

The Genetics of LMA in North American Spring Wheat (*Triticum aestivum* L.)

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Supplemental Tables 2, 3, and 4

**Supplemental Table S2.** Correlations between untreated samples (n = 251)

| A        | GH2018.1                                   | GH2018.2                                   | GH2018.3                                   |
|----------|--|--|--|
| GH2018.1 |  | $r^2 = 0.469$<br>$p < 2.2 \times 10^{-16}$ | $r^2 = 0.176$<br>$p < 1.9 \times 10^{-11}$ |
| GH2018.2 | $r^2 = 0.469$<br>$p < 2.2 \times 10^{-16}$ |  | $r^2 = 0.358$<br>$p < 2.2 \times 10^{-16}$ |
| GH2018.3 | $r^2 = 0.176$<br>$p < 1.9 \times 10^{-11}$ | $r^2 = 0.358$<br>$p < 2.2 \times 10^{-16}$ |  |

| B        | GH2019.1                                   | GH2019.2                                   | GH2019.3                                   |
|----------|--|--|--|
| GH2019.1 |  | $r^2 = 0.634$<br>$p < 2.2 \times 10^{-16}$ | $r^2 = 0.504$<br>$p < 2.2 \times 10^{-16}$ |
| GH2019.2 | $r^2 = 0.634$<br>$p < 2.2 \times 10^{-16}$ |  | $r^2 = 0.518$<br>$p < 2.2 \times 10^{-16}$ |
| GH2019.3 | $r^2 = 0.504$<br>$p < 2.2 \times 10^{-16}$ | $r^2 = 0.518$<br>$p < 2.2 \times 10^{-16}$ |  |

**Supplemental Table S3.** MLM examining sources of variation, including dpa as a fixed effect and genotype, run, and expt as random effects.

| Random Effects |                        |          |
|----------------|------------------------|----------|
| Groups         | Variance               | Std. Dev |
| Genotype       | $5.263 \times 10^{-2}$ | 0.229404 |
| Run            | $2.008 \times 10^{-2}$ | 0.141710 |
| Expt           | $2.268 \times 10^{-5}$ | 0.004762 |
| Residual       | $8.133 \times 10^{-2}$ | 0.285178 |

  

| Fixed Effects |           |            |            |        |           |
|---------------|-----------|------------|------------|--------|-----------|
|               | Estimates | Std. Error | df         | t      | Pr (> t ) |
| mean          | 0.580098  | 0.213471   | 136.273426 | 2.717  | 0.00743   |
| dpa           | -0.006257 | 0.008881   | 381.471685 | -0.705 | 0.48152   |

dpa = days past anthesis  
expt = experiment  
Std. Dev = standard deviation  
Std. Error = standard error  
df = degrees of freedom

**Supplemental Table S4.** REML (restricted maximum likelihood) output from a MLM when including genotype as a random effect and experiment (expt) as a fixed effect.

| Random Effects |          |          |  |  |  |
|----------------|----------|----------|--|--|--|
| Groups         | Variance | Std. Dev |  |  |  |
| Genotype       | 0.04049  | 0.2012   |  |  |  |
| Residual       | 0.11970  | 0.3460   |  |  |  |

  

| Fixed Effects |           |            |     |       |                     |
|---------------|-----------|------------|-----|-------|---------------------|
|               | Estimates | Std. Error | df  | t     | Pr (> t )           |
| mean          | 0.46079   | 0.02501    | 473 | 18.42 | $2 \times 10^{-16}$ |
| expt          | -0.04195  | 0.03085    | 253 | -1.36 | 0.175               |

Std. Dev = standard deviation

Std. Error = standard error

df = degrees of freedom