

Impact of pedigree depth in the Australian strawberry breeding program: variance component estimation, prediction accuracies and estimation of clonal and breeding values

Dr. Ky Mathews PhD^{1*}, Dr. Katie O'Connor PhD¹ and Dr. Jodi Neal PhD¹

^{1*}Department of Primary Industries, Queensland Government, Nambour, Australia, 4560, Queensland, Maroochy Research Facility, 47 Mayers Road.

*Corresponding author(s). E-mail(s): ky.mathews@daf.qld.gov.au;
Contributing authors: katie.oconnor@daf.qld.gov.au;
Jodi.Neal@daf.qld.gov.au;

Abstract

Key message: Less than 30 words summarising the main achievement.
Abstract should be around 200 words.

Keywords: A matrix, relatedness, yield

1 Introduction

Ways of finding the best depth. What other people have found.

Atkin 2009 and Yang 2016

Various papers analysing strawberry data have included relationship matrices constructed of somewhat arbitrary pedigree generation depths, ranging from four to 20 ([Whitaker et al, 2012](#); [Kennedy et al, 2014](#); [Osorio et al, 2014](#); [Paynter et al, 2014](#)).

Repeated measures or longitudinal analysis of yield data does not appear to be standard practice in strawberry, with research groups instead summing individual harvests to obtain a total season yield (e.g. [Cockerton et al \(2021\)](#); [Gezan et al \(2017\)](#);

Fan et al (2024); Osorio et al (2021)). Similarly for fruit weight, a common method is to divide total (marketable) yield by the total number of (marketable) fruit across a season (e.g. Sleper et al (2025); Osorio et al (2021); Whitaker et al (2012)), or average across all harvests to obtain a mean weight (e.g. Prohaska et al (2024)).

Repeated measures analyses have been used in other horticultural crops to model traits measured temporally, for example in macadamia (Hardner and De Faveri, ???), sweet cherry (Piaskowski et al, 2018) and... *Ky to insert her favourite examples (though perhaps we need to reword "horticultural")*.

Yield and fruit weight are two of the most important traits in driving profitability in strawberry (Herrington et al, 2012). Increases in yield lead to an increase in gross margin (Herrington et al, 2012). In the subtropical region, increased early season (in April and May) production is a current aim of the ASBP due to the high market punnet prices compared to the rest of the season (Herrington et al, 2012). Large fruit sizes lead to substantial increases in gross margin, as the time to pick and pack a fruit does not change with size, but larger fruit will result in more punnets and thus be more profitable for growers compared to fewer punnets of small fruit (Herrington et al, 2012). Fruit weight tends to decrease as temperatures increase (Menzel, 2021), and so average fruit weight in August in the subtropical region and in December to February in the temperate region are particularly important factors in ASBP selection decisions. It is for these reasons that the ASBP evaluates total season yield and average fruit weight as well as for individual weeks and months.

2 Materials and methods

While early studies of inheritance in octoploid strawberry found meiotic behavior to be a mix between disomic and polysomic (e.g. Lerceteanu-Köhler et al, 2003), more recent studies have found that inheritance is disomic (Rousseau-Gueutin et al, 2008; Tennessen et al, 2014; Sargent et al, 2016; Hardigan et al, 2020) and that analyses employing diploid Mendelian genetics are appropriate (Pincot et al, 2021). Therefore, we used the option `ploidy = 2` when constructing the A matrix in AGHmatrix.

If we do explore an octoploid A matrix and find that it is a better fit than diploid, then we should explore different levels of double reduction (the probability that two sister chromatids pass into the same gamete), as per Amadeu et al (2016).

Recent studies have found that a small amount of genetic variance could be attributed to dominance (Feldmann et al, 2024; Sleper et al, 2025). Thus, we also constructed a dominance relationship matrix in AGHmatrix using the option `dominance = TRUE`.

3 Results

Figures and tables are labeled with a prefix (fig or tab, respectively) plus the chunk label. Other environments such as equation and align can be labelled via the `\label{#label}` command inside or just below the `\caption{}` command. You can then use the label for cross-reference. As an example, consider the chunk label declared for Figure ?? which is fig1. To cross-reference it, use the command `\ref{fig:fig1}`, for which it comes up as "Figure ??".

To reference line numbers in an algorithm, consider the label declared for the line number 2 of Algorithm ?? is `\label{algl n2}`. To cross-reference it, use the command `\ref{algl n2}` for which it comes up as line ?? of Algorithm ??.

4 Discussion

5 Conclusions

We found. . .

Supplementary information. If your article has accompanying supplementary file/s please state so here.

Acknowledgments. The authors thank present and past members of the Australian Strawberry Breeding Program and staff at the Maroochy Research Facility for the involvement in field preparation and maintenance, plant propagation, plant maintenance, fruit harvesting and processing, data collection, etc etc.

Declarations

5.1 Funding

The Australian Strawberry Breeding Program has been funded by Hort Innovation using the strawberry research and development levy, with co-contributions from the Queensland Government through its Department of Primary Industries and funds from the Australian Government.

5.2 Competing Interests

The authors have no relevant financial or non-financial interests to disclose.

5.3 Authors' Contributions

KM and KO developed the idea. KM wrote analytical R code. etc etc.

5.4 Data Availability

The datasets generated during and/or analysed during the current study are available from the corresponding author on reasonable request.

References

Amadeu RR, Cellon C, Olmstead JW, et al (2016) Aghmatrix: R package to construct relationship matrices for autotetraploid and diploid species: a blueberry example. The plant genome 9(3):plantgenome2016.01.0009

- Cockerton HM, Karlström A, Johnson AW, et al (2021) Genomic informed breeding strategies for strawberry yield and fruit quality traits. *Frontiers in Plant Science* 12. <https://doi.org/10.3389/fpls.2021.724847>, URL <https://www.frontiersin.org/article/10.3389/fpls.2021.724847>
- Fan Z, Verma S, Lee H, et al (2024) Strawberry soluble solids qtl with inverse effects on yield. *Horticulture Research* 11(2). <https://doi.org/10.1093/hr/uhad271>, URL <https://www.scopus.com/inward/record.uri?eid=2-s2.0-85184920141&doi=10.1093%2fhr%2fuhad271&partnerID=40&md5=2076163ed5255e8c7c98638ed639a4fd>, export Date: 03 March 2024; Cited By: 0
- Feldmann MJ, Pincot DDA, Seymour DK, et al (2024) A dominance hypothesis argument for historical genetic gains and the fixation of heterosis in octoploid strawberry. *GENETICS* <https://doi.org/10.1093/genetics/iyae159>, URL <http://dx.doi.org/10.1093/genetics/iyae159>
- Gezan SA, Osorio LF, Verma S, et al (2017) An experimental validation of genomic selection in octoploid strawberry. *Horticulture Research* 4(Article 16070)
- Hardigan MA, Feldmann MJ, Lorant A, et al (2020) Genome synteny has been conserved among the octoploid progenitors of cultivated strawberry over millions of years of evolution. *Frontiers in Plant Science* 10. <https://doi.org/10.3389/fpls.2019.01789>, URL <http://dx.doi.org/10.3389/fpls.2019.01789>
- Hardner C, De Faveri J (????) Flexible linear mixed models for complex data in horticultural tree breeding. In: *Acta Horticulturae*, pp 139–146, <https://doi.org/10.17660/ActaHortic.2023.1362.19>, URL <https://www.scopus.com/inward/record.uri?eid=2-s2.0-85153536804&doi=10.17660%2fActaHortic.2023.1362.19&partnerID=40&md5=cf4c5e2580c5a46a4e1130ae1643fc76>, export Date: 23 May 2023; Cited By: 0
- Herrington ME, Wegener M, Hardner C, et al (2012) Influence of plant traits on production costs and profitability of strawberry in southeast queensland. *Agricultural Systems* 106(1):23–32. <https://doi.org/http://dx.doi.org/10.1016/j.agsy.2011.11.003>, URL <http://www.sciencedirect.com/science/article/pii/S0308521X11001636>
- Kennedy C, Osorio LF, Peres NA, et al (2014) Additive genetic effects for resistance to foliar powdery mildew in strawberry revealed through divergent selection. *Journal of the American Society for Horticultural Science* 139(3):310–316. URL <http://journal.ashspublications.org/content/139/3/310.abstract>
- Lerceteau-Köhler E, Guérin G, Laigret F, et al (2003) Characterization of mixed disomic and polysomic inheritance in the octoploid strawberry (*fragaria* × *ananassa*) using aflp mapping. *Theoretical and Applied Genetics* 107(4):619–628. <https://doi.org/10.1007/s00122-003-1300-6>, URL <http://dx.doi.org/10.1007/s00122-003-1300-6>

- Menzel C (2021) Higher temperatures decrease fruit size in strawberry growing in the subtropics. *Horticulturae* 7(2):34. URL <https://www.mdpi.com/2311-7524/7/2/34>
- Osorio LF, Pattison JA, Peres NA, et al (2014) Genetic variation and gains in resistance of strawberry to *colletotrichum gloeosporioides*. *Phytopathology* 104(1):67–74. <https://doi.org/10.1094/phyto-02-13-0032-r>, URL <http://dx.doi.org/10.1094/PHYTO-02-13-0032-R>
- Osorio LF, Gezan SA, Verma S, et al (2021) Independent validation of genomic prediction in strawberry over multiple cycles. *Frontiers in Genetics* 11(1862). <https://doi.org/10.3389/fgene.2020.596258>, URL <https://www.frontiersin.org/article/10.3389/fgene.2020.596258>
- Paynter ML, De Faveri J, Herrington ME (2014) Resistance to *fusarium oxysporum* f. sp. *fragariae* and predicted breeding values in strawberry. *Journal of the American Society for Horticultural Science* 139(2):178–184. URL <http://journal.ashspublications.org/content/139/2/178.abstract>
- Piaskowski J, Hardner C, Cai L, et al (2018) Genomic heritability estimates in sweet cherry reveal non-additive genetic variance is relevant for industry-prioritized traits. *BMC Genetics* 19(1):23–38
- Pincot DD, Ledda M, Feldmann MJ, et al (2021) Social network analysis of the genealogy of strawberry: retracing the wild roots of heirloom and modern cultivars. *G3* 11(3):jkab015
- Prohaska A, Rey-Serra P, Petit J, et al (2024) Exploration of a european-centered strawberry diversity panel provides markers and candidate genes for the control of fruit quality traits. *Horticulture Research* 11(7). <https://doi.org/10.1093/hr/uhae137>, URL <https://www.scopus.com/inward/record.uri?eid=2-s2.0-85198634207&doi=10.1093%2fhr%2fuhae137&partnerID=40&md5=6e9bd4b81a40532c5f90fee4eb696973>, export Date: 05 August 2024; Cited By: 0
- Rousseau-Gueutin M, Lerceteau-Köhler E, Barrot L, et al (2008) Comparative genetic mapping between octoploid and diploid *fragaria* species reveals a high level of colinearity between their genomes and the essentially disomic behavior of the cultivated octoploid strawberry. *Genetics* 179(4):2045–2060. <https://doi.org/10.1534/genetics.107.083840>, URL <http://dx.doi.org/10.1534/genetics.107.083840>
- Sargent D, Yang Y, Šurbanovski N, et al (2016) Haplo-snp affinities and linkage map positions illuminate subgenome composition in the octoploid, cultivated strawberry (*fragaria* × *ananassa*). *Plant Science* 242:140–150. <https://doi.org/10.1016/j.plantsci.2015.07.004>, URL <http://dx.doi.org/10.1016/j.plantsci.2015.07.004>
- Sleper J, Tapia R, Lee S, et al (2025) Within-family genomic selection in strawberry: Optimization of marker density, trial design, and training set composition. *The*

Plant Genome 18(1). <https://doi.org/10.1002/tpg2.20550>, URL <http://dx.doi.org/10.1002/tpg2.20550>

Tennessen JA, Govindarajulu R, Ashman TL, et al (2014) Evolutionary origins and dynamics of octoploid strawberry subgenomes revealed by dense targeted capture linkage maps. *Genome Biology and Evolution* 6(12):3295–3313. <https://doi.org/10.1093/gbe/evu261>, URL <http://dx.doi.org/10.1093/gbe/evu261>

Whitaker VM, Osorio LF, Hasing T, et al (2012) Estimation of genetic parameters for 12 fruit and vegetative traits in the university of florida strawberry breeding population. *Journal of the American Society for Horticultural Science* 137(5):316–324. <https://doi.org/10.21273/jashs.137.5.316>, URL <http://dx.doi.org/10.21273/JASHS.137.5.316>