Using asremlPlus, in conjunction with asreml, to do a linear mixed model analysis of a wheat experiment using hypothesis tests

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This vignette shows how to use asremlPlus (Brien, 2024), in conjunction with asreml (Butler et al., 2020), to employ hypothesis tests to select the terms to be included in a mixed model for an experiment that involves spatial variation. It also illustrates diagnostic checking and prediction production and presentation for this experiment. Here, asremlPlus and asreml are packages for the R Statistical Computing environment (R Core Team, 2024).

It is divided into the following main sections:

- 1. Set up the maximal model for this experiment
- 2. Perform a series of hypothesis tests to select a linear mixed model for the data
- 3. Diagnostic checking using residual plots and variofaces
- 4. Prediction production and presentation

1. Set up the maximal model for this experiment

```
library(knitr)
opts_chunk$set("tidy" = FALSE, comment = NA)
suppressMessages(library(asreml, quietly=TRUE))

## Offline License checked out Thu Apr 11 09:24:19 2024

packageVersion("asreml")

## [1] '4.2.0.312'
suppressMessages(library(asremlPlus))
packageVersion("asremlPlus")

## [1] '4.4.32'
suppressMessages(library(qqplotr, quietly=TRUE))
options(width = 100)
```

Get data available in asremlPlus

The data are from a 1976 spring wheat experiment and are taken from Gilmour et al. (1995). An analysis is presented in the asrem1 manual by Butler et al. (2020, Section 7.6), although they suggest that it is a barley experiment.

```
data(Wheat.dat)
```

Fit the maximal model

In the following a model is fitted that has the terms that would be included for a balanced lattice. In addition, a term WithinColPairs has been included to allow for extraneous variation arising between pairs of adjacent lanes. Also, separable ar1 residual autocorrelation has been included. This model represents the maximal anticipated model,

```
ASReml Version 4.2 11/04/2024 09:24:20
          LogLik
                        Sigma2
                                   DF
                                          wall
       -724.1213
                      23034.14
                                  124
                                        09:24:20
1
 2
      -717.4149
                                        09:24:20
                      9206.931
                                  124
                                                  ( 2 restrained)
 3
      -694.8752
                      26492.99
                                  124
                                        09:24:20
                                                  ( 2 restrained)
      -694.1600
 4
                      33101.80
                                  124
                                        09:24:20
                                                  ( 1 restrained)
 5
      -692.0020
                      36912.26
                                  124
                                        09:24:20
                                                  ( 1 restrained)
 6
      -691.7892
                                  124
                                        09:24:20
                                                  ( 2 restrained)
                      46701.51
 7
      -691.8336
                      46208.51
                                  124
                                        09:24:20
                                                  ( 1 restrained)
 8
       -691.7749
                                        09:24:20
                      47698.26
                                  124
       -691.7711
                      47041.85
                                  124
                                        09:24:20
```

Warning in asreml(yield ~ WithinColPairs + Variety, random = ~Rep/(Row + : Some components changed by more than 1% on the last iteration

The warning from asreml is probably due to a bound term.

Initialize a testing sequence by loading the current fit into an asrtests object

A label and the information criteria based on the full likelihood (Verbyla, 2019) are included in the test.summary stored in the asrtests object.

Warning in infoCriteria.asreml(asreml.obj, IClikelihood = ic.lik, bound.exclusions = bound.exclusions):
Rep

Warning in asreml(fixed = yield ~ WithinColPairs + Variety, random = ~Rep/(Row + : Log-likelihood not converged

Check for and remove any boundary terms

```
current.asrt <- rmboundary(current.asrt, IClikelihood = "full")</pre>
Warning in infoCriteria.asreml(asreml.obj, IClikelihood = ic.lik): The following bound terms were disco
ASReml Version 4.2 11/04/2024 09:24:21
          LogLik
                        Sigma2
                                  DF
                                          wall
                      47071.42
       -691.7710
                                  124
                                        09:24:21
Warning in asreml(fixed = yield ~ WithinColPairs + Variety, random = ~Rep/(Row + : Log-likelihood
not converged
summary(current.asrt$asreml.obj)$varcomp
                                                   z.ratio bound %ch
                         component
                                      std.error
Rep:Row
                      4.293282e+03 3.199458e+03 1.3418779
                                                               P 0.0
Rep:Column
                      1.575689e+02 1.480357e+03 0.1064398
                                                               P 0.7
units
                      5.742689e+03 1.652457e+03 3.4752438
                                                               P 0.0
                      4.706787e+04 2.515832e+04 1.8708669
Row:Column!R
                                                               P 0.0
Row:Column!Row!cor
                      7.920301e-01 1.014691e-01 7.8056280
                                                               U 0.0
Row:Column!Column!cor 8.799559e-01 7.370402e-02 11.9390486
                                                               U 0.0
print(current.asrt, which = "testsummary")
```

Sequence of model investigations

(If a row has NA for p but not denDF, DF and denDF relate to fixed and variance parameter numbers)

```
terms DF denDF p AIC BIC action 1 Maximal model 26 6 NA 1646.129 1742.47 Starting model 2 Rep 1 NA NA 1646.129 1742.47 Boundary
```

Rep has been removed because it has been constrained to zero. Following the recommendation of Littel et al. (2006, p. 150), the bound on all variance components is set to unconstrained (U) using setvariances.asreml so as to avoid bias in the estimate of the residual variance. Alternatively, one could move Rep to the fixed model.

Unbind Rep, Row and Column components and reload into an asrtests object

```
ASReml Version 4.2 11/04/2024 09:24:21
                       Sigma2
                                 DF
                                        wall
         LogLik
                                124
                                     09:24:21
1
      -724.1213
                     23034.14
2
      -717.4149
                     9206.931 124 09:24:21 ( 2 restrained)
                              124 09:24:21 ( 2 restrained)
 3
      -694.8752
                     26492.99
 4
      -693.9744
                    33129.65 124 09:24:21 ( 1 restrained)
5
      -692.8856
                   39662.12 124 09:24:21
                  53103.83 124 09:24:21
6
      -691.4276
                  48092.17 124 09:24:21
7
      -691.2387
8
                  47278.94 124 09:24:21
      -691.1808
9
      -691.1710
                    46850.98 124 09:24:21
      -691.1700
                    46690.46 124 09:24:21
10
Warning in asreml(fixed = yield ~ WithinColPairs + Variety, random = ~Rep/(Row + : Some components
changed by more than 1\% on the last iteration
current.asrt <- as.asrtests(current.asr, wald.tab = NULL, test.summary = current.asrt$test.summary,</pre>
                          IClikelihood = "full", label = "Max model & Unbound components")
current.asrt <- rmboundary(current.asrt)</pre>
summary(current.asrt$asreml.obj)$varcomp
                         component
                                                  z.ratio bound %ch
                                     std.error
                    -2458.3485841 1.197491e+03 -2.0529167
                                                             U 0.0
Rep
                     5008.7151486 3.401335e+03 1.4725732
                                                             U 0.0
Rep:Row
Rep:Column
                      916.4641198 1.699576e+03 0.5392309
                                                             U 0.2
units
                     5959.0220817 1.609649e+03 3.7020634
                                                             P 0.0
Row: Column! R.
                     46637.6303429 2.724392e+04 1.7118545
                                                           P 0.0
Row:Column!Row!cor
                         0.8150590 1.000281e-01 8.1483012
                                                             U 0.0
Row:Column!Column!cor
                        0.8856824 7.492514e-02 11.8208968
                                                             U 0.0
print(current.asrt, which = "testsummary")
#### Sequence of model investigations
(If a row has NA for p but not denDF, DF and denDF relate to fixed and variance parameter numbers)
                         terms DF denDF p
                                                AIC
                                                        BIC
                                                                    action
                  Maximal model 26 6 NA 1646.129 1742.470 Starting model
1
                                     NA NA 1646.129 1742.470
                           Rep 1
3 Max model & Unbound components 26
                                     7 NA 1647.193 1746.544 Starting model
print(current.asrt, which = "pseudoanova")
```

Pseudo-anova table for fixed terms

Wald tests for fixed effects.

Response: yield

```
Df denDF F.inc Pr (Intercept) 1 1.7 153.500 0.0115 WithinColPairs 1 15.6 2.545 0.1307 Variety 24 76.1 10.110 0.0000
```

Now the Rep component estimate is negative.

The test.summary output has been extended, by supplying the previous test.summary to as.asrtests, to show that there is a new starting model. The pseudo-anova table shows that Varieties are highly significant (p < 0.001)

2. Perform a series of hypothesis tests to select a linear mixed model for the data

The hypothesis tests in this section are Wald tests for fixed terms, with denominator degrees of freedom calculated using the Kenward-Rogers adjustment (Kenward and Rogers (1997), and Restricted Maximum Likelihood Ratio Tests (REMLRT) for random terms.

Check the term for within Column pairs (a post hoc factor)

The information criteria based on the full likelihood (Verbyla, 2019) is also included in the test.summary stored in the asrtests object.

Warning in asreml(fixed = yield ~ Variety, random = ~Rep/(Row + Column) + : Some components changed by more than 1% on the last iteration

Warning in asreml(fixed = yield \sim Variety, random = \sim Rep/(Row + Column) + : Some components changed by more than 1% on the last iteration

```
print(current.asrt)
```

Summary of the fitted variance parameters

```
component
                                       std.error
                                                    z.ratio bound %ch
                      -2391.9489939 1.194581e+03 -2.0023338
                                                                U 0.4
Rep
Rep:Row
                       5035.5311054 3.406006e+03 1.4784269
                                                                U 0.3
Rep:Column
                        761.9535622 1.612103e+03 0.4726458
                                                                U 1.2
units
                       5933.2133794 1.610805e+03 3.6833848
                                                                P 0.1
Row:Column!R
                      45970.8383027 2.635124e+04 1.7445415
                                                                P 0.0
Row:Column!Row!cor
                          0.8101615 9.995498e-02 8.1052641
                                                                U 0.1
                          0.8846970 7.503039e-02 11.7911827
Row:Column!Column!cor
                                                                U 0.0
```

Pseudo-anova table for fixed terms

Wald tests for fixed effects. Response: yield

```
Df denDF F.inc Pr (Intercept) 1 1.7 158.90 0.0112 Variety 24 76.8 10.27 0.0000
```

Sequence of model investigations

(If a row has NA for p but not denDF, DF and denDF relate to fixed and variance parameter numbers)

```
terms DF denDF
                                                       AIC
                                                                BIC
                                                                             action
                                                р
                   Maximal model 26
                                       6.0
                                               NA 1646.129 1742.470 Starting model
1
2
                                       NA
                                               NA 1646.129 1742.470
                             Rep 1
3 Max model & Unbound components 26
                                      7.0
                                               NA 1647.193 1746.544 Starting model
                  WithinColPairs 1
                                     15.6 0.1307 1645.325 1741.666
                                                                           Dropped
```

It is clear in the call to testranfix that the model is being changed by dropping the withinColPairs term, which could also be achieved using update.asreml. However, an asremlPlus model-changing function operates on an asrtests object, that includes an asreml object, and, except for changeTerms.asrtests, results in an asrtests object that may contain the changed model or the supplied model depending on the results of hypothesis tests or comparisons of information criteria. In addition, the result of the test or comparison will be added to a test.summary data.frame stored in the new asrtests object and, if the model was changed, the wald.tab in the new asrtests object will have been updated for the new model.

In this case, as can be seen from the summary of current.asrt after the call, the *p*-value for the withinColPairs was greater than 0.05 and so now the model stored in current.asrt does not include withinColPAirs. The wald.tab has been updated for the new model.

Test the nugget term

The nugget term represents non-spatial variance, such as random plot and measurement error. It is fitted using the asreml reserved word units.

```
current.asrt <- testranfix(current.asrt, "units", positive=TRUE, IClikelihood = "full")</pre>
```

```
Warning in asreml(fixed = yield ~ Variety, random = ~Rep + Rep:Row + Rep:Column, : Some components changed by more than 1% on the last iteration
```

Warning in asreml(fixed = yield ~ Variety, random = ~Rep + Rep:Row + Rep:Column, : Some components changed by more than 1% on the last iteration

Test Row autocorrelation

We begin testing the autocorrelation by dropping the Row autocorrelation. Because of messages about the instability of the fit, iterate.asrtests is used to execute extra iterations of the fitting process.

Warning in asreml(fixed = yield ~ Variety, random = ~Rep/(Row + Column) + : Some components changed by more than 1% on the last iteration

Warning in asreml(fixed = yield \sim Variety, random = \sim Rep/(Row + Column) + : Some components changed by more than 1% on the last iteration

```
current.asrt <- iterate(current.asrt)</pre>
```

Test Column autocorrelation (depends on whether Row autocorrelation retained)

The function getTestPvalue is used to get the p-value for the Row autocorrelation test. If it is significant then the Column autocorrelation is tested by by dropping the Column autocorrelation, while retaining the Row autocorrelation. Otherwise the model with just Row autocorrelation, whose fit is returned via current.asrt after the test, is compared to one with no autocorrelation.

```
(p <- getTestPvalue(current.asrt, label = "Row autocorrelation"))</pre>
```

[1] 4.676754e-06

Warning in DFdiff(bound.h1, bound.h0, DF = DF, bound.exclusions = bound.exclusions): There were a total The following bound terms occur in only one of the models compared and so were discounted:

Row:Column!Row!cor

Output the results

```
print(current.asrt)
```

```
\mbox{\tt \#\#\#\#} Summary of the fitted variance parameters
```

```
component std.error z.ratio bound %ch
Rep -2385.8697551 1.211207e+03 -1.9698276 U 0.0
```

```
5027.7123253 3.415391e+03 1.4720753
Rep:Row
                                                               U 0.0
Rep:Column
                       753.5913536 1.609865e+03 0.4681086
                                                               U 0.6
units
                      5920.3547038 1.611274e+03 3.6743304
                                                               P 0.0
Row:Column!R
                     45870.0971595 2.623601e+04 1.7483638
                                                               P 0.0
Row:Column!Row!cor
                         0.8098786 1.001805e-01 8.0841906
                                                               U 0.0
Row:Column!Column!cor
                         0.8845768 7.510598e-02 11.7777144
                                                               U 0.0
#### Pseudo-anova table for fixed terms
Wald tests for fixed effects.
Response: yield
           Df denDF F.inc
(Intercept) 1 1.7 159.20 0.0111
Variety
           24 76.8 10.27 0.0000
     Sequence of model investigations
(If a row has NA for p but not denDF, DF and denDF relate to fixed and variance parameter numbers)
                          terms DF denDF
                                              р
                                                     AIC
                                                              BIC
                                                                          action
1
                  Maximal model 26
                                     6.0
                                             NA 1646.129 1742.470 Starting model
                            Rep 1
                                     NA
                                             NA 1646.129 1742.470
                                                                        Boundary
3 Max model & Unbound components 26
                                     7.0
                                             NA 1647.193 1746.544 Starting model
                 WithinColPairs 1 15.6 0.1307 1645.325 1741.666
                                                                         Dropped
4
5
                                      NA 0.0006 1645.325 1741.666
                                                                        Retained
                          units 1
            Row autocorrelation 1
                                      NA 0.0000 1645.325 1741.666
6
                                                                       Unswapped
                                      NA 0.0000 1645.318 1741.658
7
            Col autocorrelation 2
                                                                       Unswapped
printFormulae(current.asrt$asreml.obj)
#### Formulae from asreml object
fixed: yield ~ Variety
random: ~ Rep/(Row + Column) + units
residual: ~ ar1(Row):ar1(Column)
print(R2adj(current.asrt$asreml.obj, include.which.random = ~ .))
ASReml Version 4.2 11/04/2024 09:24:27
                       Sigma2
                                  DF
         LogLik
                                         wall
      -694.6149
                     45855.31
                                 125
                                       09:24:27
      -694.6149
                     45854.06
                                 125
                                       09:24:27
2
[1] 44.62413
attr(,"fixed")
<environment: 0x00000241ad70f548>
attr(,"random")
```

The test.summary shows is that the model with Row and without Column autocorrelation failed to converge. The asreml.obj in current.asrt contains the model selected by the selection process, which has been printed using printFormulae.asrtests. It is clear that no changes were made to the variance terms. The adjusted R^2 value shows that the fixed and random terms in the fitted model account for 45% of the total variation in the yield.

3. Diagnosing checking using residual plots and variofaces

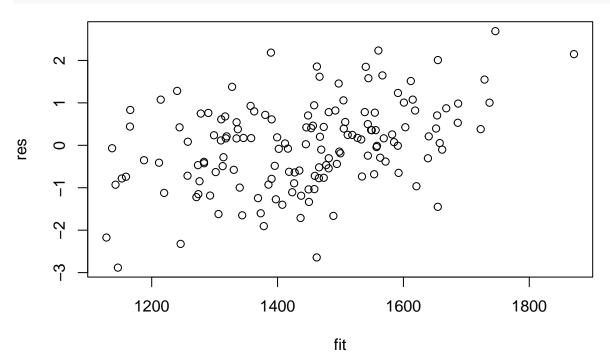
Get current fitted asreml object and update to include standardized residuals

```
current.asr <- current.asrt$asreml.obj</pre>
current.asr <- update(current.asr, aom=TRUE)</pre>
ASReml Version 4.2 11/04/2024 09:24:27
           LogLik
                          Sigma2
                                       DF
                                               wall
 1
       -694.6149
                        45855.31
                                      125
                                             09:24:27
 2
       -694.6149
                        45854.06
                                      125
                                             09:24:27
 3
       -694.6149
                        45851.09
                                      125
                                             09:24:27
Wheat.dat$res <- residuals(current.asr, type = "stdCond")</pre>
Wheat.dat$fit <- fitted(current.asr)</pre>
```

Do diagnostic checking

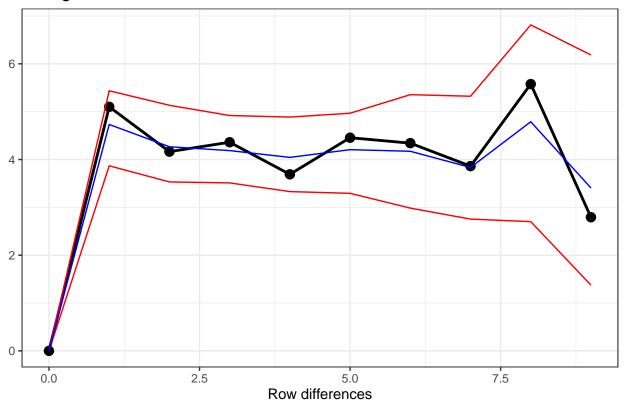
Do residuals-versus-fitted values plot

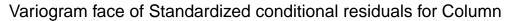
```
with(Wheat.dat, plot(fit, res))
```

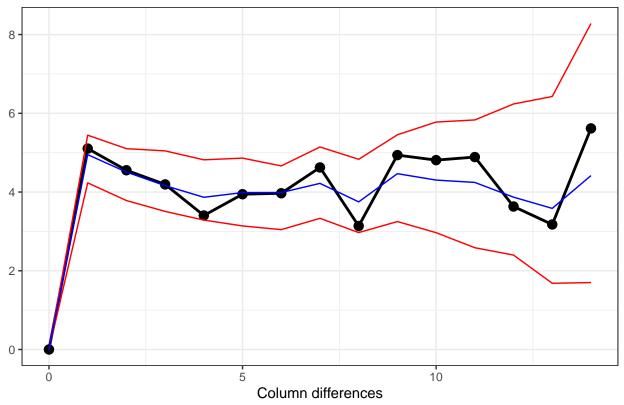


Plot variofaces

Variogram face of Standardized conditional residuals for Row





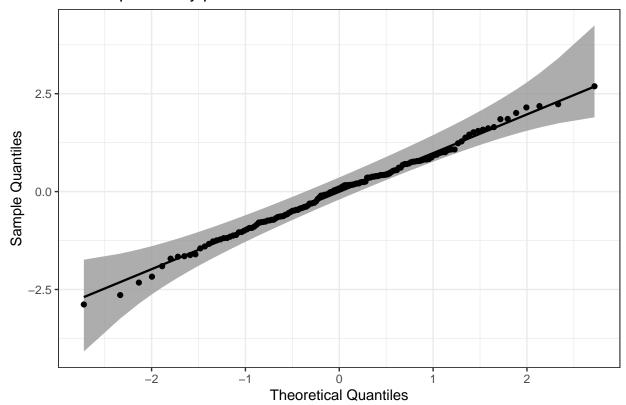


The variofaces are the lag 1 plots of the sample semivariogram with simulated confidence envelopes (Stefanova et al., 2009).

Plot normal quantile plot

The plot is obtained using the ggplot function with extensions available from the qqplotr package (Almeida, A., Loy, A. and Hofmann, H., 2023).

Normal probability plot



4. Prediction production and presentation

Get Variety predictions and all pairwise prediction differences and p-values

Predictions for yield from Variety

Notes:

- The predictions are obtained by averaging across the hypertable calculated from model terms constructed solely from factors in the averaging and classify sets.
- Use 'average' to move ignored factors into the averaging set.
- The ignored set: Rep,Row,Column,units
- Variety is included in this prediction

- (Intercept) is included in this prediction
- units is ignored in this prediction

16

17

18

19

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21

22

23

24

	Variety	nredicted value	standard (error	upper.halfLeastSignificant.limit
1	10	1168.989		.4768	1228.315
2	1	1242.750		.8104	1302.076
3	9	1257.137		.9708	1316.463
4	16	1285.718		.9400	1345.045
5	14	1293.526		.9227	1352.853
6	23	1313.653	120	. 2929	1372.979
7	11	1322.159	120	.1964	1381.485
8	7	1374.447	120	. 2407	1433.773
9	3	1394.070	120	.4032	1453.396
10	4	1410.980	120	.1055	1470.306
11	12	1444.557	120	.6034	1503.883
12	8	1453.396	120	.5940	1512.723
13	15	1458.383	120	.4346	1517.709
14	5	1473.782	120	.4455	1533.108
15	17	1487.828	120	. 2896	1547.154
16	6	1498.294	120	. 1189	1557.620
17	21	1517.121	120	. 2262	1576.447
18	2	1520.466	119	.6322	1579.792
19	24	1533.769	120	. 2995	1593.095
20	18	1541.148	120	.3664	1600.474
21	25	1575.795	120	.5142	1635.121
22	22	1610.482	120	.3281	1669.808
23	13	1610.762	120	.4575	1670.088
24	20	1627.971	120	. 2328	1687.297
25	19	1652.992		. 3435	1712.318
<pre>lower.halfLeastSignificant.limit est.status</pre>					
1			1109.663	Estim	
2			1183.424	Estim	
3			1197.811	Estin	
4			1226.392	Estim	
5			1234.200	Estin	
6			1254.327	Estim	
7			1262.832	Estim	
8			1315.120	Estim	
9			1334.743	Estim	
10			1351.653	Estim	
11			1385.231	Estim	
12			1394.070	Estim	
13			1399.057	Estin	
14			1414.456	Estin	
15			1428.501	Estim	lable

1438.968 Estimable

1457.795 Estimable

1461.140 Estimable

1474.443 Estimable

1481.821 Estimable

1516.468 Estimable

1551.156 Estimable

1551.436 Estimable 1568.645 Estimable

```
LSD values

minimum LSD = 114.0128

mean LSD = 118.6523

maximum LSD = 123.3578

(sed range / mean sed = 0.0788)
```

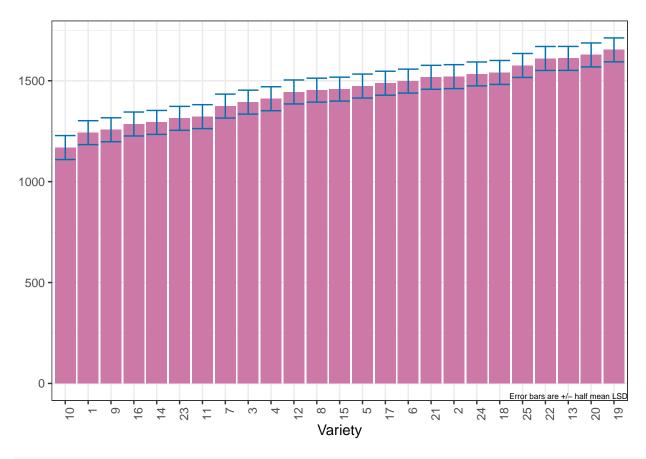
We have set error.intervals to halfLeast so that the limits for so that the limits for each prediction \pm (0.5 LSD) are calculated. When these are plotted overlapping error bars indicate predictions that are not significant, while those that do not overlap are significantly different (Snee, 1981).

Also set was sortFactor, so that the results would be ordered for the values of the predictions for Variety.

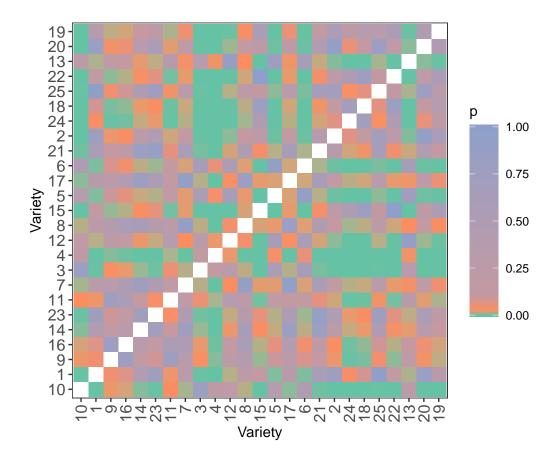
The function predictPlus returns an alldiffs object, a list consisting of the following components:

- predictions: the predictions, their standard errors and error intervals;
- vcov: the variance matrix of the predictions;
- differences: all pairwise differences between the predictions,
- p.differences: p-values for all pairwise differences between the predictions;
- sed: the standard errors of all pairwise differences between the predictions;
- LSD: the mean, minimum and maximum LSDs.

Plot the Variety predictions, with halfLSD intervals, and the p-values



plotPvalues(Var.diffs)



References

Almeida, A., Loy, A. and Hofmann, H. (2023) qqplotr: Quantile-Quantile plot extensions for 'ggplot2', Version 0.0.6. https://cran.r-project.org/package=qqplotr/ or https://github.com/aloy/qqplotr/.

Brien, C. J. (2024) asremlPlus: Augments ASReml-R in fitting mixed models and packages generally in exploring prediction differences. Version 4.4.32. https://cran.r-project.org/package=asremlPlus/ or http://chris.brien.name/rpackages/.

Butler, D. G., Cullis, B. R., Gilmour, A. R., Gogel, B. J. and Thompson, R. (2023). ASReml-R Reference Manual Version 4.2. VSN International Ltd, https://asreml.kb.vsni.co.uk/.

Gilmour, A. R., Thompson, R., & Cullis, B. R. (1995). Average Information REML: An Efficient Algorithm for Variance Parameter Estimation in Linear Mixed Models. *Biometrics*, **51**, 1440–1450.

Kenward, M. G., & Roger, J. H. (1997). Small sample inference for fixed effects from restricted maximum likelihood. *Biometrics*, **53**, 983-997.

Littell, R. C., Milliken, G. A., Stroup, W. W., Wolfinger, R. D., & Schabenberger, O. (2006). SAS for Mixed Models (2nd ed.). Cary, N.C.: SAS Press.

R Core Team (2024) R: A language and environment for statistical computing. Vienna, Austria: R Foundation for Statistical Computing. https://www.r-project.org/.

Snee, R. D. (1981). Graphical Display and Assessment of Means. Biometrics, 37, 835–836.

Stefanova, K. T., Smith, A. B. & Cullis, B. R. (2009) Enhanced diagnostics for the spatial analysis of field trials. *Journal of Agricultural, Biological, and Environmental Statistics*, **14**, 392–410.

Verbyla, A. P. (2019). A note on model selection using information criteria for general linear models estimated using REML. Australian & New Zealand Journal of Statistics, $\bf 61$, $\bf 39\text{-}50.https://doi.org/10.1111/anzs.12254/.$