

IWYP Publications (2016-20)

LEAD PI	TITLE	YEAR
Andrew Merchant	Salter, W. T., M. E. Gilbert, and T. N. Buckley. 2018. Time-Dependent Bias in Instantaneous Ceptomety caused by Row Orientation. <i>Plant Phenome J.</i> 1:180004. doi:10.2135/tppj2018.07.0004	2018
Andrew Merchant	William T. Salter, Matthew E. Gilbert, and Thomas N. Buckley (2018). A multiplexed gas exchange system for increased throughput of photosynthetic capacity measurements. <i>Plant Methods</i> 2018 14:80 https://doi.org/10.1186/s13007-018-0347-y	2018
Andrew Merchant	Salter W. T, Merchant A, Buckley T. N., (2018) PARbars: cheap, easy to build ceptometers for continuous measurement of light interception in plant canopies. <i>Journal of Visualized Experiments</i>	2019
Andrew Merchant	Salter W. T, Merchant A, Richards, RA Trethowan R, Buckley T. N. (2019) Rate of photosynthetic acclimation to fluctuating light varies widely among genotypes of wheat. <i>Journal of Experimental Botany</i>	2019
Anthony Hall	Gardiner LJ, Joynson R, Omony J, et al. Hidden variation in polyploid wheat drives local adaptation. <i>Genome Res.</i> 2018;28(9):1319-1332.	2018
Anthony Hall	Evans JR and Clarke VC. (2019) The nitrogen cost of photosynthesis. <i>J Experimental Botany</i> 70, 7-15	2019
Anthony Hall	Gardiner LJ, Wingen LU, Bailey P, et al. Analysis of the recombination landscape of hexaploid bread wheat reveals genes controlling recombination and gene conversion frequency. <i>Genome Biol.</i> 2019;20(1):69. Published 2019 Apr 15. doi:10.1186/s13059-019-1675-6	2019
Anthony Hall	Gardiner, L.-J., Brabbs, T., Akhunov, A., Jordan, K., Budak, H., Richmond, T., Singh, S., Catchpole, L., Akhunov, E., Hall, A. (2019). Integrating genomic resources to present full gene and putative promoter capture probe sets for bread wheat. <i>GigaScience</i> . doi.org/10.1093/gigascience/giz018	2019
Arnold Bloom	Bloom, A. J. and Kameritsch, P. (2017), Relative association of Rubisco with manganese and magnesium as a regulatory mechanism in plants. <i>Physiol Plantarum</i> , 161: 545-559. doi:10.1111/ppl.12616	2017
Arnold Bloom	Bloom, A. J. and K. M. Lancaster (2018) Manganese binding to Rubisco could drive a photorespiratory pathway that increases the energy efficiency of photosynthesis. <i>Nature Plants</i> 4:414-422, DOI:10.1038/s41477-018-0191-0	2018
Arnold Bloom	Bloom, A.J. (2019) Metal Regulation of metabolism. <i>Current Opinion in Chemical Biology</i> . Vol 49 33-38	2019
Arnold Bloom	Bloom, A.J., Kasemsap, P. and Rubio-Asensio, J.S. (2020), Rising atmospheric CO ₂ concentration inhibits nitrate assimilation in shoots but enhances it in roots of C ₃ plants. <i>Physiol Plantarum</i> , 168: 963-972. doi:10.1111/ppl.13040	2020
Barry Pogson	Coast, O., Shah, S., Ivakov, A., Gaju, O., Wilson, P.B., Posch, B.C., Bryant, C.J., Negrini, A.C.A., Evans, J.R., Condon, A.G., Silva-Pérez, V., Reynolds, M.P., Pogson, B.J., Millar, A.H., Furbank, R.T., and Atkin, O.K. Predicting dark respiration rates of	2019

LEAD PI	TITLE	YEAR
	wheat leaves from hyperspectral reflectance. <i>Plant Cell Environ.</i> 2019; 42: 2133– 2150. https://doi.org/10.1111/pce.13544	
Christine Raines	Driever SM, Simkin AJ, Alotaibi S, Fisk SJ, Madgwick PJ, Sparks CA, Jones HD, Lawson T, Parry MAJ, Raines CA. (2017). Increased SBPase activity improves photosynthesis and grain yield in wheat grown in greenhouse conditions. <i>Phil. Trans. R. Soc. B.</i>	2017
Christine Raines	Alotaibi, S. S., Sparks, C. A., Parry, M., Simkin, A. J., & Raines, C. A. (2018). Identification of Leaf Promoters for Use in Transgenic Wheat. <i>Plants (Basel, Switzerland)</i> , 7(2), 27. doi:10.3390/plants7020027	2018
Christine Raines	Simkin AJ, Lopez-Calcagno PE, Raines CA. (2019). Feeding the world: Improving photosynthetic efficiency for sustainable crop production. <i>J Exp Bot.</i> (doi.org/10.1093/jxb/ery445).	2019
Cristobal Uauy	Simmonds J, Scott P, Brinton J, Mestre TC, Bush M, Del Blanco A, Dubcovsky J, Uauy C (2016) A splice acceptor site mutation in TaGW2-A1 increases thousand grain weight in tetraploid and hexaploid wheat through wider and longer grains <i>Theoretical and Applied Genetics</i> 129:1099–1112.	2016
Cristobal Uauy	Krasileva KV, Vasquez-Gross H, Howell T, Bailey P, Paraiso F, Clissold L, Simmonds J, Ramirez-Gonzalez RH, Wang X, Borrill P, Fosker C, Ayling S, Phillips A, Uauy C, Dubcovsky J (2017) Uncovering hidden variation in the young polyploid wheat genomes. <i>PNAS.</i> doi: 10.1073/pnas.1619268114.	2017
Cristobal Uauy	Uauy C 2017. Wheat genomics comes of age. <i>Current Opinion Plant Biology.</i> 36:142	2017
Cristobal Uauy	Uauy C, Wulff B, Dubcovsky J. 2017. Combining Traditional Mutagenesis with New High-Throughput Sequencing and Genome Editing to Reveal Hidden Variation in Polyploid Wheat. <i>Annual Review in Genetics.</i> 51: doi.org/10.1146/annurev-genet-120116-024533	2017
Cristobal Uauy	Borrill P, Ramirez-Gonzalez R et al Uauy C. 2018. The transcriptional landscape of polyploid wheat. <i>Science</i>	2018
Cristobal Uauy	Watson A, Ghosh S et al 2018. Speed breeding is a powerful tool to accelerate crop research and breeding. <i>Nature Plants</i> 4:23-29	2018
Cristobal Uauy	Brinton, J, Uauy, C (2019) A reductionist approach to dissecting grain weight and yield in wheat. <i>J Integr Plant Biol</i> 61: 337– 358	2019
Cristobal Uauy	Wang W, Simmonds K, Pan Q, Davidson D, He F, Battal A, Akhunova A, Trick HN, Uauy C, Akhunov E. 2019. Gene editing and mutagenesis reveal inter-cultivar differences and additivity in the contribution of TaGW2 homoeologues to grain size and weight in wheat. <i>Theoretical and Applied Genetics</i>	2019
Eduard Akhunov	Wang W, Pan Q, He F, Akhunova A, Chao S, Trick HN, Akhunov E, Transgenerational CRISPR-Cas9 activity facilitates multiplex gene editing in allopolyploid wheat. <i>The CRISPR Journal</i> , 2018, 1, 65-74.	2018
Eduard Akhunov	Wang, W., Simmonds, J., Pan, Q. et al. Gene editing and mutagenesis reveal inter-cultivar differences and additivity in the contribution of TaGW2 homoeologues to grain size and weight in wheat. <i>Theor Appl Genet</i> 131, 2463–2475 <i>Theor Appl Genet</i> (2018) 131: 2463. https://doi.org/10.1007/s00122-018-3166-7	2018

LEAD PI	TITLE	YEAR
Eduard Akhunov	Wang W., Q. Pan, B. Tian, F. He, Y. Chen, G. Bai, A. Akhunova, H.N. Trick, E. Akhunov. 2019. Gene editing of the wheat homologs of TONNEAU1 recruiting motif encoding gene affects grain shape and weight in wheat. <i>Plant J.</i> 2019, 100(2):251-264, doi: 10.1111/tpj.14440.	2019
Elizabete Carmo-Silva	Perdomo JA, Degen GE, Worrall D, Carmo-Silva E. Rubisco activation by wheat Rubisco activase isoform 2 β is insensitive to inhibition by ADP. <i>The Biochemical Journal.</i> 2019 Sep;476(18):2595-2606. DOI: 10.1042/bcj20190110.	2019
Erik Murchie	Smith HL, McAusland L, Murchie EH. Don't ignore the green light: exploring diverse roles in plant processes. <i>Journal of Experimental Botany.</i> 2017 Apr;68(9):2099-2110. DOI: 10.1093/jxb/erx098.	2017
Erik Murchie	Jauregui I, Rothwell SA, Taylor SH, Parry MAJ, Carmo-Silva E, Dodd IC. 2018. Whole plant chamber to examine sensitivity of cereal gas exchange to changes in evaporative demand. <i>Plant Methods</i> 14: 97.	2018
Erik Murchie	Lawson, T. and Violet-Chabrand, S. (2019), Speedy stomata, photosynthesis and plant water use efficiency. <i>New Phytol</i> , 221: 93-98. doi:10.1111/nph.15330	2019
Erik Murchie	McAusland L., Atkinson J.A., Lawson T., Murchie E.H. (2019) High throughput procedure utilising chlorophyll fluorescence imaging to phenotype dynamic photosynthesis and photoprotection in leaves under controlled gaseous conditions. <i>Plant Methods</i> , 15 , 109. Available at: https://plantmethods.biomedcentral.com/articles/10.1186/s13007-019-0485-x	2019
Erik Murchie	Przewieslik-Allen A.M., Burrige A.J., Wilkinson P.A., Winfield M.O., Shaw D.S., McAusland L., King J., King I.P., Edwards K.J., Barker G.L.A. (2019) Developing a High-Throughput SNP-Based Marker System to Facilitate the Introgression of Traits From Aegilops Species Into Bread Wheat (<i>Triticum aestivum</i>). <i>Frontiers in Plant Science</i> , 9, 1993. Available at: https://www.frontiersin.org/article/10.3389/fpls.2018.01993/full	2019
Erik Murchie	Violet-Chabrand S., Lawson T. (2019) Dynamic leaf energy balance: deriving stomatal conductance from thermal imaging in a dynamic environment. <i>Journal of Experimental Botany</i> , 70 , 2839–2855. Available at: https://academic.oup.com/jxb/article/70/10/2839/5356941	2019
Erik Murchie	Violet-Chabrand S., Lawson T. (2020) Thermography methods to assess stomatal behaviour in a dynamic environment Ed J. Evans. <i>Journal of Experimental Botany</i> . Available at: https://academic.oup.com/jxb/advance-article/doi/10.1093/jxb/erz573/5698117	2020
Jesse Poland	Sun, J., J. A. Poland, S. Mondal, J. Crossa, P. Juliana, R. P. Singh, J. E. Rutkoski, J.-L. Jannink, L. Crespo-Herrera, G. Velu, J. Huerta-Espino and M. E. Sorrells (2019) High-throughput phenotyping platforms enhance genomic selection for wheat grain yield across populations and cycles in early stage. <i>Theoretical and Applied Genetics</i> 132(6): 1705-1720. DOI: 10.1007/s00122-019-03309-0	2019
Jesse Poland	Wang X., C. Amos, M. Lucas, G. Williams, J. Poland. (2019) Small plot identification from video streams for high-throughput phenotyping of large breeding populations with unmanned aerial systems. <i>Autonomous Air and Ground Sensing Systems for Agricultural Optimization and Phenotyping IV.</i> 11008(2019), 110080D. International Society for Optics and Photonics	2019

LEAD PI	TITLE	YEAR
Jesse Poland	Xu Wang, Hong Xuan, Byron Evers, Sandesh Shrestha, Robert Pless, Jesse Poland, High-throughput phenotyping with deep learning gives insight into the genetic architecture of flowering time in wheat, <i>GigaScience</i> , Volume 8, Issue 11, November 2019, giz120, https://doi.org/10.1093/gigascience/giz120	2019
Jorge Dubkovsky	Debernardi JM, Lin H, Chuck G, Faris JD, Dubcovsky J. microRNA172 plays a crucial role in wheat spike morphogenesis and grain threshability. <i>Development</i> . 2017;144(11):1966-1975.	2017
Jorge Dubkovsky	Blake, N. K., A. C. Varella, B. Bicego, J. M. Martin, J. P. Cook, H.-Y. Heo, R. Acharya, J. D. Sherman, D. Nash, L. E. Talbert. 2018. Maturity traits related to climate adaptation impact quality characteristics in hard red spring wheat. <i>Crop Sci</i> . 58:1954-1963	2018
Jorge Dubkovsky	El-Feki, W.M., P.F. Byrne, S.D. Reid, and S.D. Haley. 2018. Mapping quantitative trait loci for agronomic traits in winter wheat under different soil moisture levels. <i>Agronomy</i> 8: 133. doi:10.3390/agronomy8080133	2018
Jorge Dubkovsky	Gizaw S.A., J.G. Godoy, M.O. Pumphrey, and A.H. Carter. 2018. Spectral reflectance for indirect selection and genome-wide association analysis of grain yield and drought tolerance in North American spring wheat (<i>Triticum aestivum</i> L.). <i>Crop Sci</i> . 58: 2289-2301	2018
Jorge Dubkovsky	Gizaw, S.A., J.G.V. Godoy, K. Garland-Campbell, A.H. Carter. 2018. Using spectral reflectance as proxy phenotypes for genome-wide association studies of yield and yield stability in Pacific Northwest winter wheat. <i>Crop Science</i> 58:1232-1241. doi:10.2135/cropsci2017.11.0710	2018
Jorge Dubkovsky	Godoy, J., S. Gizaw, S. Chao, N. Blake, A. Carter, R. Cuthbert, J. Dubcovsky, P. Hucl, K. Kephart, C. Pozniak, P.V. V. Prasad, M. Pumphrey, and L. Talbert. 2018. Genome-wide association study of agronomic traits in a spring-planted North American elite hard red spring wheat panel. <i>Crop Science</i> 58:1838-1852	2018
Jorge Dubkovsky	Huang, M., N. Mheni, G. Brown-Guedira, A. McKendry, C. Griffey, D. Van Sanford, J. Costa, C Sneller. 2018. Genetic analysis of heading date in winter and spring wheat. <i>Euphytica</i> , 214: 128. doi:10.1007/s10681-018-2199-y	2018
Jorge Dubkovsky	Kippes, N., M. Guedira, L. Lin, G.L. Brown-Guedira and J. Dubcovsky. 2018. Single nucleotide polymorphisms in a regulatory site of VRN-A1 first intron are associated with differences in vernalization requirement in winter wheat. <i>Molecular Genetics and Genomics</i> . doi:10.1007/s0043	2018
Jorge Dubkovsky	Lozada, D.N., Mason, R.E., Sukumaran, S., Dreisigacker, S. 2018 Validation of grain yield QTL from soft winter wheat using a CIMMYT spring wheat panel. <i>Crop Science</i> . doi:10.2135/cropsci2018.04.0232	2018
Jorge Dubkovsky	Mo, J., T. Howell, H. Vasquez-Gross, L.A. de Haro, J. Dubcovsky, S. Pearce. 2018. Mapping causal mutations by exome sequencing in a wheat TILLING population: a tall mutant case study. <i>Mol. Genet. Genom.</i> 293: 463-477. doi:10.1007/s00438-017-1401-6	2018
Jorge Dubkovsky	Mo, Y., Vanzetti, L.S., Hale, I. et al. 2018. Identification and characterization of Rht25, a locus on chromosome arm 6AS affecting wheat plant height, heading time, and spike development. <i>Theor Appl Genet</i> (2018) 131: 2021. https://doi.org/10.1007/s00122-018-3130-6	2018

LEAD PI	TITLE	YEAR
Jorge Dubkovsky	Nyine, M., S. Wang, K. Kiani, K. Jordan, S.Y. Liu, P. Byrne, S. Haley, S. Baenziger, S. Chao, R. Bowden, E. Akhunov. 2018. Genotype imputation in winter wheat using first-generation haplotype map SNPs improves genome-wide association mapping and genomic predictions of traits. <i>G3</i> 9:125-133. G3/2018/200664	2018
Jorge Dubkovsky	Wang R., Y. Liu, K. Isham, W. Zhao, J. Wheeler, N. Klassen, Y. Hu, J.M. Bonman, J. Chen. 2018. QTL identification and KASP marker development for productive tiller and fertile spikelet numbers in two high-yielding hard white spring wheat cultivars. <i>Mol. Bred.</i> 38:135	2018
Jorge Dubkovsky	Zhang, J., S.A. Gizaw, E. Bossolini, J. Hegarty, T. Howell, A.H. Carter, E. Akhunov, J. Dubcovsky. 2018. Identification and validation of QTL for grain yield and plant water status under contrasting water treatments in fall-sown spring wheats. <i>Theor. Appl. Genet.</i> 131: 1741–1759. doi:10.1007/s00122-018-3111-9	2018
Jorge Dubkovsky	Bernardo, A., P.St. Amand, H.Q. Le, Z. Su, and G. Bai. 2019. Multiplex restriction amplicon sequencing: a novel next-generation sequencing-based marker platform for high-throughput genotyping. <i>Plant Biotechnol. J.</i> , https://doi.org/10.1111/pbi.13192	2019
Jorge Dubkovsky	DeWitt, N., M. Guedira, E. Lauer, M. Sarinelli, P. Tyagi, D. Fu, Q. Hao, J.P. Murphy, D. Marshall, A. Akhunova, K. Jordan, E. Akhunov, and G. Brown-Guedira. 2019. Sequence based mapping identifies AWNS1, a candidate transcription repressor underlying awn suppression at the B1 locus in wheat. <i>New Phytologist</i> . doi: 10.1111/nph.16152	2019
Jorge Dubkovsky	Hayat, H., Mason, R.E., Lozada, D.N., Acuna, A., Holder, A., Larkin, D.L., Winn, Z., Murry, J., Murphy, P., Moon, D.E., Miller, R.G. 2019. Effects of allelic variation at Rht-B1 and Rht-D1 on grain yield and agronomic traits of southern US soft red winter wheat. <i>Euphytica</i> . DOI: 10.1007/s10681-019-2478-2	2019
Jorge Dubkovsky	He F., R. Pasam, F. Shi, S. Kant, G. Keeble-Gagnere, P. Kay, K. Forrest, A. Fritz, P. Hucl, K. Wiebe, R. Knox, R. Cuthbert, C. Pozniak, A. Akhunova, P.L. Morrell, J.P. Davies, S.R. Webb, G. Spangenberg, B. Hayes, H. Daetwyler, J. Tibbits, M. Hayden, E. Akhunov. 2019. Exome sequencing highlights the role of wild relative introgression in shaping the adaptive landscape of the wheat genome. <i>Nat Genet.</i> 51:896904.	2019
Jorge Dubkovsky	Krause, M.R., L. Gonzalez-Perez, J. Crossa, P. Perez-Rodriguez, O. Montesinos-Lopez, R.P. Singh, S. Dreisigacker, J. Poland, J. Rutkoski, M.E. Sorrells, M.A. Gore, and S. Mondal. 2019. Hyperspectral reflectance-derived relationship matrices for genomic prediction of grain yield in wheat. <i>Genes Genomes Genetics.</i> 9:1231-1247	2019
Jorge Dubkovsky	Kuzay, S., Y. Xu, J. Zhang, A. Katz, S. Pearce, Z. Su, M. Fraser, J. A. Anderson, G. Brown-Guedira, N. DeWitt, A. Peters Haugrud, J.D. Faris, E. Akhunov, G. Bai, J. Dubcovsky. 2019. Identification of a candidate gene for a QTL for spikelet number per spike on wheat chromosome arm 7AL by high-resolution genetic mapping. <i>Theor. Appl. Genet.</i> 132:26892705	2019
Jorge Dubkovsky	Larkin, D.L., D.N. Lozada, and R.E Mason. 2019. Genomic selection considerations for successful implementation in wheat breeding programs. <i>Agronomy</i> . doi:10.3390/agronomy9090479	2019

LEAD PI	TITLE	YEAR
Jorge Dubkovsky	Li, C., H. Lin, A. Chen, M. Lau, J. Jernstedt, and J. Dubcovsky. 2019. Wheat VRN1, FUL2 and FUL3 play critical and redundant roles in spikelet meristem identity and spike determinacy. <i>Development</i> . 146: On line first. doi:10.1242/dev.175398	2019
Jorge Dubkovsky	Lozada, D.L., R.E. Mason, J.M. Sarinelli, and G. Brown-Guedira. 2019 Accuracy of genomic selection for grain yield and agronomic traits in soft red winter wheat. <i>BMC Genetics</i>	2019
Jorge Dubkovsky	Santantonio, N., J.L. Jannink, and M.E. Sorrells. 2019. Homeologous epistasis in wheat: the search for an immortal hybrid. <i>Genetics</i> . doi.org/10.1534/genetics.118.301851	2019
Jorge Dubkovsky	Sarinelli, J.M., J.P. Murphy, P. Tyagi, J.B. Holland, J.W. Johnson, M. Mergoum, R.E. Mason, A. Babar, S. Harrison, R. Sutton, C.A. Griffey, and G. Brown Guedira. 2019. Training population selection and use of fixed effects to optimize genomic predictions in a historical USA winter wheat panel. <i>Theor. Appl. Genet.</i> 132: 12471261	2019
Jorge Dubkovsky	Shaw, L. B. Lyu, R. Turner, C. Li, F. Chen, X. Han, D. Fu, and J. Dubcovsky. 2019. FLOWERING LOCUS T2 (FT2) regulates spike development and fertility in temperate cereals. <i>J. of Exp. Bot.</i> 70: 193-204	2019
Jorge Dubkovsky	Sun J., J.A. Poland, S. Mondal, J. Crossa, P. Juliana, R.P. Singh, J.E. Rutkoski, J-L. Jannink, L. Crespo-Herrera, G. Velu, J. Huerta-Espino, M.E. Sorrells. 2019. High-throughput phenotyping platforms enhance genomic selection for wheat grain yield across populations and cycles in early stage. <i>Theor Appl Genet.</i> 132:1705-1720	2019
Jorge Dubkovsky	Ward, B.P., G.L. Brown-Guedira, F.L. Kolb, D.A. Van Sanford, P. Tyagi, C.H. Sneller and C.A. Griffey. 2019. Genome-wide association studies for yield-related traits in soft red winter wheat grown in Virginia. <i>PLoS ONE</i> , 14, p.e0208217	2019
Jorge Dubkovsky	Ward, B.P., G.L. Brown-Guedira, P. Tyagi, F.L. Kolb, D.A. Van Sanford, C.H. Sneller and C.A. Griffey 2019. Multi-environment and multi-trait genomic selection models in unbalanced early-generation wheat yield trials. <i>Crop Science</i> . 59:491-507	2019
Jorge Dubkovsky	Debernardi, J.M., Greenwood, J.R., Jean Finnegan, E., Jernstedt, J. and Dubcovsky, J. (2020), APETALA 2-like genes AP2L2 and Q specify lemma identity and axillary floral meristem development in wheat. <i>Plant J</i> , 101: 171-187. doi:10.1111/tpj.14528	2020
Matthew Paul	Paul MJ, Watson A, Griffiths CA (2019) Linking fundamental science to crop improvement through understanding source and sink traits and their integration for yield enhancement. <i>Journal of Experimental Botany</i> , doi:10.1093/jxb/erz480	2019
Matthew Reynolds	IPPN Abstracts 2016, 4th International Plant Phenotyping Symposium. December 12-15, 2016 CDMX, Mexico: International Maize and Wheat Improvement Center (CIMMYT). http://hdl.handle.net/10883/17932	2016
Matthew Reynolds	Tattaris M, Reynolds MP, Chapman SC. A Direct Comparison of Remote Sensing Approaches for High-Throughput Phenotyping in Plant Breeding. <i>Front Plant Sci.</i> 2016;7:1131. Published 2016 Aug 3. doi:10.3389/fpls.2016.01131	2016
Matthew Reynolds	Reynolds M. et al, (2017). Proceedings of the 3rd International TRIGO (Wheat) Yield Potential Workshop 2017. CENEB, CIMMYT, Cd. Obregón, Sonora, Mexico, 22-23rd March 2017. Mexico, CDMX: CIMMYT.	2017
Matthew Reynolds	Reynolds M.P.; Pask A.J.D.; Hoppitt W.J.E.; Sonder K.; Sukumaran S.; Molero G.; Saint Pierre C.; Payne T.; Singh R.; Braun H.; Gonzalez F.; Barma N.; Hakim A.; He Z.; Fan Z.; Novoselovic D.; Maghraby M.; Gad K.; ElHusseiny Galal; Hagraas A.;	2017

LEAD PI	TITLE	YEAR
	Mohamed M.; Morad A.; Kumar U.; Singh G.; Naik R.; Kalappanavar I.; Biradar S.a; Sakuru Ve.; Chatrath R.; Sharma I.; Panchabhai K.; Sohu V.; Mavi G.; Mishra V.; Balasubramaniam A.; Jalal-Kamali M.; Khodarahmi M.; Dastfal M.; Tabib-Ghaffari S.; Jafarby J.; Nikzad A.; Moghaddam H.; Ghogh H.; Mehraban A.; Solís-Moya E.; Camacho-Casas M.; Figueroa-López P.; Moreno J.; Alvarado-Padilla J.; Borbón-Gracia A.; Torres A.; Quiche N.; Upadhyay S.; Pandey D.; Imtiaz M.; Rehman M.; Hussain M.; Hussain M.; Ud-Din R.; Qamar M.; Kundi S.; Mujahid M.Y.; Ahmad G.; Khan A.; Sial M.; Mustatea P.; von Well E.; Ncala M.; de Groot S.; Hussein A.; Tahir I.; Idris A.; Elamein H.; Joshi A. (2017). Strategic crossing of biomass and harvest index—source and sink—achieves genetic gains in wheat. <i>Euphytica</i> 213: 257. https://doi.org/10.1007/s10681-017-2040-z Print ISSN: 0014-2336. Online ISSN: 1573-5060	
Matthew Reynolds	Sukumaran S, Lopes M, Dreisigacker S, Reynolds M (2017) Genetic analysis of multi-environmental spring wheat trials identifies genomic regions for locus-specific trade-offs for grain weight and grain number. <i>Theor Appl Genet</i> 1–14. doi: 10.1007/s00122-017-3037-7	2017
Matthew Reynolds	Sukumaran S, Reynolds MP, Sansaloni CP (2018) Genome-wide association analyses identify QTL hotspots for yield and component traits in durum wheat grown under yield potential, drought, and heat stress environments. <i>Front Plant Sci</i> 9:81. doi: 10.3389/fpls.2018.00081	2017
Matthew Reynolds	Dreccer, M.F., Molero, G., Rivera-Amado, C., John-Bejai, C., Wilson, Z. 2019. Yielding to the image: How phenotyping reproductive growth can assist crop improvement and production. <i>Plant Science</i> 282 73-82 DOI: 10.1016/j.plantsci.2018.06.008	2019
Matthew Reynolds	Liu, C., Khodae, M., Lopes, M.S., Sansaloni, C., Dreisigacker, S., Sukumaran, S., Reynolds, M. 2019. Multi-environment QTL analysis using an updated genetic map of a widely distributed Seri × Babax spring wheat population. <i>Molecular Breeding</i> 39:134 DOI: 10.1007/s11032-019-1040-1	2019
Matthew Reynolds	Liu, C., Sukumaran, S., Claverie, E., Sansaloni, C., Dreisigacker, S., Reynolds, M. 2019. Genetic dissection of heat and drought stress QTLs in phenology-controlled synthetic-derived recombinant inbred lines in spring wheat. <i>Molecular Breeding</i> 39 (34) DOI 10.1007/s11032-019-0938-y	2019
Matthew Reynolds	Molero, G., Joynson R., Pinera-Chavez FJ, Gardiner LJ, Rivera-Amado C, Hall A, Reynolds MP. 2019. Elucidating the genetic basis of biomass accumulation and radiation use efficiency in spring wheat and its role in yield potential. <i>Plant Biotechnology Journal</i> 17 1276-1288 DOI: 10.1111/pbi.13052	2019
Matthew Reynolds	Reynolds, M., Borrell. A., Braun, H., Edmeades, G., Flavell, R., Gwyn, J., Jordan, D., Pixley, K., Rebetzke, G., 2019. Translational Research for Climate Resilient, Higher Yielding Crops. <i>Crop Breeding Genetics Genomics</i> 1:e190016. https://doi.org/10.20900/cbgg20190016	2019
Matthew Reynolds	Rivera-Amado, C., Trujillo-Negrellos, E., Molero, G., Reynolds, M.P., Sylvester-Bradley, R., Foulkes, M.J. 2019. Optimizing dry-matter partitioning for increased spike growth, grain number and harvest index in spring wheat. <i>Field Crops Research</i> 240 154-167 DOI: 10.1016/j.fcr.2019.04.016	2019

LEAD PI	TITLE	YEAR
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