

## PlantLink supported peer-reviewed articles

Publications supported through seed money grants are marked with an asterisk. The other publications have had bioinformatic support sponsored by PlantLink.

### 2012

Witzgall P, Proffit M, Rozpedowska E, Becher PG, Andreadis S, Coracini M, et al. "This is not an apple"-yeast mutualism in codling moth. *J Chem Ecol.* 2012;38(8):949-57.\*

### 2013

Kaki SS, Adlercreutz P. Quantitative analysis of enzymatic fractionation of multiple substrate mixtures. *Biotechnol Bioeng.* 2013;110(1):78-86.\*

### 2014

Amar D, Frades I, Danek A, Goldberg T, Sharma SK, Hedley PE, et al. Evaluation and integration of functional annotation pipelines for newly sequenced organisms: The potato genome as a test case. *BMC Plant Biology.* 2014;14(1).

Bengtsson T, Weighill D, Proux-Wéra E, Levander F, Resjö S, Burra DD, et al. Proteomics and transcriptomics of the BABA-induced resistance response in potato using a novel functional annotation approach. *BMC Genomics.* 2014;15(1).

Burra DD, Berkowitz O, Hedley PE, Morris J, Resjö S, Levander F, et al. Phosphite-induced changes of the transcriptome and secretome in *Solanum tuberosum* leading to resistance against *Phytophthora infestans*. *BMC Plant Biology.* 2014;14(1).

Crespo-Herrera LA, Akhunov E, Garkava-Gustavsson L, Jordan KW, Smith CM, Singh RP, et al. Mapping resistance to the bird cherry-oat aphid and the greenbug in wheat using sequence-based genotyping. *Theoretical and Applied Genetics.* 2014;127(9):1963-73.

Ding BJ, Hofvander P, Wang HL, Durrett TP, Stymne S, Löfstedt C. A plant factory for moth pheromone production. *Nat Commun.* 2014;5.\*

Leiva-Eriksson N, Pin PA, Kraft T, Dohm JC, Minoche AE, Himmelbauer H, et al. Differential expression patterns of non-symbiotic hemoglobins in sugar beet (*Beta vulgaris* ssp. *vulgaris*). *Plant and Cell Physiology.* 2014;55(4):834-44.\*

Turesson H, Andersson M, Marttila S, Thulin I, Hofvander P. Starch biosynthetic genes and enzymes are expressed and active in the absence of starch accumulation in sugar beet tap-root. *BMC Plant Biology.* 2014;14(1).

### 2015

Amar D, Frades I, Diels T, Zaltzman D, Ghatan N, Hedley PE, et al. The MORPH-R web server and software tool for predicting missing genes in biological pathways. *Physiol Plant.* 2015;155(1):12-20.

Andreadis SS, Witzgall P, Becher PG. Survey of arthropod assemblages responding to live yeasts in an organic apple orchard. *Front ecol evol.* 2015;3(OCT).\*

Burra DD, Mühlenbock P, Andreasson E. Salicylic and jasmonic acid pathways are necessary for defence against *Dickeya solani* as revealed by a novel method for Blackleg disease screening of in vitro grown potato. *Plant Biol.* 2015;17(5):1030-8.

Frades I, Abreha KB, Proux-Wéra E, Lankinen Å, Andreasson E, Alexandersson E. A novel workflow correlating RNA-seq data to *Phytophthora infestans* resistance levels in wild *Solanum* species and potato clones. *Frontiers in Plant Science.* 2015;6(September).

Lager I, Glab B, Eriksson L, Chen G, Banas A, Stymne S. Novel reactions in acyl editing of phosphatidylcholine by lysophosphatidylcholine transacylase (LPCT) and acyl-CoA:glycerophosphocholine acyltransferase (GPCAT) activities in microsomal preparations of plant tissues. *Planta.* 2015;241(2):347-58.\*

Manoharan L, Kushwaha SK, Hedlund K, Åhrén D. Captured metagenomics: Large-scale targeting of genes based on 'sequence capture' reveals functional diversity in soils. *DNA Res.* 2015;22(6):451-60.

Nicolia A, Proux-Wéra E, Åhman I, Onkokesung N, Andersson M, Andreasson E, et al. Targeted gene mutation in tetraploid potato through transient TALEN expression in protoplasts. *J Biotechnol.* 2015;204:17-24.

## 2016

Ding BJ, Lager I, Bansal S, Durrett TP, Stymne S, Löfstedt C. The Yeast ATF1 Acetyltransferase Efficiently Acetylates Insect Pheromone Alcohols: Implications for the Biological Production of Moth Pheromones. *Lipids.* 2016;51(4):469-75.\*

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Kuktaite R, Newson WR, Rasheed F, Plivelic TS, Hedenqvist MS, Gällstedt M, et al. Monitoring Nanostructure Dynamics and Polymerization in Glycerol Plasticized Wheat Gliadin and Glutenin Films: Relation to Mechanical Properties. *ACS Sustainable Chem Eng.* 2016;4(6):2998-3007.\*

Kushwaha SK, Chauhan P, Hedlund K, Åhren D. NBSPred: A support vector machine-based high-throughput pipeline for plant resistance protein NBSLRR prediction. *Bioinformatics.* 2016;32(8):1223-5.

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Roy A, Walker WB, Vogel H, Chattington S, Larsson MC, Anderson P, et al. Diet dependent metabolic responses in three generalist insect herbivores *Spodoptera* spp. *Insect Biochem Mol Biol*. 2016;71:91-105.

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## 2017

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Zhao M, Jones CM, Meijer J, Lundquist PO, Fransson P, Carlsson G, et al. Intercropping affects genetic potential for inorganic nitrogen cycling by root-associated microorganisms in *Medicago sativa* and *Dactylis glomerata*. *Appl Soil Ecol*. 2017;119:260-6.\*

## 2018

Abreha KB, Lankinen Å, Masini L, Hydbom S, Andreasson E. Late blight resistance screening of major wild Swedish solanum species: *S. dulcamara*, *S. nigrum*, and *S. physalifolium*. *Phytopathology*. 2018;108(7):847-57.\*

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## 2019

Anaokar S, Kodali R, Jonik B, Renne MF, Brouwers JFHM, Lager I, et al. The glycerophosphocholine acyltransferase Gpc1 is part of a phosphatidylcholine (PC)-remodeling pathway that alters PC species in yeast. *Journal of Biological Chemistry.* 2019;294(4):1189-201.\*

Eriksson NL, Reeder BJ, Wilson MT, Bülow L. Sugar beet hemoglobins: Reactions with nitric oxide and nitrite reveal differential roles for nitrogen metabolism. *BIOCHEM J.* 2019;476(14):2111-25.\*

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Odilbekov F, Armoniené R, Koc A, Svensson J, Chawade A. GWAS-Assisted Genomic Prediction to Predict Resistance to *Septoria Tritici* Blotch in Nordic Winter Wheat at Seedling Stage. *Front Genet.* 2019;10.

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## 2020

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## 2021

Juma I, Geleta M, Hovmalm HP, Nyomora A, Saripella GV, Carlsson AS, et al. Comparison of Morphological and Genetic Characteristics of Avocados Grown in Tanzania. *Genes*. 2021;12(1).