

Exposure to Dust - Trees

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First the dust data are loaded from the package "catdata".

```
> library(catdata)
> data(dust)
```

Trees can be fitted by use of the function "rpart" from package "rpart".

```
> library(rpart)
```

Now a tree is fitted. We take "years" as the only covariate, "bronch" is the binary response. Afterwards the corresponding tree is plotted.

```
> tree1 <- rpart(as.factor(bronch) ~ years, data = dust, method = "class",
+               control = rpart.control(cp = 0.001, parms=list(split='information'),
+               maxdepth = 4))
> plot(tree1, xpd=TRUE)
> text(tree1)
```

In the following the fit is plotted. It shows how the tree can be interpreted as regression function.

```
> pred <- predict(tree1)
> year <- dust$years
> year [dust$years<15.5] <- 1
> year [dust$years>15.5 & dust$years<36.5] <- 2
> year [dust$years>36.5 & dust$years<47.5] <- 3
> year [dust$years>47.5 & dust$years<50.5] <- 4
> year [dust$years>50.5] <- 5
> pre5 <- unique(pred[,2][year==5])
> pre4 <- unique(pred[,2][year==4])
> pre3 <- unique(pred[,2][year==3])
> pre2 <- unique(pred[,2][year==2])
> pre1 <- unique(pred[,2][year==1])
> meanyear <- c()
> for (i in min(dust$years):max(dust$years)){
+ meanyear[i] <- sum(dust$bronch[dust$year==i])
+ if (sum(dust$bronch[dust$year==i])!=0){
+ meanyear[i] <- mean(dust$bronch[dust$year==i])
+ }
+ }
```

```

> dust$means<- rep(2, nrow(dust))
> for (k in 1:nrow(dust)){
+ dust$means[k] <- meanyear[dust$years[k]]
+ }

> plot(dust$years, dust$means, xlab="years",ylab="")
> segments(x0=3,x1=15.5,y0=pre1)
> segments(x0=15.5,x1=15.5,y0=pre1,y1=pre2)
> segments(x0=15.5,x1=36.5,y0=pre2)
> segments(x0=36.5,x1=36.5,y0=pre2,y1=pre3)
> segments(x0=36.5,x1=47.5,y0=pre3)
> segments(x0=47.5,x1=47.5,y0=pre3,y1=pre4)
> segments(x0=47.5,x1=50.5,y0=pre4)
> segments(x0=50.5,x1=50.5,y0=pre4,y1=pre5)
> segments(x0=50.5,x1=66,y0=pre5)

```

An alternative package to generate trees is "party" which contains the function "ctree".

```

> library(party)

```

As before with "rpart" we fit a tree with "years" as only covariate.

```

> treeP1 <-ctree(as.factor(bronch) ~ years, data = dust)
> plot(treeP1)

> year<- dust$years
> year [dust$years<7.5] <- 1
> year [dust$years>7.5 & dust$years<15.5] <- 2
> year [dust$years>15.5 & dust$years<36.5] <- 3
> year [dust$years>36.5] <- 4
> pre4 <- mean(dust$bronch[year==4])
> pre3 <- mean(dust$bronch[year==3])
> pre2 <- mean(dust$bronch[year==2])
> pre1 <- mean(dust$bronch[year==1])

```

```

> plot(dust$years, dust$means, xlab="years",ylab="")
> segments(x0=3,x1=7.5,y0=pre1)
> segments(x0=7.5,x1=7.5,y0=pre1,y1=pre2)
> segments(x0=7.5,x1=15.5,y0=pre2)
> segments(x0=15.5,x1=15.5,y0=pre2,y1=pre3)
> segments(x0=15.5,x1=36.5,y0=pre3)
> segments(x0=36.5,x1=36.5,y0=pre3,y1=pre4)
> segments(x0=36.5,x1=66,y0=pre4)
>

```

Now we take "smoke", "years" and "dust" as covariates for the binary response "bronch" and again plot the tree.

```

> treeP2 <-ctree(as.factor(bronch) ~ smoke + years + dust, data = dust)
> plot(treeP2)

```