

# Ordinary Kriging in R

```
library(geoR);library(fields);library(maps)
```

```
#You will need to change the directory to load these files:
```

```
source("../\\plot.field.points.R")
```

```
load("../\\PM25.RData")
```

```
#Combine the spatial coordinates in a 84x2 matrix
```

```
s<-cbind(long,lat)
```

```
#Plot the data
```

```
plot.field.points(s,PM,map.border="county",cex=1.5)
```

```
X11()
```

```
#Estimate parameters by maximum likelihood:
```

```
ml <- likfit(data=PM,coords=s,
```

```
      fix.nugget=F,cov.model="exponential",
```

```
      ini = c(30, 5),nugget=5)
```

```
summary(ml)
```

```
#Create grid of prediction points:
```

```
sp1<-seq(min(s[,1]),max(s[,1]),length=100)
```

```
sp2<-seq(min(s[,2]),max(s[,2]),length=100)
```

```
sp<-expand.grid(sp1,sp2)
```

```
inCA<-map.where("state",x=sp[,1],y=sp[,2])
```

```
inCA[is.na(inCA)]<-"NA"
```

```
inCA<-inCA=="california"
```

```
#Perform ordinary Kriging (value of cov.pars and nugget are copied from mle output):
```

```
pred<-krige.conv(data=PM,coords=s,locations=sp,
```

```
      krige=krige.control(cov.model="exponential",
```

```
      cov.pars=c(14.73,6.144),
```

```
      nugget=4.299))
```

```
pred$predict[!inCA]<-NA
```

```
pred$krige.var[!inCA]<-NA
```

```
#Plot the predicted values:
```

```
image.plot(sp1,sp2,matrix(pred$predict,100,100),zlim=range(PM))
```

```
map("county",add=T)
```

```
#Plot the standard errors:
```

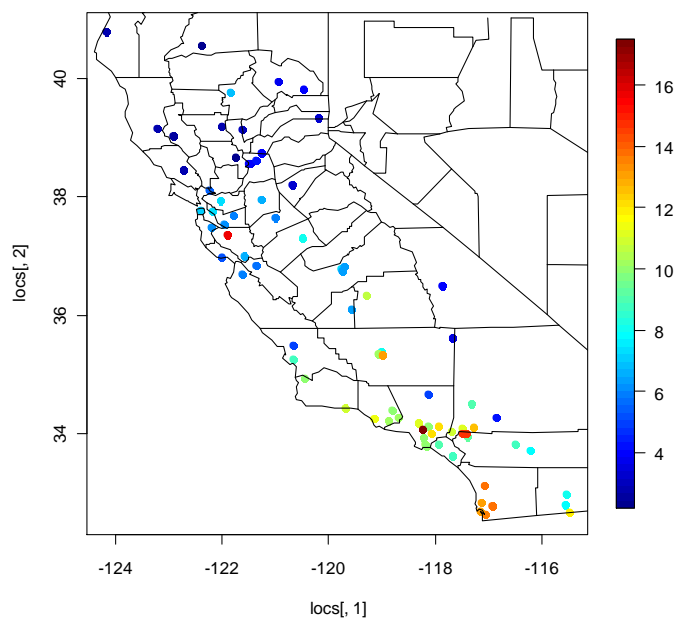
```
image.plot(sp1,sp2,matrix(sqrt(pred$krige.var),100,100),zlim=c(2,3.5))
```

```
map("county",add=T)
```

```
points(s)
```

# Output

Raw data:



```
> summary(ml)
```

Parameters of the mean component (trend):

beta

6.6346

Parameters of the spatial component:

correlation function: exponential

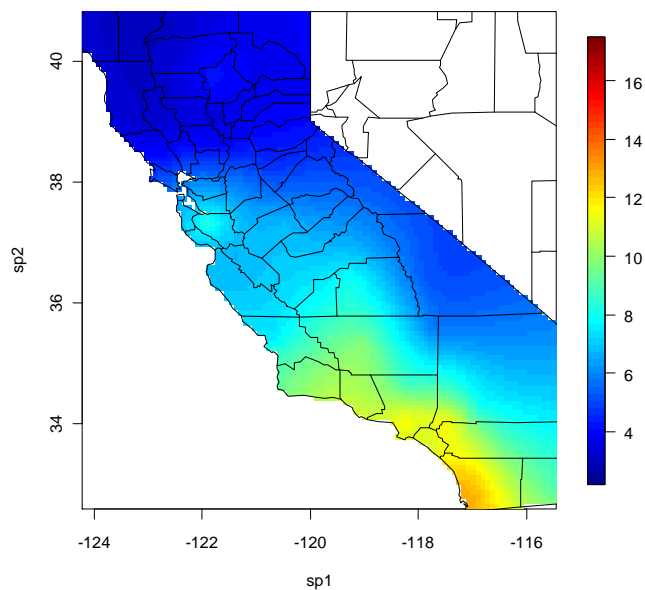
(estimated) partial sill = 14.73

(estimated) range parameter = 6.144

Parameter of the error component:

(estimated) nugget = 4.299

Predicted values



Standard deviations

