

# STATISTICS 538, Lecture #12

## From lines to curves

December 1, 2010

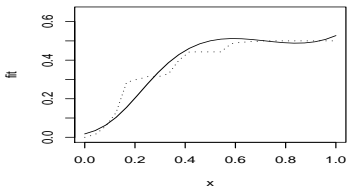
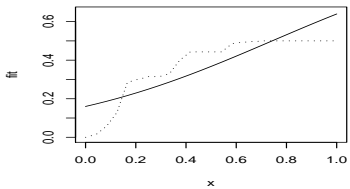
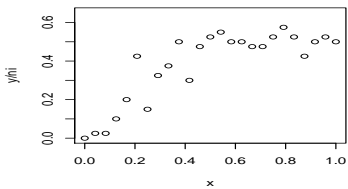
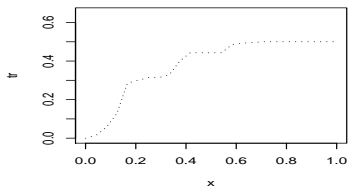
# GLM: linear predictor is linear in **parameters**, not necessarily linear in **predictors**

E.g.,  $Y$  indicates death,  $X$  indicates dose of nasty chemical. First thought

$$Pr(Y = 1|X) = \text{expit}(\beta_0 + \beta_1 X)$$

No good? Change link function? Or expand linear predictor from  $\eta = \beta_0 + \beta_1 X$  to something more complex (but still linear in parameters).

Consider this true relationship spawning these data, and two fits



# Alternative to polynomials? Splines!

**Cubic splines:** Choose **knots**  $c_1, \dots, c_q$ . Consider piecewise cubic functions (between knots) with smooth joins (at knots)

Splines have 'basis function' representations, i.e., linear in parameters

## So 'cheap homemade' spline fitting...

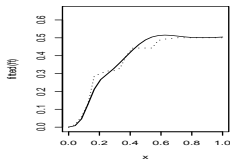
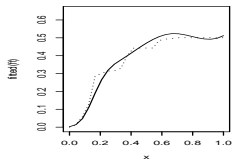
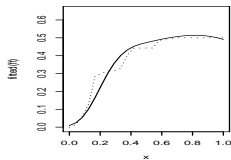
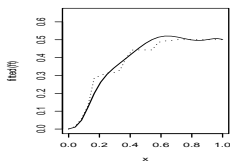
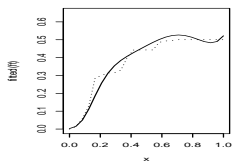
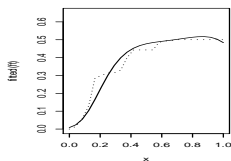
```
xmat <- cbind(x,x^2,x^3)

for (j in 1:q) {
  xmat <- cbind(xmat, pmax(x - quantile(x,j/(q+1)),0)^3)
}

ft <- glm(cbind(y,ni-y) ~ xmat, family=binomial)

plot(x, fitted(ft)) ...
```

# Add one, two, three parameters beyond the cubic model: polynomials (top) versus splines (bottom)



# Further issues

Model selection: how many knots?

More refined splines: **natural cubic splines**



## Further issues, continued

Lots of curve-fitting packages - don't need to do-it-yourself

```
> library(splines)
> ft <- glm(cbind(y,ni-y)~ ns(x,df=4), family=binomial)
```