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### List of publications:

- Badouin H, et al., (2017) The sunflower genome provides insights into oil metabolism, flowering and Asterid evolution. *Nature*. 546(7656):148-152. doi: 10.1038/nature22380.
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- Hyles J, Vautrin S, Pettolino F, MacMillan C, Stachurski Z, Breen J, Berges H, Wicker T, Spielmeier W. (2017) Repeat-length variation in a wheat cellulose synthase-like gene is associated with altered tiller number and stem cell wall composition. *J Exp Bot*. 68(7):1519-1529. doi: 10.1093/jxb/erx051
- Yousfi FE, Makhloufi E, Marande W, Ghorbel AW., Bouzayen M and Berg es H (2017) Comparative Analysis of WRKY Genes Potentially Involved in Salt Stress Responses in *Triticum turgidum* L. ssp. *durum*. *Front Plant Sci* 7:2034. doi: 10.3389/fpls.2016.02034.
- Gilles L, Khaled A, Laffaire JB, Chaignon S, Gendrot G, Laplaige J, Berg es H, Beydon G, Barret P, Comadran J, Martinant JP, Rogowsky PM and Widiez T (2017) Pollen-specific phospholipase Not Like Dad (NLD) triggers gynogenesis in maize. *The EMBO Journal*. 36(4): 389–564 doi:10.15252/embj.201796603
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- Gamas, P., F. Debellé, Bergès H., Godiard L., Niebel A., Journet E-P., Gouzy J. (2007). *Medicago truncatula* cDNA and genomic libraries. *Medicago truncatula* handbook., U. Mathesius, E. P. Journet and L. W. Sumner, ISBN 0-9754303-1-9 <http://www.noble.org/MedicagoHandbook>.

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