

MITCHELL TUINSTRA

**Professor of Plant Breeding and Genetics
Wickersham Chair of Excellence in Agricultural Research
Scientific Director – Institute for Plant Sciences**

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EDUCATION

1996 Ph.D., Purdue University, West Lafayette, IN - Plant Breeding/Genetics
1993 M.S., Purdue University, West Lafayette, IN - Molecular Genetics
1991 B.S., Calvin College, Grand Rapids, MI - Biology

PROFESSIONAL POSITIONS

2013-present Scientific Director, Institute for Plant Sciences, College of Agriculture,
Purdue University
2007-present Wickersham Chair of Excellence in Agricultural Research, Department of
Agronomy, Purdue University
2007-present Professor of Plant Breeding and Genetics, Department of Agronomy,
Purdue University
2006-2007 Professor, Department of Agronomy, Kansas State University
2001-2005 Associate Professor, Department of Agronomy, Kansas State University
1997-2001 Assistant Professor, Department of Agronomy, Kansas State University
1997 Post-Doctoral Fellow, Department of Agronomy, Purdue University
1994 Teaching Assistant, Department of Horticulture, Purdue University
1994-1996 Research Assistant, Department of Horticulture, Purdue University
1993 Teaching Assistant, Department of Horticulture, Purdue University
1993 Teaching Assistant, Department of Agronomy, Purdue University
1991-1993 Research Assistant, Department of Horticulture, Purdue University

RESEARCH/TEACHING ACTIVITIES

60% Research Plant Breeding and Genetics
10% Teaching AGRY285: World Crop Adaptation and Distribution
30% Administration Scientific Director, Institute for Plant Sciences

HONORS AND AWARDS

Lowell S. Hardin Award for Excellence in International Agriculture, College of Agriculture –
2023
Crops & Soils Merit Award In Recognition of Outstanding Contribution to Agriculture and
the Seed Industry, Indiana Crop Improvement Association – 2022

David C. Pfendler Outstanding Undergraduate Counselor Award, College of Agriculture – 2022
Agronomy Outstanding Counselor – 2017, 2018, 2021, 2022
Wickersham Chair of Excellence in Agricultural Research, Purdue University – 2007, 2015, 2021
Seed for Success, Excellence in Research, Purdue University – 2009, 2013, 2014, 2016, 2018, 2020
Agronomy Outstanding Teacher – 2020
Fellow, American Society of Agronomy – 2017
Fellow, Crop Science Society of America – 2017
Spotlight Educator – Agricultural Council Student Choice Award, College of Agriculture, Purdue University – 2016
Gamma Sigma Delta – Early Career Award – 2001

PROFESSIONAL SOCIETIES

American Society of Agronomy
Crop Science Society of America
Sorghum Improvement Conference of North America
National Association of Plant Breeders
North American Plant Phenotyping Network

PATENTS

Tuinstra, M.R. and Al-Khatib, K., Kansas State University, 2019. Acetolactate synthase herbicide resistant sorghum. U.S. Patent 10,519,461. Issue Date: December 31, 2019.
Tuinstra MR, Al-Khatib K. Kansas State University Research Foundation. Acetyl-CoA Carboxylase Herbicide Resistant Sorghum. U.S. Patent No. 9,617,530. Issue Date: April 11, 2017.
Tuinstra MR, Krothapalli K, Dilkes B, Buescher E. Genetic Mutations that Disrupt Dhurrin Production In Sorghum. U.S. Patent No. 9,512,437. Issue Date: December 6, 2016.

PUBLICATIONS

REFEREED JOURNAL ARTICLES

Wang, T., Crawford, M.M. and Tuinstra, M.R., 2023. A novel transfer learning framework for sorghum biomass prediction using UAV-based remote sensing data and genetic markers. *Frontiers in Plant Science*, 14. <https://doi.org/10.3389/fpls.2023.1138479>
Gruss, S.M., Souza, A., Yang, Y., Dahlberg, J. and Tuinstra, M.R., 2023. Expression of Stay-green Drought Tolerance in Dhurrin-free Sorghum. *Crop Science*, 2023, 1–14. <https://doi.org/10.1002/csc2.20947>
Diatta-Holgate, E., Anderson, J.S., Hatch, R., Tuinstra, M.R. and Weil, C., 2023. Rapid Determination of Protein Digestibility in Sorghum Before and After Cooking. *MethodsX*, p.102162. <https://doi.org/10.1016/j.mex.2023.102162>
Gruss, S.M., Johnson, K.D., Ghaste, M., Widhalm, J.R., Johnson, S.K., Holman, J.D., Obour, A., Aiken, R.M. and Tuinstra, M.R., 2023. Dhurrin stability and hydrogen cyanide release in

- dried sorghum samples. *Field Crops Research*, 291, p.108764.
<https://doi.org/10.1016/j.fcr.2022.108764>
- Diatta-Holgate, E., Huggis, E., Weil, C., Faye, J.M., Danquah, A., Diatta, C., Tongoona, P., Danquah, E.Y., Cisse, N. and Tuinstra, M.R., 2022. Natural variability for protein digestibility and grain quality traits in a West African Sorghum Association Panel. *Journal of Cereal Science*, p.103504. <https://doi.org/10.1016/j.jcs.2022.103504>
- Simons, J., Herbert, T., Kauffman, C., Batete, M., Simpson, A., Katsuki, Y., Le, D., Amundson, D., Buescher, E., Weil, C., Tuinstra, M.R., Addo-Quaye, C., 2022. Systematic prediction of EMS-induced mutations in a sorghum mutant population. *Plant Direct*. 6(5): e404.
<https://doi.org/10.1002/pld3.404>
- Ren, D., Engel, B. and Tuinstra, M.R., 2022. Crop improvement influences on water quantity and quality processes in an agricultural watershed. *Water Research*, p.118353.
<https://doi.org/10.1016/j.watres.2022.118353>
- Lin, M., Lynch, V., Ma, D., Maki, H., Jin, J., Tuinstra, M.R., 2022. Multi-species prediction of physiological traits with hyperspectral modeling. *Plants*, 11, 676.
<https://doi.org/10.3390/plants11050676>.
- Gruss, S.M., Ghaste, M., Widhalm, J.R., Tuinstra, M.R., 2022. Seedling growth and fall armyworm feeding preference influenced by dhurrin production in sorghum. *Theoretical and Applied Genetics*. <https://doi.org/10.1007/s00122-021-04017-4>
- Ojeda, J.J., Hammer, G., Yang, K.W., Tuinstra, M.R., DeVoil, P., McLean, G., Huber, I., Volenec, J.J., Brouder, S.M., Archontoulis, S. and Chapman, S.C., 2022. Quantifying the effects of varietal types × management on the spatial variability of sorghum biomass across US environments. *GCB Bioenergy*, 14(3), pp.411-433.
<https://doi.org/10.1111/gcbb.12919>
- Nazeri, B., Crawford, M. and Tuinstra, M.R., 2021. Estimating Leaf Area Index in Row Crops Using Wheel-Based and Airborne Discrete Return Lidar Data. *Frontiers in Plant Science*, p.2727. <https://doi.org/10.3389/fpls.2021.740322>
- Herrero, M., Meline, V., Iyer-Pascuzzi, A.S., Souza, A.M., Tuinstra, M.R. and Yang, Y., 2021. 4D Structural root architecture modeling from digital twins by X-Ray Computed Tomography. *Plant Methods* 17, 123. <https://doi.org/10.1186/s13007-021-00819-1>
- Ma, D., Rehman, T.U., Zhang, L., Maki, H., Tuinstra, M.R. and Jin, J., 2021. Modeling of Environmental Impacts on Aerial Hyperspectral Images for Corn Plant Phenotyping. *Remote Sensing*, 13, p.2520. <https://doi.org/10.3390/rs13132520>
- Tolley, S.A., Singh, A. and Tuinstra, M., 2021. Heterotic Patterns of Temperate and Tropical Maize by Ear Photometry. *Frontiers in Plant Science*, 12, p.1117.
<https://doi.org/10.3389/fpls.2021.616975>
- Perumal, R., Morris, G.P., Jagadish, S.V.K., Little, C.R., Tesso, T.T., Bean, S.R., Yu, J., Prasad, V., and Tuinstra, M.R., 2021. Registration of the Sorghum [*Sorghum bicolor* (L.) Moench] Nested Association Mapping (NAM) Population in RTx430 Background. *Journal of Plant Registration*. <https://doi.org/10.1002/plr2.20110>
- Ma, D., Rehman, T.U., Zhang, L., Maki, H., Tuinstra, M.R. and Jin, J., 2021. Modeling of diurnal changing patterns in airborne crop remote sensing images. *Remote Sensing*, 13(9), p.1719. <https://doi.org/10.3390/rs13091719>

- Herrero-Huerta, M., Meline, V., Iyer-Pascuzzi, A.S., Souza, A.M., Tuinstra, M.R. and Yang, Y., 2021. Root Phenotyping from X-Ray Computed Tomography: Skeleton Extraction. *The International Archives of Photogrammetry, Remote Sensing and Spatial Information Sciences*, 43, pp.417-422. <https://doi.org/10.5194/isprs-archives-XLIII-B4-2021-417-2021>
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- Herrero-Huerta, M., Tolley, S., Tuinstra, M.R. and Yang, Y., 2021, April. Individual maize extraction from UAS imagery-based point clouds by 3D deep learning. In *Autonomous Air and Ground Sensing Systems for Agricultural Optimization and Phenotyping VI* (Vol. 11747, p. 1174704). International Society for Optics and Photonics. <https://doi.org/10.1117/12.2587100>
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- Yang, K.W., Chapman, S., Carpenter, N., Hammer, G., McLean, G., Zheng, B., Chen, Y., Delp, E., Masjedi, A., Crawford, M. Ebert, D., Habib, A., Thompson, A., Weil, C., Tuinstra, M.R., 2021. Integrating crop growth models with remote sensing for predicting biomass yield of sorghum. *in silico Plants*, 3(1), p.diab001. <https://doi.org/10.1093/insilicoplants/diab001>
- Zhang, X., Xie, J., Chen, T., Ma, D., Yao, T., Gu, F., Lim, J., Tuinstra, M.R., Hamaker, B.R., 2021. High arabinoxylan fine structure specificity to gut bacteria driven by corn genotypes but not environment. *Carbohydrate Polymers*, 257: 117667. <https://doi.org/10.1016/j.carbpol.2021.117667>
- Griebel, S., Adedayo, A. and Tuinstra, M.R., 2021. Genetic diversity for starch quality and alkali spreading value in sorghum. *The Plant Genome*, 14(1), p.e20067. <https://doi.org/10.1002/tpg2.20067>
- Jarquín, D., de Leon, N., Romay, C., Bohn, M., Buckler, E.S., Ciampitti, I., Edwards, J., Ertl, D., Flint-Garcia, S., Gore, M.A., Graham, C., Hirsch, C.N., Holland, J.B., Hooker, D., Kaeppler, S.M., Knoll, K., Lee, E.C., Lawrence-Dill, C.J., Lynch, J.P., Moose, S.P., Murray, S.C., Nelson, R., Rocheford, T., Schnable, J.C., Schnable, P.S., Smith, M., Springer, N., Thomison, P., Tuinstra, M.R., Wisser, R.J., Xu, W., and Lorenz, A., 2020. Utility of Climatic Information via Combining Ability Models to Improve Genomic Prediction for Yield within the Genomes to Fields Maize Project. *Frontiers in Genetics*, 11, p.1819. <https://doi.org/10.3389/fgene.2020.592769>

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- Cotrozzi, L., Peron, R., Tuinstra, M.R., Mickelbart, M.V. and Couture, J.J., 2020. Spectral phenotyping of physiological and anatomical leaf traits related with maize water status. *Plant Physiology*, 184(3), pp.1363-1377. <https://doi.org/10.1104/pp.20.00577>
- McFarland, B.A., AlKhalifah, N., Bohn, M., Bubert, J., Buckler, E.S., Ciampitti, I., Edwards, J., Ertl, D., Gage, J.L., Falcon, C.M., Flint-Garcia, S., Gore, M.A., Graham, C., Hirsch, C.N., Holland, J.B., Hood, E., Hooker, D., Jarquin, D., Kaeppler, S.M., Knoll, J., Kruger, G., Lauter, N., Lee, E.C., Lima, D.C., Lorenz, A., Lynch, J.P., McKay, J., Miller, N.D., Moose, S.P., Murray, S.C., Nelson, R., Poudyal C., Rocheford T., Rodriguez, O., Romay, M.C., Schnable, J.C., Schnable, P.S., Scully, B., Sekhon, R., Silverstein, K., Singh, M., Smith, M., Spalding, E.P., Springer, N., Thelen, K., Thomison, P., Tuinstra, M.R., Wallace, J., Walls, R., Wills, D., Wisser, R.J., Wenwei, X., Yeh, C.-T., and de Leon, N., 2020. Maize genomes to fields (G2F): 2014–2017 field seasons: genotype, phenotype, climatic, soil, and inbred ear image datasets. *BMC Research Notes*, 13(1), pp.1-6. <https://doi.org/10.1186/s13104-020-4922-8>
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- Wang, L., Jin, J., Song, Z., Wang, J., Zhang, L., Rehman, T.U., Ma, D., Carpenter, N.R. and Tuinstra, M.R., 2020. LeafSpec: An accurate and portable hyperspectral corn leaf imager. *Computers and Electronics in Agriculture*, 169, p.105209. <https://doi.org/10.1016/j.compag.2019.105209>
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- Wu, Y., Guo, T., Mu, Q., Wang, J., Li, X., Wu, Y., Tian, B., Wang, M.L., Bai, G., Perumal, R., Trick, H.N., Bean, S.R., Dweikat, I.M., Tuinstra, M.R., Morris, G., Tesso, T.T., Yu, J., Li, X., 2019. Allelochemicals targeted to balance competing selections in African agroecosystems. *Nature Plants*, pp.1-8. <https://doi.org/10.1038/s41477-019-0563-0>
- Griebel, S., Westerman, R.P., Adeyanju, A., Addo-Quaye, C., Craig, B.A., Weil, C.F., Cunningham, S.M., Patel, B., Campanella, O.H. and Tuinstra, M.R., 2019. Mutations in sorghum SBEIIb and SSIIa affect alkali spreading value, starch composition, thermal properties and flour viscosity. *Theoretical and Applied Genetics*, 132(12), pp.3357-3374. <https://doi.org/10.1007/s00122-019-03430-0>

- Ma, D., Carpenter, N., Amatya, S., Maki, H., Wang, L., Zhang, L., Neeno, S., Tuinstra, M.R. and Jin, J., 2019. Removal of greenhouse microclimate heterogeneity with conveyor system for indoor phenotyping. *Computers and Electronics in Agriculture*, 166, p.104979. <https://doi.org/10.1016/j.compag.2019.104979>
- Ma, D., Carpenter, N., Rehman, T., Maki, H., Tuinstra, M.R. and Jin J., 2019. Greenhouse Environment Modeling and Simulation for Microclimate Control. *Computers and Electronics in Agriculture*, 162, pp.134-142. <https://doi.org/10.1016/j.compag.2019.04.013>
- Griebel, S., Webb, M.M., Campanella, O.H., Craig, B.A., Weil, C.F. and Tuinstra, M.R., 2019. The alkali spreading phenotype in Sorghum bicolor and its relationship to starch gelatinization. *Journal of cereal science*, 86, pp.41-47. <https://doi.org/10.1016/j.jcs.2019.01.002>
- Al Khalifah, N., Campbell, D.A., Falcon, C.M., Gardiner, J.M., Miller, N.D., Cinta Romay R, Walls, R., Walton, R., Yeh, C.T., Bohn, M., Bubert, J., Buckler, E.S., Ciampitti, I., Flint-Garcia, S., Gore, M.A., Graham, C., Hirsch, C., Holland, J.B., Hooker, D., Kaeppler, S., Knoll, J., Lauter, N., Lee, E.C., Lorenz, A., Lynch, N.P., Moose, S.P., Murray, S.C., Nelson, R., Rocheford, T., Rodriguez, O., Schnable, J.C., Scully, B., Smith, M., Springer, N., Thomison, P., Tuinstra, M.R., Wisser, R.J., Xu, W., Ertl, D., Schnable, P., De Leon, N., Spalding, E.P., Edwards, J., Lawrence-Dill, C.J. 2018. Maize Genomes to Fields: 2014 and 2015 field season genotype, phenotype, environment, and inbred ear image datasets. *BMC research notes*, 11(1), p.452. <https://doi.org/10.1186/s13104-018-3508-1>
- Balzan, S., Carraro, N., Salleres, B., Dal Cortivo, C., Tuinstra, M.R., Johal, G. and Varotto, S., 2018. Genetic and phenotypic characterization of a novel brachytic2 allele of maize. *Plant Growth Regulation*, pp.1-12. <https://doi.org/10.1007/s10725-018-0412-6>
- Addo-Quaye, C., Tuinstra, M., Carraro, N., Weil, C. and Dilkes, B.P., 2018. Whole genome sequence accuracy is improved by replication in a population of mutagenized sorghum. *G3: Genes, Genomes, Genetics*, pp.g3-300301. <https://doi.org/10.1534/g3.117.300301>
- Bouchet, S., Olatoye, M.O., Marla, S.R., Perumal, R., Tesso, T., Yu, J., Tuinstra, M. and Morris, G.P., 2017. Increased power to dissect adaptive traits in global sorghum diversity using a nested association mapping population. *Genetics*, 206(2), pp.573-585. <https://doi.org/10.1534/genetics.116.198499>