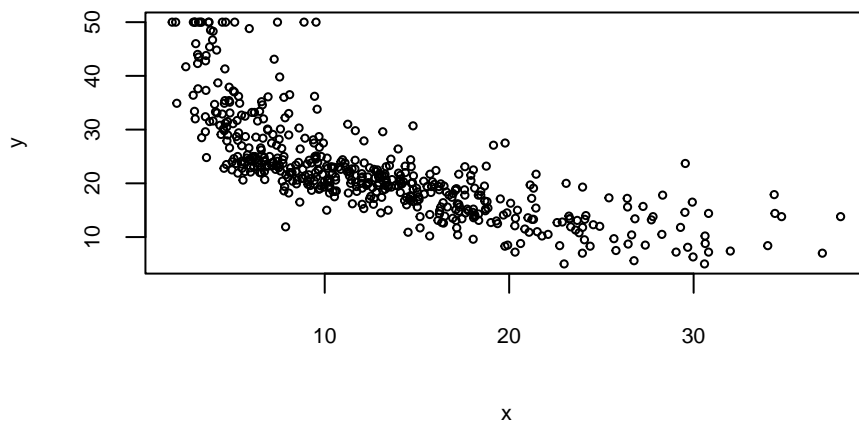
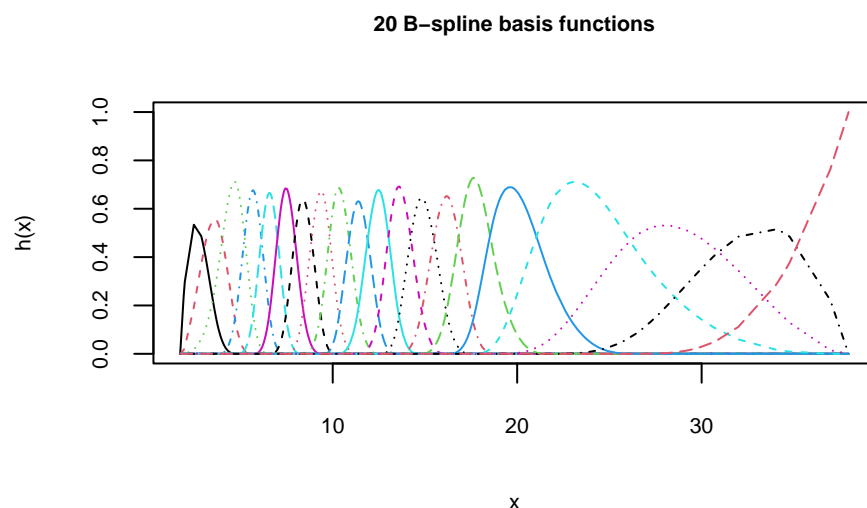


B-spline bases

```
library("MASS")
data(Boston)
plot(Boston$lstat, Boston$medv, xlab = "x", ylab = "y", cex = .5, cex.axis=.7,
      cex.lab = .7)
```



```
library("splines")
bs_x <- bs(Boston$lstat, df = 20)
matplot(Boston$lstat[order(Boston$lstat)], bs_x[order(Boston$lstat), ],
        type = "l", xlab = "x", ylab = "h(x)",
        main = "20 B-spline basis functions",
        cex.main = .7, cex.axis=.7, cex.lab = .7)
```



Note that by default, knots are placed based on quantiles (so more knots in areas with higher density).

Estimate coefficients (in an unpenalized manner) and inspect the fitted spline curve:

```
library("gam")
mod_df20 <- gam(medv ~ bs(lstat, df = 20), data = Boston)
coef(mod_df20)
```

```
##      (Intercept) bs(lstat, df = 20)1 bs(lstat, df = 20)2
##      47.084153    -7.542724         -1.107027
## bs(lstat, df = 20)3 bs(lstat, df = 20)4 bs(lstat, df = 20)5
##     -15.832894    -20.156822         -21.295745
## bs(lstat, df = 20)6 bs(lstat, df = 20)7 bs(lstat, df = 20)8
##     -17.657272    -27.771313         -18.138844
## bs(lstat, df = 20)9 bs(lstat, df = 20)10 bs(lstat, df = 20)11
##     -29.350675    -21.835623         -29.433491
## bs(lstat, df = 20)12 bs(lstat, df = 20)13 bs(lstat, df = 20)14
##     -24.612153    -28.927923         -30.660864
## bs(lstat, df = 20)15 bs(lstat, df = 20)16 bs(lstat, df = 20)17
##     -30.524051    -30.998140         -35.464397
## bs(lstat, df = 20)18 bs(lstat, df = 20)19 bs(lstat, df = 20)20
##     -34.369267    -36.984612         -34.627202
```

```
plot(mod_df20, residuals = TRUE, col = "blue", cex = .5,
      cex.axis=.7, cex.lab = .7)
```

