

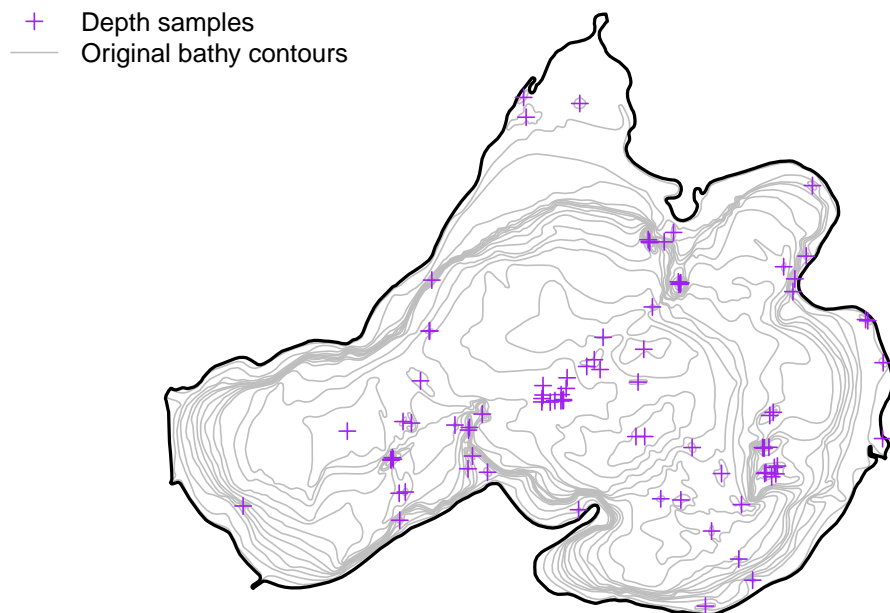
# Loken\_\_HW8

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Dear Dr. Dugan:

Create a spatial interpolated map of Mendota bathymetry using the Mendota-Bathy.csv file



```
#Create spatial lines to calculate distance to shore
MEline<-as(MEoutline2, "SpatialLines")
MEcsv$DistToShore<-gDistance(MEcsv, MEline, byid=T)[1,]

# Make polygrid - This is each location to make predictions
bbox<-MEoutline2@bbox
pixelsize=200
data1.grid<-polygrid(seq(bbox[1,1], bbox[1,2], pixelsize),
                     seq(bbox[2,1], bbox[2,2], pixelsize),
                     borders=MEoutline2@polygons[[1]]@Polygons[[1]]@coords)
```

```

# Set names of coordinates and match projection
coordinates(data1.grid)<--x+y
proj4string(data1.grid) <- crs(MEoutline2)

data1.grid$DistToShore<-gDistance(data1.grid, MEline, byid=T)[1,]

# Convert to gridded (pixels) rather than points
gridded(data1.grid) <- TRUE

# plot data grid, boundary, and observations
# plot(data1.grid, border="grey")
# plot(MEoutline2, add=TRUE, col=NA, lwd=3)
# plot(MEcsv, add=TRUE, col="red", cex=0.5)

#Create variogram for interpolation
gs <- gstat(formula=depth_m~1, locations=MEcsv)
v <- variogram(gs, width=100 )
v.fit<-fit.variogram(v, vgm(c('Lin', 'Sph', 'Exp', 'Gau', 'Nug')), fit.method=2)

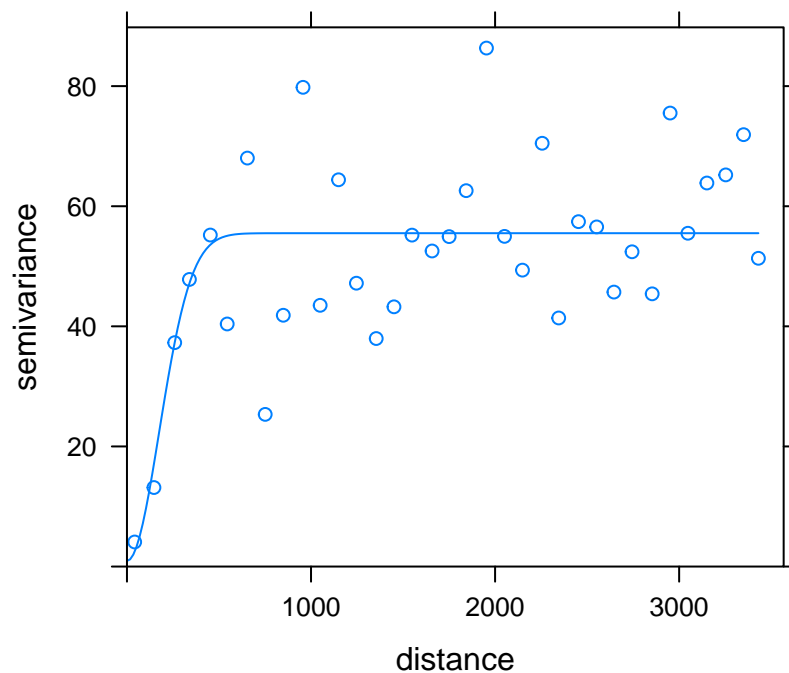
## Warning in fit.variogram(object, x, fit.sills = fit.sills, fit.ranges =
## fit.ranges, : No convergence after 200 iterations: try different initial
## values?

v.fit

##   model    psill   range
## 1   Nug 0.9390482  0.0000
## 2   Gau 54.5682175 247.5283

plot(v, v.fit)

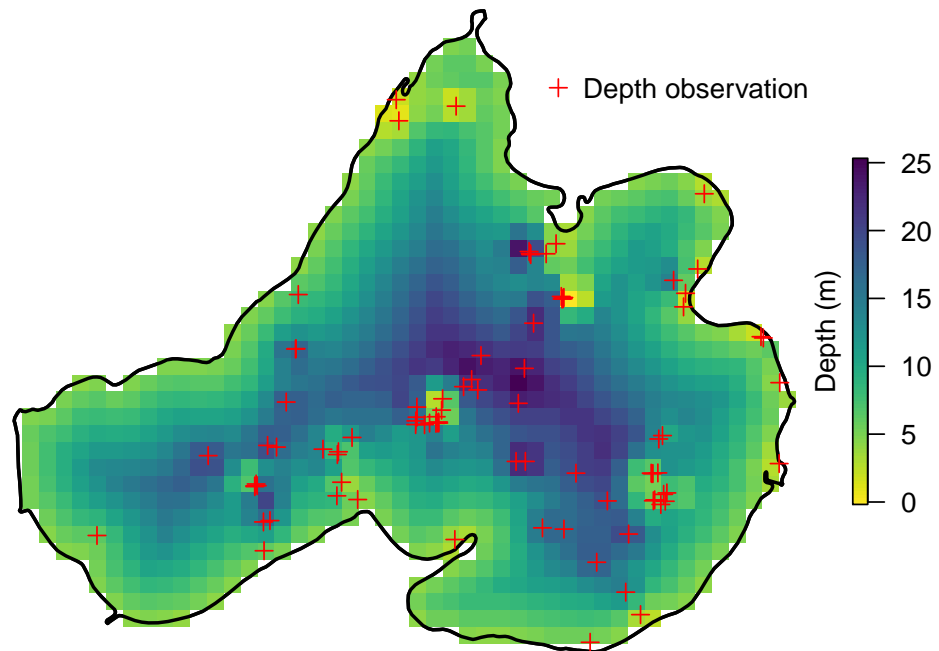
```



```
#Krig to datagrid using Distance to shore and semivariance model
krig1<- krige(depth_m~DistToShore, MEcsv, data1.grid, v.fit)

## [using universal kriging]
krig_r<-raster(krig1)
```

Predicted depth using universal Kriging and distance to shore



Nice trick changing the depths of the first 7 observations in the middle of the lake...Dr. Sneaky