

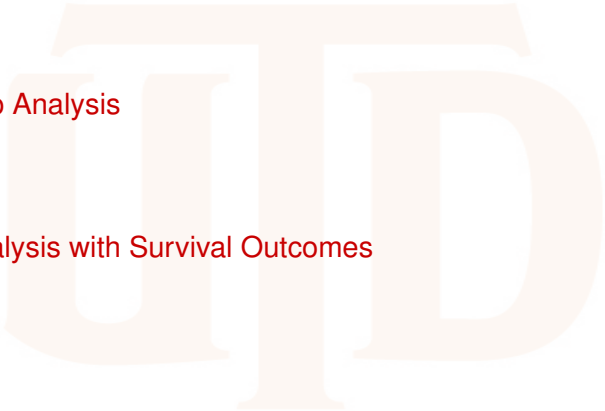
# Meta-Analysis with R

## Meta-Analysis of Other Outcomes

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# Subgroup Analysis

- From time to time we need to work with subgroups of studies in a meta-analysis. The various “R” commands for meta-analysis in the “R” package meta support a “byvar” option, i.e. conduct a subgroup analysis by a variable.

```
> # 1. Read in the data:
> data3 <- read.csv("dataset03.csv")
> # 2. As usual, to view an object, type its name:
> data3
```

	author	year	Ne	Me	Se	Nc	Mc	Sc	duration
1	Bontognali	1991	30	0.70	3.76	30	1.27	4.58	<= 3 months
2	Castiglioni	1986	311	0.10	0.21	302	0.20	0.29	<= 3 months
3	Cremonini	1986	21	0.25	0.23	20	0.71	0.29	<= 3 months
4	Grassi	1994	42	0.16	0.29	41	0.45	0.43	<= 3 months
5	Jackson	1984	61	0.11	0.00	60	0.13	0.00	<= 3 months
6	Allegra	1996	223	0.07	0.11	218	0.11	0.14	> 3 months
7	Babolini	1980	254	0.13	0.18	241	0.33	0.27	> 3 months
8	Boman	1983	98	0.20	0.27	105	0.32	0.30	> 3 months
9	Borgia	1981	10	0.05	0.08	9	0.15	0.17	> 3 months
10	Decramer	2005	256	0.10	0.11	267	0.11	0.16	> 3 months
11	Grassi	1976	35	0.14	0.15	34	0.27	0.21	> 3 months
12	Grillage	1985	54	0.10	0.00	55	0.12	0.00	> 3 months
13	Hansen	1994	59	0.11	0.15	70	0.16	0.19	> 3 months
14	Malerba	2004	115	0.06	0.08	119	0.07	0.08	> 3 months
15	McGavin	1985	72	0.42	0.34	76	0.52	0.35	> 3 months
16	Meister	1986	90	0.15	0.15	91	0.20	0.19	> 3 months
17	Meister	1999	122	0.06	0.15	124	0.10	0.15	> 3 months
18	Moretti	2004	63	0.12	0.14	61	0.17	0.17	> 3 months
19	Nowak	1999	147	0.03	0.06	148	0.06	0.12	> 3 months
20	Olivieri	1987	110	0.18	0.31	104	0.33	0.41	> 3 months
21	Parr	1987	243	0.18	0.21	210	0.21	0.21	> 3 months
22	Pela	1999	83	0.17	0.18	80	0.29	0.32	> 3 months
23	Rasmussen	1988	44	0.13	0.21	47	0.14	0.19	> 3 months

# Subgroup Analysis

- The result of the meta-analysis is given by

```
> mc3 <- metacont(Ne, Me, Se, Nc, Mc, Sc, data=data3,
+               studlab=paste(author, year))
Warning message:

> print(summary(mc3), digits=2)
Number of studies combined: k=21

              MD              95%-CI              z  p-value
Fixed effect model  -0.05 [-0.05; -0.04] -10.06 < 0.0001
Random effects model -0.08 [-0.11; -0.05]  -5.82 < 0.0001

Quantifying heterogeneity:
tau^2 = 0.0027; H = 2.63 [2.19; 3.15]; I^2 = 85.5% [79.1%; 89.9%]

Test of heterogeneity:
      Q d.f.  p-value
138.08   20 < 0.0001
Details on meta-analytical method:
- Inverse variance method
- DerSimonian-Laird estimator for tau^2
```

- The results indicate significant between-study heterogeneity ( $Q = 138, p < 0.0001$ ) with  $I^2 = 85.5\%$ . Here subgroup information is available for study duration: studies whose duration was greater or less than three months.

# Subgroup Analysis

- A subgroup analysis can be done by using argument “byvar” in the original call of the “metacont” function:

```
> mc3s <- metacont(Ne, Me, Se, Nc, Mc, Sc, data=data3,  
+                  studlab=paste(author, year),  
+                  byvar=duration, print.byvar=FALSE)
```

- Another more convenient way is to update the original meta-analysis by using the “update.meta” function from “R” package “meta”.

```
> mc3s <- update(mc3, byvar=duration, print.byvar=FALSE)
```

# Subgroup Analysis

```
> print(summary(mc3s), digits=2)
Number of studies combined: k=21

              MD              95%-CI              z  p-value
Fixed effect model  -0.05 [-0.05; -0.04] -10.06 < 0.0001
Random effects model -0.08 [-0.11; -0.05] -5.82 < 0.0001

Quantifying heterogeneity:
tau^2 = 0.0027; H = 2.63 [2.19; 3.15]; I^2 = 85.5% [79.1%; 89.9%]

Test of heterogeneity:
      Q d.f.  p-value
138.08   20 < 0.0001

Results for subgroups (fixed effect model):
      k  MD      95%-CI      Q    tau^2  I^2
<= 3 months   4 -0.13 [-0.17; -0.09] 22.43  0.035 86.6%
> 3 months   17 -0.04 [-0.05; -0.03] 94.92  0.002 83.1%

Test for subgroup differences (fixed effect model):
      Q d.f.  p-value

Between groups  20.73   1 < 0.0001
Within groups 117.35  19 < 0.0001

Results for subgroups (random effects model):
      k  MD      95%-CI      Q    tau^2  I^2
<= 3 months   4 -0.28 [-0.50; -0.05] 22.43  0.035 86.6%
> 3 months   17 -0.06 [-0.09; -0.04] 94.92  0.002 83.1%

Test for subgroup differences (random effects model):
      Q d.f.  p-value
Between groups  3.41   1  0.0647

Details on meta-analytical method:
```

# Subgroup Analysis

- The results for the fixed effect model show that between-group heterogeneity is highly statistically significant ( $Q = 20.73$  on 1 degrees of freedom) as well as within-group heterogeneity ( $Q = 117.35$ , 19 degrees of freedom).
- Further, the fixed effect estimates ( $-0.13$ , short duration  $-0.04$ , long duration) are not that different. While short duration studies seem to have far fewer patients, the effect appears similar; study duration does not appear to be the source of the high degree of heterogeneity in these data.
- This observation is supported by the results for the random effects model (between-study heterogeneity:  $Q = 3.41$ , 1 degrees of freedom).

# Meta-Analysis with Survival Outcomes

- A meta-analysis with survival time outcomes is typically based on the hazard ratio as measure of treatment effect.
- Accordingly, the logarithm of the hazard ratio and its standard error are the basic quantities utilised in meta-analysis.
- The generic inverse variance method can be used straightforward with log hazard ratio  $\hat{\theta}_k$  and its standard error  $S.E.(\hat{\theta}_k)$ , for study  $k$ ,  $k = 1, \dots, K$ .



# Meta-Analysis with Survival Outcomes

- Using these quantities, all methods described before can be used for meta-analysis. In the following example we consider the most basic case, i.e. fixed effect and random effects meta-analysis using the DerSimonian-Laird method to estimate the between-study variance  $\hat{\tau}^2$ .

```
> # 1. Read in the data
> data4 <- read.csv("dataset04.csv")
> # 2. Print data
> data4
```

	author	year	Ne	Nc	logHR	selogHR
1	FCG on CLL	1996	53	52	-0.5920	0.3450
2	Leporrier	2001	341	597	-0.0791	0.0787
3	Rai	2000	195	200	-0.2370	0.1440
4	Robak	2000	133	117	0.1630	0.3120

# Meta-Analysis with Survival Outcomes

- Columns “logHR” and “selogHR” correspond to the log hazard ratio and its standard error.

```
> mgl <- metagen(logHR, selogHR,
+               studlab=paste(author, year), data=data4,
+               sm="HR")
```

```
> print(mgl, digits=2)
```

	HR	95%-CI	%W(fixed)	%W(random)
FCG on CLL 1996	0.55 [0.28; 1.09]		3.68	5.85
Leporrier 2001	0.92 [0.79; 1.08]		70.70	59.76
Rai 2000	0.79 [0.59; 1.05]		21.12	27.32
Robak 2000	1.18 [0.64; 2.17]		4.50	7.08

Number of studies combined: k=4

	HR	95%-CI	z	p-value
Fixed effect model	0.89 [0.78; 1.01]		-1.82	0.0688
Random effects model	0.87 [0.74; 1.03]		-1.58	0.1142

Quantifying heterogeneity:

$\tau^2 = 0.0061$ ;  $H = 1.1$  [1; 2.81];  $I^2 = 17.2\%$  [0%; 87.3%]

# Meta-Analysis with Survival Outcomes

Test of heterogeneity:

Q	d.f.	p-value
3.62	3	0.3049

Details on meta-analytical method:

- Inverse variance method
- DerSimonian-Laird estimator for  $\tau^2$