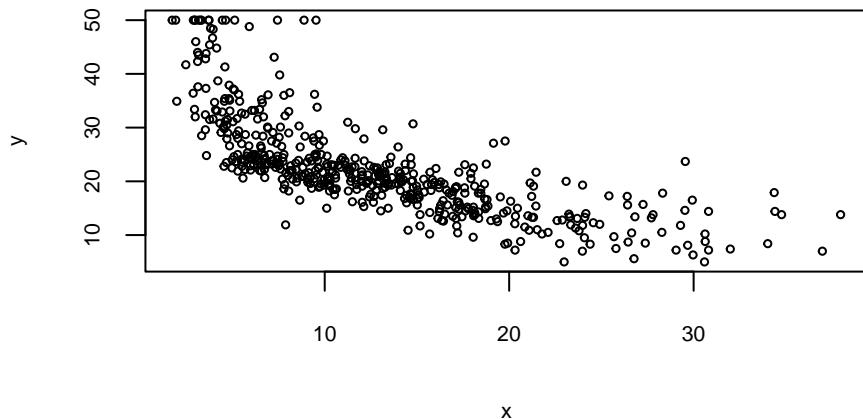


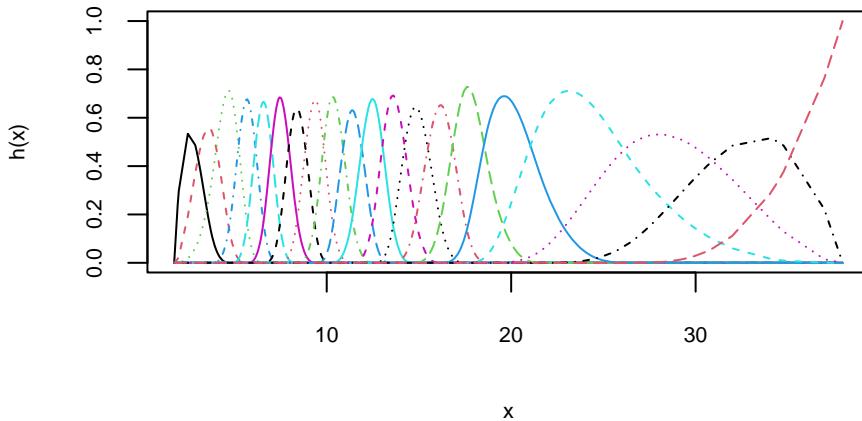
B-spline bases

```
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)
```

20 B-spline basis functions



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)
```

