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Development and Characterization of a Multiparent Advanced Generation Inter-Cross (MAGIC) Population of Jute (*Corchorus olitorius*)

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Abstract

Intermating of multiple founder lines is increasingly becoming a popular breeding method of choice for enhancing genetic diversity, developing mapping populations and breaking negative linkage drags. Low genetic diversity in the breeding populations is a major bottleneck for genetic improvement in jute (*Corchorus olitorius* L.). Here, we report the development of the first multiparent advanced generation inter-cross (MAGIC) population of jute. This MAGIC panel comprising 341 recombinant inbred lines was created by intermating 20 geographically isolated founder lines for four generations followed by six generations of inbreeding. We investigated the extent of phenotypic variability in four economically important traits using this permanent panel across two diverse locations. Significant trait variability, high broad-sense heritability (0.76–0.83) and transgressive segregation (up to 17.60%) revealed the potential of this MAGIC panel for genetic mapping and breeding for economic traits. Moreover, robust correlations of plant height and green biomass with bast fibre yield ($r = 0.67–0.72$) reinforced the importance of these two traits in selection for fibre yield in jute. While the major variance components were attributed to genotypic differences, significant non-heritable blockvariance for all the four traits indicated rather high influence of micro-environments on jute. Taken together, our study envisages the utility of the MAGIC population in jute and analogous crops constrained with a narrow gene pool.

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References

1. Bandillo N, Raghavan C, Muyco PA, Sevilla MAL, Lobina IT et al (2013) Multi-parent advanced generation inter-cross (MAGIC) populations in rice: progress and potential for genetics research and breeding. *Rice* 6:11
2. Basak SL, Chaudhuri BB (1966) Genetic variation and phenotypic plasticity of roots in two cultivated species of jute (*Corchorus olitorius* L. and *C. capsularis* L.). *Biol Plant* 9:292–300
3. Benor S, Demissew S, Hammer K, Blattner FR (2012) Genetic diversity and relationships in *Corchorus olitorius* (Malvaceae s. l.) inferred from molecular and morphological data. *Genet Resour Crop Evol* 59:1125–1146
4. Bernardo R (2020) Reinventing quantitative genetics for plant breeding: something old, something new, something borrowed, something BLUE. *Heredity* 125:375–385
5. Casler MD (2015) Fundamentals of experimental design: guidelines for designing successful experiments. *Agron J* 107:692–705
6. Dube SP, Marais D, Mavengahama S, Jaarsveld CMV, Gerrano AS (2018) Characterisation of agro-morphological traits of *Corchorus* accessions. *Acta Agric Scand Sec B Soil Plant Sci* 69:126–134
7. Fry JD (1992) The mixed-model analysis of variance applied to quantitative genetics: biological meaning of the parameters. *Evolution* 46:540–550
8. Gardner KA, Wittern LM, Mackay IJ (2016) A highly recombined, high-density, eight-founder wheat MAGIC map reveals extensive segregation

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9. González-Barrios P, Díaz-García L, Gutiérrez L (2018) Mega-environmental design: using genotype environment interaction to optimize resources for cultivar testing. *Crop Sci* 59:1899–1915

10. Hammer Ø, Harper DAT, Ryan PD (2001) Past: paleontological statistics software package for education and data analysis. *Palaeontol Electron* 4:1–9

11. Huang BE, Verbyla KL, Verbyla AP, Raghavan C, Singh VK et al (2015) MAGIC populations in crops: current status and future prospects. *Theor Appl Genet* 128:999–1017

12. Jiménez-Galindo JC, Malvar RA, Butrón A et al (2019) Mapping of resistance to corn borers in a MAGIC population of maize. *BMC Plant Biol* 19:431

13. Kover PX, Valdar W, Trakalo J (2009) A multiparent advanced generation inter-cross to fine-map quantitative traits in *Arabidopsis thaliana*. *PLoS Genet* 5:7

14. Kundu A, Chakraborty A, Mandal NA, Das D, Karmakar PG, Singh NK, Sarkar D (2015) A restriction-site-associated DNA (RAD) linkage map, comparative genomics and identification of QTL for histological fibre content coincident with those for retted bast fibre yield and its major components in jute (*Corchorus olitorius* L., Malvaceae s. l.). *Mol Breed* 35:19

15. Li Y, Ruperao P, Batley J, Edwards D, Khan T, Colmer TD, Pang J, Siddique KHM, Sutton T (2018) Investigating drought tolerance in chickpea using genome-wide association mapping and genomic selection based on whole-genome resequencing data. *Front Plant Sci* 9:190

16. Mathew B, Léon J, Sannemann W, Sillanpää MJ (2018) Detection of epistasis for flowering time using Bayesian multilocus estimation in a barley MAGIC population. *Genetics* 208:525–536

PDF

Help

17. Morrison GD, Linder CR (2014) Association mapping of germination traits in *Arabidopsis thaliana* under light and nutrient treatments: searching for G × E effects. *G3* 4:1465–1478

18. Nyadanu D, Amoah RA, Kwarteng AO, Akromah R, Aboagye LM, Adu-Dapaah H, Dansi A, Lotsu F, Tsama A (2017) Domestication of jute mallow (*Corchorus olitorius* L.): ethnobotany, production constraints and phenomics of local cultivars in Ghana. *Genet Resour Crop Evol* 64:1313–1329

19. Ongom PO, Ejeta G (2019) Mating design and genetic structure of a multi-parent advanced generation intercross (MAGIC) population of sorghum (*Sorghum bicolor* (L.) Moench). *G3* 8:331–341

20. Palit P, Sasmal BC, Bhattacharyya AC (1996) Germplasm diversity and estimate of genetic advance of four morpho-physiological traits in a world collection of jute. *Euphytica* 90:49–58

21. Pascual L, Desplat N, Huang BE, Desgroux A, Bruguier L et al (2015) Potential of a tomato MAGIC population to decipher the genetic control of quantitative traits and detect causal variants in the resequencing era. *Plant Biotechnol J* 13:565–577

22. Piepho HP, Möehring J (2007) Computing heritability and selection response from unbalanced plant breeding trials. *Genetics* 177:1881–1888

23. R Core Team (2017) R: a language and environment for statistical computing. R Foundation for Statistical Computing, Vienna

24. Sarkar D, Kundu A, Das D, Chakraborty A et al (2019) Resolving population structure and genetic differentiation associated with RAD-SNP loci under selection in tossa jute (*Corchorus olitorius* L.). *Mol Genet Genom* 294:479–492

25. Sarkar D, Satya P, Mandal NA, Das D, Karmakar PG, Singh NK (2016) Jute genomics: emerging resources and tools for molecular breeding. In: Ramawat KG, Ahuja MR (eds) *Fiber plants. Biology, biotechnology and applications*. Springer International Publishing AG, Cham, pp 155–200

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26. Sarker A, Singh M (2015) Improving breeding efficiency through application of appropriate experimental designs and analysis models: a case of lentil (*Lens culinaris* Medikus subsp. *culinaris*) yield trials. *Field Crops Res* 179:26–34

27. Satya P, Banerjee R, Biswas C, Karan M, Ghosh S, Ali N (2014) Genetic analysis of population structure using peroxidase gene and phenylalanine ammonia-lyase gene-based DNA markers: a case study in jute (*Corchorus* spp.). *Crop Sci* 54:1609–1620

28. Satya P, Maiti RK (2013) Bast and leaf fibre crops: kenaf, hemp, jute, *Agave*, etc. In: Singh BP (ed) *Biofuel crops: production, physiology and genetics*. CABI, UK, pp 292–311

29. Singh SR, Kundu DK, Tripathi MK, Dey P, Saha AR, Kumar M, Singh I, Mahapatra BS (2015) Impact of balanced fertilization on nutrient acquisition, fibre yield of jute and soil quality in New Gangetic alluvial soils of India. *Appl Soil Ecol* 92:24–34

30. Smith AB, Cullis BR, Thompson R (2005) The analysis of crop cultivar breeding and evaluation trials: an overview of current mixed models approach. *J Agric Sci* 143:449–462

31. Stadlmeier M, Hartl L, Mohler V (2018) Usefulness of a multiparent advanced generation intercross population with a greatly reduced mating design for genetic studies in winter wheat. *Front Plant Sci* 9:1825

32. Valdar W, Flint J, Mott R (2006) Simulating the collaborative cross: power of QTL detection and mapping resolution in large sets of recombinant inbred strains of mice. *Genetics* 172:1783–1797

33. Verbyla AP, Cavanagh CR, Verbyla KL (2014) Whole-genome analysis of multi-environment or multitrait QTL in MAGIC. *G3* 4:1569–1584

34. Wei J, Xu S (2016) A random-model approach to QTL mapping in multiparent advanced generation intercross (MAGIC) populations. *Genetics* 202:471–486

35. Wrońska-Pilarek D, Wiatrowska B, Bocianowski J (2019) Pollen morphology and variability of invasive *Spiraea tomentosa* L. (Rosaceae) from

PDF

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-
36. Yamamoto E, Iwata H, Tanabata T, Mizobuchi R, Yonemaru J et al (2014) Effect of advanced intercrossing on genome structure and on the power to detect linked quantitative trait loci in a multiparent population: a simulation study in rice. *BMC Genet* 15:50
-
37. Zhang L, Ibrahim AK, Niyitanga S, Zhang L, Qi J (2019) Jute (*Corchorus* spp.) breeding. In: Al-Khayri JM, Jain SM, Johnson DV (eds) *Advances in plant breeding strategies: industrial and food crops*, vol 6. Springer International Publishing AG, Cham, pp 86–113

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DS and NKS contributed to conceptualization and supervision; AK, DD, and NAM contributed to methodology; DS, PS, and CSK contributed to data generation and analysis; PS contributed to writing—original draft preparation; DS, JM, GK, and NKS contributed to writing—review and editing; DS and NKS contributed to funding acquisition.

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Ethics declarations

Conflict of interest

The authors disclose that they have no conflict of interest.

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