# Statistics 360: Advanced R for Data Science MARS, part V

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### More details on the implementation

- So far we have discussed the formula and data inputs, and pre-processing.
- ► Today we'll talk about the mars.control object input, some hints about the forward algorithm, and packaging the output as a mars object.
- You will implement the forward and backward algorithms yourself.

#### mars.control object as input

- ► From the mars4 lecture we noted that mars() should take an object of class mars.control as input.
- ► The elements of a mars.control object are Mmax, d and trace.
- Write a constructor, validator and helper for this class.
- Helper:
  - Use default values Mmax=2, d=3, trace=FALSE.
  - ► Mmax should be an even integer ≥ 2. If not, coerce user's input and throw a warning.
- Validator:
  - ► Check that Mmax is an even integer ≥ 2, d is numeric, and trace is logical.
- Constructor:
  - Its input is a list, its output is an object of class mars.control.

#### Hints on fwd\_stepwise()

- ➤ You should be working on a first draft of the fwd\_stepwise() function, as a modification of your recpart\_fwd() algorithm from lab 5.
- ► The key modifications are
  - Replace the indicator functions H() with mirror-image hinge functions h().
  - Do not remove parent basis function  $B_m(x)$  after it is split. We are therefore adding **pairs** of basis functions in each outer-loop iteration.
  - Only split basis function  $B_m(x)$  into  $B_m(x)h(t-x_v)$  and  $B_m(x)h(x_v-t)$  for variables  $x_v$  not already involved in  $B_m$ .

### Indicators to hinge functions

- ► This modification is relatively straightforward.
- Your hinge function h() should take arguments s (sign), x (variable) and t (knot) and return the maximum of 0 and s\*(x-t) for each value of x.
- ▶ You may find the pmax() function useful.

#### Adding pairs of basis functions

- ► Can initialize B as in recpart\_fwd(), but we'll initialize Bfuncs to be of length Mmax+1 to match B.
  - ▶ Bfuncs <- vector(mode="list",length=Mmax+1)
  - ▶ Recall: B has Mmax+1 columns, including the intercept B0.
- ► Replace the for loop over basis functions M from 1 to Mmax with a loop over **pairs** 1 to Mmax/2.
- When adding pair i, there are currently M=2\*i−1 basis functions to consider splitting.
  - ► E.G., when you add pair 1 there is 2\*1-1=1 (the constant basis function), when you add pair 2 there are currently 2\*2-1=3 (the constant and the first pair), etc.
- As in recpart\_fwd() when you loop over basis functions, variables and split points, keep track of lof\_best and the best m, v and t. When you finish the three loops, construct left-and right-split data frames from Bfuncs[[m]] and add this pair to the B matrix and the Bfuncs list.
  - The indices of pair i in B and Bfuncs are 2\*i and 2\*i+1.

### Restriction on splitting

- Only split basis function  $B_m(x)$  into  $B_m(x)h(+(x_v-t))$  and  $B_m(x)h(-(x_v-t))$  for variables  $x_v$  not already involved in  $B_m$ .
- ▶ You will need to consult the Bfuncs[[m]] to see which basis functions make up  $B_m$ .

## Value/output

- object of S3 class mars.
- inherits from class lm and includes all of the components of the lm() from the final fit
  - Use c() to combine these with any of your own components.
  - Note: bwd\_selection() will select the best model but will not return the fit. You will need to call lm() after bwd selection() to obtain the final fit.
- include Bfuncs data structure from final fit.
- write a constructor for class mars no need for a validator or helper since you are the only one who will call the constructor.

#### Methods

- Use methods() to find a list of methods implemented for the S3 class lm.
- Write more informative print and summary methods for mars objects
- Write a plot method.
  - ► The details are up to you, but you should consult Section 3.5 of the Friedman paper (ANOVA decomposition).
  - ➤ Two sources of inspiration are the plot.earth method for earth objects (see the earth package), and plot.Gam for plotting generalized additive model components (see the gam package).
- Write a predict method with the same interface as predict.lm.
- You can use the residuals(), fitted(), hatvalues() and other methods for lm objects for methods that depend only on the final lm.