

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 $M_{\max}+1$ columns, including the intercept `B0`, and `Bfuncs` has M_{\max} elements
- ▶ Replace the for loop over basis functions `M` from 1 to M_{\max} with a loop over **pairs** 1 to $M_{\max}/2$.
- ▶ 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 `B` are $2*i$ and $2*i+1$. and 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 `lm` 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`.