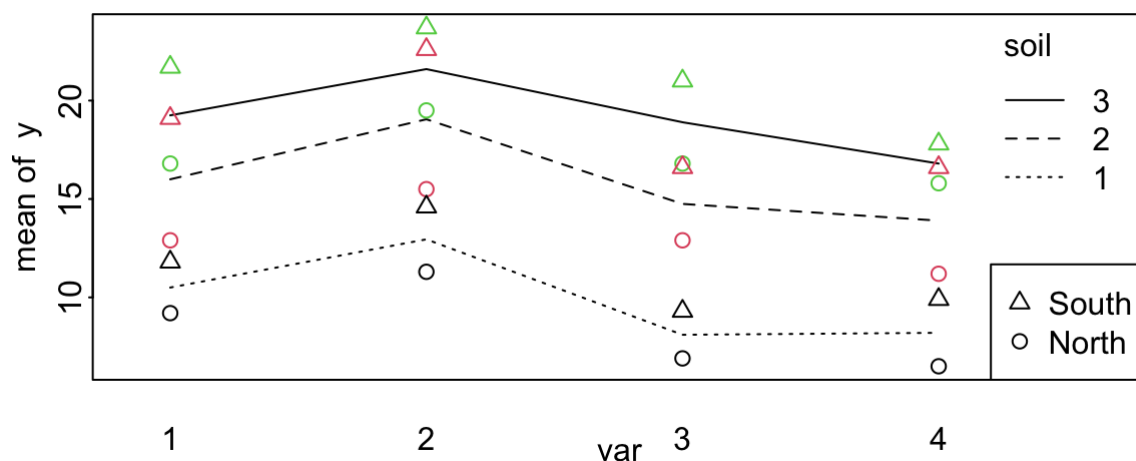


# Split Plot analysis with R

```
dat = read.table("splitplot.txt", header=T,
                 colClasses=c("factor","factor","factor","numeric"))
dat$pan = with(dat, block:soil) # or call it "bench"
with(dat, interaction.plot(var, soil, y, ylim=range(y)))
with(dat, points(var, y, col=soil, pch=as.numeric(block)))
legend("bottomright", pch=2:1, legend=levels(dat$block)[2:1])
```



```
library(lme4); library(lmerTest)
fit = lmer(y ~ block + soil + var + soil:var + (1 | pan), dat)
print(VarCorr(fit), comp=c("Variance","Std.Dev.)) # summary(fit)
```

Groups	Name	Variance	Std.Dev.
pan	(Intercept)	0.76281	0.87339
Residual		0.65417	0.80881

```
anova(fit)
```

```
Type III Analysis of Variance Table with Satterthwaite's method
```

	Sum Sq	Mean Sq	NumDF	DenDF	F value	Pr(>F)
block	17.951	17.9512	1	2	27.4414	0.03456
soil	61.583	30.7916	2	2	47.0699	0.02080
var	81.530	27.1767	3	9	41.5440	1.343e-05
soil:var	3.743	0.6238	6	9	0.9535	0.50444

```
ls_means(fit, which="soil")
```

Least Squares Means table:

	Estimate	Std. Error	df	t value	lower	upper	Pr(> t )
soil1	9.93750	0.68057	2	14.602	7.00924	12.86576	0.004657
soil2	15.92500	0.68057	2	23.399	12.99674	18.85326	0.001821
soil3	19.13750	0.68057	2	28.120	16.20924	22.06576	0.001262

Confidence level: 95%

Degrees of freedom method: Satterthwaite

```
lsm = ls_means(fit, which="soil", pairwise=TRUE); lsm
```

Least Squares Means table:

	Estimate	Std. Error	df	t value	lower	upper	Pr(> t )
soil1 - soil2	-5.98750	0.96247	2	-6.2210	-10.12869	-1.84631	0.02488
soil1 - soil3	-9.20000	0.96247	2	-9.5587	-13.34119	-5.05881	0.01077
soil2 - soil3	-3.21250	0.96247	2	-3.3378	-7.35369	0.92869	0.07924

Confidence level: 95%

Degrees of freedom method: Satterthwaite

```
show_tests(lsm) # shows which contrasts were used above
```

\$soil

	(Intercept)	blockSouth	soil2	soil3	var2	var3	var4	soil2:var2	soil3:var2
soil1 - soil2	0	0	-1	0	0	0	0	-0.25	0.00
soil1 - soil3	0	0	0	-1	0	0	0	0.00	-0.25
soil2 - soil3	0	0	1	-1	0	0	0	0.25	-0.25

	soil2:var3	soil3:var3	soil2:var4	soil3:var4
soil1 - soil2	-0.25	0.00	-0.25	0.00
soil1 - soil3	0.00	-0.25	0.00	-0.25
soil2 - soil3	0.25	-0.25	0.25	-0.25

```
ls_means(fit, which="var")
```

Least Squares Means table:

	Estimate	Std. Error	df	t value	lower	upper	Pr(> t )
var1	15.25000	0.48597	4.4	31.381	13.94828	16.55172	2.385e-06
var2	17.86667	0.48597	4.4	36.765	16.56495	19.16839	1.190e-06
var3	13.91667	0.48597	4.4	28.637	12.61495	15.21839	3.563e-06
var4	12.96667	0.48597	4.4	26.682	11.66495	14.26839	4.857e-06

Confidence level: 95%

Degrees of freedom method: Satterthwaite

Why this standard error, and why this degree of freedom of 4.4?

Mean for variety 1 (say): from average over 2 blocks and 3 soils. It's standard error is then:

```
sqrt((0.7628 + 0.6542)/6) # sqrt of (s2_epsilon + s2_delta) * 1/(2*3)
```

```
[1] 0.4859698
```

This estimated variance is a rescaled version of

$MSWPE + (4-1) MSSPE = 4(s2\_epsilon + s2\_delta)$ , so the Satterthwaite approximation will use:

```
MSWPE = 0.6542 + 4*0.7628 # df = 2
MSSPE = 0.6542 # df = 9
dfsat = (MSWPE + 3*MSSPE)^2 / ( MSWPE^2/2 + (3*MSSPE)^2/9)
dfsat # 4.4 degrees of freedom to approximate by a chi-square distribution
```

```
[1] 4.405093
```

```
head(ls_means(fit, which = "soil:var"), n=2)
```

Least Squares Means table:

	Estimate	Std. Error	df	t value	lower	upper	Pr(> t )
soil1:var1	10.50000	0.84172	4.4	12.475	8.24535	12.75465	0.0001328
soil2:var1	16.00000	0.84172	4.4	19.009	13.74535	18.25465	2.139e-05

Confidence level: 95%

Degrees of freedom method: Satterthwaite

```
head(ls_means(fit, which = "soil:var", pairwise=TRUE), n=4)
```

Least Squares Means table:

	Estimate	Std. Error	df	t value	lower	upper	Pr(> t )
soil1:var1 - soil2:var1	-5.50000	1.19037	4.4	-4.6204	-8.68855	-2.31145	0.007833
soil1:var1 - soil3:var1	-8.75000	1.19037	4.4	-7.3507	-11.93855	-5.56145	0.001243
soil1:var1 - soil1:var2	-2.45000	0.80881	9.0	-3.0292	-4.27965	-0.62035	0.014266
soil1:var1 - soil2:var2	-8.55000	1.19037	4.4	-7.1826	-11.73855	-5.36145	0.001367

Confidence level: 95%

Degrees of freedom method: Satterthwaite

**Warning:** an alternative is the `multcomp` package. It uses a normal approximation to test contrasts and pairwise comparisons. This is very liberal for `soil` comparisons, for which `df` should be taken to be 2 (not infinite).

```
library(multcomp)
summary(glht(fit, linfct=mcp(soil="Tukey", interaction_average=T)))
```

### Simultaneous Tests for General Linear Hypotheses

Multiple Comparisons of Means: Tukey Contrasts

```
Fit: lmer(formula = y ~ block + soil + var + soil:var + (1 | pan),
  data = dat)
```

Linear Hypotheses:

	Estimate	Std. Error	z value	Pr(> z )
2 - 1 == 0	5.9875	0.9625	6.221	<1e-04
3 - 1 == 0	9.2000	0.9625	9.559	<1e-04
3 - 2 == 0	3.2125	0.9625	3.338	0.0024

(Adjusted p values reported -- single-step method)