refitME: Tutorial for fitting MCEM models with measurement error in covariates

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Example 1: A simple GLM example taken from Carroll et al. (2006).

The Framingham heart study data set.

Load data and R-packages.

```
suppressMessages(library(refitME));
suppressMessages(library(simex));
epsilon<-0.00001;
B<-100;
family<-"binomial";
data.Fram<-as.matrix(read.table(file="Framinghamdata.txt"));</pre>
```

Setup all variables.

The construction below follows the Carroll et al. (2006) monograph.

```
Y<-data.Fram[,10];
n<-length(Y);
z1<-(data.Fram[,9]); # Cholesterol.
z2<-(data.Fram[,2]); # Age.
z3<-data.Fram[,7]; # Smoke.
w1<-(log((data.Fram[,3]+data.Fram[,4]+data.Fram[,5]+data.Fram[,6])/4-50)); # Mean exam 2 and 3.
dat<-data.frame(cbind(Y,z1,z2,z3,w1));
sigma.sq.u<-0.01259/2 # ME variance, obtained from Carroll et al. (2006) monograph.</pre>
```

Fit the naive model.

```
mod_naiv1<-glm(Y~w1+z1+z2+z3,x=TRUE,family=binomial,data=dat);</pre>
```

Fit the SIMEX model.

Fit the MCEM model.

```
start<-Sys.time();
est<-refitME(mod_naiv1,sigma.sq.u,B);

## [1] "convergence :-)"
## [1] 5
end<-Sys.time();
t2<-difftime(end,start,units="secs");
comp.time<-c(comp.time,t2);</pre>
```

Report and compare times and model estimates.

```
est.beta<-rbind(coef(mod_naiv1),coef(mod_simex1),est$beta);</pre>
est.beta.se<-rbind(sqrt(diag(vcov(mod_naiv1))),</pre>
                   sqrt(diag(mod_simex1$variance.jackknife)),
                   est$beta.se2);
round(est.beta,digits=3);
        (Intercept)
                                    z2
##
                       w1
                              z1
## [1,]
            -14.951 1.707 0.008 0.055 0.592
## [2,]
            -15.710 1.897 0.008 0.054 0.601
## [3,]
            -16.114 1.966 0.008 0.055 0.597
round(est.beta.se,digits=3);
##
        (Intercept)
                       w1
                              z1
## [1,]
             1.900 0.418 0.002 0.012 0.250
## [2,]
              2.102 0.473 0.002 0.012 0.251
## [3,]
              2.185 0.486 0.002 0.012 0.251
comp.time;
## Time differences in secs
## [1] 14.435396 8.071469
```

Example 2: A GAM example taken from Ganguli et al. (2005).

The Milan mortality air pollution data set.

Load data and R-packages.

```
rm(list=ls());
suppressMessages(library(refitME));
suppressMessages(library(SemiPar));
epsilon<-0.00001;
B<-2;
family<-"poisson";
data(milan.mort);
dat.air<-milan.mort;</pre>
```

Setup all variables.

```
Y<-dat.air[,6];
n<-length(Y);
w1<-log(dat.air[,9]);
w1<-scale(w1);
colnames(w1)<-"w1";
z1<-(dat.air[,1]);
z2<-(dat.air[,4]);
z3<-(dat.air[,5]);
dat<-data.frame(cbind(Y,z1,z2,z3,w1));
## Reliability ratio.
sigma.sq.u<-0.1; # Rel. ratio of 0.9.
#sigma.sq.u<-0.2; # Rel. ratio of 0.8.
#sigma.sq.u<-0.3; # Rel. ratio of 0.7.
rel.rat<-(1-sigma.sq.u/var(dat$w1))*100;</pre>
```

Fit the naive model.

```
mod_naiv1 < -gam(Y \sim s(w1, k=5) + s(z1, bs='cc', k=25) + s(z2, k=5) + s(z3, k=5), family="poisson", data=dat);
```

Fit the MCEM model.

```
This models can take a while to fit.
```

```
est<-refitME(mod_naiv1,sigma.sq.u,B);</pre>
```

```
## [1] "convergence :-)"
## [1] 6
```

Plots.

MCEM (Poisson GAM) fitted to the air pollution data

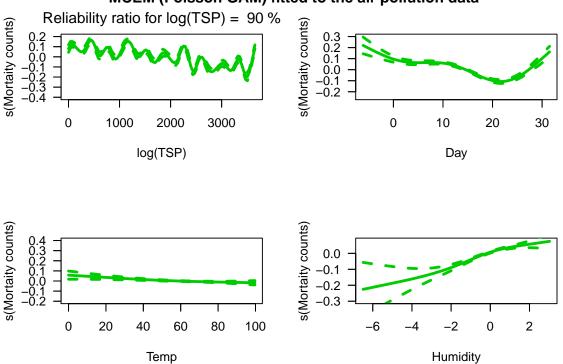


Figure 1: Plots of smooths against covariate.