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` and `Bfuncs` as in `recpart_fwd()`.
  - Note: `B` has `Mmax+1` columns, including the intercept `B0`, and `Bfuncs` has `Mmax` elements
- 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 `Bfuncs` list.
  - The indices of pair `i` in `Bfuncs` are `2*i-1` and `2*i`.

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`.