

Single trial analysis for

1 General information

| | |
|--------------------|-------------------|
| Analysis done on | 18-03-26 14:41:25 |
| Pipeline version | 0.0.0.9000 |
| Mixed model engine | SpATS |

2 Description of the fitted model

| | |
|------------------------------|---|
| Trait | yield |
| Experimental design features | replicate (F) + row(replicate) (R) + col(replicate) (R) |
| Spatial model | 2 dimensional P-splines |
| Genotype | fixed |

3 Summary of the results

Table 1: Size of the data set and eight number descriptives

| | |
|--------------------------|---------|
| Number of observations | 338.00 |
| Number of missing values | 0.00 |
| Mean | 184.16 |
| Median | 179.81 |
| Min | 54.58 |
| Max | 341.77 |
| Lower quartile | 148.56 |
| Upper quartile | 220.50 |
| Variance | 2569.23 |

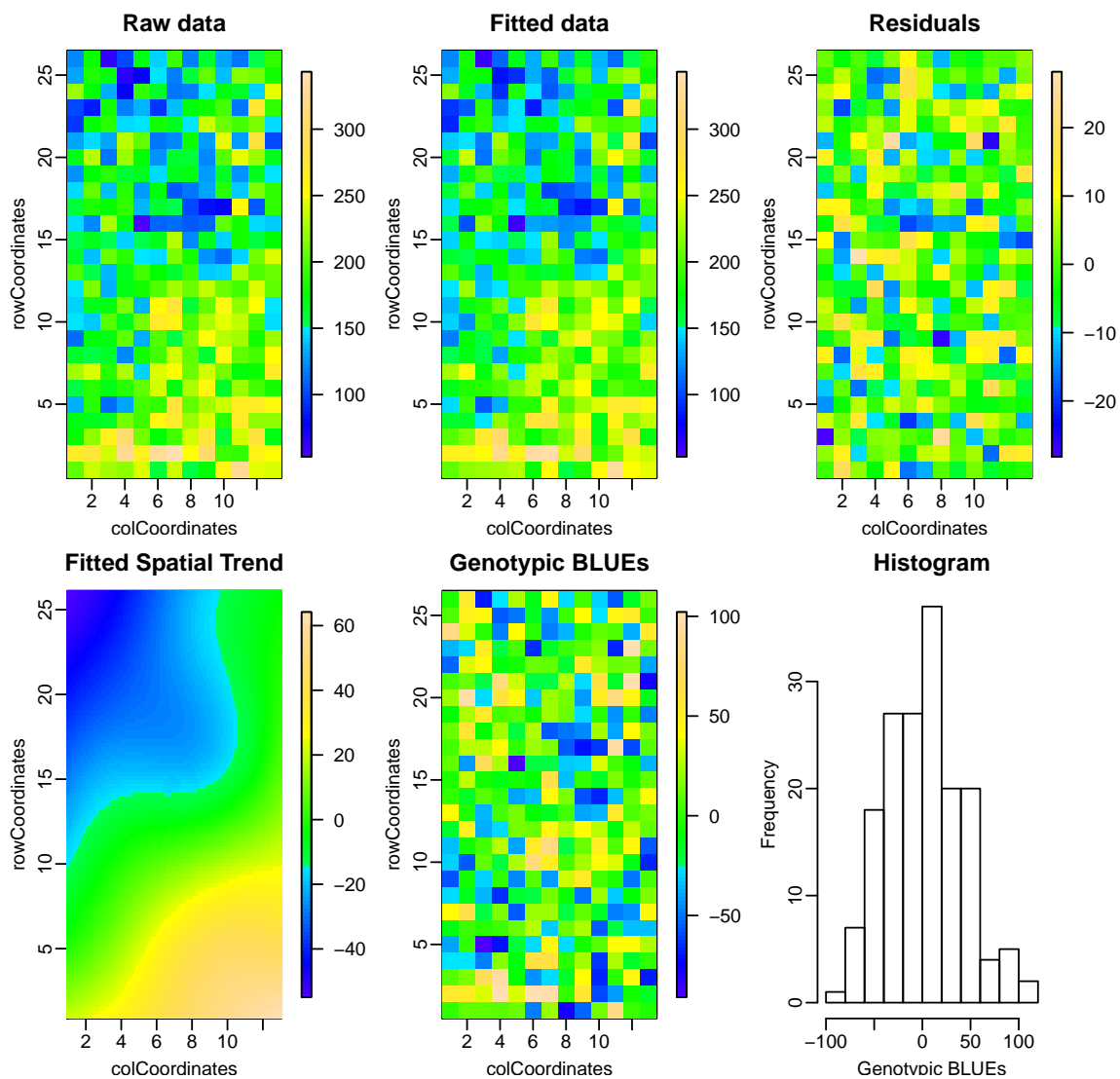
Table 2: Effective dimensions

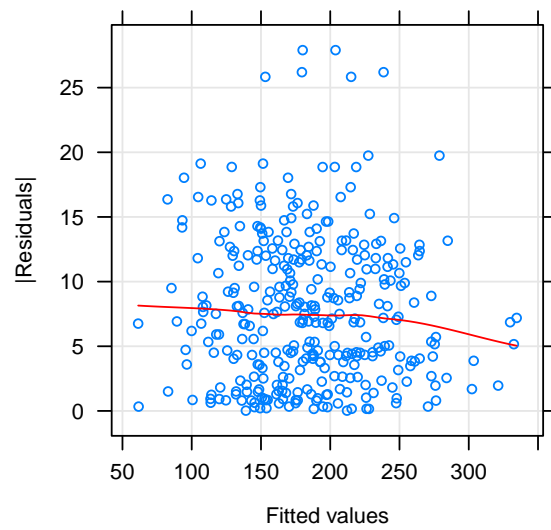
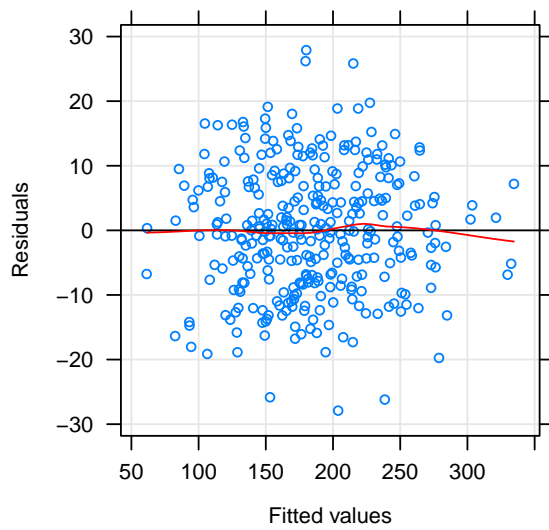
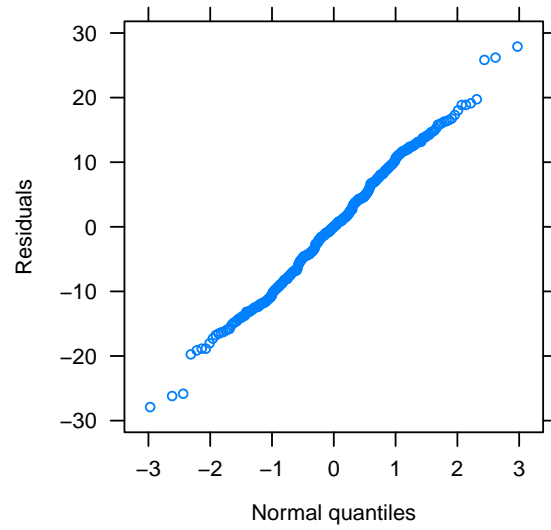
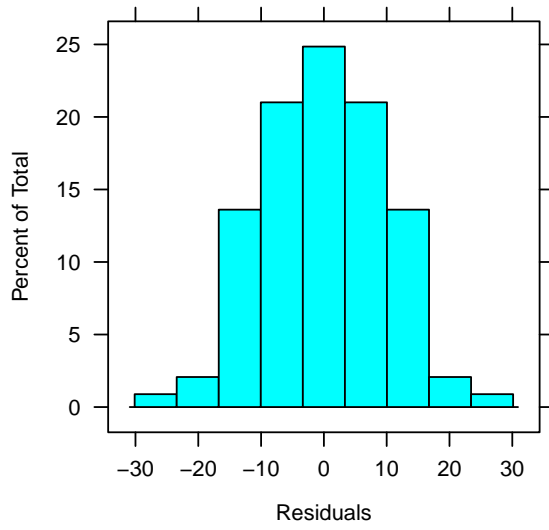
| | Effective | Model | Nominal | Ratio | Type |
|---|-----------|-------|---------|-------|------|
| Genotype | 168.0 | 168 | 168 | 1.00 | F |
| Intercept | 1.0 | 1 | 1 | 1.00 | F |
| Replicate | 1.0 | 1 | 1 | 1.00 | F |
| Row(replicate) | 10.5 | 52 | 51 | 0.21 | R |
| Col(replicate) | 0.0 | 26 | 25 | 0.00 | R |
| Linear trend along cols | 1.0 | 1 | 1 | 1.00 | S |
| Linear trend along rows | 1.0 | 1 | 1 | 1.00 | S |
| Linear trend along rows and cols | 1.0 | 1 | 1 | 1.00 | S |
| Smooth trend along cols | 0.0 | 8 | 8 | 0.00 | S |
| Smooth trend along rows | 1.6 | 14 | 14 | 0.12 | S |
| Linear trend in rows changing smoothly along cols | 0.0 | 8 | 8 | 0.00 | S |
| Linear trend in cols changing smoothly along rows | 0.1 | 14 | 14 | 0.00 | S |
| Smooth-by-smooth interaction trend over rows and cols | 5.7 | 28 | 28 | 0.20 | S |
| Total | 190.8 | 323 | 321 | 0.59 | |
| Residual | 147.2 | | | | |
| Number of observations | 338 | | | | |

Table 3: Variance components

| | Variance |
|---|----------|
| Row(replicate) | 34.27 |
| Col(replicate) | 0.00 |
| Smooth trend along cols | 0.00 |
| Smooth trend along rows | 78.89 |
| Linear trend in rows changing smoothly along cols | 0.00 |
| Linear trend in cols changing smoothly along rows | 0.59 |
| Smooth-by-smooth interaction trend over rows and cols | 1732.00 |
| Residual | 213.80 |

4 Diagnostic plots





5 List of best 10% genotypes with highest BLUEs

| genotype | yield | SE |
|----------|--------|-------|
| SB023 | 288.69 | 11.81 |
| SB025 | 287.90 | 11.75 |
| SB022 | 285.49 | 11.86 |
| SB032 | 282.39 | 11.92 |
| SB011 | 274.11 | 12.03 |
| SB156 | 270.44 | 11.78 |
| SB144 | 268.80 | 11.89 |
| SB031 | 256.97 | 11.99 |
| SB093 | 251.19 | 12.26 |
| SB055 | 250.36 | 11.99 |
| SB054 | 250.21 | 12.05 |
| SB020 | 244.95 | 11.87 |
| SB028 | 242.94 | 12.12 |
| SB194 | 241.81 | 11.57 |
| SB128 | 241.31 | 11.97 |
| SB155 | 239.79 | 11.76 |
| SB027 | 239.54 | 11.88 |

