatty acid synthases. *Nature Chem. Biol.* 13:360-362
- 473. L. Wenning; T. Yu; F. David; J. Nielsen; V. Siewers (2016) Establishing very long-chain fatty alcohol and wax ester biosynthesis. *Biotechnol. Bioeng.* 114:1025-1035
- 474. Y. Wei; V. Siewers; J. Nielsen (2017) Cocoa butter-like lipid production ability of non-oleaginous and oleaginous yeasts under nitrogen limited culture conditions. *Appl. Microbiol. Biotechnol.* **101**:13577-3585
- 475. A. Nilsson; A. Mardinoglu; J. Nielsen (2017) Predicting growth of the healthy infant using a genome-scale metabolic model. *npj Systems Biol. Appl.* **3**:3
- 476. A. Mardinoglu; E. Bjornson; C. Zhang; M. Klevstig; S. Söderlund; M. Ståhlman; M. Adiels; A. Hakkarainen; N. Lundbom; M. Kilicarslan; B.M. Hallström; J. Lundbom; B. Verges; P.H.R. Barret; G.F. Watts; M. Serlie; J. Nielsen; M. Uhlen; U. Smith; H.-U. Marschall; M.-R. Taskine; J. Boren (2017) Personalized genome-scale modeling identifies NAD+ and glutathione metabolism as a target for treatment of NAFLD. *Mol. Sys. Biol.* 13:916
- 477. Y. Wei; M. Gossing; D. Bergenholm; V. Siewers; J. Nielsen (2017) Increasing cocoa butter-like lipid production of *Saccharomyces cerevisiae* by expression of selected cocoa genes. *AMB Express* **7**:34
- 478. S. Tippmann; R. Ferreira; V. Siewers; J. Nielsen; Y. Chen (2017) Effects of acetoacetyl-CoA synthase expression on production of farnesene in *Saccharomyces cerevisiae*. J. Ind. Biotechnol. Microbiol. 44:911-922
- 479. S. Tippmann; J. Anfelt; F. David; J.M. Rand; V. Siewers; M. Uhlen; J. Nielsen; E.P. Hudson (2017) Affibody scaffold improve sesquiterpene production in *Saccharomyces cerevisiae*. ACS Syn. Biol. 6:19-28
- 480. E.L. Bredeweg; K.R. Pomraning; Z. Dai; J. Nielsen; E. Kerkhoven; S.E. Baker (2017) A molecular genetic toolbox for *Yarrowia lipolytica*. *Biotechnol*. *Biofuels* 10:2
- 481. Y. Chen; Y. Wang; J. Nielsen (2017) Systematic inference of functional phosphorylation events in yeast metabolism. *Bioinformatics* 33:1995-2001
- 482. P.-J. Lahtvee; B.J. Sanchez; A. Smialowska; S. Kasvandik; I. Elsemman; F. Gatto; **J. Nielsen** (2017) Absolute quantification of protein and mRNA abundances demonstrate variability in gene-specific translation efficiency in yeast. *Cell Systems* **4**:495-504
- 483. Y. Hu; J. Zhou; J. Bao; L. Huang; **J. Nielsen**; A. Krivoruchko (2017) Metabolic engineering of *Saccharomyces cerevisiae* for production of germacreme A, a precursor of beta-elemene. *J. Ind. Microbiol. Biotechnol.* **44**:1065-1072
- 484. T. Yu; Y.J. Zhou; L. Wenning; Q. Liu; A. Krivoruchko; V. Siewers; J. Nielsen; F. David (2017) Metabolic engineering of *Saccharomyces cerevisiae* for production of very long chain fatty acid-derived chemicals. *Nature Com.* 8:15587

- 485. P.G. Teixeira; R. Ferreira; Y.J. Zhou; V. Siewers; J. Nielsen (2017) Dynamic regulation of fatty acid pools for improved production of fatty alcohols in *Saccharomyces cerevisiae*. *Microb. Cell Fac.* **16**:45
- 486. E.D. Jensen; R. Ferreira; T. Jakociunas; D. Arsovska; L. Ding; J.D. Smith; F. David; **J. Nielsen**; M.K. Jensen; J.D. Keasling (2017) Transcriptional reprogramming in yeast using dCas9 and combinatorial gRNA strategies. *Microb. Cell Fac.* **16**:46
- 487. R. Babazadeh; P.-J. Lahtvee; C. Beck; M. Goksör; J. Nielsen; S. Hohmann (2017) The yeast osmostress response is carbon source dependent. *Sci. Rep.* **7**:990
- 488. M.-K. Kang; Y. Zhou; N.A. Buijs; J. Nielsen (2017) Functional screening of aldehyde decarboxylases for long-chain alkane production by *Saccharomyces cerevisiae*. *Microb. Cell Fac.* **16**:74
- 489. L. Väremo; T.I. Henriksen; C. Scheele; C. Broholm; M. Pedersen; M. Uhlen; B.K. Pedersen; J. Nielsen (2017) Type 2 diabetes and obesity are independently associated with similar inherent transcriptional reprogramming in human myocytes. *Genome Med.* **9**:47
- 490. J.C. Nielsen; S. Grijseels; S. Prigent; B. Ji; J. Dainat; K.F. Nielsen; J.C. Frisvad; M. Workman; **J. Nielsen** (2017) Global analysis of biosynthetic gene clusters reveals vast potential of secondary metabolite production in *Penicillium* species. *Nature Microbiol.* **2**:17044
- 491. J. Bao; M. Huang; D. Petranovic; **J. Nielsen** (2017) Moderate expression of *SEC16* increases protein secretion by *Saccharomyces cerevisiae*. *Appl. Environ. Microbiol.* **83**:e03400-16
- 492. E.J. Kerkhoven; Y.-M. Kim; S. Wei; C.D. Nicora; T.L. Filllmore; S.O. Purvine; B.-J. Webb-Robertson; R.D. Smith; S.E. Baker; T.O. Metz; **J. Nielsen** (2017) Leucine biosynthesis is involved in regulating high lipid accumulation in *Yarrowia lipolytica. mBio* **6**:e00857-17
- 493. P. Sen; A. Mardinoglu; **J. Nielsen** (2017) Selection of complementary foods based on optimal nutrition values. *Sci. Rep.* **7**:5413
- 494. A. Rodriguez; T. Strucko; S.G. Stahlhut; M. Kristensen; D.K. Svenssen; J. Forster; J. Nielsen; I. Borodina (2017) Metabolic engineering of yeast for fermentative production of flavonoids. *Biores. Technol.* **245**:1645-1654
- 495. A. Feizi; F. Gatto; M. Uhlen; J. Nielsen (2017) Human protein secretory pathway genes are expressed in a tissue-specific pattern to match secretion demand. *Systems Biol. Appl.* **3**:22
- 496. B.J. Sanchez; C. Zhang; A- Nilsson; P.-J. Lahtvee; E. Kerkhoven; J. Nielsen (2017) Improving the phenotype predictions of a yeast genome-scale metabolic model by incorporating enzymatic constraints. *Mol. Systems Biol.* 13:935
- 497. K. Thorell; J. Bengtsson-Palme; O. Liu; R.V. Gonzales; I. Nookaew; L. Rabeneck; L. Paszat; D. Graham; J. Nielsen; S, Lundin; Å. Sjöling (2017) *In vivo* analysis of the viable microbiota and *Helicobacter pylori* transcriptome in gastic infection and early stages of carcinogenesis. *Infect. Immun.* 85:e00031-17
- 498. S. Lee; C. Zhang; Z. Liu; M. Klevstig; B. Mukhopadhyay; M. Bergentall; R. Cinar; M. Ståhlman; N. Sikanic; J.K. Park; S. Deshmukh; A.M. Pour; T. Kuijpers; M. Grøtli; S.J. Elsässer; B.D. Piening; M. Snyder; J. Nielsen; F. Bäckhed; G. Kunos; M. Uhlen; J. Boren; A. Mardinoglu (2017) Network analysis identify liver-specific targets for treating liver diseases. *Mol. Sys. Biol.* 13:938
- 499. M. Huang; J. Bao; B.M. Hallström; D. Petranovic; J. Nielsen (2017) Efficient protein production by yeast requires global tuning of metabolism. *Nature Com.* 8:1131
- 500. G. Liu; Y. Chen; N. Færgeman; **J. Nielsen** (2017) Elimination of the last reactions in ergosterol biosynthesis alters the reistance of *Saccharomyces cerevisiae* to multiple stress. *FEMS Yeast Res.* **17**:fox063
- 501. R. Ferreira; F. Gatto; J. Nielsen (2017) Exploiting off-targeting in guide-RNAs for CRISPR systems for editing of multiple genes. FEBS Let. 591:3288-3295
- 502. Z. Zhu; Y.J. Zhou; M.-K. Kang; A. Krivoruchko; N.A. Buijs; J. Nielsen (2017) Enabling the synthesis of medium chain alkanes and 1-alkenes in yeast. *Met. Eng.* 44:81-88
- 503. S. Grijseels; J.C. Nielsen; **J. Nielsen**; T.O. Larsen; J.C. Frisvad; K.F. Nielsen; R. Frandsen; M. Workman (2017) Physiological characterization of secondary metabolite producing *Penicillium* cell factories. *Fungal Biol. Biotechnol.* **4**:8
- 504. A. Rodriguez; Y. Chen; S. Khoomrung; E. Özdemir; I. Borodina; J. Nielsen (2017) Comparison of the metabolic response to over-production of p-coumaric acid in two yeast strains. *Met. Eng.* 44:265-272

- 505. S. Lee; C. Zhang; A. Muhammad; L. Zhengtao; R. Benfeitas; G. Bidkhor; S. Deshmukh; M.A. Shobky; A. Lovric; J. Boren; J. Nielsen; M. Uhlen; A Mardinoglu (2018) TCSBN: a database of tissue and cancer specific biological networks. *Nucl. Acid. Res.* 46:D595-D600
- 506. E. Brunk; S. Sahoo; D.C. Zielinski; A. Altunkaya; A. Dräger; N. Mih; F. Gatto; A. Nilsson; G.A.P. Gonzalez; M.K. Aurich; A. Prlic; A. Sastry; A.D. Danielsdottir; A. Heinken; A. Noronha; P.W. Rose; S.K. Burley; R.M.T. Fleming; J. Nielsen; I. Thiele; B.O. Palsson (2018) Recon3D enables a three-dimensional view of gene variation in human metabolism. *Nature Biotechnol.* 36:272-281
- 507. R. Ferreira; C. Skrekas; J. Nielsen; F. David (2018) Multiplexed CRISPR/Cas9 genome editing and gene regulation using Csy9 in *Saccharomyces cerevisiae*. ACS Syn. Biol. 7:10-15
- 508. Y. Hu; Z. Zhu; J. Nielsen; V. Siewers (2018) Transporter engineering for improved fatty alcohol secretion in yeast. *Meg. Eng.* 45:51-58
- 509. D. Bergenholm; M. Gossing; Y. Wei; V. Siewers; J. Nielsen (2018) Modulation of saturation and chain length of fatty acids in *Saccharomyces cerevisiae* for production of cocoa butter-like lipids. *Biotechnol. Bioeng.* 115:932-942
- 510. A. Mardinoglu; H. Wu; E. Bjornson; C. Zhang; A. Hakkarainen; S.M. Räsänen; S. Lee; R.M. Mancina; M. Bergentall; K.H. Pietiläinen; S. Söderlund; N. Matikainen; M. Ståhlman; P.-O. Bergh; M. Adiels; B.D. Piening; M. Graner; N. Lundbom; K.J. Williams; S. Romeo; J. Nielsen; M. Snyder; M. Uhlen; G. Bergström; R. Perkins; H.-U. Marschall; F.

- Bäckhed; M.-R. Taskinen; J. Boren (2018) An integrated understanding of the rapid metabolic benefits of a carbohydrate restricted diet on hepatic steatosis in humans. *Cell Metab.* **27**:559-571
- 511. Y. Zhou; Y. Hu; Z. Zhu; V. Siewers; J. Nielsen (2018) Engineering of 1-alkene biosynthesis and secretion by dynamic regulation in yeast. ACS Syn. Biol. 7:584-590
- 512. R. Ferreira; P.G. Teixeira; V. Siewers; J. Nielsen (2018) Redirection of lipid flux towards phospholipids in yeast increases fatty acid turnover and secretion. *Proc. Nat. Acad. Sci. USA* 115:1262-1267
- 513. P. Jenjaroenpun; T. Wongsurawat; R. Pereira; P. Patumcharoenpol; D.W. Ussery; J. Nielsen; I. Nookaew (2018) Complete genomic and transcriptional landscape analysis using third generation sequencing: A case study of Saccharomyces cerevisiae CEN.PK113-7D. Nuc. Acid Res. 46:e38
- 514. A. Azimi; S. Caramuta; R. Jafari; B.S. Ludlow; J. Boström; J. Robinson; F. Edfors; R. Tuominen; **J. Nielsen**; J. Hansson; S.E. Brage; M. Altun; J. Lehtiö; M. Uhlen; G. Maddalo (2018) Targeting CDK2 overcomes melanoma reistance against BRAF and Hsp90 inhibitors. *Mol. Systems Biol.* **14**:e7858
- 515. Y. Wei; D. Bergenholm; M. Gossing; V. Siewers; J. Nielsen (2018) Expression of cocoa genes in *Saccharomyces cerevisiae* improves cocoa butter production. *Microb. Cell Fac.* 17:11
- 516. R. Ferreira; P.G. Teixeira; M. Gossing; F. David; V. Siewers; J. Nielsen (2018) Metabolic engineering of *Saccharomyces cerevisiae* for overproduction of triacylglycerols. *Met. Eng. Com.* **6**:22-27
- 517. D. Carmona-Gutierrez; M.A. Bauer; A. Zimmermann; A. Aguilera; N. Austriaco; K. Ayscough; R. Balzan; S. Bar-Nun; A. Barrientos; P. Belenky; M. Blondel; R.J. Braun; M. Breitenbach; W.C. Burhans; S. Büttner; D. Cavalieri; M. Chang; K.F. Cooper; M. Côrte-Real; V. Costa; C. Cullin; I. Dawes; J. Dengjel; M.B. Dickman; T. Eisenberg; B. Fahrenkrog; N. Fasel; K.-U. Fröhlich; A. Gargouri; S. Giannattasio; P. Goffrini; C.W. Gourlay; C.M. Grant; M.T. Greenwood; N. Guaragnella; T. Heger; J. Heinisch; E. Herker; J.M. Herrmann; S. Hofer; A. Jiménez-Ruiz; H. Jungwirth; K. Kainz; D.P. Kontoyiannis; P. Ludovico; S. Manon; E. Martegani; C. Mazzoni; L.A. Megeney; C. Meisinger; J. Nielsen; T. Nyström; H.D. Osiewacz; T.F. Outeiro; H.-O. Park; T. Pendl; D. Petranovic; S. Picot; P. Polčic; T. Powers; M. Ramsdale; M. Rinnerthaler; P. Rockenfeller; C. Ruckenstuh; R. Schaffrath; M. Segovia; F.F. Severin; A. Sharon; S.J. Sigrist; C. Sommer-Ruck; M.J. Sousa; J.M. Thevelein; K. Thevissen; V. Titorenko; M.B. Toledano; M. Tuite; F.-N. Vögtle; B. Westermann; J. Winderickx; S. Wissing; S. Wölfl; Z.J. Zhang; R.Y. Zhao; B. Zhou; L. Galluzzi; G. Kroemer; F. Madeo (2018) Guidelines and recommendations on yeast cell death nomenclature. *Microb. Cell* 5:4-31
- 518. J. Bao; M. Huang; D. Petranovic; **J. Nielsen** (2018) Balanced trafficking between the ER and the Golgi apparatus increases protein secretion in yeast. *AMB Express* **8**:37
- 519. P. Das; B. Ji; P. Kovatcheva-Datchary; F. Bäckhed; J. Nielsen (2018) *In vitro* co-cultures of human gut bacterial species as predicted from co-occurrence network analysis. *PLOS One* **13**:e0195161
- 520. F. Gatto; K.A. Blum; S.S. Hosseini; M. Ghanaat; M. Kashan; F. Maccari; F. Galeotti; J. Hsieh; N. Volpi; A.A. Hakimi; **J. Nielsen** (2018) Plasma glycosaminoglycans as diagnostic and prognostic biomarkers in surgically treated renal cell carcinoma. *Eur. Urol. Onc.* 1:364-377
- 521. J. Maury; S. Kannan; N.B. Jensen; F.K. Öberg; K.R. Kildegaard; J. Förster; J. Nielsen; C.T. Workman; I. Borodina (2018) Glucose-dependent promoters for dynamic regulation of metabolic pathways. *Fron. Bioeng. Biotechnol.* **6**:63
- 522. P. Teixeira; F. David; V. Siewers; **J. Nielsen** (2018) Engineering lipid droplet assembly mechanisms for improved triacylglycerols accumulation in *Saccharomyces cerevisiae*. *FEMS Yeast Res.* **18**:foy060
- 523. S. Prigent; J.C. Nielsen; J.C. Frisvad; J. Nielsen (2018) Reconstruction of 24 *Penicillium* genome-scale metabolic models shows diversity based on their secondary metabolism. *Biotechnol. Bioeng.* 115:2604-2612
- 524. Z. Dai; M. Huang; Y. Chen; V. Siewers; J. Nielsen (2018) Global rewiring of cellular metabolism renders Saccharomyces cerevisiae Crabtree-negative. Nature Com. 9:3059
- 525. G. Bidkhori; R. Benfeitas; E. Elmas; M.N. Karadoudi; M. Arif; M. Uhlen; J. Nielsen; A. Mardinoglu (2018) Metabolic network-based identification and prioritization of anti-cancer targets based on expression data in hepatocellular carcinoma. *Fron. Phys.* 9:916
- 526. T. Yu; Y. Zhou; M. Huang; Q. Liu; R. Pereira; F. David; **J. Nielsen** (2018) Reprogramming yeast metabolism from alcoholic fermentation to lipogenesis. *Cell* **174**:1-10
- 527. D. Bergenholm; G. Liu; P. Holland; **J. Nielsen** (2018) Reconstruction of a global transcriptional regulatory network for control of lipid metabolism in yeast using ChIP-exo. *mSystems* **3**:e00215-17
- 528. P. Babaei; S. Shoaei; B. Ji; **J. Nielsen** (2018) Challenges in modeling the human gut microbiome. *Nature Biotechnol.* **16**:682-686
- 529. M. Kumar; B. Ji; P. Babaei; P. Das; D. Lappa; G. Ramakrishnan; T.E. Fox; R. Haque; W.A. Petri Jr.; F. Bäckhed; J. Nielsen (2018) Gut microbiota dysbiosis is associated with malnutrition and reduced plasma amino acid levels: Lessons from genome-scale metabolic modeling. *Met. Eng.* **49**:128-142
- 530. T.M. Souza-Moreira; C. Navarrete; X. Chen; C.F. Zanelli; S.R. Valentini; M. Furlan; **J. Nielsen**; A. Krivoruchko (2018) Screening of 2A peptides for polycistronic gene expression in yeast. *FEMS Yeast Res.* **18**:foy036
- 531. L. Ouyang; P. Holland; H. Lu; D. Bergenholm; J. Nielsen (2018) Integrated analysis of the yeast NADPH-regulator Stb5 reveals distinct differences in NADPH requirements and regulation in different states of yeast metabolism. FEMS Yeast Res. 18:foy091
- 532. M. Gossing; A. Smialowska; J. Nielsen (2018) Impact of forced fatty acid synthesis on metabolism and physiology of *Saccharomyces cerevisiae*. FEMS Yeast Res. 18:foy096
- 533. H. Wang; S. Marcisauskas; D. Hermansson; R. Agren; **J. Nielsen**; E. Kerkhoven (2018) The RAVEN Toolbox 2.0: A versatile platform for metabolic network reconstruction and a case study on *Streptomyces coelicolor*. *PLOS Comp. Biol.* **14**:e1006541
- 534. M. Huang; G. Wang; J. Qin; D. Petranovic; J. Nielsen (2018) Engineering the protein secretory pathway of *Saccharomyces cerevisiae* enables improved protein production. *Proc. Nat. Acad. Sci. USA* **115**:E11025-E11032

- 535. K. Pomraning; E. Bredeweg; E. Kerkhoven; K. Berrie; S. Haridas; H. Hundley; K. LaButti; A. Lipzen; M. Yan; J. Magnuson; B. Simmons; I. Grigoriev; **J. Nielsen**; S. Baker (2018) Regulation of yeast to hyphae transition in *Yarrowia lipolytica. mSphere* 3:e00541-18
- 536. G. Bidkhori; R. Benfeitas; M. Klevstig; J. Nielsen; M. Uhlen; J. Boren; A. Mardinoglu (2018) Metabolic network stratification of the hepatocellular carcinoma reveals three distrinct tumor subtypes. *Proc. Nat. Acad. Sci. USA* 115:E11874-E11883
- 537. Z.P. Guo; S. Khoomrung; **J. Nielsen**; L. Olsson (2018) Changes in lipid metabolism convey acid tolerance in *Saccharomyces cerevisiae*. *Biotechnol. Biofuels* **11**:297
- 538. S. Grijseels; C. Pohl; J.C. Nielsen; Z. Wasil; Y. Nygård; J. Nielsen; J.C. Frisvad; K.F. Nielsen; M. Workman; T.O. Larsen; A.J.M. Driessen; R.J.N. Frandsen (2018) Identification of the decumbenone biosynthetic gene cluster in *Penicillium decumbens* and the importance for production of calbistrin. *Fun. Biol. Biotechnol.* 5:18

- 539. T. Ma; B. Shi; Z. Ye; X. Li; M. Liu; Y. Chen; J. Xia; J. Nielsen; Z. Deng; T. Liu (2018) Lipid engineering combined with systematic metabolic engineering of *Saccharomyces cerevisiae* for high-yield production of lycopene. *Met. Eng.* 52:134-142
- 540. P. Holland; J. Nielsen; T.D.G.A. Mondeel; M. Barberis (2019) Coupling cell division to metabolic pathways through transcription. *Encyclopedia Bioinf. Comp. Biol.* **3**:74-92
- 541. B.J. Sanchez; F. Li; E. Kerkhoven; J. Nielsen (2019) SLIMEr: Probing flexibility of lipid metabolism in yeast with an improved constraint-based modeling framework. *BMC Systems Biol.* 13:4
- 542. C. Börlin; N. Cvetesic; P. Holland; D. Bergenholm; V. Siewers; B. Lenhard; J. Nielsen (2019) Saccharomyces cerevisiae displays a stable transcription start site landscape in multiple conditions. FEMS Yeast Res. 19:foy128
- 543. R. Benfeitas; G. Bidkhori; B. Mukhopadhyay; M. Klevstig; M. Arif; C. Zhang; S. Lee; R. Cinar; J. Nielsen; M. Uhlen; J. Boren; G. Kunos; A. Mardinoglu (2019) Characterization of heterogeneous redox responses in hepatocellular carcinoma patients using network analysis. *E Biomed.* 40:471-487
- 544. Z. Liu; C. Zhang; S. Lee; W. Kim; M. Klevstig; A.M. Harzandi; N. Sikanic; M. Arif; M. Stählman; **J. Nielsen**; M. Uhlen; J. Boren; A. Mardinoglu (2019) Pyruvate kinase L/R is a regulator of lipid metabolism and mitochondrial function. *Met. Eng.* **52**:263-272
- 545. D. Bergenholm; G. Liu; D. Hansson; J. Nielsen (2019) Construction of mini-chemostats for high-throughput strain characterization. *Biotechnol. Bioeng.* 116:1029-1038
- 546. A. Bergman; J. Hellgren; T. Moritz; V. Siewers; **J. Nielsen**; Y. Chen (2019) Heterologous phosphoketolase expression redirects flux towards acetate, perturbs sugar phosphate pools and increases respiratory demand in *Saccharomyces cerevisiae*. *Microb. Cell Fac.* **18**:25
- 547. P. Kovatcheva-Datchary; S. Shoaie; S. Lee; A. Wahlström; I. Nookaew; A. Hallen; R. Perkins; **J. Nielsen**; F. Bäckhed (2019) Simplified intestinal microbiota to study microbe-diet-host interactions in a mouse model. *Cell Rep.* **26**:3772-3783
- 548. J.L. Robinson; A. Feizi; M. Uhlen; J. Nielsen (2019) A systematic investigation of the malignant functions and diagnostic potential of the cancer secretome. *Cell Rep.* 10:2622-2635
- 549. Y. Wei; B. Ji; V. Siewers; D. Xu; B.A. Halkier; J. Nielsen (2019) Identification of genes involved in shea butter biosynthesis from *Vitellaria paradoxa* fruits through transcriptomics and functional heterologous expression. *Appl. Mic. Biotechnol.* 103:3727-3736
- 550. L. Wenning; C.S. Ejsing; F. David; R.R. Sprenger; J. Nielsen; V. Siewers (2019) Increasing jojoba-like wax ester production in *Saccharomyces cerevisiae* by enhancing very long-chain, monosaturated fatty acid synthesis. *Microb. Cell Factories* 18:49
- 551. P. Das; P. Babaei; **J. Nielsen** (2019) Metagenomic analysis of microbe-mediated vitamin metabolism in the human gut microbiome. *BMC Genom.* **20**:208
- 552. J.C. Nielsen; S. Prigent; S. Grijseels; M. Workman; B. Ji; **J. Nielsen** (2019) Comparative transcriptome analysis shows conserved metabolic regulations during production of secondary metabolites in filamentous fungi. *mSystems* **4**:e00012-19
- 553. G. Wang; G. S.M. Björk; M. Huang; Q. Liu; K. Campbell; J. Nielsen; H.N. Jönsson; D. Petranovic (2019) RNAi expression tuning, microfluidic screening, and genome recombineering for improved protein production in *Saccharomyces cerevisiae*. *Proc. Nat. Acad. Sci. USA* 116:9324-9332
- 554. A. Bergman; D. Vitay; J. Hellgren; Y. Chen; J. Nielsen; V. Siewers (2019) Effects of overexpression of *STB5* in *Saccharomyces cerevisiae* on fatty acid biosynthesis, physiology and transcriptome. *FEMS Yeast Res.* 19:foz027
- 555. P. Holland; D. Bergenholm; C. Börlin; G. Liu; J. Nielsen (2019) Predictive models of eukaryotic transcriptional regulation reveals changes in transcription factor roles and promoter usage between metabolic conditions. *Nuc. Acid Res.* 47:4986-5000
- 556. R. Pereira; P. Vilaca; P. Maia; **J. Nielsen**; I. Rocha (2019) TDPS Turnover dependent phenotypic simulation: a quantitative constraint-based simulation method that accommodates all main strain design strategies. *ACS Syn. Biol.* **8**:976-988
- 557. I.A. Tiukova; M.E. Pettersson; M.P. Hoeppner; R.-A. Olsen; M. Käller; J. Nielsen; J. Dainat; H. Lantz; J. Söderberg; V. Passoth (2019) Chromosomal genome assembly of the ethanol production strain CBS 11270 indicates a highly dynamic genome structure in the yeast species *Bretanomyces bruxellensis*. *PLOS One* 14:e0215077
- 558. I.A. Tiukova; J. Brandenburg; J. Blomqvist; S. Samples; N. Mikkelsen; M. Skaugen; M.Ø. Arntzen; **J. Nielsen**; M. Sandgren; E.J. Kerkhoven (2019) Proteome analysis of xylose metabolism in *Rhodotorula toruloides* during lipid production. *Biotechnol. Bioeng.* **12**:137
- 559. T.I. Henriksen; L.V. Wigge; J. Nielsen; B.K. Pedersen; M. Sandri; C. Scheele (2019) Dysregulated autophagy in muscle precursor cells from humans with type 2 diabetes. *Sci. Rep.* **9**:8169

- 560. P. Das; S. Marcisauskas; B. Ji; J. Nielsen (2019) Metagenomic analysis of bile salt biotransformation in the human gut microbiome. *BMC Genom.* 20:517
- 561. T.D.G.A. Mondeel; P. Holland; J. Nielsen; M. Barberis (2019) ChIP-exo analysis highlights Fkh1 and Fkh2 transcription factors as hubs that integrate multi-scale networks in budding yeast. *Nuc. Acid Res.* 47:7825-7841
- 562. H. Lu; F. Li; B.J. Sanchez; Z. Zhu; G. Li; I. Domenzain; S. Marcisauskas; P.M. Anton; D. Lappa; C. Lieven; M.E. Beber; N. Sonnenschein; E.J. Kerkhoven; J. Nielsen (2019) A consensus *S. cerevisiae* metabolic model Yeast8 and its ecosystem for comprehensively probing cellular metabolism. *Nature Com.* 10:3586
- 563. Y. Dabirian; P.G. Teixeira; J. Nielsen; V. Siewers; F. David (2019) FadR-based biosensor assisted screening for genes enhancing fatty acyl-CoA pools in *Saccharomyces cerevisiae*. ACS Syn. Biol. 8:1788-1800
- 564. Y. Chen; J. Nielsen (2019) Energy metabolism controls phenotypes by protein efficiency and allocation. *Proc. Nat. Acad. Sci. USA* 116:17592-17597
- 565. C. Börlin; D. Bergenholm; P. Holland; **J. Nielsen** (2019) A bioinformatics pipeline to analyze ChIP-exo datasets. *Biol. Meth. Protoc.* **4**:bpz011
- 566. Y. Dabirian; X. Li; Y. Chen; F. David; J. Nielsen; V. Siewers (2019) Expanding the dynamic range of a transcription factor-based biosensor in *Saccharomyces cerevisiae*. ACS Syn. Biol. 8:1968-1975
- 567. I. Tiukova; S. Prigent; J. Nielsen; M. Sandgren; E.J. Kerkhoven (2019) Genome-scale model of *Rhodotorula toruloides* metabolism. *Biotechnol. Bioeng.* **116**:3396-3408
- 568. R. Pereira; Y. Wei; E. Mohamed; M. Radi; C. Malina; M.J. Herrgård; A.M. Feist; J. Nielsen; Y. Chen (2019) Adaptive laboratory evolution of tolerance to dicarboxylic acids in *Saccharomyces cerevisiae*. *Met. Eng.* **56**:130-141
- 569. G. Li; B. Ji; J. Nielsen (2019) The pan-genome of Saccharomyces cerevisiae. FEMS Yeast Res. 19:foz064
- 570. R. Ferreira; C. Skrekas; A. Hedin; B. Sanchez; V. Siewers; J. Nielsen; F. David (2019) Model-assisted fine-tuning of central carbon metabolism in yeast through dCas9-based regulation. ACS Syn. Biol. 8:2457-2463
- 571. G. Li; K.S. Rabe; J. Nielsen; M.K.M. Engqvist (2019) Machine learning applied to predicting microorganism growth temperatures and enzyme catalytic optima. ACS Syn. Biol. 8:1411-1420
- 572. Q. Liu; T. Yu; X. Li; Y. Chen; K. Campbell; J. Nielsen; Y. Chen (2019) Rewiring carbon metabolism in yeast for high level production of aromatic chemicals. *Nature Com.* **10**:4976
- 573. S. Marcisauskas; B. Ji; J. Nielsen (2019) Reconstruction and analysis of a *Klyveromyces marxianus* genome-scale metabolic model. *BMC Bioinfor*. **20**:551
- 574. I. Tiukova; I Møller-Hansen; Z.M. Belew; D. Darbani: E. Boles; E. Nour; H. Hussam; T. Linder; **J. Nielsen**; I. Borodina (2019) Identification and characterization of two high-affinity glucose transporters from the spoilage yeast *Brettanomyces bruxellensis. FEMS Lett.* **366**:fnz222
- 575. I.A. Tiukova; H. Jiang; J. Dainat; M.P. Hoeppner; H. Lantz; J. Piskur; M. Sandgren; **J. Nielsen**; Z. Gu; V. Passoth (2019) Assembly and analysis of the genome sequence of the yeast *Brettanomyces naardenensis* CBS7540. *Microorganisms* 7:489
- 576. G.M. Borja; A. Rodriguez; K. Campbell; I. Borodina; Y. Chen; **J. Nielsen** (2019) Metabolic engineering and transcriptomic analysis of *Saccharomyces cerevisiae* producing p-coumaric acid from xylose. *Microb. Cell Fac.* **18**:191
- 577. M. Uhlen; H. Tegel; Å. Sivertsson; C.-C. Kuo; J.M. Gutierrez; N. Lewis; B. Forsström; M. Dannemeyer; L. Fagerberg; J. Rockberg; M. Malm; H. Vunk; F. Edfors; A. Hober; E. Sjöstedt; J. Mulder; A. Mardinoglu; J. Schwenk; P. Nilsson; M. Zwahlen; K. von Feilitzen; C. Lindskog; F. Ponten; J. Nielsen; B.G. Voldborg; B.O. Palsson; A.-L.R. Volk; M. Lundqvist; A. Berling; A.-S. Svensson; S. Kanje; H. Enstedt; D. Afshari; S. Ekblad; J. Scheffel; L. Xu; R. Mihai; L. Bremer; M. Westin; M. Muse; L. Mayr; S. Knight; S. Göpel; R. Davies; P. Varley; D. Hatton; J.O. Takanen; L.H. Schiavone; S. Hober (2019) The human secretome. *Science Signaling* 12:eaaz0274
- 578. L. Ye; P. Das; P. Li; B. Ji; J. Nielsen (2019) Carbohydrate active enzymes are affected by diet transition from milk to solid food in infant gut microbiota. *FEMS Microb. Eco.* **95**:fiz159
- 579. G. Othoum; S. Prigent; A. Derouiche; L. Shi; A. Bokhari; S. Alamoudi; S. Bougouffa; X. Gao; R. Hoehndorf; S.T. Aroid; T. Gojobori; H. Hirt; F.F. Lafi; **J. Nielsen**; V.B. Bajic; I. Mijakovic; M. Essack (2019) Comparative genomics study reveals Red Sea *Bacillus* with characteristics associated with potential microbial cell factories (MCFs). *Sci. Rep.* 9:19254
- 580. A. Nilsson; E. Björnson; M. Flockhart; F.J. Larsen; J. Nielsen (2019) Complex I is bypassed during high intensity exercise. *Nature Com.* 10:5072

- 581. N. Raethong; H. Wang; J. Nielsen; W. Vongsangnak (2020) Optimizing cultivation of *Cordyceps militaris* for fast growth and cordyceptin overproduction using rational design of synthetic media. *Comp. Struc. Biotechnol. J.* 18:1-8
- 582. J. Gutierrez; A. Feizi; S. Li; T. Kallehauge; G.M. Lee; B. Palsson; J. Nielsen; N. Lewis (2020) Genome-scale reconstruction of the mammalian secretory pathway predicts metabolic costs and limitations of protein secretion.

 Nature Com. 11:68
- 583. F. Gatto; R. Ferreira; J. Nielsen (2020) Pan-cancer analysis of the metabolic reaction network. Met. Eng. 57:51-62
- 584. Y. Liu; Q. Liu; A. Krivoruchko; S. Khoomrung; J. Nielsen (2020) Engineering yeast phospholipid metabolism for *de novo* oleoylethanolamide production. *Nature Chem. Biol.* 16:197-205
- 585. S. Khoomrung; I. Nookaew; P. Sen; T. Olafsdottir; J. Persson; T. Moritz; P. Andersen; A. Harandi; J. Nielsen (2019) Metabolic profiling and compound-class identification reveal alterations in serum triglyceride levels in mice immunized with human vaccine adjuvant Alum. J. Prot. Res. 19:269-278
- 586. C. Lieven; M.E. Beber; B.G. Olivier; F.T. Bergmann; M. Ataman; P. Babaei; J.A. Bartell; L.M. Blank; S. Chauhan; K. Correia; C. Diener; A. Dräger; B.E. Ebert; J.N. Edirisinghe; J.P. Faria; A.Feist; G. Fengos; R.M.T. Fleming; B. García-Jiménez; V. Hatzimanikatis; W. van Helvoirt; C.S. Henry; H. Hermjakob; M.J. Herrgård; A. Kaafarani; H.U. Kim; Z. King; J.J. Koehorst; M. König; S. Klamt; E. Klipp; M. Lakshmanan; D.-Y. Lee; S.Y. Lee; S. Lee; N.E.

- Lewis; F. Liu; H. Ma; D. Machado; R. Mahadevan; P. Maia; A. Mardinoglu; G.L. Medlock; J.M. Monk; J. Nielsen; L.K. Nielsen; J. Nogales; I. Nookaew; O. Resendis-Antonio; B.O. Palsson; J.A. Papin; K.R. Patil; M. Poolman; N.D. Price; A. Richelle; I. Rocha; B.J. Sánchez; P.J. Schaap; R.S.M. Sheriff; S. Shoaie; N. Sonnenschein; B. Teusink; P. Vilaça; J.O. Vik; J.A.H. Wodke; J.C. Xavier; Q. Yuan; M. Zakhartsev; C. Zhang (2019) Memote for standardized genome-scale metabolic model testing. *Nature Biotechnol.* 38:272-276
- 587. Z. Zhu; Y. Hu; P.G. Teixeira; R. Pereira; Y. Chen; V. Siewers; J. Nielsen (2020) Multidimensional engineering of *Saccharomyces cerevisiae* for efficient synthesis of medium-chain fatty acids. *Nature Cat.* **3**:64-74
- 588. S. Lam; S. Doran; H. Yuksel; O. Altay; H. Turkez; J. Nielsen; J. Boren; M. Uhlen; A. Mardinoglu (2020) Addressing the heterogeneity in liver diseases using biological networks. *Brief. Bioinform.* Bbaa002
- 589. X. Li; B. Turanli; K. Juszczak; W. Kim; M. Arif; Y. Sato; S. Ogawa; H. Turkez; **J. Nielsen**; J. Boren; M. Uhlen; C. Zhang; A. Mardinoglu (2020) Classification of clear renal cell carcinoma based on *PKM* alternative splicing. *Heliyon* 6:e03440
- 590. K. Campbell; J. Westholm; S. Kasvendik; F. Di Bartolomei; M. Mormino; J. Nielsen (2020) Building blocks are synthesized on demand during the yeast cell cycle. *Proc. Nat. Acad. Sci. USA* 117:7575-7583
- 591. F. Di Bartolomeo; C. Malina; K. Campbell; M. Mormini; J. Fuchs; E. Vorontsov; C.M. Gustafsson; J. Nielsen (2020) Absolute yeast mitochondrial proteome quantification reveals trade-off between biosynthesis and energy generation during diauxic shift. *Proc. Nat. Acad. Sci. USA* 117:7524-7535
- 592. C. Zhang; E. Björnson; M. Arif; A. Tebani; A. Lovric; R. Benfeitas; M. Ozcan; K. Juszczak; W. Kim; J.T. Kim; G. Bidkhori; M. Ståhlman; P.-O. Bergh; M. Adiels; H. Turkez; M.-R. Taskinen; J. Bosley; H.-U. Marschall; J. Nielsen; M. Uhlen; J. Boren; A. Mardinoglu (2020) The acute effect of naturally occurring metabolic cofactor supplementation. *Mol. Systems Biol.* 16:e9495
- 593. D.J. Cook; J. Kallus; R. Jörnsten; J. Nielsen (2020) Molecular natural history of breast cancer: leveraging transcriptomics to predict breast cancer diagnosis and aggressiveness. *Cancer Med.* 9:3551-3562
- 594. K.S. Pedersen; F. Gatto; B. Zerahn; J. Nielsen; B.K. Pedersen; P. Hojman; J. Gehl (2020) Exercise-mediated lowering of glutamine availability suppresses tumor growth and attenuates muscle wasting. iScience 23:100978
- 595. J.L. Robinson; P. Kocabas; H. Wang; P.-E. Cholly; D. Cook; A. Nilsson; M. Anton; R. Ferreira; I. Domenzain; V. Billa; A. Limeta; A. Hedin; J. Gustafsson; E.J. Kerkhoven; L.T. Svensson; B.O. Palsson; A. Mardinoglu; L. Hansson; M. Uhlen; J. Nielsen (2020) An Atlas of Human Metabolism. *Science Signal.* 13:eaaz1482
- 596. A. Nilsson; J.R. Haanstra; M. Engqvist; A. Gerding; B.M. Bakker; U. Klingmüller; B. Teusink; **J. Nielsen** (2020) Quantitative analysis of amino acid metabolism in liver cancer links glutamate excretion to nucleotide synthesis. *Proc. Nat. Acad. Sci. USA* **117**:10294-10304
- 597. R. Yu; K. Campbell; R. Pereira; J. Björkeroth; Q. Qi; E. Vorontsov; C. Sihlbom; J. Nielsen (2020) Nitrogen limitation reveals large reserves in metabolic and translational capacities of yeast. *Nature Com.* **11**:1881
- 598. T:W. Doughty; I. Domenzain; A. Millan-Oropeza; N. Montini; P.A. de Groot; R. Pereira; **J. Nielsen**; C. Henry; J.-M.G. Daran; V. Siewers; J.P. Morriseey (2020) Stress-induced expression is enriched for evolutionarily young genes in diverse budding yeasts. *Nature Com.* **11**:2144
- 599. L. Dzanaeva; B. Kruk; J. Ruchala; J. Nielsen; A. Sibirny; K. Dmytruk (2020) The role of peroxisomes in xylose alcoholic fermentation in the engineered *Saccharomyces cerevisiae*. *Cell Biol. Int.* **44**:1606-1615
- 600. S. Vieira-Silva; G. Falony; E. Belda; T. Nielsen; J. Aron-Wisnewsky; R. Chakaroun; S.K. Forslund; K. Assmann; M. Valles-Colomer; T.T.D. Nguyen; S. Proost; E. Prifti; V. Tremaroli; N. Pons: E. Le Chatelier; F. Andreelli; J.-P. Bastard; L.P. Coelho; N. Galleron; T.H. Hansen; J.-S. Hulot1; C. Lewinter; H.K. Pedersen; B. Quinquis; C. Rouault; H. Roume; J.-E. Salem; N.B. Søndertoft; S. Touch; MetaCardis Consortium*, M.-E. Dumas; S.D. Ehrlich; P. Galan; J.P. Gøtze; T. Hansen; J.J. Holst; L. Køber; I. Letunic; J. Nielsen; J.-M. Oppert; M. Stumvoll; H. Vestergaard; J.-D. Zucker; P. Bork; O. Pedersen; F. Bäckhed; K. Clément; J. Raes (2020) Statin therapy is associated with a lower prevalence of gut microbiota dysbiosis. *Nature* 581:310-315
- 601. J. Hou; C. Gao; L. Guo; J. Nielsen; Q. Ding; W. Tang; G. Hu; X. Chen; L. Liu (2020) Rewiring carbon flux in *Escherichia coli* using a bifunctional molecular switch. *Met. Eng.* **61**:47-57
- 602. C. Börlin; J. Nielsen; V. Siewers (2020) The transcription factor Leu3 shows differential binding behavior in response to changing leucine availability. *FEMS Lett.* **367**:fnaa107
- 603. M. Babaei; G.B. Zamfir; X. Chen; H.B. Christensen; M. Kristensen; J. Nielsen; I. Borodina (2020) Metabolic engineering of *Saccharomyces cerevisiae* for rosmarinic acid production. *ACS Syn. Biol.* **9**:1978-1988
- 604. J. Björkeroth; K. Campbell; C. Malina; R. Yu; F. Di Bartolomeo; J. Nielsen (2020) Proteome re-allocation from amino acid biosynthesis to ribosomes enables yeast to grow faster in rich media. *Proc. Nat. Acad. Sci. USA* 117:21804-21812
- 605. A.A.-Haleem; V. Ravikumar; B. Ji; K. Mineta; X. Gao; **J. Nielsen**; T. Gojobori; I. Mijakovic (2020) Integrated metabolic modeling, culturing and transcriptomics explains enhanced virulence of *V. cholerae* during co-infection with ETEC. *mSystems* **5**:e00491-20
- 606. J. Zhang; S. Petersen; T. Radivojevic; A. Ramirez; A. Perez-Manriquez; E. Abeliuk; B. Sanchez; Z. Costello; Y. Chen; M. Fero; M.G. Martin; **J. Nielsen**; J. Keasling; M. Jensen (2020) Combining mechanistic and machine learning models for predictive engineering and optimization of tryptophan metabolism. *Nature Com.* **11**:4880
- 607. Y. Hu; Z. Zhu; D. Gradischnig; M. Winkler; **J. Nielsen**; V. Siewers (2020) Engineering carboxylic acid reductase for selective synthesis of medium-chain fatty alcohols in yeast. *Proc. Nat. Acad. Sci. USA* **17**:22974-22983
- 608. S. Vaga; S. Lee; B. Ji; A. Andreasson; N.J. Talley; L. Agreus; G. Bidkhori; P. Kovatcheva-Datchary; J. Park; D. Lee; G. Proctor; D. Eherlich: J. Nielsen; L. Engstrand; S. Shoaie (2020) Composition and functional differences of the mucosal microbiota along the intestine of healthy individuals. *Sci. Rep.* 10:14977
- 609. J. Hellgren; A. Godina; O. Vidalin; **J. Nielsen**; V. Siewers (2020) Promiscuous phosphoketolase and metabolic rewiring enables non-oxidative glycolysis in yeast for high yield production of acetyl-CoA products. *Met. Eng.* **62**:150-160

- 610. J. Gustafsson; F. Held; J. Robinson; E. Björnson; R. Jörnsten; **J. Nielsen** (2020) Sources of variation in cell-type RNA-seq profiles. *PLOS One* **15**:e0239495
- 611. Q. Qi; F. Li; R. Yu; M.K.M. Engqvist; V. Siewers; J. Fuchs; J. Nielsen (2020) Different routes of protein folding converge on improved protein production in *Saccharomyces cerevisiae*. mBio 11:e02743-20
- 612. R. Pereira; E. Mohamed; M. Radi; M.J. Herrgård; A.M. Feist; J. Nielsen; Y. Chen (2020) Elucidating aromatic acid tolerance at low pH in *Saccharomyces cerevisiae* using adaptive laboratory evolution. *Proc. Nat. Acad. Sci. USA* 117:27954-27961
- 613. Y. Zhang; M. Su; N. Qin; J. Nielsen; Z. Liu (2020) Expressing a cytosolic pyruvate dehydrogensase complex to increase free fatty acid production in *Saccharomyces cerevisiae*. *Microb. Cell Fac.* 19:226
- 614. A. Limeta; B. Ji; M. Levin; F. Gatto; J. Nielsen (2020) Meta-analysis of the gut microbiota in predicting response to cancer immunotherapy in metastatic melanoma. *J. Clin. Inv. Insight* 5:e140940
- 615. M. Malm; R. Saghaleyni; M. Lundqvist; M. Giudici; V. Chotteau; R. Field; P.G. Varley; D. Hatton; L. Grassi; T, Svensson; J. Nielsen; J. Rockberg (2020) Evolution from adherent to suspension systems biology of HEK293 cell line development. *Sci. Rep.* 10:18996
- 616. A. Molinaro; P.B. Lassen; M. Henricsson; H. Wu; S. Adriouch; E. Belda; R. Chakaroun; T. Nielsen; P.-O. Bergh; C. Rouault; S. André; F. Marquet; F. Andreelli; J.-E. Salem; K. Assmann; J.-P. Bastard; S. Forslund; E. Le Chatelier; G. Falony; N. Pons; E. Prifti; B. Quinquis; H. Roume; S. Vieira-Silva; T.H. Hansen; H.K. Pedersen; C. Lewinter; N.B. Sønderskov; MetaCardis Consortium; L. Køber; H. Vestergaard; T. Hansen; J.-D. Zucker; P. Galan; M.-E. Dumas; J. Raes; J.-M. Oppert; I. Letunic; J. Nielsen; P. Bork; S. Dusko Ehrlich; M. Stumvoll; O. Pedersen; J. Aron-Wisnewsky; K. Clément; F. Bäckhed (2020) Imidazole propionate is increased in diabetes and associated with dietary patterns and altered microbial ecology. *Nature Com.* 11:5881
- 617. M. Zhao; Y. Zhao; Q. Hu; H. Iqbal; M. Yao; H. Liu; B. Qiao; C.A.E. Skovbjerg; J.C.F. Nielsen; J. Nielsen; R.J.N. Frandsen; Y. Yuan; J.D. Boeke (2020) Pathway engineering in yeast for synthesizing the complex polyketide bikaverin. *Nature Com.* 11:6197
- 618. J. Zrimec; C. Börlin; F. Buric; A.S. Muhammad; R. Chen; V. Siewers; V. Verendel; **J. Nielsen**; M. Töpel; A. Zelezniak (2020) Deep learning suggests that gene expression is encoded in all parts of a co-evolving interacting gene regulatory structure. *Nature Com.* **11**:6141
- 619. J. Gustafsson; J. Robinson; J.I. Diaz; E. Björnsson; R. Jörnsten; **J. Nielsen** (2020) DSAVE: Detection of misclassified cells in single-cell RNA-seq data. *PLOS ONE* **15**:e0243360
- 620. N. Qin; L. Li; X. Ji; X. Li; Y. Zhang; C. Larsson; Y. Chen; J. Nielsen; Z. Liu (2020) Rewiring central carbon metabolism ensures increased provision of acetyl-CoA and NADPH required for 3-OH-propionic acid production. *ACS Syn. Biol.* **9**:3236-3244
- 621. C. Ye; N. Xu; C. Gao; G. Liu; J. Xu; W. Zhang; X. Chen; J. Nielsen; L. Liu (2020) Comprehensive understanding of *Saccharomyces cerevisiaei* with whole-cell model WM_S288C. *Biotechnol. Bioeng.* 117:1562-1574

- 622. C.C. van Olden; A.W. van de Laar; A.S. Meijnikman; O. Aydin; N. van Olst; J. Hoozemans; L.M. de Brauw; S.C. Bruin; Y.I.Z. Acherman; J. Verheij; M. Hoogendoorn; J. Nielsen; T. Schwartz; A.K. Groen; V.E.A. Gerdes; F. Bäckhed; M. Nieuwdorp (2020) Systems Biology and Gut Microbiota in understanding the metabolic effects of Bariatric Surgery: The BARIA Longitudinal Cohort Study. J. Int. Med. 289:340-354
- 623. G. Li; Y. Hu; J. Zrimec; H. Lou; H. Wang; A. Zelezniak; B. Ji; J. Nielsen (2021) Bayesian genome scale modeling identifies termal determinants of yeast metabolism. *Nature Com.* **12**:190
- 624. H.G. Blitzblau; A. Consiglio; P. Teixeira; D.V. Crabtree; S. Chen: O. Konzokck; G. Chifamba; A. Su; A. Kamineni; K.M.S. MacEwen; M. Hamilton; V. Tsakraklides; **J. Nielsen**; V. Siewers; J. Shaw (2021) Production of 10-methyl branched fatty acids in yeast. *Biotechnol. Biofuels* 14:12
- 625. S. Blasche; Y. Kim; R. Mars; D, Machado; M. Maansson; E. Kafkia; A. Milanese; G. Zeller; B. Teusink; J. Nielsen; V. Benes; R. Neves; U. Sauer; K.R. Patil (2021) Metabolic cooperation and spatiotemporal niche partitioning in a kefir microbial community. *Nature Microbiol.* 6:196-208
- 626. X. Li: W. Kim; M. Arif; C. Gao; A. Hober; D. Kotol; L. Strandberg; B. Forsström; A. Sivertsson; P. Oksvold; H. Turkez; M. Grottli; Y. Sato; H. Kume; S. Ogawa; J. Boren; J. Nielsen; M. Uhlen; Z. Chang; A. Mardinoglu (2021) Dicovery of functional alternatively spliced PKM transcripts in human cancers. *Cancers* 13:348
- 627. B.J. Sanchez; P.-J. Lahtvee; K. Campbell; S. Kasvandik; R. Yu; I. Domenzain; A. Zelezniak; **J. Nielsen** (2021) Benchmarking accuracy and precision of intensity-based absolute quantification of protein abundances in *Saccharomyces cerevisiae*. *Proteomics* **21**:2000093
- 628. J. Qin, A. Krivoruchko; B. Ji; Y. Chen; M. Kristensen, E. Özdemir; J. D. Keasling; M.K. Jensen; **J. Nielsen** (2021) Engineering yeast metabolism for the discovery and production of polyamines and polyamine analogues. *Nature Cat.* **4**:498-509
- 629. L. Baumann; T. Doughty; V. Siewers; J. Nielsen; E. Boles; M. Oreb (2021) Transcriptomic response of Saccharomyces cerevisiae to octanoic acid production. FEMS Yeast Res. 21:foab011
- 630. Y. Zhao; Y. Zhao; **J. Nielsen**; Z. Liu (2021) Production of β-carotene in *Saccharomyces cerevisiae* through altering yeast lipid metabolism. *Biotechnol. Bioeng.* **118**:2043-2052
- 631. Y. Chen; F. Li; J. Mao; Y. Chen; J. Nielsen (2021) Yeast optimizes metal utilization based on metabolic network and enzyme kinetics. *Proc. Nat. Acad. Sci. USA* 118:e2020154118
- 632. G.B. Ferraro; A. Ali; A. Luengo; D.P. Kodack; A. Deik; K.K. Abbott; D. Bezwada; L. Blanc; B. Prideaux; X. Jin; J.M. Possada; J. Chen; C.R. Chin; Z. Amoozgar; R. Ferreira; I. Chen; K. Nexerova; C. Ng; A.M. Westermark; M. Duquette; S. Roberge; N.I. Lindeman; C.A. Lyssiotis; J. Nielsen; D.E. Housman; D.G. Duda; E. Brachtel; T.R. Golub; L.C. Cantley; J.M. Asara; S.M. Davidson; D. Fukumura; V.A. Dartois; C.B. Clish; R.K. Jain; M.G. Vander Heiden (2021) Fatty acid synthesis is required for breast cancer brain metastasis. *Nature Cancer* 2:414-428

- 633. P. Li; D. Sundh; B. Ji; D. Lappa; L. Ye; **J. Nielsen**; M. Lorentzon (2021) Metabolic alterations in older women supplemented with *Lactobacillus reuteri* or with low bone mineral density. *JBMR Plus* **5**:e10478
- 634. J. Geng; B. Ji: G. Li; F. Lopez-Isunza; **J. Nielsen** (2021) CODY enables quantitatively spatiotemporal predictions on in vivo gut microbial variability induced by diet-intervention. *Proc. Nat. Acad. Sci. USA* **118**:e2019336118
- 635. Y. Chen, E. van Pelt-KleinJan, B. van Olst, S. Douwenga, S. Boeren, H. Bachmann, D. Molenaar, J. Nielsen, B. Teusink (2021) Proteome constraints reveal targets for improving microbial fitness in nutrient-rich environments. *Mol. Systems Biol.* 17:e10093
- 636. R. Yu; E. Vorontsov; C. Sihlbom; **J. Nielsen** (2021) Quantifying absolute gene expression profiles reveals distinct regulation of central carbon metabolism genes in yeast. *eLife* **10**:e65722
- 637. L. Österberg; I. Domenzain; J. Münch; J. Nielsen; S. Hohmann; M. Cvijovic (2021) A novel hybrid modeling framework integrating Boolean and enzyme constrained networks enables exploration of the interplay between signaling and metabolism. *PLOS Comp. Biol.* 17:e1008891
- 638. R. Saghaleyni; M.A. Sheikh; P. Bangalore; J. Nielsen; J.L. Robinson (2021) Machine learning-based investigation of the cancer protein secretory pathway. *PLOS Comp. Biol.* 17:e1008898
- 639. H.O. Altay; C. Zhang; H. Turkez; **J. Nielsen**; M. Uhlen; A. Mardinoglu (2021) Revealing the metabolic alterations during biofilm development of *Burkholderia cenocepacia* based on genome-scale metabolic modeling. *Metabolites* 11:221
- 640. S. Marcisauskas; Y. Kim; S. Blasche; K.R. Patil; B. Ji; J. Nielsen (2021) Draft genome sequences of five fungal strains isolated from Kefir. *Microb. Res. Announcements* 10:e0019521
- 641. D. Tamburro; S. Bratulic; S.A. Shameh; N.K. Soni; A. Bacooni; F. Maccari; F. Galeotti; K. Mattsson; N. Volpi; J. Nielsen; F. Gatto (2021) Analytical performance of a standardized kit for mass spectrometry-based measurements of human glycosaminoglycans. *J. Chromat. B* **1177**:122761
- 642. J. Gustafsson; J. Robinson; J. Nielsen; L. Pachter (2021) BUTTERFLY: Addressing the pooled amplification paradox with unique molecular identifiers in single-cell RNA-seq. *Genom. Biol.* 22:174
- 643. O. Altay; M. Arif; X. Li; H. Yang; M. Aydin; G. Alkurt; W. Kim; D. Akyol; C. Zhang; G. Dinler-Doganay; H. Turkez; S. Shoaie; **J. Nielsen**; J. Boren; O. Olmuscelik; L. Doganay; M. Uhlen; A. Mardinoglu (2021) Combined metabolic activators (CMA) accelerates recovery in mild-to-moderate COVID-19. *Adv. Science* **8**:2101222
- 644. D. Bergenholm; Y. Dabirian; R. Ferreira; V. Siewers; F. David; J. Nielsen (2021) Rational gRNA design based on transcription factor binding data (2021) Syn. Biol. 6:1-7
- 645. G. Gong; Y. Zhang; Z. Wang; L. Liu; S. Shi; V. Siewers; Q. Yuan; **J. Nielsen**; X. Zhang; Z. Liu (2021) GTR 2.0: gRNA-tRNA Array and Cas9-NG based genome disruption and single-nucleotide conversion in *Saccharomyces cerevisiae*. ACS Syn. Biol. **10**:1328-1337
- 646. G. Li; J. Zrimec; B. Ji; J. Geng; J. Larsbrink; A. Zelezniak; J. Nielsen; M.K.M. Engqvist (2021) Performance of regression models as a function of experiment noise. *Bioinf. Biol. Insights* 15:1-10
- 647. S. Alseekh; A. Aharoni; Y. Brotman; K. Contrepois; J. D'Auria; J. Ewald, J. Ewald; P. Frase; P. Giavalisco; R. Hall; M. Heinemann; H. Link; J. Luo; S. Neumann; J. Nielsen; L.P. de Souza; K. Saito; U. Sauer; F. Schroeder; S. Schuster; G. Siuzdak; A. Skirycz; L. Sumner; M. Snyder; H. Tang; T. Tohge; Y. Wang; W. Wen; S. Wu; G. Xu; N. Zamboni (2021) Mass spectrometry-based metabolomics: A guide for annotation, quantification and best report practices. *Naure Met.* 18:747-756
- 648. X. Li; W. Kim; K. Juszczak; M. Arif; Y. Sato; H. Kume; S. Ogawa; H. Turkez; J. Boren; **J. Nielsen**; M. Uhlen; C. Zhang; A. Mardinoglu (2021) Stratification of clear cell renal cell carcinoma patients to facilitate drug repositioning. *iScience* 24:102772
- 649. H. Wang; J.L. Robinson; P. Kocabas; J. Gustafsson; M. Anton; P.-E. Cholley; S. Huang; J. Gobom; T. Svensson; M. Uhlen; H. Zetterberg; **J. Nielsen** (2021) Genome-scale metabolic network reconstruction of model animals as a platform for translational research. *Proc. Nat. Acad. Sci. USA* **118**:e2102344118
- 650. Y. Wang; X. Li; X. Chen; **J. Nielsen**; D. Petranovic; V. Siewers (2021) Expression of antibody fragments in *Saccharomyces cerevisiae* strains evolved for enhanced protein secretion. *Mic. Cell Fac.* **20**:134
- 651. X. Li; Y. Wang; G. Lo; Q. Liu; R. Pereira; Y. Chen; J. Nielsen (2021) Metabolic network remodeling enhances yeast's fitness on xylose using aerobic glycolysis. *Nature Cat.* **4**:783-796
- 652. Y. Chen; **J. Nielsen** (2021) In vitro turnover numbers do not reflect in vivo activities of yeast enzymes. *Proc. Nat. Acad. Sci. USA* **118**:e2108391118:
- 653. T.W. Doughty; R. Yu; L. F.-I. Chao; Z. Qin; V. Siewers; **J. Nielsen** (2021) A single chromosome strain of *S. cerevisiae* exhibits dimished ethanol metabolism and tolerance. *BMC Genom.* **22**:688
- 654. L. Caspeta; E.J. Kerkhoven; A. Martinez; J. Nielsen (2021) The yeastGemMap: A process diagram to assist yeast systems-metabolic studies. *Biotechnol. Bioeng.* 118:4800-4814
- 655. C. Malina; F. Di Barolomeo; E. Kerkhoven; J. Nielsen (2021) Constraint-based modeling of yeast mitochondria reveals the dynamics of protein impot and iron-sulfur cluster biogenesis. *iScience* 24:103294
- 656. M. Zeybel; H. Yang; O. Altay; M. Arif; C. Fredolini; M. Akyildiz; B. Saglam; M.G. Gonenli; D. Ural; W. Kim; X. Li; J.M. Schwenk; C. Zhang; S. Shoaie; J. Nielsen; M. Uhlen; J. Boren; A. Mardinoglu (2021) Combined metabolic cofactor supplementation ameliorates liver fat in NAFLD. *Mol. Systems Biol.* 17:e10459
- 657. H. Lu; F. Li; L. Yuan; I. Domenzain; R. Yu; H. Wang; G. Li; Y. Chen; B. Ji; E.J. Kerkhoven; **J. Nielsen** (2021) Yeast metabolic innovations emerged via expanded metabolic network and gene positive selection. *Mol. Systems Biol.* **17**:e10427
- 658. Q. Liu; Y. Liu; G. Li; O. Savolaiinen; Y. Chen; J. Nielsen (2021) *De novo* biosynthesis of bioactive isoflavonoids by engineered yeast cell factories. *Nature Com.* **2**:6085
- 659. S.K. Forslund; R. Chakaroun; M. Zimmermann-Kogadeeva; L. Markó; J. Aron-Wisnewsky; T. Nielsen; L. Moitinho-Silva; T.S.B. Schmidt; G. Falony; S. Vieira-Silva; S. Adriouch; R.J. Alves; K. Assmann; J.-P. Bastard; T. Birkner; R. Caesar; J. Chilloux; L.P. Coelho; L. Fezeu; N. Galleron; G. Helft; R. Isnard; B. Ji; M. Kuhn; E. Le Chatelier; A.

Myridakis; L. Olsson; N. Pons; E. Prifti; B. Quinquis; H. Roume; J.-E. Salem; N. Sokolovska; V. Tremaroli; M. Valles-Colomer; C. Lewinter; N.B. Søndertoft; H.K. Pedersen; T.H. Hansen; The MetaCardis Consortium; J.P. Gøtze; L. Køber; H. Vestergaard; T. Hansen; J.-D. Zucker; S. Hercberg; J.-M. Oppert; I. Letunic; J. Nielsen; F. Bäckhed; S.D. Ehrlich; M.-E. Dumas; J. Raes; O. Pedersen; K. Clément; M. Stumvol; P. Bork (2021) Combinatorial, additive and dose-dependent drug-microbiome associations. *Nature* 600:500-505

- 660. X. Liu; M. Liu; J. Zhang; Y. Chang; Z. Cui; B. Ji; J. **Nielsen**; Q. Qi; J. Hou (2022) Mapping of non-homologous end joining mediated integration facilitates genome-scale trackable mutagenesis in *Yarrowia lipolytica*. *ACS Syn. Biol.* 11:216-227
- 661. S. Bratulic; A. Limeta; F. Maccari; F. Galeotti; M. Levin; N. Volpi; J. Nielsen; F. Gatto (2022) Analysis of normal levels of free glycoaminoglycans in urine and plasma in adults. *J. Biol. Chem.* **298**:101575
- 662. S. Fromentin; S.K. Forslund; K. Chechi; J. Aron-Wisnewsky; R. Chakaroun; T. Nielsen; V. Tremaroli; B. Ji; E. Prifti; A. Myridakis; J. Chilloux; A. Petros; Y. Fan; M. T. Olanipekun; R. Alves; S. Adiouch; N. Bar; Y. Talmor-Barkan; E. Belda; R. Caesar; L.P. Coelho; G. Falony; S. Fellahi; P. Galan; N. Galleron; G. Helft; L. Hoyles; R. Isnard; E. Le Chatelier; H. Julienne; L. Olsson; H.K. Pedersen; N. Pons; B. Quinquis; C. Rouault; H. Roume; J.-E. Salem; T.S.B. Schmidt; S. Vieira-Silva; P. Li; M. Zimmermann-Kogadeeva; C. Lewinter; N.B. Søndertoft; T.H. Hansen; D. Gauguier; J.P. Gøtze; L. Køber; R. Kornowski; H. Vestergaard; T. Hansen; J.-D. Zucker; S. Hercberg; I. Letunic; F. Bäckhed; J.-M. Oppert; J. Nielsen; J. Raes; P. Bork; M. Stumvoll; E. Segal; K. Clément; M.-E. Dumas; S.D. Ehrlich; O. Pedersen (2022) Microbiome and metabolome features of the cardiometabolic disease spectrum. Nature Med. 28:303-314
- 663. M. Zeybel; M. Arif; X. Li; O. Altay; M. Shi; M. Akyildiz; B. Saglam; M.G. Gonenli; B. Yigit; B. Ulukan; D. Ural; S. Shoaie; H. Turkez; J. Nielsen; C. Zhang; M. Uhlen; J. Boren; A. Mardinoglu (2022) Multi-omics analysis reveals the impact of microbiota on host metabolism in hepatic steatosis. *Adv. Sci.* 9:2104373
- 664. N. Montini; T.W. Doughty; I. Domenzain; D.A. Fenton; P.V. Baranov; R. Harrington; J. Nielsen; V. Siewers; J.P. Morrissey (2022) Identification of novel gene required for competitive growth at high temperature in the thermotolerant yeast *Klyveromyces marxianus*. *Microbiol.* **168**:001148
- 665. I. Elsemman; A.R. Prado; P. Grigaitis; M. Garcia-Abomoz; V. Harman; S. Holman; J. Van Heerden; F. Bruggemann; M. Bisschops; N. Sonnenschein; S. Hubbard; R. Beynon; P. Daran-Lapujade; J. Nielsen; B. Teusink (2022) Wholecell modelling in yeast predicts compartment-specific proteome constraints that drive metabolic strategies. *Nature Com.* 13:801
- 666. E. Belda; L. Voland; V. Tremaroli; G. Falony; S. Adriouch: K.E. Assmann; E. Prifti; J. Aron-Wisnewsky; J. Debedat; T. Le Roy; T. Nielsen; C. Amouyal; S. Andre; F. Andreelli; M. Blüher; R. Chakaroun; J. Chilloux; L.P. Coelho; M.C. Dao; P. Das; S. Fellahi; S. Forslund; N. Galleron; T.H. Hansen; B. Holmes; B. Ji; H.K. Pedersen; P. Le; E. Le Chatelier; C. Lewinter; L. Mannerås-Holm; F. Marquet; A. Myridakis; V. Pelloux; N. Pons; B. Quinquis; C. Rouault; H. Roume; J.-E. Salem; N. Sokolovska; N.B. Søndertoft; S. Touch; S. Vieira-Silva; MetaCardis Consortium; P. Galan; J. Holst; J.P. Gøtze; L. Køber; H. Vestergaard; T. Hansen; S. Hercberg; J.-M. Oppert; J. Nielsen; I. Letunic; M.-E. Dumas; M. Stumvoll; O.B. Pedersen; P. Bork; S.D. Ehrlich; J.-D. Zucker; F. Bäckhed; J. Raes; K. Clement (2022) Impairment of gut microbial biotin metabolism and host biotin status in severe obesity: effect of biotin and prebiotic supplementation on improved metabolism. *Gut* 71:2463-2480
- 667. R. Saghaleyni; M. Malm; N. Moruzzi; J. Zrimec; R. Razavi; N. Wistbacka; H. Thorell; A. Pintar; A. Hober; F. Edfors; V. Chotteau; P.-O. Berggren; L. Grassi; A. Zelezniak; T. Svensson; D. Hatton; J. Nielsen; J. Robinson; J. Rockberg (2022) Enhanced metabolism and negative regulation of ER stress support higher erythropoietin production in HEK293 cells. *Cell Rep.* 39:110936
- 668. X. Li; K. Shong; W. Kim; M. Yuan; H. Yang; Y. Sato; H. Kume; S. Ogawa; H. Turkez; S. Shoaie; J. Boren; J. Nielsen; M Uhlen; C. Zhang; A. Mardinoglu (2022) Prediction of drug candidates for clear cell renal carcinoma using a systems biology-based drug repositioning approach. *eBioMedicine* 78:103963
- 669. A.S. Meijnikman; D. Lappa; H. Herrema; O. Aydin; K.A. Krautkramer; V. Tremaroli; L.E. Olofsson; A. Lundqvist; S. Bruin; Y. Acherman; Dutch liver Pathology Panel; J. Verheij; S. Hjort; V.E.A. Gerdes; T.W. Schwartz; M. Nieuwdorp; A.K. Groen; J. Nielsen; F. Bäckhed (2022) A systems biology approach to study non-alcoholic fatty liver (NAFL) in women with obesity. *iScience* 25:104828
- 670. F. Li; L. Yuan; H. Lu; G. Li; Y. Chen; M.K.M. Engqvist; E.J. Kerkhoven; **J. Nielsen** (2022) Deep learning based *k*_{cat} prediction enables improved enzyme constrained model reconstruction. *Nature Cat.* **5**:662-672
- 671. F. Li; Y. Chen; Q. Qi; Y. Wang; L. Yuan; M. Huang; I.E. Elsemman; A. Feizi; E.J. Kerkhoven; J. Nielsen (2022) Improving recombinant protein production by yeast through genome-scale modelling using proteome constraints. *Nature Com.* **13**:2969
- 672. J. Xia; B.J. Sanchez; Y. Chen; K. Campbell; S. Kasvandik; J. Nielsen (2022) Proteome allocation change linearly with specific growth rate of *Saccharomyces cerevisiae* under glucose-limitation. *Nature Com.* 13:2819
- 673. T. Xie; M. Chen; **J. Nielsen**; J. Xia (2022) Multi-omics analyses of the transition to the Crabtree effect in *S. cerevisiae* reveals a key role for the citric acid shuttle. *FEMS Yeast Res.* **22**:foac030
- 674. O.P. Ishchuk; I. Domenzain; B.J. Sanchez; F. Muniz-Paredes; J.L. Martinez; J. Nielsen; D. Petranovic (2022) Genome-scale modelling drives 70-fold improvement of intracellular heme production in *Saccharomyces cerevisiae*. *Proc. Nat. Acad. Sci. USA* 119:e2108245119
- 675. I. Domenzain; B. Sanchez; M. Anton; E.J. Kerkhoven; A. Millan-Oropeza; C. Henry; V. Siewers; J.P. Morrissey; N. Sonnenschein; J. Nielsen (2022) Reconstruction of a catalogie of genome-scale metabolic models with enzymatic constraints using GECKO2.0. *Nature Com.* 13:3766

- 676. F. Gatto; S. Dabestani; S. Bratulic; A. Limeta; F. Maccari; F. Galeotti; N. Volpi; U. Steiner; **J. Nielsen**; S. Lundstam (2022) Plasma and urine free glucosaminoglycans as monitoring biomarkers in nonmetastatic renal cell carcinoma A prospective cohort study. *Eur. Urol. Open Sci.* **42**:30-39
- 677. N. Begum; S. Lee; T.J. Portlock; A. Pellon; S.D.S. Nasab; J. Nielsen; M. Uhlen; D.L. Moyes; S. Shoaie (2022) Integrative functional analysis uncovers metabolic differences between *Candida* species. *Comms. Bio.* 5:1013
- 678. J. Zrimec; X. Fu; A.S. Muhammad; C. Skrekas; V. Jauniskis; N.K. Speicher; C.S. Börlin; V. Verendel; M.H. Chehreghani; D. Dubhashi; V. Siewers; F. David; J. Nielsen; A. Zelezniak (2022) Controlling gene expression with deep generative design of regulatory DNA. *Nature Com.* 13:5099
- 679. C. Zhang; M. Shi; W. Kim; M. Arif; M. Klevstig; X. Li; H. Yang; C. Bayram; I. Bolat; Ö.Ö. Tozlu; H. Hacimuftuoglu; S. Yildirim; J. Sebhaoui; S. Iqbal; Y. Wei; X. Shi; J. Nielsen; H. Turkez; M. Uhlen; J. Boren; A. Mardinoglu (2022) Discovery of therapeutic agents targeting *PKLR* for NAFLD using drug repositioning. *eBioMed*. 83:104214
- 680. G. Zhang; C. Zhang; Z. Wang; Q. Wang; J. Nielsen; Z. Dai (2022) Dual β-oxidation pathway and transcription factor engineering methyl ketones production in *Saccharomyces cerevisiae*. *Met. Eng.* **73**:225-234
- 681. T. Yu; Q. Liu; X. Wang; X. Liu; V. Siewers; Y. Chen; J. Nielsen (2022) Metabolic reconfiguration enable synthetic reductive metabolism. *Nature Met.* 4:1551-1559
- 682. A.S. Meijnikman; M. Davids; H. Herrema; O. Aydin; V. Tremaroli; M. Rios-Morales; H. Levels; S. Bruin; M. de Brauw; J. Verheij; M. Kemper; A.G. Holleboom; M. Tushuizen; T.W. Schwartz; **J. Nielsen**; D. Brandjes; E. Dirinck; J. Weyler; A. Verrijken; C. De Block; L. Vonghia; S. Francque; U. Beuers; V.E.A. Gerdes; F. Bäckhed; A.K. Groen; M. Nieuwdorp (2022) Microbiome-derived ethanol in non-alcoholic fatty liver disease. *Nature Med.* **28**:2100-2106
- 683. P. Li; B. Ji; H. Lou; D. Sundh; M. Lorentzon; **J. Nielsen** (2022) One-year supplementation with *Lactobacillus reuteri* ATCC PTA 6475 counteracts a degradation of gut microbiota in older women with low bone mineral density. *NPJ Biofilms & Microbiomes* **8**:84
- 684. G. Li; F. Buric; J. Zrimec; S. Viknander; J. Nielsen; A. Zelezniak; M.K.M. Engqvist (2022) Learning deep representations of enzyme thermal adaptation. *Prot. Sci.* 31:e4480
- 685. H. Lu; E. Kerkhoven; J. Nielsen (2022) A pan-draft metabolic model reflects evolutionary diversity across 332 yeast species. *Biomolecules* 12:1632
- 686. F. Gatto; S. Bratulic; E. Jonasch; A. Limeta; F. Maccari; F. Galeotti; N. Volpi; S. Lundtam; J. Nielsen; U. Steiner (2022) Plasma and urine free glycosaminoglycans as monitoring and predictive biomarkers in metastatic renal cell carcinoma: A prospective cohort study. *JCO Precision Oncol.* **42**:30-39
- 687. M.S.S. Guman; J.B. Hoozemans; S. Haal; P.A. de Jonge; Ö. Aydin; D. Lappa; A.S. Meijnikman; F. Westerink; Y. Acherman; F. Bäckhed; M. de Brauw; J. Nielsen; M. Nieuwdorp; A.K. Groen; V.E.A. Gerdes (2022) Adipose Tissue, Bile Acids, and Gut Microbiome Species Associated With Gallstones After Bariatric Surgery. J Lipid Res. 63:100280
- 688. S. Bratulic; A. Limeta; S. Dabestani; H. Birgisson; G. Enblad; K. Stålberg; G. Hesselager; M. Häggman; M. Höglund; O.E. Simonson; P. Stålberg; H. Lindman; A. Bång-Rudenstam; M. Ekstrand; G. Kumar; I. Cavarretta; M. Alfano; F. Pellegrino; T. Mandel-Clausen; A. Salanti; F. Maccari; F. Galeotti; N. Volpi; M. Daugaard; M. Belting; S. Lundstam; U. Steiner; J. Nyman; B. Bergman; P.-H. Edqvist; M. Levin; A. Salonia; H. Kjölhede; E. Jonasch; J. Nielsen; F. Gatto (2022) *Proc. Nat. Acad. Sci. USA* 119:e2115328119

- 689. Y. Zhao; K. Meng; J. Fu; S. Xu; G. Cai; G. Meng; J. Nielsen; Z. Liu; Y. Zhang (2023) Protein engineering of invertase for enhancing yeast dough fermentation under high sucrose concentrations. *Folia Microbiol.* **68**:207-217
- 690. F. Li; Y. Chen; M. Anton; **J. Nielsen** (2023) GotEnzymes: An extensive database of enzyme parameter predictions. *Nuc. Acid Res.* **51**:D583-D586
- 691. X. Cao; W. Yu; Y. Chen; S. Yang; Z.K. Zhao; J. Nielsen; H. Luan; Y.J. Zhou (2023) Engineering yeast for high-level production of diterpenoid sclareol. *Met. Eng.* 75:19-28
- 692. D. Lappa; A.S. Meijnikman; K.A. Krautkramer; L.M. Olsson; Ö. Aydin; A.-S. van Rijswijk; Y.I.Z. Acherman; M.L. De Brauw; V. Tremaroli; L.E. Olofsson; A- Lundqvist; S.A. Hjorth; B. Ji; V.E.A. Gerdes; A.K. Groen; T.W. Schwartz; M. Nieuwdorp; F. Bäckhed; J. Nielsen (2023) Self-organized metabotyping of obese individuals identifies clusters responding differently to bariatric surgery. *PLOS One*, in press
- 693. H. Lou; P. Li; B. Ji; J. Nielsen (2023) Modeling the metabolic dynamics at the genome-scale by optimizaed yield analysis. *Met. Eng.*, in press
- 694. N. Qin; L. Li; X. Ji; R. Pereira; Y. Chen; S. Yin; C. Li; X. Wan; D. Qiu; J. Jiang; H. Lou; Y. Zhang; G. Dong; Y. Zhang; S. Shi; H.J. Jessen; J. Xia; Y. Chen; C. Larsson; T. Tan; Z. Liu; J. Nielsen (2023) Flux regulation through glycolysis and respiration is balanced by inositol pyrophosphates. *Cell* 186:748-763
- 695. S. Rahimi; D. van Leeuwen; F. Roshanzamir; S. Pandit; L. Shi; N. Sasanian; **J. Nielsen**; E.K. Esbjörner; I. Mijakovic (2023) Ginsenoside Rg3 reduces the toxicity of graphene oxide used for pH-responsive delivery of doxorubicin to liver and breast cancer cells. *Pharmaceutics* **15**:391
- 696. Z. Yao; Y. Guo; H. Wang: Y. Chen; Q. Wang; Z. Dai; J. Nielsen (2023) A highly efficient transcriptome-based biosynthesis of non-ethanol chemicals in Crabtree negative Saccharomyces cerevisiae. Biotechnol. Biofuels, in press
- 697. J.O. Valle; V. Siewers; J. Nielsen; S. Shi (2023) Directed evolution of a wax ester synthase for production of fatty acid ethyl esters in *Saccharomyces cerevisiaei*. *Appl. Microbiol. Biotechnol.* **107**:2921-2932
- 698. K. Wang; Y. Da; H. Bi; Y. Liu: B. Chen; M. Wang; Z. Liu; J. Nielsen; T. Tan (2023) A one-carbon chemicals conversion strategy to produce precursor of biofuels with *Saccharomyces cerevisiae*. *Renewable Energy* 208:331-340
- 699. P. Adrikopoulos; J. Aron-Wisnewsky; R. Charkaroun; A: Myridakis; S.K. Forslund; T. Nielsen; S. Adriouch; B. Holmes; J. Chilloux; S. Vieira-Silva; G. Falony; J.-E. Salem; F. Adnreelli; E. Belda; J. Kieswich; K. Chechi; F. Puig-Castellvi; M. Chevalier; E. Le Chatelier; M.T. Olanipekun; L. Hoyles; R. Alves; G. Helft; R. Isnard; L. Køber; L.P. Coelho; C. Rouault; D. Gauguir; J.P. Gøtze; E. Prifti; MetaCardis Consortium; J.-D. Zucker; F. Bäckhed; H.

- Vestergaard; T. Hansen; J.-M. Oppert; M. Blüher; **J. Nielsen**; J. Ras; P. Bork; M.M. Yaqoob; M. Stumvoli; O. Pedersen; S.D. Ehrlich; K. Clement; M.-E. Dumas (2023) Evidence of causal and modifiable relationship between kidney function and circulating trimethylamine N-oxide. *Nature Com.*, in press
- 700. L. Yuan; H. Lu; F. Li; J. Nielsen; E.J. Kerkhoven (2023) HGTphyloDetect facilitating the identification of phylogenetic analysis of horizontal gene transfer. *Brif. Bioinfor.* 24:bbad035
- 701. M. Chen; T. Xie; H. Li; Y. Zhuang; J. Xia; J. Nielsen (2023) Increased glycolytic flux in yeast is accompanies by decreased metabolite regulation and lower protein phosphorylation. *Proc. Nat. Acad. Sci. USA*, in press
- 702. H. Yang; X. Li; H. Jin; H. Turkez; G. Ozturk; H.L. Doganay; C. Zhang; J. Nielsen; M. Ühlen; J. Boren; A. Mardinoglu (2023) Longitudinal metabolomics analysis reveals the acute effect of cysteine and NAC included in the combined metabolic activators. Free Radical Biol. Med. 204:347-358
- 703. M.-E. Dumas; P. Andrikopoulos; J. Aron-Wisneswky: R. Chakaroun; A. Myridakis; S. Forslund; T. Nielsen; S. Adiouch; B. Holmes; J. Chilloux; S. Vieira-Silva; G. Falony; J.-E. Salem; F. Andreelli; E. Belda; J. Kieswich; K. Chechi; F. Puig-Castellví; M. Chevalier; E. Le Chatelier; M. Olanipekun; L. Hoyles; R. Alves; G. Helft; R. Isnard; L. V. Køber; L.P. Coelho; C. Rouault; D. Gauguier; J. Gøtze; E. Prifti; J.-D. Zucker; F. Bäckhed; H. Vestergaard; T. Hansen; J.-M. Oppert; M. Blüher; J. Nielsen; J. Raes; P. Bork; M. Yaqoob; M. Stumvoll; O. Pedersen; S. Ehrlich; K. Clement; MetaCardis Consortium (2023) Evidence of a causal and modifiable relationship between kidney function and circulating trimethylamine N-oxide. *Nature Com.*, in press
- 704. J. Mao; M.T. Mohedano; J. Fu; X. Li; Q. Liu; J. Nielsen; V. Siewers; Y. Chen (2023) Fine-tuning of p-coumaric acid synthesis to increase (2S)-naringenin production in yeast. *Met. Eng.* **79**:192-202
- 705. Y. Chen; J. Gustafsson; A.T. Rangel; M. Anton; I. Domenzain; C. Kittikunapong; F. Li; L. Yuan; J. Nielsen; E. Kerkhoven (2023) Reconstruction, simulation and analysis of enzyme-constrained metabolic models using GECKO toolbox 3.0. *Nature Prot.*, in press
- 706. B. Yulug; O. Altay; X. Li; L. Hanoglu; S. Cankaya; S. Lam; H.A. Velioglu; H. Yang; E. Coskun; E. Idil; R. Nogaylar; A. Ozsimsek; C. Bayram; I. Bolat; S. Oner; O.O. Tozlu; M.E. Arslan; A. Hacimuftuoglu; S. Yildirim; M. Arif; S. Shoaie; C. Zhang; J. Nielsen; H. Turkez; J. Borén; M. Uhlén; A. Mardinoglu (2023) Combined metabolic activators improve cognitive functions in Alzheimer's disease patients: a randomised double-blinded, placebo-controlled phase-II trial. *Transl. Neurodegener.* 12:4
- 707. K. Wang; C. Su; H. Bi; C. Zhang; D. Cai; Y. Liu; M. Wang; B. Chen; J. Nielsen; Z. Liu; T. Tan (2023) The transition from 2G to 3G-feedstocks enabled efficient production of fuels and chemicals. *Green Energy & Environ.*, in press

Reviews in international journals with peer review

- 1. J. Nielsen; J. Villadsen (1992) Modelling of microbial kinetics. Chem. Eng. Sci. 47:4225-4270
- 2. J. Nielsen (1996) Modelling the morphology of filamentous microorganisms. Trends Biotechnol. 14:438-443
- 3. L. Olsson; **J. Nielsen** (1997) Studies of on-line and *in-situ* measuring methods for biomass concentration. *Trends Biotechnol* **15**:517-522
- 4. C. Klein; L. Olsson; **J. Nielsen** (1998) Glucose control in *Saccharomyces cerevisiae*. The role of *MIG1* on metabolic functions. *Microbiol* **144**:13-24
- L. Olsson; U. Schulze; J. Nielsen (1998) On-line bioprocess monitoring an academic discipline or an industrial tool? Trends Anal. Chem. 17:88-95
- C. M. Henriksen; J. Nielsen; J. Villadsen (1998) Cyclization of α-aminoadipic acid into the δ-lactam 6-oxopiperidine-2-carboxylic acid by *Penicillium chrysogenum*. J. Antibio. 51:99-106
- J. Nielsen (1998) The role of metabolic engineering in the production of secondary metabolites. Curr. Opion Microbiol. 1:330-336
- 8. S. Ostergaard; L. Olsson; J. Nielsen (2000) Metabolic engineering of Saccharomyces cerevisiae. Microb. Mol. Biol. Rev. 64:34-50
- 9. A. K. Gombert; J. Nielsen (2000) Mathematical modelling of metabolism. Curr. Opinion Biotechnol. 11:180-186
- 10. J. Nielsen (2001) Metabolic engineering. Appl. Microbiol. Biotechnol. 55:263-283
- 11. J. Zaldivar; J. Nielsen; L. Olsson (2001) Fuel ethanol production from lignocellulose: A challenge for metabolic engineering and process integration. *Appl. Microbiol. Biotechnol.* **56**:17-34
- 12. J. Thykaer; J. Nielsen (2003) Metabolic engineering of β-lactam production. *Metabolic Eng.* **5**:56-69
- G. Hofmann; M. McIntyre; J. Nielsen (2003) Fungal genomics beyond Saccharomyces cerevisiae. Curr. Op. Biotechnol. 14:226-231
- K. Patil; M. Åkesson; J. Nielsen (2004) Use of genome scale microbial models for metabolic enginereing. Curr. Op. Biotechnol. 15:64-69
- 15. I. Borodina; J. Nielsen (2005) From genomes to in silico cells via metabolic networks. Curr. Op. Biotechnol. 16:1-6
- S. G. Villas-Bôas; S. Mas; M. Åkesson; J. Smedsgaard; J. Nielsen (2005) Mass spectrometry in metabolome analysis. Mass Spec. Rev. 24:613-646
- 17. J. Nielsen; S. Oliver (2005) The next wave in metabolome analysis. Trends Biotechnol. 23:544-546
- 18. M. C. Jewett; G. Hofmann; J. Nielsen (2006) Fungal metabolite analysis in genomics and phenomics. *Curr. Op. Biotechnol.* 17:1-7
- 19. S. Rokem; A. Eliasson; J. Nielsen (2007) Systems biology of secondary metabolism in microorganisms. *Nat. Prod. Rep.* **24**:1262-1287
- 20. **J. Nielsen**; M. Jewett (2008) Impact of systems biology on metabolic engineering of *Saccharomyces cerevisiae*. *FEMS Yeast Res.* **8**:122-131

- 21. V. Mapelli; L. Olsson; **J. Nielsen** (2008) Metabolic footprinting in microbiology: methods and applications in functional genomics and biotechnology. *Trends Biotechnol.* **26**:490-497
- 22. D. Petranovic; **J. Nielsen** (2008) Can yeast systems biology contribute to the understanding of human disease? *Trends Biotechnol.* **26**:584-590
- 23. M. R. Andersen; J. Nielsen (2009) Current status of systems biology in Aspergilli. Fun. Gen. Biol. 46:S180-S190
- G. Panagiotou; J. Nielsen (2009) Nutritional systems biology: Definitions and approaches. Annual Rev. Nutr. 29:329-339
- 25. J. Nielsen (2009) Systems biology of lipid metabolism: From yeast to human. FEBS Lett. 583:3905-3913
- 26. J. M. Otero; J. Nielsen (2010) Industrial Systems Biology. Biotechnol. Bioeng. 105:439-460
- K. Tyo; K. Kocharin; J. Nielsen (2010) Toward Design-based Engineering of Industrial Microbes. Curr. Op. Microbiol. 13:255-262
- 28. L. Liu; R. Agren; S. Bordel; **J. Nielsen** (2010) Use of genome-scale metabolic models for understanding microbial physiology. *FEBS Lett.* **584**:2556-2564
- J. Zhang; G. Vemuri; J. Nielsen (2010) Systems biology of energy homeostasis in yeast. Curr. Op. Microbiol. 13:382-388
- 30. D. Petranovic; K. Tyo; G.N. Vemuri; **J. Nielsen** (2010) Prospects of yeast systems biology for human health: integrating lipid, protein and energy metabolism. *FEMS Yeast Res.* **10**:1046-1059
- 31. A. Krivoruchko; V. Siewers; J. Nielsen (2011) Opportunities for yeast metabolic engineering: Lessons from synthetic biology. *Biotechnol. J.* 6:262-276
- 32. S. Shi; J.O. Valle-Rodriguez; V. Siewers; **J. Nielsen** (2011) Prospects for microbial biodiesel production. *Biotechnol. J.* **6**:277-285
- 33. F. H. Karlsson; I. Nookaew; D. Petranovic; J. Nielsen (2011) Prospects for systems biology and modeling of the gut microbiome. *Trends Biotechnol.* 29:251-258
- 34. M. Cvijovic; S. Bordel; J. Nielsen (2011) Mathematical models of cell factories: Moving towards the cores of industrial biotechnology. *Microbiol. Biotechnol.* **4**:572-584
- 35. T. Österlund; I. Nookaew; **J. Nielsen** (2012) Fifteen years of large scale metabolic modeling of yeast: Developments and impacts. *Biotechnol. Adv.* **30**:979-988
- 36. B. de Jong; V. Siewers; J. Nielsen (2012) Systems biology of yeast: enabling technology for development of cell factories for production of advanced biofuels. *Curr. Opion. Biotechnol.* 23:1-7
- 37. I.-K. Kim; A. Roldao; V. Siewers; **J. Nielsen** (2012) A system-level approach for metabolic engineering of yeast cell factories. *FEMS Yeast Res.* **12**:228-248
- 38. A. Mardinoglu; J. Nielsen (2012) Systems medicine and metabolic modeling. J. Int. Med. 271:142-154
- 39. K.-K. Hong; J. Nielsen (2012) Metabolic engineering of Saccharomyces cerevisiae: A key cell factory platform for future biorefineries. Cell. Mol. Life Sci. 16:2671-2690
- 40. J. L. Martinez; L. Liu; D. Petranovic; J. Nielsen (2012) Pharmaceutical protein production by yeast: towards production of human blood proteins by microbial fermentation. *Curr. Opionion Biotechnol.* 23:965-971
- 41. J. Hou; K. E. J. Tyo; Z. Liu; D. Petranovic; J. Nielsen (2012) Metabolic engineering of recombinant protein secretion by Saccharomyces cerevisiae. FEMS Yeast Res. 12:491-510
- 42. C. Knuf; J. Nielsen (2012) Aspergilli: Systems biology and industrial applications. Biotechnol. J. 7:1147-1155
- 43. **J. Nielsen** (2013) Production of biopharmaceutical proteins by yeast. Advances through metabolic engineering. *Bioengineered* **4**:207-211
- 44. Y. Chen; J. Nielsen (2013) Advances in metabolic pathway and strain engineering paving the way for sustainable production of chemical building blocks. *Curr. Op. Biotechnol.* 24:965-972
- 45. A. Mardinoglu; F. Gatto; J. Nielsen (2013) Genome-scale modeling of human metabolism. Biotechnol. J. 8:985-996
- 46. L. Caspeta; **J. Nielsen** (2013) Toward systems metabolic engineering of *Aspergillus* and *Pichia* species for the production of chemicals and biofuels. *Biotechnol. J.* **8**:534-544
- 47. J. Nielsen; C. Larsson; A. van Maris; J. Pronk (2013) Metabolic engineering of yeast for production of fuels and chemicals. *Curr. Op. Biotechnol.* 24:398-404
- 48. N. A. Buijs; V. Siewers; J. Nielsen (2013) Advanced biofuel production by the yeast *Saccharomyces cerevisiae*. *Curr. Op. Chem. Biol.* 17:480-488
- F. Karlsson; V. Tremaroli; J. Nielsen; F. Bäckhed (2013) Assessing the human gut microbiota in metabolic diseases. Diabetes 62:3341-3349
- M. A. Garcia-Albornoz; J. Nielsen (2013) Application of genome-scale metabolic models in metabolic engineering. Ind. Biotechnol. 9:203-214
- 51. S. Tippmann; Y. Chen; V. Siewers; **J. Nielsen** (2013) From flavors and pharmeuticals to advanced biofuels: Production of isoprenoids in *Saccharomyces cerevisiae*. *Biotechnol. J.* **8**:1435-1444
- 52. L. Väremo; I. Nookaew; J. Nielsen (2013) Novel insights into obesity and diabetes through genome-scale metabolic modeling. *Fron. Physiology* **4**:92
- 53. M. Huang; J. Bao; J. Nielsen (2014) Biopharmaceutical protein production by *Saccharomyces cerevisiae*: Current state and future prospects. *Pharmaceut. Bioprocessing* 2:167-182
- 54. I. Borodina; J. Nielsen (2014) Advances in metabolic engineering of yeast *Saccharomyces cerevisiae* for production of chemicals. *Biotechnol. J.* **9**:609-620
- 55. J. Almquist; M. Cvijovic; V. Hatzimanikatis; J. Nielsen; M. Jirstrand (2014) Kinetic models in industrial biotechnology improving cell factory performance. *Met. Eng.* 24:38-60
- 56. S. Shoai; **J. Nielsen** (2014) Elucidating the interactions between the human gut microbiota and its host through metabolic modeling. *Fron. Genetics* **5**:86
- 57. G. Liu; A. Marras; J. Nielsen (2014) The future of genome-scale modeling of yeast through integration of a transcriptional regulatory network. *Quan. Biol.* 2:30-46

- 58. Y.J. Zhou; N.A.Buijs; V. Siewers; **J. Nielsen** (2014) Fatty acid-derived biofuels and chemicals production in *Saccharomyces cerevisiae*. Front. Synth. Biol. **2**:32
- 59. E.J. Kerkhoven; P.-J. Lahtvee; J. Nielsen (2014) Application of computational modeling in metabolic engineering of Saccharomyces cerevisiae. FEMS Yeast Res. 15:1-13
- 60. B. Ji; J. Nielsen (2015) New insight into the gut microbiome through metagenomes. Adv. Genom. Genet. 5:77-91
- A. Krivoruchko; Y. Zhang; V. Siewers; J. Nielsen (2015) Microbial acetyl-CoA metabolism and metabolic engineering. Met. Eng. 28:28-42
- 62. A. Krivoruchko; J. Nielsen (2015) Production of natural products through metabolic engineering of Saccharomyces cerevisiae. Curr. Op. Biotechnol. 35:7-15
- 63. A. Mardinoglu; **J. Nielsen** (2015) New paradigms for metabolic modeling of human cells. *Curr. Op. Biotechnol.* **34**:91-97
- B. Pfleger; M. Gossing; J. Nielsen (2015) Metabolic engineering strategies for microbial production of oleochemicals. Met. Eng. 29:1-11
- 65. D. Jullesson; F. David; B. Pfleger; **J. Nielsen** (2015) Impact of Synthetic Biology and Metabolic Engineering on Industrial Production of Fine Chemicals. *Biotechnol. Adv.* **33**:1395-1402
- B. J. Sanchez; J. Nielsen (2015) Genome-scale models of yeast: Towards standardized evaluation and consistent omic integration. *Integrative Biol.* 7:846-858
- 67. O. Kayikci; J. Nielsen (2015) Glucose repression in Saccharomyces cerevisiae. FEMS Yeast Res. 15:fov068
- 68. Z. Dai; **J. Nielsen** (2015) Advancing metabolic engineering through systems biology of industrial microorganisms. *Curr. On. Biotechnol.* **36**:8-15
- P.N. Ghaffari; A. Mardinoglu; J. Nielsen (2015) Cancer metabolism: A modeling perspective. Front. Physiol. 6:Article 382
- L. Caspeta; T. Castilio; J. Nielsen (2015) Modifying yeast tolerance to inhibitory conditions of ethanol production. Fron. Bioeng. Biotechnol. 3:Article 184
- 71. B. Ji; **J. Nielsen** (2015) From next-generation sequencing to systematic modeling of the gut microbiome. *Front. Genet.* **6**:219
- 72. E. Fletcher; A. Krivoruchko; **J. Nielsen** (2016) Industrial systems biology and its impact on synthetic biology of yeast cell factories. *Biotechnol. Bioeng.* **113**:1164-1170
- 73. J.L. Martinez; D. Petranovic; J. Nielsen (2016) Heme metabolism in stress regulation and protein production: From cinderalla to key player. *BioEngineered* 7:112-115
- 74. F. Gatto; J. Nielsen (2016) In search for symmetries in the metabolism of cancer. WIREs Systems Biol. Med. 8:23-35
- 75. J. Nielsen; J. Keasling (2016) Engineering Cellular Metabolism. Cell 164:1185-1197
- M. Uhlen; B.M. Hallström; C. Lindskog; A. Mardinoglu; F. Ponten; J. Nielsen (2016) Transcriptomics resources of human tissues and organs. Mol. Sys. Biol. 12:862
- 77. M. Kumar; P. Babaei; B. Ji; J. Nielsen (2016) Human gut microbiota and healthy ageing: Recent developments and future perspective. *Nutr. Healthy Ageing* **4**:3-16
- J. Robinson; J. Nielsen (2016) Integrative analysis of human omics data using biomolecular networks. Mol. BioSystems 12:2953-2964
- Y. Chen; J. Nielsen (2016) Biobased organic acid production by metabolically engineered microorganisms. Curr. Op. Biotechnol. 37:165-172
- 80. Y. Chen; J. Nielsen (2016) Flux control through protein phosphorylation in yeast. FEMS Yeast Res. 16:fow096
- M.-K. Kang; J. Nielsen (2017) Biobased production of alkanes and alkenes through metabolic engineering of microorganisms. J. Ind. Microbiol. Biotechnol. 44:613-622
- 82. A. Nilsson; J. Nielsen (2017) Genome-scale metabolic modeling of cancer. Met. Eng. 43:103-112
- 83. L.J. Sweetlove; **J. Nielsen**; A.R. Fernie (2017) Engineering central metabolism a grand challenge for plant biologists. *Plant J.* **90**:749-763
- 84. J. Geng; J. Nielsen (2017) *In silico* analysis of human metabolism: Reconstruction, contextualization and application. *Curr. Op. Systems Biol.* **2**:28-37
- J. Nielsen (2017) Systems Biology of Metabolism: A Driver for Developing Personalized and Precision Medicine. Cell Met. 25:572-579
- 86. J.C. Nielsen; **J. Nielsen** (2017) Development of fungal cell factories for the production of secondary metabolites: linking genomics and metabolism. *Synth. Systems Biotechnol.* **2**:5-12
- 87. G. Wang; M. Huang; J. Nielsen (2017) Exploring the potential of *Saccharomyces cerevisiae* for biopharmaceutical protein production. *Curr. Opion. Biotechnol.* **48**:77-84
- 88. J. Nielsen (2017) Systems Biology of Metabolism. Ann. Rev. Biochem. 86:245-275
- 89. J.L. Robinson; J. Nielsen (2017) Anticancer drug discovery through genome-scale metabolic modeling. *Curr. Opion. Syst. Biol.* **4**:1-8
- 90. **J. Nielsen**; J. Archer; V. Bajic; T. Gojobori; I. Mijakovic (2017) Building a biobased industry in the Middle East through harnessing the potential of the Red See biodiversity. *Appl. Microbiol. Biotechnol.*, in press
- 91. D. Cook; **J. Nielsen** (2017) Genome-scale metabolic models applied to human health and disease. *WIRE Systems Biol.*Med. 9:e1393
- 92. X. Chen; C. Gao; L. Guo; G. Hu; Q. Luo; J. Liu; J. Nielsen; J. Chen; L. Liu (2018) DCEO Biotechnology: Tools to Design, Construct, Evaluate and Optimize the Metabolic Pathway for Biosynthesis of Chemicals. *Chem. Rev.* 118:4-72
- 93. R. Benfeitas; M. Uhlen; J. Nielsen; A. Mardinoglu (2017) New challenges to study heterogeneity in cancer redox metabolism. *Fron. Cell Develop. Biol.* **5**:65
- 94. J. Bosley; C. Boren; S. Lee; M. Grøtli; J. Nielsen; M. Uhlen; J. Boren; A. Mardinoglu (2017) Improving economics of NASH/NAFLD treatment through the use of systems biology. *Drug Discov. Today* 22:1532-1538
- 95. Z. Gong; J. Nielsen; Y. Zhou (2017) Engineering robustness of microbial cell factories. Biotechnol. J. 12:1700014

- K. Campbell; J. Xia; J. Nielsen (2017) The impact of systems biology on bioprocessing. Trends Biotechnol. 35:1156-1168
- 97. Y. Zhang; **J. Nielsen**; Z. Liu (2017) Engineering yeast metabolism for production of terpenoids for use a perfume ingredients, pharmaceuticals, and biofuels. *FEMS Yeast Res.* **17**:fox080
- 98. A. Mardinoglu; J. Boren; U. Smith; M. Uhlen; J. Nielsen (2018) Systems biology in hepatology: Approaches and applications *Nature Rev. Gastro. Hep.* **15**:365-377
- 99. R. Ferreira; F. David; **J. Nielsen** (2018) Advancing biotechnology with CRISPR/Cas9: Recent applications and patent landscape. *J. Ind. Microbiol. Biotechnol.* **45**:467-480
- 100. F.R. Pinu; N. Granucci; J. Daniell; T.-L. Han; S. Carneiro; I. Rocha; J. Nielsen; S.G. Villas-Boas (2018) Metabolite secretion in microorganisms: The theory of metabolic overflow put to the test. *Metabolomics* 14:43
- 101. C. Malina; C. Larsson; J. Nielsen (2018) Yeast mitochondria: An overview of mitochondrial biology and the potential of mitochondrial systems biology. FEMS Yeast Res. 18:foy040
- 102. B. Turanli; M. Grøtli; J. Boren; J. Nielsen; M. Uhlen; K.Y. Arga; A. Mardinoglu (2018) Drug repositioning for effective prostate cancer treatment. Fron. Physiol. 9:500
- 103. Y. Zhang; J. Nielsen; L. Zihe (2018) Metabolic engineering of *Saccharomyces cerevisiae* for production of fatty acid derived hydrocarbons. *Biotechnol. Bioeng.* 115:2139-2147
- 104. Y. Zhou; E. Kerkhoven; **J. Nielsen** (2018) Barriers and opportunities in bio-based production of hydrocarbons. *Nature Energy* **3**:925-935
- 105. R. Ferreira; A. Limeta; J. Nielsen (2019) Tackling cancer with yeast based technologies. Trends Biotechnol. 37:592-603
- 106. L. Zihe; Y. Zhang; J. Nielsen (2019) Synthetic biology of yeast. Biochem. 58:1511-1520
- 107. X. Li; Y. Chen; J. Nielsen (2019) Harnessing xylose pathways for biofuels production. *Curr. Opion. Biotechnol.* 57:56-65
- 108. J. Nielsen (2019) Yeast Cell Factory: Model Organism and Cell Factory. Biotechnol. J. 14:1800421
- 109. T. Yu; Y. Dabirian; Q. Liu; V. Siewers; J. Nielsen (2019) Challenges and strategies for metabolic rewiring. Curr. Op. Systems Biol. 15:30-38
- 110. Y. Hu; Z. Zhu; J. Nielsen; V. Siewers (2019) Engineering Saccharomyces cerevisiae cells for production of fatty acid derived biofuels and chemicals. Open Biol. 9:190049
- 111. M. Kumar; B. Ji; K. Zengler; J. Nielsen (2019) Modeling approaches for studying the gut microbiota. *Nature Microbiol.* 4:1253-1267
- 112. Y. Liu; **J. Nielsen** (2019) Recent trends in metabolic engineering of microbial chemical factories. *Curr. Op. Biotechnol.* **60**:188-197
- 113. B. Turanli; O. Altay; J. Boren; H. Turkez; J. Nielsen; M. Uhlen; K.Y. Arga; A. Mardinoglu (2019) Systems biology based drug repositioning for development of cancer therapy. *Sem. Cancer Biol.* 10.1016
- 114. R. Yu; J. Nielsen (2019) Big data in yeast systems biology. FEMS Yeast Res. 19:foz070
- 115. M. Palmnäs; C. Brunius; L. Shi; A. Rostgaard-Hansen; N.E. Torres; R. Gonzalez-Dominguez; R. Zamra-Ros; Y. Lingqun; J. Halkjær; A. Tjønneland; G. Riccardi; T. Giacco; G. Costabile; C. Vetrani; J. Nielsen; C. Andres-Lacueva; R. Landberg (2019) Metabotyping: A potential personalized nutrition strategy for precision prevention of cardiometabolic disease. Adv. Nutrition 11:524-532
- 116. S. Bratulic; F. Gatto; **J. Nielsen** (2019) The translational status of cancer liquid biopsies. *Reg. Eng. Transl. Med.* S40883-019-00141-2
- 117. R. Yu; J. Nielsen (2019) Yeast systems biology in understanding principles of physiology underlying complex human diseases. Curr. Op. Biotechnol. 121:1-9
- 118. Z. Liu; K. Wang; Y. Chen; T. Tan; J. Nielsen (2020) Third-generation biorefineries as a mean to produce fuels and chemicals from CO₂. *Nature Cat.* **3**:274-288
- 119. Q. Liu; Y. Liu; Y. Chen; J. Nielsen (2020) Current state of aromatics production using yeast: Achievements and challenges. *Curr. Opion. Biotechnol.* **65**:65-74
- 120. Z. Lin; J. Nielsen; Z. Liu (2020) Bioprospecting through cloning of whole natural product biosynthetic gene clusters. Fron. Bioeng. Biotechnol. Fbio.2020.00526
- 121. E. Mohammadi; R. Benfeitas; H. Turkez, J. Boren; J. Nielsen; M. Uhlen; A. Mardinoglu (2020) Applications of genome-wide screening and systems biology approaches in drug repositioning. *Cancers* 12:2694
- 122. Y. Zhang; Z. Liu; **J. Nielsen** (2021) Yeast based biorefineries for oleochemical production. *Curr. Op. Biotechnol.* **67**:26-34
- 123. Z. Liu; Z. Liu; J. Nielsen (2021) Expression of fungal biosynthetic gene clusters in S. cerevisiae for natural product discovery. Syn. Systems Biotechnol. 6:20-22
- 124. Y. Chen; J. Nielsen (2021) Mathematical modeling of proteome constraints within metabolism. *Curr. Op. Systems Biol.* **25**:50-56
- 125. C. Zhan; X. Li; Y. Yang; J. Nielsen; Z. Bai; Y. Chen (2021) Strategies and challenges with the microbial conversion of methanol to high-value chemicals. *Biotechnol. Bioeng.* 118:3655-3668
- 126. J. Wang; J. Nielsen; Z. Liu (2021) Synthetic biology advanced natural product discovery. Metabolites 11:785
- 127. H. Lu; E.J. Kerkhoven; J. Nielsen (2022) Multiscale models quantifying yeast physiology: towards a whole-cell model. *Trends Biotechnol.* **40**:291-305
- 128. S. Shi; N. Qi; J. Nielsen (2022) Microbial production of chemicals driven by CRISPR-Cas systems. *Curr. Op. Biotechnol.* **73**:34-42
- 129. Z. Liu; J. Wang; **J. Nielsen** (2022) Yeast synthetic biology advances biofuel production. *Curr. Op. Microbiol.* **67**:33-39
- 130. Y. Chen; F. Li; J. Nielsen (2022) Genome-scale modeling of yeast metabolism: retrospective and perspectives. *FEMS Yeast Res.*, in press

- 131. X. Tan; J. Nielsen (2022) The integration of bio-catalysis and electrocatalysis to produce fuels and chemicals from carbon dioxide. *Chem. Soc. Rev.* **51**:4753
- 132. J. Nielsen; C.B. Tillegreen; D. Petranovic (2022) Innovation trends in industrial biotechnology. *Trends Biotechnol.*, in press
- 133. P. Li; H. Lou; B. Ji; J. Nielsen (2022) Machine learning for data integration in human gut microbiome. *Microb. Cell Fac.* 21:241
- 134. G. Zou; J. Nielsen; Y. Wei (2022) Harnessing synthetic biology for mushroom farming. Trends. Biotechnol. 41:480-483
- 135. P. Li; S. Roos; H. Lou; B. Ji; J. Nielsen (2023) Metabolic engineering in human gut microbiome: recent developments and future perspectives. *Met. Eng.*, in press
- 136. A. Limeta; F. Gatto; M.J. Herrgård; B. Ji; J. Nielsen (2023) Leverating high-resolution omics data for predicting response and adverse events to immune checkpoint inhibitors. *Comp. Struc. Biotechnol. J.*, in press
- 137. A. Limeta; F. Gatto; M.J. Herrgård; B. Ji; J. Nielsen (2023) Leveraging high-resolution omics dat for predicting responses and adverse events to immune checkpoint inhibitors. *Comp. Struc. Biotechnol. J.* 21:3912-3919
- 138. P. Li; S. Roos; H. Lou; B. Ji; J. Nielsen (2023) Metabolic engineering of human gut microbiome: Recent developments and future perspectives. *Met. Eng.* **79**:1-13
- 139. Z. Liu; S. Shi; Y. Ji; K. Wang; T. Tan; J. Nielsen (2023) Opportunities of CO₂-based biorefineries for production of fuels and chemicals. *Green Carbon* 1:75-84

Commentaries, Editorials and Short Papers

- 1. B. Hahn-Hägerdal; K. van Dam; I. Spencer-Martins; J. Nielsen (1998) Le dédale du métabolisme. Biofutur 184:81-82
- J. Thykaer; B. Christensen; J. Nielsen (2001) The application of metabolic network analysis in metabolic engineering. BIOforum International 5:228-231
- 3. J. M. Otero; L. Olsson; J. Nielsen (2007) Industrial biotech meets systems biology. Genetic Eng. News 1:28-31
- J. Nielsen (2007) Principles of optimal metabolic network operation. Mol. Sys. Biol. 3:126
- 5. G. Vemuri; J. Nielsen (2008) Systems biology: Is the hope worth the hype. SIM News 58:178-188
- 6. **J. Nielsen** (2009) Vintage paper Introduction. *Biotechnol. Bioeng.* **103**:1
- 7. J. Nielsen (2009) Quantification of flux control in metabolic pathways, Biotechnol. Bioeng. 104: 1-2
- 8. J. Nielsen; M. Vidal (2010) Systems biology of microorganisms. Current Opp. Microbiol. 13:335-336
- 9. J. Nielsen (2011) Transcriptional control of metabolic fluxes. Mol. Systems Biol. 7:478
- 10. J. Nielsen (2011) Chimeric Synthetic Pathways. Nature Chem. Biol. 7:195-196
- 11. **J. Nielsen**; J. Keasling (2011) Synergies between synthetic biology and metabolic engineering. *Nature Biotechnol.* **29**:693-695
- 12. J. Nielsen (2012) Translational and systems medicine. J. Int. Med. 271:108-110
- 13. J. Nielsen; S. Y. Lee (2012) Systems Biology: The "new biotechnology". Current Opinion Biotechnol. 23:583-584
- M. Cvijovic; J. Almquist; J. Hagmar; M. Heinemann; S. Hohmann; H.-M. Kaltenbach; E. Klipp; M. Krantz; P. Mendes; S. Nelander; J. Nielsen; A. Pagnani; N. Przulj; A. Raue; J. Stelling; S. Stoma; F. Tobin; J. Wodke; R. Zecchina; M. Jirstrand (2014) Bridging the gaps in systems biology. *Mol. Gen. Genom.* 289:727-734
- 15. J. Nielsen; M. Fussenegger; J. Keasling; S.Y. Lee; J.C. Liao; K. Prather; B. Palsson (2014) Engineering synergy in biotechnology. *Nature Chem. Biol.* 10:319-322
- 16. **J. Nielsen** (2014) Maintaining a strong yeast research community. *FEMS Yeast Res.* **14**:527-528
- 17. J. Nielsen (2014) Synthetic Biology for Engineering Acetyl Coenzyme A Metabolism in Yeast. mBio 5:e02153-14
- 18. J. Pronk; S.Y. Lee; J. Lievense; J. Pierce; B. Palsson; M. Uhlen; J. Nielsen (2015) How to set up collaborations between academia and industrial biotech companies. *Nature Biotechnol.* 33:237-240
- 19. L. Väremo; J. Nielsen (2015) Networking in metabolism and human disease. Oncotarget 6:18
- 20. A. Ebrahim; E. Almaas; E. Bauer; A. Bordbar; A.P. Burgard; R.L. Chang; A. Dräger; I. Famili; A.M. Feist; R.M.T. Fleming; S.S. Fong; V. Hatzimanikatis; M.J. Herrgård; A. Holder; M. Hucka; D. Hyduke; N. Jamshidi; S.Y. Lee; N. Le Novère; J.A. Lerman; N.E. Lewis; D. Ma; R. Mahadevan; C. Maranas; H. Nagarajan; A. Navid; J. Nielsen; L.K. Nielsen; J. Nogales; A. Noronha; C. Pal; B.O. Palsson; J.A. Papin; K.R. Patil; N.D. Price; J.L. Reed; M. Saunders; R.S. Senger; N. Sonnenschein; Y. Sun; I. Thiele (2015) Do genome-scale models need exact solvers or clearer standards? *Mol. Sys. Biol.* 11:831
- 21. J. Nielsen (2015) Yeast cell factories on the horizon. Science 349:1050-1051
- 22. D. Fraenkel; **J. Nielsen** (2016) Trehalose-6-phosphate synthase and stabilization of yeast glycolysis. *FEMS Yeast Res.* **16**:fov100
- 23. J. Nielsen (2016) All you need is mentorship: A journey of equals. Cell 164:1093
- 24. A. Mardinoglu; J. Nielsen (2016) The impact of systems medicine on human health and disease. Fron. Physiol. 7:552
- 25. J. Nielsen (2017) Built on stable catalysts. Nature Microbiol. 2:17085
- 26. A. Nilsson; **J. Nielsen**; B. Palsson (2017) Metabolic models of protein allocation call for the Kinetome. *Cell Systems* **5**:538-541
- 27. A. Nilsson; J.R. Haanstra; B. Teusink; **J. Nielsen** (2018) Metabolite depletion affects flux profiling of cell lines. *Trends Biochem. Sci.* **43**:395-397
- 28. J. Nielsen (2019) Designer Microbes Serving Society. Cell Met. 29:509
- 29. J. Nielsen (2019) Cell factory engineering for improved production of natural products. Nat. Prod. Rep. 36:1233-1236
- 30. **J. Nielsen** (2019) Antibiotic lethality is impacted by nutrient availabilities: New insights from machine learning. *Cell* **177**:1373-1374

- 31. J. Nielsen (2019) A stress-coping strategy for yeast cells. Nature 572:184-185
- 32. J. Nielsen (2022) Bioactive metabolites: The double-edged sword in your food. Cell 185:4469-4471
- 33. J. Nielsen (2023) Engineering yeast to produce plant-derived anti-obesity agent. Nature Chem. 15:1204-1205

Books

- 1. M. Y. Andersen *et al.* (1991) Bioreaktionsteknik. Metoder til fortolkning af data fra bioreaktorer. Akademiet for Tekniske Videnskaber, Lyngby
- 2. J. Nielsen; J. Villadsen (1994) Bioreaction Engineering Principles. Plenum Press, New York
- 3. J. Nielsen (1997) Physiological Engineering Aspects of Penicillium chrysogenum. World Scientific Publishing Co., Singapore
- 4. G. Stephanopoulos; A. Aristodou; J. Nielsen (1998) Metabolic Engineering. Academic Press, San Diego
- 5. **J. Nielsen** (2001) (editor) Metabolic Engineering. Advances in Biochemical Engineering/Biotechnology, Vol. 73, Springer Verlag, Heidelberg
- 6. J. Nielsen; J. Villadsen; G. Liden (2003) Bioreaction Engineering Principles, 2. ed., Kluywer Plenum, New York
- 7. **J. Nielsen** (2006) (editor) Frontiers in Biotechnology. Advances in Biochemical Engineering/Biotechnology, Vol. 100, Springer Verlag, Heidelberg
- 8. S. G. Villas-Boas; U. Roessner; M. A. E. Hansen; J. Smedsgaard; J. Nielsen (2007) Metabolome Analysis. An Introduction, Wiley, Hoboken
- 9. J. Nielsen; M. C. Jewett (2008) (editors) Metabolomics. Topics in Current Genetics, Vol. 18, Springer Verlag, Heidelberg
- 10. J. Villadsen; J. Nielsen; G. Liden (2011) Bioreaction Engineering Principles, 3. ed., Springer, New York
- 11. J. Nielsen; S. Hohmann (2017) Systems Biology. Advanced Biotechnology, Vol. 5, Wiley VCH, Mannheim

Book Chapters

- 1. J. Nielsen (1992) Modelling the growth of filamentous fungi. Adv. Biochem. Eng./Biotechnol. 46:187-223
- 2. **J. Nielsen**; J. Villadsen (1993) Bioreactors: Description and modelling. pp. 77-104, Chap. 5 in Biotechnology Vol. 3 (2. ed.). Eds. H.-J. Rehm; G. Reed (volume editor G. Stephanopoulos), VCR Verlag
- 3. **J. Nielsen**; M. Carlsen (1996) Fungal Pellets. pp. 273-293, Chap. 13. In: Immobilised living cell systems: Modelling and experimental methods. Eds. G. Baron, R. Willaert; L. De Backer, John Wiley & Sons, Chichester
- 4. P. Krabben; **J. Nielsen** (1998) Modeling the mycelium morphology of *Penicillium* species in submerged cultures. Adv. Biochem. Eng./Biotechnol. **60**:125-152
- A. B. Spohr, T. Agger; M. Carlsen; J. Nielsen (1998) Quantitative morphology of filamentous microorganisms. pp. 373-410, Chap. 14. In: Digital analysis of Microbes. Eds. M. H. F. Wilkinson; F. Schut, John Wiley & Sons
- J. Nielsen (1999) Fermentation monitoring, design and optimization. pp. 1147-1157. In: The Encyclopedia of Bioprocess Technology: Fermentation, Biocatalysis & Bioseparations. Eds. M. C. Flickinger; S. W. Drew, John Wiley & Sons
- 7. **J. Nielsen** (1999) Fermentation kinetics. pp. 69-119, Chap. 4. In: Modern Concepts in Fermentation Biotechnology. Eds. M. El-Mansi, Bryce, Taylor and Francis, London
- B. Christensen; J. Nielsen (1999) Metabolic network analysis powerful tool in metabolic engineering. Adv. Biochem. Eng./Biotechnol. 66:209-231
- J. Nielsen (2001) Microbial Process Kinetics. pp. 127-149, Chap. 6. In: Basic Biotechnology. 2. Edition. Eds. C. Ratledge, B. Kristiansen, Cambridge University Press, Cambridge
- 10. **J. Nielsen** (2002) Metabolic Turnover. In: Encyclopaedia of Life Sciences. John Wiley and Sons, Ltd., London (www.els.net)
- T. Agger; J. Nielsen (2001) Mathematical modelling of microbial processes Motivation and means. pp. 61-75. In: Engineering and Manufacturing for Biotechnology. Eds. M. Hofman, P. Thonart, Kluwer Academic Publishers, Dordrecht
- 12. S. Ostergaard; L. Olsson; **J. Nielsen** (2001) Metabolic Pathway Analysis of *Saccharomyces cerevisiae*. pp. 75-85. In: Applied Microbiology. Focus on Biotechnology Vol. 2. Eds. M. Hofman, P. Thonart, Kluwer Academic Publishers, Dordrecht
- 13. M. McIntyre; C. Müller; J. Dynesen; **J. Nielsen** (2001) Metabolic engineering of the morphology of *Aspergillus*. Adv. Biochem. Eng./Biotechnol. **73**:103-128
- J. Nielsen (2002) Metabolic Engineering. pp. 391-406. In: Encyclopedia of Physical Science and Technology, Vol. 9. Academic Press, San Diego
- M. M. dos Santos; M. Åkesson; J. Nielsen (2003) Metabolic Flux Analysis in the post genomics area. pp. 89-105. In: Metabolic Engineering in the Post Genomic Era. Eds. B. Kholodenko; H. Westerhoff, Horizon Scientific Press, Norfolk
- 16. C. Bro; B. Regenberg; **J. Nielsen** (2003) Yeast Functional Genomics and Metabolic Engineering; Past, present and future. pp. 331-360. In: Functional Genetics of Industrial Yeast. Ed. H. de Winde, Springer, Heidelberg
- A. Gombert; J. Nielsen (2003) Quantification of metabolic fluxes. In: Encyclopaedia of Life Sciences. Nature Publishing Group, London (www.els.net
- N. Gunnarsson, A. Eliasson, J. Nielsen (2004) Control of fluxes towards antibiotics and the role of primary metabolism in production of antibiotics. Adv. Biochem. Eng./Biotechnol. 88:137-178

- J. Nielsen, A. Eliasson (2005) From glucose to antibiotics: What controls the fluxes? pp. 195-214. In: Biocombinatorial Approaches for Drug Finding. Ed. W. Wohlleben, T. Spelling, B. Müller-Tiemann, Springer, Heidelberg
- H. David, J. Nielsen (2005) Modelling of fungal metabolism. Pp. 195-214. In: Metabolome Analysis. Strategies for Systems Biology. Ed. S. Vaidyanathan, G. G. Harrigan, R. Goodacre, Springer, New York
- 21. J. Maury, M. A. Asadollahi, K. Møller, A. Clark, **J. Nielsen** (2005) Microbial isoprenoid production: An example of green chemistry through metabolic engineering. Adv. Biochem. Eng./Biotechnol. **100**:19-51
- J. Højer-Pedersen; J. Smedsgaard; J. Nielsen (2006) Elucidating the mode-of-action of compounds from metabolite profiling studies, Pp. 103-130. In: Systems Biological Approaches in Infectious Diseases. Progress in Drug Research Vol. 64. Ed. H. I. Boshoff, C. E. Barry III, Birkhäuser, Berlin
- 23. **J. Nielsen** (2006) Microbial Process Kinetics. pp. 155-180, Chap. 6. In: Basic Biotechnology. 3. Edition. Eds. C. Ratledge, B. Kristiansen, Cambridge University Press, Cambridge
- J. Nielsen (2006) Fermentation Kinetics. pp. 69-120. Chap. 4. In: Fermentation Microbiology and Biotechnology. Eds.
 E. M. T. El-Mansi, C. F. A. Bryce, A. L. Demain, A. R. Allman, Taylor and Francis, London
- I. Rocha; J. Förster, J. Nielsen (2007) Design and application of genome-scale reconstructed metabolic models, Pp. 409-431. In: Methods in Molecular Biology, Vol. 416: Gene Essentiality. Ed. S. Y. Gerdes, A. L. Osterman, Humana Press Inc., Totowa, USA
- W. de Jongh, J. Nielsen (2007) Filamentous fungi as cell factories for metabolite production, Pp. 163-182. In: Food Mycology. A multifaceted approach to fungi and food. Ed. J. Dijksterhuis, R. A. Samson, CRC Press, Boca Raton, USA
- 27. A. P. Oliveira; M. C. Jewett; J. Nielsen (2007) From gene expression to metabolic fluxes, Pp. 37-66. In: Introduction to Systems Biology. Ed. S. Choi, Humana Press, Totowa, USA
- 28. M. C. Jewett; J. Nielsen (2008) The role of Metabolomics in systems biology. Pp. 1-10. In: Metabolomics. Topics in Current Genetics. Eds. J. Nielsen and M. C. Jewett, Vol. 18, Springer Verlag, Heidelberg
- M. C. Jewett; M. A. E. Hansen; J. Nielsen (2008) Data acquisition, analysis, and mining: Integrative tools for discerning metabolic function in *Saccharomyces cerevisiae*. Pp. 159-188. In: Metabolomics. Topics in Current Genetics. Eds. J. Nielsen and M. C. Jewett, Vol. 18, Springer Verlag, Heidelberg
- W. Vongsangnak; J. Nielsen (2009) Bioinformatics and systems biology of Aspergillus. Pp. 61-84. In: Aspergillus. Molecular Biology and Genomics. Eds. M. Machida and K. Gomi. Caister Academic Press, UK
- 31. G. Vemuri; **J. Nielsen** (2009) Yeast as a prototype for systems biology. Pp. 287-354. In: Systems biology and synthetic biology. Eds. P. Fu and S. Panke, Wiley, Hoboken, USA
- 32. V. Siewers; U. H. Mortensen; **J. Nielsen** (2010) Genetic engineering tools for *Saccharomyces cerevisiae*. Pp. 287-301. In: Manual of Industrial Microbiology and Biotechnology (3.ed). Eds. R. H. Baltz, A. L. Demain and J. E. Davies. ASM Press, Washington, USA
- 33. J. M. Otero; J. Nielsen (2010) Industrial Systems Biology. Pp. 79-148. In: Industrial Biotechnology. Eds. W. Soetaert and E. J. Vandamme. Wiley VCH, Weinheim, Germany
- 34. M. R. Andersen; K. Rucksomtawin; G. Hofmann; J. Nielsen (2010) Metabolic Engineering of filamentous fungi. Pp. 25.1-25.30. In: The Metabolic Pathway Engineering Handbook. Ed. C. D. Smolke. CRC Press, Boca Raton, USA
- 35. I. Borodina; A. Eliasson; J. Nielsen (2010) Metabolic Engineering of Streptomyces. Pp. 24.1-24.30. In: The Metabolic Pathway Engineering Handbook. Ed. C. D. Smolke. CRC Press, Boca Raton, USA
- 36. J. Nielsen (2010) Developing appropriate hosts for metabolic engineering. Pp. V.1-V.8. In: The Metabolic Pathway Engineering Handbook. Ed. C. D. Smolke. CRC Press, Boca Raton, USA
- 37. K. R. Patil; P. M. Bapat; J. Nielsen (2010) Structure and flux analysis of metabolic networks. Pp. 17.1-17.18. In: The Metabolic Pathway Engineering Handbook. Ed. C. D. Smolke. CRC Press, Boca Raton, USA
- 38. M. Papini; M. Salazar; **J. Nielsen** (2010) Systems Biology of Industrial Microorganisms. *Adv. Biochem. Eng./Biotechnol.* **120**:51-99
- V. Siewers; J. Nielsen; U. Mortensen (2010) Genetic engineering tools for Saccharomyces cerevisiae. Chap. 20, Pp. 287-301. In Manual of Industrial Microbiology and Biotechnology, 3rd ed., Eds. R. H. Baltz, J. E. Davies, A. Demain, American Society for Microbiology, USA
- 40. A. Roldao; I.-K. Kim; **J. Nielsen** (2012) Bridging Omics Technologies with Synthetic Biology in Yeast Industrial Biotechnology. Chap. 9, Pp. 271-327. In Systems Metabolic Engineering, Eds. C. Wittmann and S. Y. Lee, Springer, Dordrecht, The Netherlands
- 41. J. Maury; M. Asadollahi; R. L. Formenti; M. Schalk; **J. Nielsen** (2013) Metabolic engineering of isoprenoid production: Reconstruction of multi step heterologous pathways in tractable hosts. Chap. 6. In Isoprenoid synthesis in plants and microorganisms. Eds. T. J. Bach and M. Rohmer, Springer, Germany
- 42. W. Vongsangnak; J. Nielsen (2013) Systems biology methods and developments of filamentous fungi in relation to the production of food ingredients. Chap. 2. In Microbial production of food ingredients, enzymes and nutraceuticals. Eds. B. McNeil, D. Archer, I. Giavasis and L. Harvey, Woodhead Publishing, Cambridge, UK
- 43. R. Kumar; P.-J. Lahtvee; J. Nielsen (2014) Systems Biology: Developments and Applications. Chap. 4. In Molecular Mechanisms in Yeast Carbon Metabolism. Eds. Jure Piskur and Concetta Compagno, Springer, Berlin, Germany
- 44. **J. Nielsen**; S. Bordel; I. Nookaew (2014) Genome-scale metabolic models: A link between bioinformatics and systems biology. Chap. 6.11. Pp. 165-173. In Comprehensive Biomedical Physics. Ed. A. Brahme, Elsevier, Amsterdam, The Netherlands
- 45. Y. Chen; Y.J. Zhou; V. Siewers; J. Nielsen (2015) Enabling technologies to advance microbial isoprenoid production. *Adv. Biochem. Eng./Biotechnol.* 148:143-160
- 46. J. Kim; J. Nielsen (2017) Bioproduction of Fuels: An Introduction. Pp. 1-22. In Handbook of hydrocarbons and lipid microbiology series. Consequences of microbial interactions with hydrocarbons, oils and lipids. Ed. S. Y. Lee

- M. Huang; H. Joensson J. Nielsen (2018) High-throughput microfluidics for screening of yeast libraries. Methods Mol. Biol. 1671:307-317
- 48. Q. Liu; T. Yu; K. Campbell; J. Nielsen; Y. Chen (2018) Modular pathway rewiring of yeast for amino acid production. Meth. Enz. 608:417-439
- Y. Chen; G. Li; J. Nielsen (2019) Genome-scale metabolic modeling from yeast to human cell models of complex diseases: Latest advances and challenges. *Methods Mol. Biol.* 2049:329-345

Papers in Conference Proceedings

- 1. **J. Nielsen**; K. Nikolajsen; J. Villadsen (1989) Computer controlled system for on-line monitoring of a fermentation process. pp. 53-57. In Computer Applications in Fermentation Technology. Modelling and Control of Biotechnological Processes, eds. N. M. Fish; R. I. Fox; N. F. Thornhill, Elsevier Applied Science, London
- J. Villadsen; J. Nielsen (1990) Modelling of fermentation kinetics. pp. 259-266. Proc. V'th European Congress on Biotechnology, Lyngby
- J. Nielsen (1991) Application of structured fermentation models. pp. 843-848. Proc. European Simulation Multiconference, Copenhagen
- S. Benthin; J. Nielsen; J. Villadsen (1992) Transmembrane transport systems, studied on actively growing cells. pp. 351-355. In Harnessing Biotechnology for the 21st Century (Proc. IX'th International Biotechnology Symposium, Crystal City), eds. M. R. Ladisch; A. Rose,
- J. Nielsen (1993) Modelling of filamentous microorganisms. pp. 45-52. In Computer Applications in Fermentation Technology. Modelling and Control of Biotechnological Processes, eds. M. N. Karim; G. Stephanopoulos, Pergamon Press
- 6. H. S. Jørgensen; H. Møllgaard; **J. Nielsen**; J. Villadsen (1993) Identification of rate controlling enzymes in the metabolic pathway of penicillin in a high yielding strain of *P. chrysogenum*. pp. 251-254. In Computer Applications in Fermentation Technology. Modelling and Control of Biotechnological Processes, eds. M. N. Karim; G. Stephanopoulos, Pergamon Press
- 7. **J. Nielsen** (1993) Modelling the morphology of filamentous fungi. pp. 131-142. Proc. Bioreactor Performance, eds. U. Mortensen; H. J. Noorman, Elsingore
- 8. S. Frandsen; J. Nielsen; J. Villadsen (1993) Application of regimen analysis of yeast fermentation for down-scaling. pp. 171-179. Proc. Bioreactor Performance, eds. U. Mortensen; H. J. Noorman, Elsingore
- 9. S. Frandsen; J. Nielsen; J. Villadsen (1993) Dynamics of *Saccharomyces cerevisiae* in continuous culture. pp. 887-890. In Progress in Biotechnology, Vol. 9. Proc. VI'th European Congress on Biotechnology, Florence. Elsevier
- A. G. Pedersen; J. Nielsen; J. Villadsen (1993) Characterization of bioreactors using isotope techniques. pp. 931-934.
 In Progress in Biotechnology, Vol. 9. Proc. VI'th European Congress on Biotechnology, Florence. Elsevier
- J. Nielsen (1994) Physiological Engineering Towards a new science. pp. 30-38. In Proc. The 1994 IChemE Research Event, London, Vol. I
- 12. M. Carlsen; A. Spohr; R. Mørkeberg; **J. Nielsen**; J. Villadsen (1994) Growth and protein formation of recombinant Aspergillus: Utility of morphological characterization by image analysis. pp. 197-202. In Proc. Advances in Bioprocess Engineering, Cuernavaca. Kluwer Academic Publishers
- 13. **J. Nielsen** (1994) Physiological aspects of *Penicillium chrysogenum*. pp. 15-18. Proc. Modelling of filamentousfungi, Otocek.
- 14. G. Lidén; U. Schulze; J. Nielsen; J. Villadsen (1995) On the optimization of ethanol production in nitrogen limited yeast fermentations. In Proc. VII'th European Congress on Biotechnology, Nice
- 15. **J. Nielsen**; H. Jørgensen (1995) A kinetic model for the penicillin biosynthetic pathway in *Penicillium chrysogenum*. pp. 136-141. In Proc. Computer Application in Biotechnology 6, Garmisch-Partenkirchen
- P. N. Pissarra; M. J. Bazin; K. Schmidt; J. Nielsen (1995) A user-friendly and interactive methodology for simulation and non-linear parameter estimation of biotechnological process models using SIMULINK. Proc. Nordic Matlab Conference '95, Stockholm
- 17. K. Schmidt: P. N. Pissarra; J. Nielsen (1995) The evolutionary strategy: A simple and straightforward optimization algorithm for parameter estimation in structured biochemical models. Proc. Nordic Matlab Conference '95, Stockholm
- J. Nielsen (1995) Metabolic control analysis of the penicillin biosynthetic pathway. Proc. National Biotechnology and Bioengineering Conference, Ixtapa, Mexico
- 19. U. Schulze; T. L. Nissen; J. Nielsen; J. Villadsen (1996) Application of Metabolic Flux Analysis in physiological studies. Proc. V'th World Congress on Chemical Engineering, San Diego, USA
- P. de N. Pissarra; J. Nielsen (1996) Kinetic and thermodynamic analysis of the penicillin biosynthetic pathway. Proc. V'th World Congress on Chemical Engineering, San Diego, USA
- D. L. Bogle et al. (1996) Process synthesis, design and simulation of integrated biochemical processes. Proc. V'th World Congress on Chemical Engineering, San Diego, USA
- 22. J. Nielsen; I. Spencer Martins (1997) Metabolic Engineering. Proc. European Cell Factory Conference, Lund, Sweden
- 23. S. Østergaard, L. Olsson; J. Nielsen (1998) Metabolic control analysis of the Leloir pathway in *Saccharomyces cerevisiae*. BioThermoKinetics in the post genomic era. Proc. 8'th BTK Meeting, Fiskebäckskil, Sweden
- 24. H. Aae Theilgaard; J. Nielsen (1998) The influence of the LLD-ACV:bisACV ration on a MCA of the penicillin biosynthetic pathway in *Penicillium chrysogenum*. BioThermoKinetics in the post genomic era. Proc. 8'th BTK Meeting, Fiskebäckskil, Sweden
- 25. **J. Nielsen** (1998) Mathematical modelling of biochemical pathways. BioThermoKinetics in the post genomic era. Proc. 8'th BTK Meeting, Fiskebäckskil, Sweden

Other Publications

- 1. **J. Nielsen** (1990) Post doc på Universität Hannover. Dansk Kemi **5**:178-179
- 2. J. Nielsen (1990) On-line måling af biomasse i bioreaktorer. Dansk Kemi 8:260-265
- 3. **J. Nielsen**; J. Gram; L. Foldager (1990) Bioreaktionsteknik: Kendte metoder anvendt på nye processer. Dansk Kemi **12**:430-431
- 4. **J. Nielsen** (1993) Physiological Engineering. Dansk Kemi **11**:14-17
- 5. **J. Nielsen**; A. G. Pedersen; M. Bundgaard-Nielsen; H. Andersen (1994) Anvendelse af radioaktive isotoper til karakterisering af bioreaktorer. Dansk Kemi **4**:28-31
- 6. **J. Nielsen** (1995) Doktorafhandling om industriel penicillinproduktion. Dansk Kemi 11:20-23
- 7. J. Nielsen (1995) Penicillinets historie. Dansk Kemi 12:24-28
- 8. U. Schulze; J. Nielsen (1997) Massachusetts Institute of Technology. Dansk Kemi
- 9. T. Agger; J. Nielsen (1999) Optimering af enzymproduktion med skimmelsvampe. Dansk Kemi 1:14-15
- B. Christensen; J. Nielsen (1999) Metabolsk flux analyse en kvantitativ beskrivelse af mikroorganismers primære metabolisme. Dansk Kemi 1:16-18
- 11. **J. Nielsen** (1999) Functional genomics og dens rolle i udviklingen af fremtidige bioteknologiske processer. Dansk Kemi **1**:20-27
- 12. C. Bro; B. Regenberg; J. Nielsen (2001) DNA Arrays. Dansk Kemi 1:18-20
- 13. M. R. Andersen; M. L. Nielsen; J. Nielsen (2006) Genomet systembiologiens rygrad. Dansk Kemi 3:15-17
- J. Højer-Pedersen; J. Smedsgaard; J. Nielsen (2006) Metabolomet: Et indirekte produkt af genomet. Dansk Kemi 3:18-20
- 15. F. Gatto; J. Nielsen (2014) Särskild metabolism i klarcellig njurcellscancer. Onkologi i Sverige 3:76-80
- 16. A.W. Lykke; B. Palsson; J. Nielsen (2017) Why microbes rule the world and our industries. BioZoom 2:4-5
- 17. B. Palsson; **J. Nielsen** (2017) Biobased production of chemicals A transformation of a USD3T industry. BioZoom 2:6-9

List of Presentations

Invited conference presentations

- 1. Modelling of fermentation kinetics, European Congress on Biotechnology 5, Lyngby (1990)
- 2. Modelling of the lactic acid fermentation, European Congress on Biotechnology 5, Lyngby (1990)
- 3. Modelling of fermentation processes, European Simulation Multiconference, Copenhagen (1991)
- 4. On-line monitoring of microbial processes, FACSS, Anaheim, California (1991)
- 5. Modelling the growth of filamentous microorganisms, IFAC/ICCAFT, Keystone, Colorado (1992)
- 6. On-line monitoring of fermentation processes, Bioreactor Performance, Annual Conference of a Nordic Industrial programme, Stockholm, Sweden (1992)
- 7. On-line måling af bioprocesser, 2. Danske Symposium i Analytisk Kemi, Lyngby, Denmark (1992)
- 8. On-line monitoring of penicillin fermentations, AnaBiotec'92, Noordwijkerhout, Holland (1992)
- 9. Modelling the morphology of filamentous fungi, Bioreactor Performance, Elsingore, Denmark (1993)
- 10. Simulation of bioreactions, ESCAPE-3, Graz, Austria (1993)
- Pellet formation, Process Integration in Biochemical Engineering, European Science Foundation Workshop, Sitges, Spain (1993)
- 12. Physiological Engineering Towards a new science, The 1994 IChemE Research Event, London (1994)
- 13. Modelling the growth and product formation of Penicillium chrysogenum, IUMS Congresses '94, Prague (1994)
- 14. Mathematical models: An excellent tool in physiological studies of filamentous fungi, International conference on Modelling of filamentous fungi, Otocek, Slovenien (1994)
- 15. In-situ and on-line analysis of cultivation processes, International seminar on Analyses in biomass conversion to ethanol, Lund, Sweden (1994)
- 16. A kinetic model for the penicillin biosynthetic pathway in Penicillium chrysogenum, 6'th International conference on Computer Application in Biotechnology, Garmisch-Partenkirchen, Germany (1995)
- 17. Modelling the growth of filamentous fungi, National Biotechnology and Bioengineering Conference, Ixtapa, Mexico (1995)
- 18. Metabolic Control Analysis of the penicillin biosynthetic pathway, National Biotechnology and Bioengineering Conference, Ixtapa, Mexico (1995)
- 19. Metabolic Control Analysis of the penicillin biosynthetic pathway, Pacifichem'95, Honolulu, USA (1995)
- Fungal Morphology. Measurements and modelling, 6th Netherlands Biotechnology Congress, Amsterdam, The Netherlands (1996)
- 21. Metabolic Flux Analysis of filamentous fungi, DECHEMA Jahrestagungen'96, Wiesbaden, Germany (1996)
- 22. Metabolic Engineering: The analysis part, rDNA Biotechnology: Focus on Metabolic Engineering, Engineering Foundation, Danvers, USA (1996)
- 23. What can we learn from metabolic flux analysis? VW-symposium on Metabolic fluxes, Hannover, Germany (1997)
- 24. Metabolic engineering, Danish Biotechnology Conference III, Vejle, Denmark (1997)
- Metabolic control analysis of the penicillin biosynthetic pathway based on a kinetic model and on a thermokinetic description of reaction rates, ESF Conference on Control of Metabolic Flux, Giens, France (1997)
- 26. Physiological Engineering, European Congress on Biotechnology 8, Budapest, Hungary (1997)
- 27. Mathematical modelling of biochemical pathways, 8'th BTK meeting, Fiskebäckskil, Sweden (1998)
- 28. Enzyme production by Aspergillus, SGM meeting, Norwich, UK (1998)
- 29. Metabolic engineering of *Saccharomyces cerevisiae* for the improvement of industrial processes, Yeast as a cell factory, Vlaardingen, The Netherlands (1998)
- 30. The role of mathematical models in kinetic studies of hyphal growth, Analysis of microbial cells at the single cell level, Como, Italy (1999)
- 31. The role of mathematical models in microbial physiology, Annual Meeting of Swedish Society of Microbiology, Lund, Sweden (1999)
- 32. Modeling of cellular processes, European Congress on Biotechnology 9, Brussels, Belgium (1999)
- 33. The application of mathematical models in molecular physiology, European Congress on Biotechnology 9, Brussels, Belgium (1999)
- 34. Metabolic engineering of *Saccharomyces cerevisiae* for the improvement of ethanol production, IEA Bioenergy Workshop, Itala Game Reserve, South Africa (1999)
- 35. Yeast mixed sugar metabolism, Cell Factory Area. Grand Finale, Graz, Austria (1999)
- 36. The role of metabolic engineering in the improvement of industrial processes, APBioChEC'99, Phuket, Thailand (1999)
- Transgenic microorganisms and industrial strain enhancement, Perspectives and limitations of biotechnology in developing countries, San José, Costa Rica (2000)
- 38. Metabolic engineering of enzyme production by filamentous fungi, 4th International Congress on Biochemical Engineering, Stuttgart, Germany (2000)
- 39. Metabolic Engineering of Saccharomyces cerevisiae, XIII SINAFERM, Teresopolis, Brasilien (2000)
- 40. Metabolome analysis: A powerful tool in metabolic engineering and functional genomics, Metabolic Engineering III, Colorado Springs, USA (2000)

- 41. Metabolome analysis: A powerful tool in functional genomics, WFCS Symposium on Genomics, Wageningen, The Netherlands (2000)
- 42. Genome wide expression monitoring of metabolically engineered strains of *S. cerevisiae* with improved ethanol yield, PacificChem2000, Honolulu, USA (2000)
- 43. Metabolome analysis: A powerful tool in metabolic engineering and functional genomics, MTBio Workshop, Dresden, Germany (2001)
- 44. The role of functional genomics in metabolic engineering, SIM Annual Meeting, St. Louis, USA (2001)
- 45. Metabolic engineering for improved β-lactam production, Recent Advances in Fermentation Technology IV, Long Beach, USA (2001)
- 46. The role of yeast in modern biotechnology, International Specialized Symposium on Yeast 2002, Pilansberg, South Africa (2002)
- 47. Metabolic engineering and functional genomics: Moving towards systems biology, SGM Annual Meeting, Warwick, UK (2002)
- 48. Impact of Systems Biology on Modern Biotechnology, Danish Biotechnology Conference VIII, Vejle, Denmark (2002)
- 49. Metabolic engineering for improvement of β-lactam production, Genetics of Industrial Microorganisms 2002, South Korea (2002)
- 50. Carbon metabolism in Aspergillus and Penicillium, International Mycology Congress 7, Oslo, Norway (2002)
- 51. From the Genome to the Fluxome: A Metabolic Engineering challenge, ASM Annual Meeting, Washington DC, USA (2003)
- 52. Systems Biology of glucose repression in *S. cerevisiae*, Yeast Genetics and Molecular Biology, Gothenburg, Sweden (2003)
- 53. From Genomics to Industrial Bioprocesses: A Metabolic Engineering challenge, European Congress on Biotechnology 11, Basel, Switzerland (2003)
- 54. Systems Biology of S. cerevisiae, First International Workshop on Yeast Systems Biology, St. Louis, USA (2003)
- 55. From Glucose to Antibiotics what controls the flux, Ernst Schering Foundation Research Workshop, Berlin, Germany (2004)
- 56. Systems Biology of S. cerevisiae, South Africa Microbial Society's annual meeting, Stellenbosch, South Africa (2004)
- 57. Aspergillus the Ultimate Cell Factory for Production of Chemicals, European Congress on Fungal Genetics 7, Copenhagen, Denmark (2004)
- 58. Microorganisms the Chemical Factories of the Future, ETIF Conference, Lund, Sweden (2004)
- 59. Grøn Kemi, Konference om Ansvarlig Bioteknologi, Copenhagen, Denmark (2004)
- 60. Integration of Metabolic Models and Ome Data: Lessons from S. cerevisiae, ASM Conference on Integration of Metabolism and Genomics, Montreal, Canada (2004)
- 61. Green Chemistry the New S-curve in Biotechnology, 35th R³ Nordic Symposium and Exibition, Elsingore, Denmark (2004)
- 62. Metabolic Engineering, European Symposium on Biochemical Engineering Science 5, Stuttgart, Germany (2004)
- 63. Metabolic Engineering: Impacts of Functional Genomics, Metabolic Engineering V, Lake Tahoe, USA (2004)
- 64. From gene expression to metabolic fluxes, International Congress on Systems Biology 2004, Heidelberg, Germany (2004)
- 65. Integration of the metabolism of *S. coelicolor* through genome-scale modelling, Streptomyces Dissemination Meeting, University of Surrey, UK (2005)
- 66. Design af cellefabrikker til production af nye levnedsmiddelingredienser, LMC Congress, Lyngby, Denmark (2005)
- 67. White Biotechnology: From gene expression to metabolic fluxes, DECHEMA Bioperspectives, Wiesbaden, Germany (2005)
- 68. Identification of global regulatory structures in cellular metabolism, Biochemical Engineering XIV, Harrison Hot Springs, Canada (2005)
- 69. Genome-scale models of fungi, 13th International Meeting on Microbial Genomes (2005)
- 70. The role of chemical engineering in modern biotechnology, CHEMPOR 2005, Braga, Portugal (2005)
- 71. Systems biology of the yeast *Saccharomyces cerevisiae*, The Norwegian Biochemical Society, 42. Contact Meeting, Storefjell, Norway (2006)
- 72. Model driven data integration in yeast systems biology, Genomes to Systems Conference 2006, Manchester, UK (2006)
- 73. Systems Biology of Industrial Microbes, American Society for Microbiology 106th General Meeting, Orlando, USA (2006)
- 74. Impacts of systems biology on industrial biotechnology, Danish Conference on Molecular Biology and Biotechnology, Munkebjerg, Denmark (2006)
- 75. Reporter features: A tool for mapping of global control in metabolism through model driven analysis of ome data, ISSY25 Systems Biology of Yeasts from Models to Applications, Helsinki, Finland (2006)
- 76. Yeast as a versatile cell factory, Genetics of Industrial Microorganisms, Prague, Czech Republic (2006)
- 77. Metabolism of *Aspergillus*: Lessons from Genomics, Society for Industrial Microbiology Annual Meeting, Baltimore, USA (2006)
- 78. Production of engineered haemoglobin from yeast, International Visions on Blood Substitutes, Parma, Italy (2006)
- 79. Systems biology of lipid metabolism in Saccharomyces cerevisiae: Mapping of global regulatory structures, Keystone conference on Bioactive Lipids, Taos, New Mexico, USA (2007)
- 80. Yeast as a versatile chemical factory, ASM Annual Meeting, Toronto, Canada (2007)
- 81. Integrated analysis of yeast metabolism, ASM Annual Meeting, Toronto, Canada (2007)
- 82. Systems biology of the cell factory Aspergillus niger, International Conference on Biorefinery, Beijing, China (2007)

- 83. The role of chemical engineering in modern biotechnology, European Congress on Chemical Engineering 6, Copenhagen, Denmark (2007)
- 84. Integrated analysis of yeast metabolism, FOSBE2007, Stuttgart, Germany (2007)
- 85. Integrated analysis of yeast metabolism, 9th Functional Genomics Conference: Synthetic Biology, Gothenburg, Sweden
- 86. Systems biology of the cell factory Aspergillus niger, RAFT VII, St. Petersburg, Florida, USA (2007)
- 87. Systems Biology as a driver for industrial biotechnology, 2008 BERN meeting, University College London, UK (2008)
- 88. Comparative metabolic analysis of Aspergilli at the genome level, European Conference on Fungal Genetics 9, Edinburgh, UK (2008)
- 89. The role of Snf1/AMPK on regulation of lipid metabolism: Lessons from yeast, Danish Conference on Molecular Biology and Biotechnology II, Munkebjerg, Denmark (2008)
- 90. Modeling of microbial metabolic networks, Tutzing Symposium 2008, Tutzing, Germany (2008)
- 91. Systems Biology of the cell factories A. niger and A. oryzae, FEBS2008, Athens, Greece (2008)
- 92. Systems Biology of the yeast *Saccharomyces cerevisiae*, International Congress on Genetics 2008, Berlin, Germany (2008)
- 93. Systems Biology of lipid and energy metabolism in yeast, International Conference on Yeast, Kiev, Ukraine (2008)
- 94. Industrial systems biology: Yeast and Aspergilli as cell factories for sustainable production of chemicals, International Conference on Systems Biology 2008, Gothenburg, Sweden (2008)
- 95. Production of fine chemicals by yeast and Aspergilli, International Biotechnology Symposium 2008, Dalian, China (2008)
- Integrated analysis of yeast metabolism, 61st Annual Symposium on Cancer Research: Systems Biology of Cancer, Houston, USA (2008)
- 97. The metabolic networks of Aspergillii, Asperfest, Asilomar, USA (2009)
- 98. Metabolism of Aspergilli at the genome level, 25th Fungal Genetics Conference, Asilomar, USA (2009)
- 99. Systems biology of lipid metabolism: Mapping of global regulatory structures, 9th Yeast Lipid Conference, Berlin, Germany (2009)
- 100. Integrated analysis of metabolism; From yeast to human, Nobel Symposium on Systems Biology, Stockholm, Sweden (2009)
- 101. Industrial Systems Biology: Identification of metabolic engineering targets using metabolic networks, Foundation of Engineering in Systems Biology 2009, Denver, USA (2009)
- 102. From omics data to phenotype through integrative systems biology, ISSY 27, Paris, France (2009)
- 103. Systems biology of metabolism: From yeast to mammals, European Congress on Biotechnology 14, Barcelona, Spain (2009)
- 104. Yeast Metabolomics, Metabolomics Conference, University of Copenhagen, Copenhagen, Denmark (2009)
- 105. Prospects of Systems Biology for advancing our understanding of global regulation of metabolism, Annual INRA Conference, Genoscope, France (2010)
- 106. Genome-Scale Modeling of Fungi: Expansion from Metabolism to Protein Production, PYFF4, Rotterdam, The Netherlands (2010)
- 107. Yeast as a Platform Cell Factory for the Production of Fuels and Chemicals, MEVIII, Jeju Island, South Korea (2010)
- 108. Metabolism of Filamentous Fungi at the Genome Level, FEBS2010, Gothenburg, Sweden (2010)
- 109. Insight into Metabolic Diseases through Systems Biology, 8th Aegian Conference on Pathways, Networks and Systems, Rhodes, Greece (2010)
- 110. Genome-Scale Metabolic Models: The Core of Industrial Systems Biology, Industrial Systems Biology 2010, Gothenburg, Sweden (2010)
- 111. Yeast as a platform cell factory for production of fuels and chemicals. International Biotechnology Symposium 2010, Rimini, Italy (2010)
- 112. Yeast as a platform cell factory. Power of Microbes in Industry and Environment 2010, Island Krk, Croatia (2010)
- 113. The application of genome-scale metabolic models in industrial biotechnology, ICSB2010, Edinburgh, UK (2010)
- 114. Synthetic biology and industrial biotechnology, Green Chem Conference, Lund, Sweden (2010)
- 115. Development of yeast as a platform cell factory, 3rd International Conference on Biomolecular Engineering, San Fransisco, USA (2011)
- 116. Engineering of the Biocatalyst in Biorefineries, 3rd International Symposium for Innovation Bioproduction Kobe, Kobe, Japan (2011)
- 117. The Art of Making Yeast to Produce Biodiesel, Chalmers Energy Conference 2011, Gothenburg, Sweden (2011)
- 118. Yeast as Platform Cell Factories in Future Biorefineries, AAAS Annual Symposium 2011, Washington, USA (2011)
- 119. Yeast as Platform Cell Factories in Future Biorefineries, Asian Congress on Biotechnologyn 2011, Shanghai, China (2011)
- 120. Nutritional Systems Biology, Experimental Biology 2011, Washington, USA (2011)
- 121. Yeast as Platform Cell Factories in Future Biorefineries, Danish Conference on Biotechnology 6, Munkebjerg, Denmark (2011)
- 122. Towards the Human Metabolic Atlas, 8th Key Symposium, Stockholm, Sweden (2011)
- 123. Application of Genome Scale Metabolic Models in Industrial Biotechnology, 1st COBRA Conference, Reykavik, Iceland (2011)
- 124. Metabolic Engineering, Synthetic Biology, Systems Biology, ... what is the role of Biochemical Engineering, Biochemical and Biomolecular Engineering XVII, Seattle, USA (2011)
- 125. Integration of Metabolomics Data into Metabolic Networks, Metabomeeting 2011, Helsinki, Finland (2011)
- 126. Gut Metagenomics and its relation to Systems Medicine, Science for Life Laboratory Annual Conference 2012, Uppsala, Sweden (2012)

- 127. Synthetic Biology and Metabolic Engineering of Yeast, EFB Applied Synthetic Biology in Europe, Barcelona, Spain (2012)
- 128. Yeast as a Platform Cell Factory for Biorefineries, APCChE2012, Singapore (2012)
- 129. Prospects for systems biology and modelling of the gut microbiome, MetaHit, Paris, France (2012)
- 130. Systems Biology of Metabolism: Impact of Human Health and Industrial Biotechnology, Molecular Frontiers Symposium, Singapore (2012)
- 131. Systems biology of industrially important filamentous fungi, 11th European Congress on Fungal Genetics, Marburg, Germany (2012)
- 132. Modeling and Engineering of Protein Secretion by Yeast, Microbial Stress: From Molecules to Systems, Belgirate, Italy (2012)
- 133. Systems Biology of Metabolism: Enabling Technologies for Metabolic Engineering of Yeast, Metabolic Engineering IX, Biarritz, France (2012)
- 134. Genome-Scale Metabolic Models: A Bridge Between Bioinformatics and Systems Biology, Nordic Bioinformatics Conference, Stockholm (2012)
- 135. Towards a Human Metabolic Atlas, Nordic Clinical Chemistry Conference, Reykavic, Iceland (2012)
- 136. Genome-Scale Metabolic Models: A Bridge Between Bioinformatics and Systems Biology, ICSB2012, Toronto, Canada (2012)
- 137. Systems Biology of Saccharomyces cerevisiae Metabolism: Impact on Cell Factory Design, ICY2012, Madison, USA, 2012.
- 138. Systems Biology and its Application in Nutritional Studies, IVA Symposium "Mat som Medicin", Gothenburg, Sweden (2012)
- 139. The Sense of Biofuels, IVA Symposium "Technologies for the Future Visions and Opportunities", Stockholm, Sweden (2012)
- 140. Yeast as a Platform Cell Factory in Future Biorefineries, 4th International Conference on Biomolecular Engineering, Fort Lauderdale, USA (2013)
- 141. Systems Biology: Yeast as a platform cell factory, Annual B-Basic Conference, Nordweijkerhout, The Netherlands (2013)
- 142. Yeast as a platform cell factory in future biorefineries, Copenhagen Bioscience Conference on Biosustainability, Hillerød, Denmark
- 143. Advancing metabolic engineering of fungal cell factories through systems biology, PYFF5, Montpeiller, France (2013)
- 144. Developing yeast cell factories for sustainable production of chemicals, Biochemical and Molecular Engineering XVIII, Beijing, China (2013)
- 145. Role of Systems and Synthetic Biology in Metabolic Engineering, Tsinghua Mini-Symposium on Synthetic Biology, Beijing, China (2013)
- 146. Yeast as a platform cell factory in future biorefineries, Genetics of Industrial Microorganisms 2013, Cancun, Mexico (2013)
- 147. Engineering of yeast cell factories for the production of advanced biofuels, Genetics of Industrial Microorganisms 2013, Cancun, Mexico (2013)
- 148. From metagenomics to systems biology of the gut ecosystem, Metagenopolis Conference 2013, Jouy-en-Josas, France (2013)
- 149. Genome scale modelling for unravelling the mechanisms of cancer, Chalmers Conference on Life Science Engineering, Gothenburg, Sweden (2013)
- 150. Regulation of the yeast metabolome, ICSB2013, Copenhagen, Denmark (2013)
- 151. Engineering of yeast cell factories for production of advanced biofuels, Lignofuels 2013, London, UK (2013)
- 152. Metabolic engineering of yeast for production of fuels and chemicals, Industrial Biotechnology: Meeting the challenges, Lund, Sweden (2013)
- 153. Use of genome-scale metabolic models for integrative analysis, SILS Conference, Öresundsbro, Sweden (2013)
- 154. Systems Biology: From microbe to man. Putting microbial genomes to work, Delft, The Netherlands (2013)
- 155. Yeast as a platform cell factory for production of fuels and chemicals. AMBC2014, Bangkok, Thailand (2014)
- 156. Systems biology for identification of novel metabolic engineering targets. Metabolic Engineering X, Vancouver, Canada (2014) (also available on-line at www.aiche.org/sbe/)
- 157. Systems biology and metabolic engineering of yeast. IUMS2014, Montreal, Canada (2014)
- 158. Genome wide reprogramming of human metabolism to obesity and cancer. International Summer Symposium on Systems Biology, INMEGEN, Mexico City, Mexico (2014)
- 159. Combining systems biology and adaptive laboratory evolution for mapping genotype-phenotype relaionsships in yeast. Frontiers in Fungal Systems Biology, EMB, Heidelberg, Germany (2014)
- 160. Studies of metabolic diseases and cancer through genome-scale metabolic modelling of human metabolism, Cell Symposia: Systems Approach to Metabolic Diseases, Chicago, USA (2014)
- 161. Aquiring novel phenotypes of yeast through adaptive laboratory evolution, ISSY31, Nova Gorcia, Slovenia (2014)
- 162. Impacts of systems biology on synthetic biology of yeast, International Symposium on Synthetic Biology, Beijing, China (2014)
- 163. Integrative analysis of lipid metabolism through genome-scale metabolic modelling, Keystone Symposium on Systems Biology of Lipid Metabolism, Breckenridge, USA (2015)
- 164. New insights into cancer and obesity related diseases through systems biology of human metabolism, Systems Biology: Networks, Cold Spring Harbor Laboratory, Cold Spring Harbor, USA (2015)
- 165. Identifying genome-level metabolic reprogremming in cancer cells, DFG Hinterzartener Kreis für Krebsforschung, Lago de Como, Italy (2015)

- 166. Yeast as a platform cell factory for production of fatty acid derived products, Copenhagen Bioscience Conference on Biosustainability, Hillerød, Denmark (2015)
- 167. Metagenome analysis of the human gut microbiome, Nobel Symposium, Gothenburg, Sweden (2015)
- 168. Combining systems biology and adaptive laboratory evolution for mapping genotype-phenotype relations in yeast, Ho-Am Forum, Seoul, South Korea (2015)
- 169. Impact of synthetic biology on metabolic engineering of yeast, Eukaryotic Synthetic Biology, EMBO/EMBL Symposium, Heidelberg, Germany (2015)
- 170. Studies of metabolic diseases and cancer through genome-scale metabolic modelling of human metabolism, 11th
 International Conference on Pathways, Networks and Systems Medicine, Aegean Conferences, Crete, Greece (2015)
- 171. Acquiring novel phenotypes of yeast through adaptive laboratory evolution, Adler Symposium, University of Gothenburg, Gothenburg, Sweden (2015)
- 172. Mathematical modelling of yeast: A driver for innovation in biotechnology and medicine. 27th International Conference on Yeast Genetics and Molecular Biology, Levico Terme, Italy (2015)
- 173. Global regulation of yeast metabolism. 27th International Conference on Yeast Genetics and Molecular Biology, Levico Terme, Italy (2015)
- 174. Metabolism of the gut microbiome: New insights through genome-scale metabolic modelling. 4th Conference on Construction and Analysis, Heidelberg, Germany (2015)
- 175. Systems biology of yeast metabolism. IMYA11, Porto, Portugal (2015)
- 176. Metabolism of the gut microbiome: New insights through genome-scale metabolic modelling. Copenhagen Bioscience Conference on Metabolism, Hillerød, Denmark (2015)
- 177. Integrative analysis of omics data for studying metabolism in human disease. BILS Annual Meeting, Stockholm, Sweden
- 178. Meabolism of the gut microbiome: New insights through genome-scale metabolic modelling. The First KSSEA Workshop, Stockholm, Sweden (2015)
- 179. Advancing the Design-Build-Test Cycle for Metabolic Engineering. Metabolic Engineering Summit 2015, Beijing, China (2015)
- 180. Systems biology for mapping genotype-phenotype relations in yeast. KAUST Research Conference on Computationa and Experimental Interfaces of Big Data and Biotechnology, KAUST, Saudi Arabia (2016)
- 181. Systems Biology of yeast metabolism. Jahrestagung der Vereinigung für Allgemeine und Angewandte Mikrobiologie (VAAM), Jena, Germany (2016)
- 182. Metabolic engineering of yeast. Biosystems Design 2.0, A-Star, Singapore (2016)
- 183. Yeast as a platform cell factory. Gaden Award Lecture, ACS Annual Meeting, San Diego, USA (2016)
- 184. Assessing the human gut microbiota in metabolic disease. FEAM Workshop on Precission Medicine, Bern, Switzerland (2016)
- 185. Metabolic engineering of yeast. Metabolic Engineering 12, Awaji Island, Japan (2016)
- 186. Engineering yeast metabolism for production of fuels and chemicals. European Congress on Biotechnology, Krakow, Poland (2016)
- 187. Systems biology of yeast metabolism. ICY2016, Awaji Island, Japan (2016)
- 188. Systems biology of yeast metabolism. FISV2016, Rome, Italy (2016)
- 189. Systems biology of industrial microorganisms. 13th International Symposium on the Genetics of Industrial Microorganisms, Wuhan, China (2016)
- 190. Genome-scale metabolic modelling of yeast. The International Conference on Metabolic Science, Shanghai, China (2016)
- 191. Biobased production of fuels and chemicals: Barriers and opportunities. International Forum on Innovation and Emerging Industries Development (IEID), Shanghai, China (2016)
- 192. A systems biology approach to measuring human-microbiome interactions. Microbiome Drug Development Summit Europe, Paris (2017)
- 193. Biomarker and diagnostics development using omics integration into human metabolic networks. 12th Biomarker Congress, Manchester, UK (2017)
- 194. Assessing the human gut microbiota in metabolic diseases. Symbiosis in Evolution, Biology and Human Health. IBPS International Symposium, Paris, France (2017)
- 195. Systems Biology of Metabolism. BioSB 2017, Lunteren, The Netherlands (2017)
- 196. Systems Biology of Yeast Metabolism. 12th International Meeting on Yeast Apoptosis, Bari, Italy (2017)
- 197. Systems Biology of Yeast Lipid Metabolism. 13th Yeast Lipid Conference, Paris, France (2017)
- 198. Controlling Metabolism: From Microorganisms to Human. Molecular Frontiers Symposium, Royal Swedish Academy of Science, Stockholm, Sweden (2017)
- 199. Metabolic engineering of yeast for production of fuels and chemicals. 14th International Conference on Renewable Resources and Biorefineries, Wroclaw, Poland (2017)
- 200. Solving environmental challenges using life science, Where is Life Science, Stockholm, Sweden (2017)
- 201. Quantifying global rewiring of metabolism in cancer through genome scale modelling. Feeding the beast the metabolic landscape of the tumour and its host, Glasgow, Scotland (2017)
- 202. Systems metabolic engineering of yeast. 28th International Conference on Yeast Genetics and Molecular Biology, Prague, Czech Republic (2017)
- 203. Systems biology of yeast metabolism. Emerging Topics in Biological Networks and Systems Biology, Uppsala, Sweden (2017)
- 204. Synthetic biology of yeast. 6th International Conference of the International Chemical Biology Society, Shanghai, China (2017)
- 205. Systems biology of yeast metabolism. Metabolic Engineering Summit 2017, Beijing, China (2017)

- 206. Engineering yeast for production of advanced biofuels. Fuel Choices and Smart Mobility Summit 2017. Tel Aviv, Israel (2017)
- 207. Quantifying global rewiring of metabolism in cancer through genome-scale modelling. The 1st International Symposium for Trans-Omics, University of Tokyo, Japan (2017)
- 208. Metabolic engineering of yeast for production of fuels and chemicals. SynBio UK 2017, Manchester, UK (2017)
- 209. Metabolic engineering of yeast. Sino-Swedish Workshop, Shandong University, Qingdao, China (2018)
- 210. Systems biology of yeast metabolism. Advancing Synthetic Biology, Beijing University of Chemical Technology, Beijing, China (2018)
- 211. Systems biology of yeast metabolism. Metabolic Engineering 12, Munich, Germany (2018)
- 212. Genome-scale modelling of yeast. COBRA5, Seattle, USA (2018)
- 213. Systems biology of yeast metabolism. EMBO Workshop on Experimental Approaches to Evolution and Ecology Using Yeast and Other Model Systems, Heidelberg, Germany (2018)
- 214. Engineering yeast metabolism using synthetic biology. 2018 World Life Science Conference, Beijing, China (2018)
- 215. Systems biology of yeast metabolism. Nature Conference on Cellular Metabolism, Xiamen, China (2019)
- 216. Entrepreneurship in Biotechnology. Danish Biotechnology Conference 14, Munkebjerg, Denmark (2019)
- 217. Systems Biology of Yeast Metabolism. Emerging Applications of Microbes, VIB Conferences, Leuven, Belgium (2019)
- 218. Synthetic Biology of Yeast. XXIX International Conference on Yeast Genetics and Molecular Biology, Gothenburg, Sweden (2019)
- 219. Metabolic Engineering of Yeast. Metabolic Engineering Summit 2019, Tianjin, China (2019)
- 220. Synthetic Biology of Yeast. Green Manufacturing International Conference, Beijing, China (2019)
- 221. Systems Biology of Metabolism: Role in Production of Advanced Biofuels, Obesity and Cancer. James E. Bailey Award lecture, AIChE Annual Meeting, Orlando, USA (2019)
- 222. Systems Biology of Metabolism. Trumping the Trumps. Symposium in honor of Prof. Hans Westerhof. Free University of Amsterdam, The Netherlands (2020)
- 223. The power of microbes. Global Summit, Hello Tomorrow, Paris (2020)
- 224. Synthetic Biology of yeast. Biocatalysis Open Day 2020, Virtual (2020)
- 225. Systems Biology of Metabolism. 7th Conference on Constraint Based Reconstruction and Analysis. IMES/AIChE, Virtual (2021)
- 226. Metabolic Engineering of Yeast. Symposium on Synthetic Biology and Natural Product Biosynthesis, Chinese Academy of Chinese Medical Sciences, Beijing, China (2021)
- 227. Engineering of yeast for production of food ingredients. International Summit Forum on Future Food and Biotechnology 2021, Wuxi, China (2021)
- 228. Metabolic Engineering of Yeast. 43rd Symposium on Biomaterials, Fuels and Chemicals, SIMB, Virtual (2021)
- 229. From Science to Market. Metabolic Engineering 14, Viirtual (2021)
- 230. Systems Biology of Yeast Metabolism. Metabolic Engineering 14, Virtual (2021)
- 231. Systems Biology of Yeast Metabolism. 61st meeting of Italian Society for Biochemistry and Molecular Biology, Viirtual (2021)
- 232. Metabolic engineering of yeast. Brazilian Conference on Biotechnology and Bioengineering. Virtual (2021)
- 233. Synthetic Biology of Yeast for Production of Food Ingredients. 2021 MiFFI, Copenhagen, Denmark (2021)
- 234. Systems Biology of Yeast Metabolism. Major Ideas in Quantitative Biology, Copenhagen, Denmark (2022)
- 235. Systems Biology of Metabolism. EMBO Annual Conference, Heidelberg, Germany (2022)
- 236. BioInnovation Institute. MiXii Conference, Jerusalem, Israel (2022)
- 237. Innovation trends in metabolic engineering. ME13, Singapore (2023)
- 238. Systems biology of yeast metabolic. European Congress on Chemical Engineering, Berlin, Germany (2023)
- 239. Metabolic engineering of yeast. Korean Society for Biotechnology and Bioengineering Fall Meeting, Busan, Korea (2023)

Webinars and other web-based talks (not complete)

- 1. Industrial Systems Biology, Society for Biological Engineering, 2009. www.aiche.org/sbe/
- 2. *Impact of systems biology on metabolic engineering*, in H. Kitano (Ed), Systems Biology: The Biomedical and Life Science Collection, Henry Stewart Talks Ltd., London, 2014. http://hstalks.com/?t=BL1893791-Nielsen
- 3. Advancing the design-build-test cycle for metabolic engineering of yeast. SBE Webinar, AIChE, USA
- 4. Systems Biology of Yeast Metabolism. CHASSY Webinar, University of Cork, Ireland

Invited seminar presentations (not complete)

- 1. Design of an on-line monitoring system for lactic acid fermentations, Ciba Geigy, Basel, Switzerland (1987)
- 2. Automation of laboratory fermentors, Technical University of Denmark, Lyngby (1987)
- 3. On-line monitoring of lactic acid fermentation, Department of Chemical Engineering, Caltech, Pasedena, California (1988)
- 4. Application of FIA for on-line monitoring of fermentation processes, Technion, Lund, Sweden (1988)
- 5. Structured modelling of microbial systems, Technical University of Denmark, Lyngby (1989)
- 6. Experimentally verification of fermentation models, University of Lund, Sweden (1990)

- 7. Structural models for fermentation processes, Departmental seminar, Institut für Biotechnologie, Technische Universität Graz, Austria (1990)
- 8. Modelling and on-line monitoring of fermentation processes, Departmental seminar, Institut für Technische Chemie, Universität Hannover, Germany (1990)
- 9. Verification of structured models for fermentation processes, Departmental seminar, Kemisk Reaktionsteknik, Chalmars Tekniska Högskola, Gothenburg, Sweden (1991)
- Modelling of filamentous microorganisms, Department of Chemical Engineering, University of Michigan, Ann Arbor, Michigan (1992)
- 11. On-line monitoring and modelling of microbial processes, Departmental seminar, Department of Biochemical Engineering, Indian Institute of Technology, New Delhi (1992)
- 12. Growth of filamentous fungi An engineers perspective, Seminar at University of Manchester, Manchester (1992)
- 13. On-line monitoring of filamentous fungi fermentations, Meeting of EFB Working Party on "Measurement and Control", Florence (1993)
- Modelling of filamentous growth, Meeting of DECHEMA Working Party on "Messung und Regelung in Biotechnologie", Frankfurt (1993)
- 15. Industriel anvendelse af skimmelsvampe, Dansk Ingeniørforening, Copenhagen (1994)
- 16. Metabolic flux analysis of the penicillin fermentation, Institute of Chemical Metalurgy, Chinese Academy of Science, Beijing (1994)
- 17. Mathematical models An excellent tool in physiological studies of filamentous fungi, Department of Chemical Engineering, University of Minnesota, Minneapolis (1994)
- 18. Physiological Engineering of filamentous fungi, Departmental Seminar, Department of Chemical Engineering, MIT, Cambridge (1994)
- Physiological Engineering. The integration of microbial physiology and chemical engineering, Department of Chemical Engineering, MIT, Cambridge, USA (1996)
- 20. Modelling the growth of filamentous microorganisms, Departmental seminar, Department of Chemical Engineering, Tufts University, Medford, USA (1996)
- 21. Modelling the growth of filamentous microorganisms, Departmental seminar, Department of Chemical and Biochemical Engineering, University of Western Ontario, London, Canada (1996)
- 22. Modelling the growth of filamentous fungi, Department of Chemical Engineering, MIT, Cambridge (1996)
- 23. Metabolic Engineering, University of Shandong, Jinan (1996)
- 24. Metabolic flux analysis of filamentous fungi, BASF, Ludwigshafen (1996)
- 25. Metabolic Flux Analysis, Departmental seminar, Department of Chemical Engineering, Xinghua University, Beijing, China (1996)
- Biochemical analysis of the penicillin biosynthetic pathway, Departmental Seminar, Department of Microbiology, Technical University of Denmark, Lyngby, Denmark (1997)
- Biochemical characterization of the penicillin biosynthetic pathway, Departmental Seminar, Institut für Biotechnologie, ETH, Zürich, Switzerland (1997)
- 28. Metabolic engineering. Methods and applications, Institute seminar, Institut für Biotechnologie, Forschungszentrum Jülich, Germany (1998)
- 29. Enzyme production by Aspergillus. Biochemical engineering methods for fundamental research and process optimisation, Seminar, Carlsberg Laboratory, Copenhagen, Denmark (1999)
- 30. Metabolomics, Seminar, Biologisk Selskab, Copenhagen, Denmark (2000)
- 31. Metabolic Engineering and Functional Genomics, Seminar, Novo Nordisk, Bagsværd, Denmark (2000)
- 32. Metabolic Engineering and Functional Genomics, Seminar, Chr. Hansen, Hørsholm, Denmark (2000)
- 33. Metabolic Engineering and Functional Genomics, Bjerrum-Brøndsted-Lang lecture, Carlsberg Laboratory, Denmark (2000)
- 34. Metabolic engineering of Penicillium chrysogenum for improved β-lactam production, Seminar, Microbia, Cambridge, USA (2001)
- 35. Analysis of metabolism: Control of fluxes, Seminar, BASF, Ludwigshafen, Germany (2001)
- 36. The role of Functional Genomics in Metabolic Engineering, Seminar, EPFL, Lausanne, Switzerland (2001)
- 37. Metabolomics, Seminar, Genomics in Food Science, KVL, Denmark (2002)
- 38. From quantitative physiology to metabolic engineering and systems biology, Sunner Memorial Lecture, Lund University, Lund, Sweden (2002)
- 39. Hough Memorial Lecture, University of Birmingham, Birmingham, UK (2004)
- 40. Metabolic engineering: Impacts of functional genomics, Seconda Università degli Studi di Napoli, Naples, Italy (2005)
- 41. Linking the transcriptome and the metabolome through genome-scale metabolic models, University Milano-Bicocca, Milan, Italy (2005)
- 42. Systems Biology: Current status and future challenges, NTNU, Trondheim, Norway (2005)
- 43. Systems Biology of Yeasts: Impacts on Metabolic Engineering and Basic Sciences, The National Hellenic Research Foundation, Athens, Greece (2006)
- 44. Impacts of systems biology on biotech process based on yeast, Merck, West Point, USA (2006)
- 45. The role of metabolic engineering in the improvement of industrial processes, BIOTEC, Bangkok, Thailand (2006)
- 46. Yeast systems biology: A vehicle for medical and biotechnological research, Department of Chemical and Biological Engineering. Chalmers University of Technology, Gothenburg, Sweden (2007)
- 47. Yeast as a model organism for studying nutragenomic, Department of Systems Biology, ETH Zürich, Switzerland (2007)
- 48. Systems Biology and Synthetic Biology, Beijing University of Chemical Technology, Beijing, China (2007)
- 49. Systems Biology of lipid metabolism in yeast, Department of Biotechnology, University of Graz, Austria (2008)

- 50. Systems Biology as a driver for industrial biotechnology, Department of Chemical Engineering, Catholic University of Chile, Santiago, Chile (2008)
- 51. Systems Biology of Lipid Metabolismi, Departmental Seminar, Institute for Molecular Biosciences, Universität Graz, Austria (2008)
- 52. Industrial systems biology: Yeast and Aspergillus as cell factories for sustainable production of chemicals, Sandoz, Kundl, Austria (2008)
- 53. Industrial Systems Biology: Yeast and Filamentous Fungi as Cell Factories for Sustainable Production of Chemicals, Departmental Seminar, Department of Chemical Engineering, Rice University, Houston, Texas, USA (2008)
- Yeast systems biology: Does this have any medical relevance? Wallenberg Laboratory, Sahlgrenska Academy, Gothenburg, Sweden (2008)
- 55. Industrial systems biology, Amyris, Berkeley, USA (2009)
- 56. Understanding the function of biological networks through systems biology, Faculty of Science and Mathematics, University of Zagreb, Zagreb, Croatia (2009)
- 57. Systems biology of metabolism: From yeast to mammals, Department of Bioengineering, UC San Diego, California, USA (2009)
- 58. Industrial systems biology, Genomatica, San Diego, USA (2009)
- Industrial systems biology: Yeast and Aspergilli as cell factories for sustainable production of chemicals. CJ Company, Seoul, South Korea (2010)
- 60. Engineering the metabolism of yeast cell factories, Novo Nordisk Foundation, Denmark (2010)
- 61. Development of yeast as a platform cell factory for production of fuels and chemicals through industrial systems biology, Tufts University, Boston, USA (2010)
- 62. Impact of systems biology on synthetic biology, Yale University, New Haven, USA (2010)
- 63. Yeast Systems Biology, Carlsberg Laboratory, Copenhagen, Denmark (2010)
- 64. Biorefinery. Cell Factory Design and Implementation, World Council of Industrial Biotechnology meeting, Beijing, China (2010)
- 65. Development of Yeast as a Platform Cell Factory, Joint Bioenergy Institute, Berkeley, USA (2011)
- 66. Insights into Metabolic Diseases through Systems Biology, SomaLogic, Boulder, USA (2011)
- 67. Systems Biology: Integrated Analysis of Human Metabolism through the Human Metabolic Atlas, Rigshospitalet, Copenhagen, Denmark (2011)
- 68. Yeast as a Cell Factory Platform for Production of Fuels and Chemicals, University of Umeå, Umeå, Sweden (2011)
- 69. Biochemical Engineering as the Foundation of Systems Biology, Synthetic Biology and Metabolic Engineering, Seminar in connection with Prof. Reuss, University of Stuttgart, Germany (2011)
- 70. Systems Biology, Synthetic Biology and Metabolic Engineering of Yeast, Genomatica, San Diego, USA (2011)
- 71. The role of DNA and RNA sequencing in systems biology, University of Gothenburg, Gothenburg, Sweden (2012)
- 72. Metabolic modelling applied to metabolic engineering and systems medicine, Departmental Seminar, Department of Chemical Engineering, MIT, Cambridge, USA (2012)
- 73. Systems biology of yeast, Gevo, Denver, USA (2012)
- 74. Production of recombinant proteins by yeast, Department of Chemical and Biological Engineering, Chalmers University of Technology, Gothenburg, Sweden (2012)
- 75. The Human Metabolic Atlas: A novel resource for studying cancer metabolism, Dana Faber Center for Cancer Systems Biology, Boston, USA (2012)
- 76. Genome-scale metabolic models as a scaffold for integrative analysis of metabolomics data, Umeå University, Umeå, Sweden (2013)
- 77. Systems biology: Yeast as a platform cell factory, Shanghai Jiao Tong University, Shanghai, China (2013)
- 78. The Human Metabolic Atlas: A novel resource for studying metabolic diseases and cancer, Astra-Zeneca Bioinformatics Seminar, Mölndal, Sweden (2014)
- 79. Yeast systems biology, Odense University, Odense, Denmark (2014)
- 80. Yeast as a platform cell factory for production of fuels and chemicals, Genomics Science, UNAM, Mexico City, Mexico (2014) (Webinar)
- 81. Metabolism and its role in production of advanced biofuels, obesity and cancer. KVVS, Gothenburg, Sweden (2014)
- 82. Systems Biology: From microbe to man. Department of Genetics and Microbiology, Trinity College Dublin, Ireland (2014)
- 83. Systems Biology: From Microbe to Man. Department of Chemistry, EPFL, Lausanne, Switzerland (2014)
- 84. Systems Biology of Metabolism: A Journey from Yeast to Cancer. Svenska Kemistsamfundet, Gothenburg, Sweden (2014)
- 85. Impact of Systems Biology on Industrial Biotechnology. Systems Biology and Systems Medicine, Lake Como School of Advanced Studies, Como, Italy (2014)
- 86. Genome wide reprogramming of human metabolism in response to obesity and cancer. Systems Biology and Systems Medicine, Lake Como School of Advanced Studies, Como, Italy (2014)
- 87. Genome wide reprogramming of human metabolism in response to obesity and cancer. Kemiska Institut, Ljubljana, Slovenia (2014)
- 88. From metagenomics to systems biology of the gut ecosystem. MetaCardis Annual Meeting, Gothenburg, Sweden (2014)
- 89. Metabolic Engineering and Systems Biology of Yeast. School of Life Sciences, Tsinghua University, Beijing, China (2014)
- 90. Yeast as a platform cell factory in future biorefineries. Institute of Biotechnology, Wuhan University, Wuhan, China (2014)
- 91. Metabolic engineering and synthetic biology of yeast. Energy Biosciences Institute, UC Berkeley, USA (2014)

- 92. Prospects for systems biology and modelling of the gut microbiome. EPFL, Switzerland (2015)
- 93. The Human Metabolic Atlas: A resource for studying metabolic diseases. SciLifeLab Day, Karolinska Institute, Stockholm, Sweden (2015)
- 94. Metabolic engineering and synthetic biology of yeast. Chung-Ang University, Seoul, South Korea (2015)
- 95. Genome scale modelling of human metabolism. University of Ljubljana, Ljubljana, Slovenia (2015)
- 96. Mapping Genotype Phenotype Relations in Yeast through Systems Biology. Novozymes Symposium, Copenhagen, Denmark (2015)
- 97. Yeast as a Platform Cell Factory for Production of Fuels and Chemicals. Zhang Dayu Lectureship, Dalian Institute for Chemical Physics, Chinese Academy of Science, Dalian, China (2015)
- 98. Impacts of Systems Biology on Metabolic Engineering. Department of Chemical Engineering, University of Princeton, Princeton, USA (2015)
- 99. Big data in life science: Cancer and gut microbiome research. Big Data Seminar, Chalmers University of Technology, Gothenburg, Sweden (2016)
- 100. Yeast as a platform cell factory for the production of fuels and chemical, Novozymes Prize Lecture, Chalmers University of Technology, Gothenburg, Sweden (2016)
- 101. Metabolic engineering, synthetic biology and microbiome, Ajinomoto, Tokyo, Japan (2016)
- 102. Metabolic engineering of yeast, Beijing University of Chemical Technology, Beijing, China (2016)
- 103. Metabolic engineering of yeast, Tian Gong Symposium, Tianjin Institute of Industrial Biotechnology, Chinese Academy of Sciences, Tianjin, China (2016)
- 104. Systems Biology of Metabolism. Volterra Lecture, Norwegian University of Life Sciences, Ås, Norway (2017)
- 105. Studies of metabolic diseases and cancer using gneome-scale metabolic models for human metabolism. Astra Zeneca, Mölndal, Sweden (2017)
- 106. Systems biology of yeast metabolism. Manchester Institute for Biotechnology, Manchester University, UK (2017)
- 107. Yeast as a platform cell factory for the production of fuels and chemicals. Shanghai Institute for Plant Physiology and Ecology, Chinese Academy of Science, Shanghai, China (2017)
- 108. Yeast as a platform cell factory for the production of fuels and chemicals. East China University of Science and Technology, Shanghai, China (2017)
- 109. Yeast as a platform cell factory for the production of fuels and chemicals. Jiangnan University, Wuxi, China (2017)
- 110. Metabolic engineering of yeast for production of fuels and chemicals. Department of Life Science, Swedish Agricultural University, Uppsala, Sweden
- 111. Towards biobased production of fuels and chemicals. Politechnico di Torino, Torino, Italy (2017)
- 112. Towards biobased production of fuels and chemicals. Universita degli Studi della Basillicata, Portenza, Italy (2017)
- 113. Metabolic engineering of yeast for production of fuels and chemicals. Department of Molecular Sciences, Swedish University of Agricultural Sciences, Uppsala, Sweden (2017)
- 114. What does biofuels and cancer have in common? Royal Academy of Engineering Sciences, Stockholm, Sweden (2017)
- 115. Assessing the human gut microbiota in metabolic disease. Royal Academy of Engineering Sciences, Stockholm, Sweden (2017)
- 116. Research leadership, mentoring, publishing and innovation. Copenhagen Bioscience Lectures, the Novo Nordisk Foundation, Copenhagen, Denmark (2018)
- 117. Systems biology of yeast metabolism. Tartu University, Tartu, Estonia (2018)
- 118. Systems biology of yeast metabolism. EPLF, Lausanne, Switzerland (2018)
- 119. Systems biology of yeast metabolism. Technical University of Denmark, Lyngby, Denmark (2018)
- 120. Systems biology of yeast metabolism. China National Gene Bank, Shenzhen, China (2018)
- 121. Metabolic engineering of yeast. Chinese Academy of Traditional Chinese Medicines, Beijing, China (2018)
- 122. Metabolic engineering of yeast. Beijing Technology and Management University, Beijing, China (2018)
- 123. Systems biology of yeast metabolism. Zymergen, Emmeryville, USA (2018)
- 124. Metabolic modelling of the human gut microbiome. SERES Pharmaceuticals, Cambridge, USA (2018)
- 125. Systems biology of yeast metabolism. Ginko Bioworks, Boston, USA (2018)
- 126. Quantifying cancer metabolism through genome-scale metabolic modelling, Koch Institute, MIT, Cambridge, USA (2018)
- 127. Systems biology of yeast metabolism. Joint BioEnergy Institute, Emmeryville, USA (2018)
- 128. Systems biology of metabolism. California Institute of Technology, Pasadena, USA (2018)
- 129. Systems biology of yeast metabolism. Provivi, Santa Monica, USA (2018)
- 130. Systems biology of yeast metabolism. Jiangnan University, Wuxi, China (2019)
- 131. Systems biology of yeast metabolism. Steve Fest, University of Cambridge, Cambridge, UK (2019)
- 132. Systems Biology of Yeast Metabolism. Peiyang Lecture, Tianjin University, Tianjin, China (2019)
- 133. Metabolic Engineerin of Yeast. Nankai University, Tianjin, China (2019)
- 134. How studying metabolism can impact production of biofuels and identifying novel cancer biomarkers. PhD student Day, Department of Biology, University of Copenhagen, Denmark (2019)
- 135. Systems Biology of Metabolism. NYU School of Medicine, Institute for Systems Genetics, USA (2020)
- Systems Biology: A Driver for Metabolic Engineering. Department of Chemical and Biomolecular Engineering, KAIST, South Korea (2020)
- 137. Systems biology of human metabolism. Department of Chemical and Biomolecular Engineering, KAIST, South Korea (2020)
- 138. Systems Biology of Yeast Metabolism. Unviersity of Texas, Virtual (2021)
- 139. Systems Biology of Yeast. Pacific Northern National Laboratory, Virtual (2021)
- 140. Systems biology of yeast metabolism. NNF Center for Biosustainability annual meeting, Elsinore, Denmark (2021)
- 141. Life Science Innovation. NNF Center for Biosustainability, DTU, Lyngby, Denmark (2021)

- 142. Systems biology of Human metabolism. FEBS Course, Stockholm, Sweden (2022)
- 143. Systems Biology of Metabolism. Sahlgrenska Academy, Gothenburg, Sweden (2023) 144. Systems Biology of Metabolism. Center for Basic Metabolic Research, University of Copenhagen, Denmark (2023)
- 145. Innovation trends in industrial biotechnology. KAIST, Daejon, Korea (2023)
- 146. Life science innovation. Center for Basic Metabolic Research, University of Copenhagen, Denmark (2023)

List of Conferences co-organized

Chair/Co-Chair of Organizing Committee (not complete)

- 1. Danish Biotechnology Conference I, Munkebjerg, Denmark (1995) (130 participants) Chair
- 2. Danish Biotechnology Conference II, Munkebjerg, Denmark (1996) (130 participants) Chair
- 3. Danish Biotechnology Conference III, Munkebjerg, Denmark (1997) (180 participants) Chair
- 4. Danish Biotechnology Conference IV, Munkebjerg, Denmark (1998) (130 participants) Chair
- 5. Danish Biotechnology Conference V, Munkebjerg, Denmark (1999) (120 participants) Chair
- European Symposium on Biochemical Engineering Science 3, Copenhagen, Denmark (2000) (300 participants) Chair
- 7. Metabolic Engineering IV, Il Ciocco, Italy (2002) (250 participants) Chair
- 8. 7th European Conference on Fungal Genetics, Copenhagen, Denmark (800 participants) Co-Chair
- 9. European Congress on Biotechnology 12, Copenhagen, Denmark (2005) Chair of Scientific Committee
- 10. Swedish Bioinformatics Workshop, Gothenburg, Sweden (2010) (100 participants) Chair
- 11. Gothenburg Life Science Conference XI, Gothenburg, Sweden (2010) (150 participants) Chair
- 12. Key Sympoisum on Translational and Systems Medicine, Stockholm, Sweden (2011) (150 participants)

 Chair
- 13. Copenhagen Bioscience Conference on Biosustainability, Hilleröd, Denmark (2013) (150 participants). **Co-chair**
- 14. Copenhagen Bioscience Conference on Biosustainability, Hilleröd, Denmark (2015) (150 participants). Chair
- Copenhagen Bioscience Conference on Biosustainability, Hilleröd, Denmark (2019) (150 participants)
 Chair

Member of Organizing Committee (not complete)

- 1. Danish Biotechnology Conference VI, Munkebjerg, Denmark (2000)
- 2. Danish Biotechnology Conference VII, Munkebjerg, Denmark (2001)
- 3. Symposium on Physiology of Yeast and Filamentous Fungi, Hindsgayl, Denmark (2001)
- 4. Danish Biotechnology Conference VIII, Munkebjerg, Denmark (2002)
- 2nd International Conference on Analysis of Microbial Cells at the Single Cell Level, Munkebjerg, Denmark (2002)
- 6. Danish Biotechnology Conference IX, Munkebjerg, Denmark (2003)
- 7. Danish Biotechnology Conference X, Munkebjerg, Denmark (2004)
- 8. The 9th International Conference on Systems Biology, Gothenburg, Sweden (2008)
- 9. 35th FEBS Congress, Gothenburg, Sweden (2010)
- 10. Industrial Systems Biology 2010, Gothenburg, Sweden (2010)
- 11. Key Symposium on Systems Medicine, Salsjöbaden, Sweden (2011)
- 12. Copenhagen Bioscience Conference, Hillerød, Denmark (2013)
- 13. Eukaryotic Synthetic Biology, Heidelberg, Germany (2015)
- 14. Copenhagen Bioscience Conference, Hillerød, Denmark (2015)
- 15. Metabolic Engineering Summit, Beijing, China (2017)

Member of Scientific/Advisory Committee (not complete)

- 1. Modeling for Improved Bioreactor Performance II, Otocec, Slovenia (1994)
- 2. Yeast as a Cell Factory, Vlaardingen, The Netherlands (1998)
- 3. ISSY22: Yeast fermentations and other yeast bioprocesses, Pilansberg, South Africa (2002)
- 4. European Symposium on Biochemical Engineering Science 4, Delft, The Netherlands (2002)
- 5. Metabolic Engineering V, Taos, USA (2004)
- 6. European Symposium on Biochemical Engineering Science 5, Stuttgart, Germany (2004)
- 7. Biochemical Engineering XIV, Harrison Hot Springs, Canada (2005)
- 8. 8th European Conference on Fungal Genetics, Vienna, Austria (2006)
- 9. Metabolic Engineering VI, Noordwijkerhout, The Netherlands (2006)

- 10. ISSY25: Systems Biology of Yeasts from Models to Applications, Espoo, Finland (2006)
- 11. FOSBE 2007, Stuttgart, Germany (2007)
- 12. 9th European Conference on Fungal Genetics, Edinburgh, UK (2008)
- 13. Metabolic Engineering VII, Puerto Vallarta, Mexico (2008)
- 14. 13th International Biotechnology Symosium and Exhibition, Dalian, China (2008)
- 15. 12th International Congress on Yeasts, Kiev, Ukraine (2008)
- 16. FOSBE 2009, Englewood, USA (2009)
- 17. ISSY27: Yeast for health and biotechnologies, Paris, France (2009)
- 18. PYFF4, Rotterdam, The Netherlands (2010)
- 19. Metabolic Engineering VIII, Jeju Island, South Korea (2010)
- 20. 10th European Conference on Fungal Genetics, Leiden, The Netherlands (2010)
- 21. 14th International Biotechnology Symposium and Exhibition, Rimini, Italy (2010)
- 22. Microbial Stress: From Molecules to Systems, Beligrate, Italy (2012)
- 23. Metabolic Engineering IX, Biarritz, France (2012)
- 24. Metabolic Engineering X, Vancouver, Canada (2014)
- 25. Metabolic Engineering Summit, Beijing, China (2015)
- 26. Metabolic Engineering 11, Ajawi Island, Japan (2016)
- 27. ISSY33, Cork, Ireland (2017)
- 28. Metabolic Engineering 12, Munich, Germany (2018)

Research Supervision

Current Post Doctoral Researchers

- Peishun Li, Chalmers (2022-)
- Hao Wang (2017-)

Former Post Doctoral Researchers

- Lars Højlund Christensen, DTU (1992-1994)
- Gunnar Liden, DTU (1993-1994) 2.
- 3. Lisbeth OIsson, DTU (1994-1996)
- Morten Carlsen, DTU (1995-1997)
- Anne Santerre Henriksen, DTU (1995-1999) 5.
- Aradhana Srivastava, DTU (1996-1998) 6.
- Hans Peter Smits, DTU (1996-1999; 2000-2001) 7.
- Alexei Aleksenko, DTU (1997-2001)
- Philippe Duboc, DTU (1997-1999) 9
- 10. Fernando Bautista, DTU (1998-1999)
- 11. Uffe Mortensen, DTU (1999-2002)
- 12. Ana Borges, DTU (1999-2001)
- Biarke Christensen, DTU (1999-2001)
- Mhairi Workman, DTU (1999-2004)
- 15. Birgitte Regenberg, DTU (1999-2005)
- 16. Anna Eliasson Lantz, DTU (2000-2002)
- 17. Christian Müller, DTU (2001-2002)
- 18. Vsevolod Serebrianyi, DTU (2001-2002)
- 19. Mats Åkesson, DTU (2001-2003)
- 20. Kasper Møller, DTU (2001-2003)
- 21. Per Bruheim, DTU (2002)
- 22. Vasimon Ruanglek, DTU (2002)
- 23. Tamay Seker, DTU (2002-2003)
- 24. Sandrine Mas, DTU (2002-2005)
- 25. Gerald Hofmann, DTU (2004-2006)
- 26. Roberta Mustachi, DTU (2004-2007)
- 27. Jerome Maury, DTU (2004-2008)
- 28. Isabel Rocha, DTU (2004)
- 29. Jette Thykær, DTU (2005-2007)
- 30. Dongmei Bai, DTU (2005-2007) 31. Michael Jewett, DTU (2005-2008)
- 32. Goutham Vemuri, DTU (2006-2007)
- Manuel Quiros Asensio, DTU (2006-2008)
- 34. Prashant Bapat, DTU (2006-2008)
- 35. Sven Even Borgos, DTU (2007)
- 36. Dina Petranovic, DTU (2007-2008)
- 37. Subir Kumar Nandy, Chalmers (2008-2010)
- 38. Keith Tyo, Chalmers (2008-2010)
- 39. Marija Cvijovic, Chalmers (2008-2010)
- 40. Andrea Neiss, Chalmers (2009-2010)
- 41. Wanwipa Vongsangnak, Chalmers (2010)
- 42. Liming Liu, Chalmers (2009-2010)
- 43. Sergio Bordel Velasco (2008-2010)
- 44. Intawat Nookaew (2008-2012)
- 45. Jin Hou, Chalmers (2010-2012)
- Fredrik Öberg, DTU (2011-2012)
- 47. Shuobo Shi, Chalmers (2009-2012) 48. Luis Caspeta, Chalmers (2009-2013)
- 49. Il-Kwon Kim, Chalmers (2010-2013)
- 50. Rahul Kumar, Chalmers (2010-2013)
- 51. Antonio Roldao, Chalmers (2010-2013)
- Adil Mardinoglu, Chalmers (2010-2015) Marina Sanchez Martinez, Chalmers (2011-2013)
- 54. Manuel Garcia, Chalmers (2012-2014)

- 55. Clara Navarrete, Chalmers (2013-2014)
- 56. Zheng Wang, Chalmers (2013-2014)
- 57. Subazini Thankaswarmy, Chalmers (2013-2014)
- 58. Martin Engqvist, Chalmers (2014-2015)
- 59. Anastasia Krivoruchko, Post doc, Chalmers (2010-2016)
- 60. Jose L. Martinez, Chalmers (2011-2015)
- 61. Agata Smialowska, Chalmers (2014-2015)
- 62. Sunjae Lee, Chalmers (2014-2015)
- 63. Petri-Jaan Lahtvee, Chalmers (2013-2015)
- 64. Eugene Fletcher, Chalmers (2014-2015)
- 65. Hülya Karaca Gencer, Chalmers (2014-2015)
- 66. Silveira Wendel, Chalmers (2015-2016)
- 67. Min-Kyoung Kang, Chalmers (2014-2016)
- 68. Guodong Liu, Chalmers (2013-2016)
- 69. Mark Bisschops, Chalmers (2014-2016)
- 70. Ömür Kayikci, Chalmers (2014-2016)
- 71. Tatiana Moreira, Chalmers (2015-2016)
- 72. Yongjin Zhou, Chalmers (2012-2016)
- 73. Yongjun Wei, Chalmers (2014-2016)
- 74. Amir Feizi, Chalmers (2016)
- 75. Mingtao Huang, Chalmers (2012-2017)
- Zongjie Dai, Chalmers (2013-2017)
- 77. Partho Sarathi Sen, Chalmers (2014-2017)
- 78. Sakda Khoomrung, Chalmers (2011-2017)
- 79. Michael Gossing, Chalmers (2014-2017)
- 80. Francesco Gatto, Chalmers (2015-2017)
- 81. Yi Liu, Chalmers (2015-2020)
- 82. JinHo Kim, Chalmers (2016-2018)
- 83. Manish Kumar, Chalmers (2015-2018)
- 84. Petter Holland, Chalmers (2016-2018)
- 85. Sylvain Prigent, Chalmers (2015-2018)
- 86. Ibrahim El-Semman, DTU (2015-2018)
- 87. Francesca Di Bartolomeo, Chalmers (2017-2019)
- 88. Lucy Fang-I Chao, Chalmers (2017-2019)
- 89. Zhiwei Zhu, Chalmers (2014-2019)
- 90. Tao Yu, Chalmers (2014-2019)
- 91. Tyler Doughty, Chalmers (2017-2019)
- 92. Ling-Qun Ye, Chalmers (2017-2019)
- 93. Yeping Zhang, BUCT (2017-2019)
- 94. Kate Campbell, Chalmers (2016-2020) 95. Jonathan Robinson, Chalmers (2016-2020)
- 96. Rui Pereira, Chalmers (2016-2020)
- 97. Xiaowei Li, Chalmers (2016-2020)
- 98. Quanli Liu, Chalmers (2016-2020)
- 99. Daniel Cook, Chalmers (2017-2020)
- 100. Ievgeniia Tiukova, Chalmers (2018-2020)
- 101. Hongzhong Lu, Chalmers (2017-2020)
- 102. Pinar Kocabas, Chalmers (2017-2020)
- 103. Jun Geng, Chalmers (2015-2020)
- 104. Jiufu Qin, Chalmers & DTU (2015-2020)
- 105. Boyang Ji, Chalmers (2013-2020)
- 106. Yiming Zhang, BUCT (2016-2021)
- 107. Rosemary Yu, Chalmers (2017-2021)
- 108. Sinisa Bratulic, Chalmers (2018-2022)
- 109. Yu Chen, Chalmers (2018-2022)
- 110. Rasool Saghaleyni, Chalmers (2021-2022)
- 111. Feiran Li, Chalmers (2021-2022)
- 112. Yu Chen, Chalmers (2018-2023)
- 113. Rasool Saghaleyni, Chalmers (2021-2023)
- 114. Feiran Li, Chalmers (2021-2023)

Current PhD Students (main supervisor)

- Angelo Limeta, Chalmers (2019-)
- Mihail Petre Anton, Chalmers (2021-)

Graduated PhD Students (main supervisor)

- 1. Henrik Jørgensen, DTU (1991-1993)
- 2. Claus Lindvad Johansen, DTU (1991-1993)
- 3. Morten Carlsen, DTU (1992-1995)
- 4. Rong Wei Min, DTU (1992-1995)
- 5. Anders Spohr, DTU (1993-1996)
- 6. Preben Krabben, DTU (1993-1997)
- 7. Claus Maxel Henriksen, DTU (1993-1996)
- 8. Christoffer Klein, DTU (1995-1998)
- 9. Karsten Schmidt, DTU (1995-1998)
- 10. Torben Nissen, DTU (1995-1998)
- 11. Henrik Pedersen, DTU (1996-1999)
- 12. Bjarke Christensen, DTU (1996-1999)
- 13. Teit Agger, DTU (1996-1999)
- 14. Hanne Theilgaard, DTU (1996-1999)
- 15. Simon Østergaard, DTU (1997-2000)
- 16. Wai Prathumpai, DTU (1998-2001)
- 17. Tina Lübbehüsen, DTU (1998-2001)
- 18. Jens Dynesen, DTU (1998-2001)
- 19. Torben Christiansen (1998-2001)
- 20. Christian Müller, DTU (1998-2001)
- 21. Jarno Robin, DTU (1998-2002)
- 22. Jochen Förster, DTU (1999-2002)
- 23. Michael Lynge Nielsen, DTU (1999-2003)
- 24. Jette Thykær, DTU (2000-2005)
- 25. Nina Gunnarsson, DTU (2000-2003)
- 26. Margarida Moreira dos Santos, DTU (1999-2003)
- 27. Christoffer Bro, DTU (2000-2003)
- 28. Gerald Hofmann, DTU (2001-2004)
- 29. Helga David, DTU (2001-2005)
- 30. Thomas Grotkjær, DTU (2001-2004)
- 31. Nadine Eckert-Boulet, DTU (2001-2004)
- 32. Vijay Raghevendrabn, DTU (2001-2005)
- 33. Steen Lund Westergaard, DTU (2002-2005)
- 34. Wian de Jongh, DTU (2002-2005)
- 35. Silas Granato Villas-Boas, DTU (2002-2005)
- 36. Kiran Patil, DTU (2003-2006)
- 37. Songsak Wattanachaisaereekul, DTU (2003-2007)
- 38. Audrey Diano, DTU (2003-2007)
- 39. Torsten Bak Reguira, DTU (2003-2007)
- 40. Irina Borodina, DTU (2004-2008)
- 41. Susan Meijer, DTU (2004-2007)
- 42. Ana Paula Oliveira, DTU (2004-2008)
- 43. Mohammad Asahollahi, DTU (2004-2008)
- 44. Mikael Rørdam Andersen, DTU (2004-2008)
- 45. Jesper Højer Pedersen, DTU (2004-2008)
- 46. Kjeld Kjeldsen, DTU (2005-2008)
- 47. Wanwipa Vongsangnak, Chalmers (2005-2009)
- 48. Jose Manuel Otero, Chalmers (2005-2009)
- 49. Margarita Salazar Pena, Chalmers (2006-2010)
- 50. Jie Zhang, Chalmers (2007-2011)
- 51. Roberto Olivares, Chalmers (2006-2011)
- 52. Pramote Chumnapuen, Chalmers (2008-2012)
- 53. Marta Papini, Chalmers (2008-2012)
- 54. Gionata Scalcinati, Chalmers (2008-2012)
- 55. Siavash Partow, Chalmers (2008-2012)
- 56. Kuk-Ki Hong, Chalmers (2008-2012)
- 57. Liu Zihe, Chalmers (2008-2012)
- 58. Kanokarn Kocharin, Chalmers (2009-2013)
- 59. Rasmus Ågren, Chalmers (2008-2013)
- 60. Tobias Österlund, Chalmers (2009-2014)
- 61. Fredrik Karlsson, Chalmers (2009-2014)
- 62. Christoph Knuf, Chalmers (2009-2014)
- 63. Natapol Pornputtapong, Chalmers (2010-2014)
- 64. Lifang Liu, Chalmers (2010-2014)
- 65. Bouke de Jong, Chalmers (2010-2015)

- 66. Yiming Zhang, Chalmers (2010-2015)
- 67. Francesco Gatto, Chalmers (2012-2015)
- 68. Saeed Shoaie, Chalmers (2011-2015)
- 69. Amir Feizi, Chalmers (2011-2016)
- 70. Leif Väremo, Chalmers (2011-2016)
- 71. Nicolaas Buijs, Chalmers (2011-2016)
- 72. Mingji Li, DTU (2013-2016)
- 73. Edith Angelica Rodriguez Prado, DTU (2013-2016)
- 74. Stefan Tippmann, Chalmers (2012-2016)
- 75. Pouyan Ghaffari Nouran, Chalmers (2013-2017)
- 76. Jens Christian F. Nielsen, Chalmers (2014-2018)
- 77. Paulo Teixeira, Chalmers (2013-2018)
- 78. Jichen Bao, Chalmers (2013-2018)
- 79. Alexandra Bergman, Chalmers (2013-2019)
- 80. Avlant Nilsson, Chalmers (2014-2019)
- 81. Benjamín José Sánchez Barja, Chalmers (2014-2019)
- 82. David Bergenholm, Chalmers (2014-2020)
- 83. Parizad Babaei, Chalmers (2015-2019)
- 84. Promi Das, Chalmers (2015-2019)
- 85. Raphael Ferreira, Chalmers (2015-2019)
- 86. Gang Li, Chalmers (2016-2020)
- 87. Simonas Marcišauskas, Chalmers (2015-2020)
- 88. Christoph Börlin, Chalmers (2016-2020)
- 89. Rasool Saghaleyni, Chalmers (2016-2021)
- 90. Carl Melina, Chalmers (2016-2021)
- 91. Ning Qin, BUCT (2016-2021)
- 92. Qi Qi, Chalmers (2017-2021)
- 93. Feiran Li, Chalmers (2017-2021)
- 94. Yijin Zhao, BUCT (2017-2021)
- 95. Peishun Li, Chalmers (2018-2022)
- 96. Johan Gustafsson, Chalmers (2017-2022)
- 97. Hao Lou, Chalmers (2018-2022)
- 98. Dimitra Lappa, Chalmers (2016-2023)
- 99. Lingyun Li, BUCT (2018-2023)
- 100. Ivan Domenzain Del Castillo Cerecer (2017-2023)
- 101. Juan Octavo Valle (2012-2023)

Graduated PhD Students (Examiner)

- 1. Joachim Almquist, FCC, Sweden (2010-2017)
- Yasaman Dabirian, Chalmers (2016-2021)
- 3. Oliver Konzock, Chalmers (2018-2022)
- 4. Christos Skrekas, Chalmers (2018-2022)
- 5. Maximillian Otto, Chalmers (2018-2022)
- 6. Dany Liu, Chalmers (2019-2023)
- 7. Marta Tous Mohedano, Chalmers (2019-2023)

Graduated PhD Students (Co-supervisor)

- 1. Lars Højlund Christensen, DTU (1989-1992)
- 2. Stig Benthin, DTU (1989-1992)
- 3. Annemarie Gade Pedersen, DTU (1990-1992)
- 4. Simone de Jong Frandsen, DTU (1991-1993)
- 5. Ulrik Schultze, DTU (1992-1995)
- 6. Teresa Zangorilami, DTU (1994-1998)
- 7. Kasper Møller, DTU (1998-2001)
- 8. Mikkel Nordkvist, DTU (2001-2005)
- 9. Renata Usaite, DTU (2004-2008)
- 10. Lasse Pedersen, DTU (2007-2010)
- 11. Kanchana Rueksomtawin Kildegaard, DTU (2004-2011)
- 12. Xiao Chen, DTU (2007-2011)
- 13. Kwanjeera Wanichthanarak, Chalmers (2010-2014)
- 14. Kaisa Thorell, Gothenburg University, Sweden (2010-2014)
- 15. Leonie Wenning, Chalmers (2014-2018)
- 16. Yating Hu, Chalmers (2015-2019)

- 17. Elias Björnson, Gothenburg University (2014-2021)
- 18. John Hellgren, Chalmers (2017-2022)

Former Visiting PhD Students (co-supervisor)

- 1. Robert Lejeune, Free University of Brussels (1994-1995)
- 2. Pedro N. Pissarra, Kings College London (1994-1995)
- Einar Jonsbu, NTNU (1999-2000)
- Andreas Karoly Gomberg, Sao Paulo University (1998-2000)
- 5. Joel Forest Moxley, MIT (2003-2005)
- 6. Tunahan Cakir, Bogazici University (2004-2005)
- 7. Kazim Yalcin Arga, Bogacizi University (2004-2005)
- 8. Intawat Nookaew, KMUTT (2005-2006)
- 9. Goutham Vemuri, University of Georgia (2005-2006)
- 10. Donatella Cimini, University of Napoli (2006)
- 11. Valeria Mapelli, University Milan-Bicocca (2006-2007)
- 12. Paula Jouhten, Technical University of Helsinki (2008-2009)
- 13. Raphael Aggio, University of Auckland (2009)
- 14. William Alfonso Rodriguez Limas, National University of Mexico (2009-2010)
- 15. Kantida Kusonmano, UMIT (2011)
- 16. Akarin Boonsombuti, Mahasarakham University (2011-2012)
- 17. Jorge Alberto Vasquez Castillo, Antioquia University (2011-2012)
- 18. Josh Michener, Caltech/Stanford (2011)
- 19. Ibrahim E. El-Semman, Assiut University (2012-2014)
- 20. Jiufu Qin, Jiangnan University, China (2011-2014)
- 21. Rui Pereira, University of Minho, Portugal (2012-2014)
- 22. Cheng Zhang, East China University of Science and Technology, China (2013-2015)
- 23. John Casey, University of Hawaii, USA (2014)
- 24. Jacqueline Rand, University of Wisconsin, USA (2014)
- 25. Zahra Azim Zadeh Irani, Tarbiat Modares University, Iran (2014-2015)
- 26. Yu Chen, East China University of Science and Technology, China (2015-2017)
- 27. Chinh Bkrong Nguyen, Oslo University, Norway (2017-2018)
- 28. Zhengming Zhu, Jiangnan University, China (2017-2018)
- 29. Chunjun Zhan, Jiangnan University, China (2017-2018)
- 30. Zeinab Hefny, Katholiek University of Leuven, Belgium (2017-2019)

Other Former Affiliated Researchers and Staff

- 1. Tina Johansen, Research Engineer, DTU (1990-2008)
- 2. Lene Christiansen, Research Engineer, DTU (1992-2008)
- 3. Martin Hjortso, Visiting Professor, DTU (1992-1993, 2002-2003)
- 4. Susanne Sloth Larsen, Head of Administration, DTU (1995-1999)
- 5. Jette Mortensen, Laboratory Technician, DTU (1996-2008)
- 6. Birgitte Karsbøl, Administrative Assistant, DTU (1996-2008)
- 7. Kirsten Nielsen, Administrative Assistant, DTU (1998-2005)
- 8. Trine Bro, Head of Administration, CMB, DTU (1999-2008)
- 9. Lars K. Nielsen, Visiting Professor, DTU (2004)
- 10. Stefan Rokem, Visiting Professor, DTU (2004-2005)
- 11. Eduardo Agosin, Visiting Professor, DTU (2005)
- 12. Verena Siewers, Senior Researcher & Docent, DTU & Chalmers (2006-2020)
- 13. Yun Chen, Resarcher & Docent, Chalmers (2008-2020)
- 14. Marie Nordqvist, Research Engineering and Lab Manager (2008-2017)
- 15. Marie-Louise Wennerhag, Financial Officer, Chalmers (2008-2018)
- 16. Martina Butorac (2008-2020)
- 17. Malin Nordvall, Research Engineer, Chalmers (2009-2014)
- 18. Stefan Rokem, Visiting Professor, Chalmers (2010)
- 19. Pegah Khorramzadeh, Research Engineering, Chalmers (2010-2011)
- 20. Sergio Bordel Velasco, Assistant Professor (2010-2014)
- 21. Suwanee Jansa-Ard, Research Engineer, Chalmers (2011-2014)
- 22. Martin Markström, Co-Director AoA, Chalmers (2011-2014)
- 23. Ximena Rozo Sevilla, Research Engineer, Chalmers (2011-2015)
- 24. Danilo Porro, Visiting Professor, Chalmers (2012)
- 25. Emma Ribbenhed, Research Engineer, Chalmers (2013-2015)
- 26. Antonio Marras, Research Assistant Chalmers (2013-2015)
- 27. Eduard Kerkhoven, Researcher, Chalmers (2013-2020)

- 28. Shaghayegh Hosseini, Data Manager, Chalmers (2013-2019)
- 29. Gheorge Manuel Borja Zamfir, PhD student, DTU (2013-2019)
- 30. Brian Pfleger, Visiting Professor, Chalmers (2014)
- 31. Julia Karlsson, Research Engineer, Chalmers (2014-2016)
- 32. Matthias Nilsson, Research Engineer, Chalmers (2015-2016)
- 33. Daniel Hermansson, Research Engineer, Chalmers (2016)
- 34. Xiaojun Ji, Visiting Researcher (2016-2017)
- 35. Jianye Xia, Visiting Professor, Chalmers (2016-2017)
- 36. Liming Quyang, Visiting Professor, Chalmers (2016-2017)
- 37. Takayoshi Fujii, Visiting Researcher, Chalmers (2016-2018)
- 38. Fredrik Schubert, Chalmers (2016-2019)
- 39. Joakim Norbeck, Researcher & Docent, Chalmers (2015-2020)
- 40. Johan Björkeroth, PhD student, Chalmers (2016-2020)
- 41. Erica Dahlin, Chalmers (2008-2020)
- 42. Angelica Ardehed (Lab Manager), Chalmers (2016-2020)
- 43. Emilie Lindquist, Chalmers (2017-2020)
- 44. Pierre-Etienne Cholley, Chalmers (2017-2020)
- 45. Virinchi Billa, Chalmers (2017-2019)
- 46. Mihail Anton, Chalmers (2017-2020)
- 47. Felipe Lopez-Isunza, Visiting Professor, Chalmers (2019)