# InterLymph meta-analysis

August 31, 2016

Load libraries:

```
library(mvmeta)
## This is mumeta 0.4.7. For an overview type: help('mumeta-package').
library(xtable)
```

#### 1 Load study-level results

Load the effect sizes and covariance matrices from the RData file:

```
load("InterLymph_summary_stats.RData")

##names of variables
Var1 <- "Wine"
Var2 <- "Liquor"
Var3 <- "Beer"
Var4 <- "Smoking"</pre>
```

## 2 Fixed effects meta-analyses

We fit the MVMA and UVMA fixed effects models:

Save all the results in a table:

Here is the table in a form that can be copied/pasted into a LaTeX document from the .tex file:

	Variable	Method	Estimate	Pvalue	CI
1	Wine	MVMA	-0.3300	0.06	(-0.67, 0.01)
2	Liquor	MVMA	-0.3100	0.55	(-1.31, 0.69)
3	Beer	MVMA	-0.8200	< 0.001	(-1.38, -0.27)
4	Smoking	MVMA	0.3200	< 0.001	(0.17, 0.46)
5	Wine	UVMA	-0.4900	0.01	(-0.86, -0.12)
6	Liquor	UVMA	-0.1800	0.75	(-1.26, 0.9)
7	Beer	UVMA	-0.9600	< 0.001	(-1.53, -0.4)
8	Smoking	UVMA	0.3200	< 0.001	(0.18, 0.47)

#### 3 Random effects meta-analyses

We fit the MVMA and UVMA random effects models (using REML):

```
##MVMA
MVMA.RE <- coef(summary(mvmeta(EffSizes, S = CovMats,
                                 method="reml")))
MVMA.RE[, -c(2,4)] \leftarrow round(MVMA.RE[, -c(2,4)], 2)
MVMA.RE[, 4] <- round(MVMA.RE[, 4], 2)
##UVMAs have to be fit separately for RE case
UVMA.RE1 <- mvmeta(EffSizes[,1,drop=FALSE],</pre>
                    S = sapply(CovMats, function(x) \{x[1,1,drop=FALSE]\}),
                    method="reml")
UVMA.RE2 <- mvmeta(EffSizes[,2,drop=FALSE],</pre>
                    S = sapply(CovMats, function(x){x[2,2,drop=FALSE]}),
                    method="reml")
UVMA.RE3 <- mvmeta(EffSizes[,3,drop=FALSE],</pre>
                    S = sapply(CovMats, function(x){x[3,3,drop=FALSE]}),
                    method="reml")
UVMA.RE4 <- mvmeta(EffSizes[,4,drop=FALSE],</pre>
                    S = sapply(CovMats, function(x) \{x[4,4,drop=FALSE]\}),
                    method="reml")
UVMA.RE <- rbind(coef(summary(UVMA.RE1)), coef(summary(UVMA.RE2)),</pre>
                  coef(summary(UVMA.RE3)), coef(summary(UVMA.RE4)))
UVMA.RE[, -c(2,4)] \leftarrow round(UVMA.RE[, -c(2,4)], 2)
UVMA.RE[, 4] <- round(UVMA.RE[, 4], 2)
```

Save all the results in a table:

Here is the table in a form that can be copied/pasted into a LaTeX document from the .tex file:

	Variable	Method	Estimate	Pvalue	CI
1	Wine	MVMA	-0.5900	0.01	(-1.01, -0.18)
2	Liquor	MVMA	0.0600	0.94	(-1.56, 1.68)
3	Beer	MVMA	-0.7500	0.05	(-1.48, -0.01)
4	Smoking	MVMA	0.2600	0.01	(0.06, 0.47)
5	Wine	UVMA	-0.4900	0.01	(-0.86, -0.12)
6	Liquor	UVMA	0.2600	0.76	(-1.39, 1.92)
7	Beer	UVMA	-0.9200	0.03	(-1.75, -0.09)
8	Smoking	UVMA	0.2600	0.01	(0.06, 0.46)

## 4 Make table combining results from fixed- and random-effects

Make combined table for manuscript:

Here is the table in a form that can be copied/pasted into a LaTeX document from the .tex file:

FE: MVMA	FE: UVMA	RE: MVMA	RE: UVMA	NA
Wine	-0.33 (-0.67, 0.01)	-0.49 (-0.86, -0.12)	-0.59 (-1.01, -0.18)	-0.49 (-0.86, -0.12)
Liquor	-0.31 (-1.31, 0.69)	-0.18 (-1.26, 0.9)	0.06 (-1.56, 1.68)	0.26 (-1.39, 1.92)
Beer	-0.82 (-1.38, -0.27)	-0.96 (-1.53, -0.4)	-0.75 (-1.48, -0.01)	-0.92 (-1.75, -0.09)
Smoking	0.32 (0.17, 0.46)	0.32 (0.18, 0.47)	0.26 (0.06, 0.47)	0.26 (0.06, 0.46)