Package ‘sealasso’

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Title Standard Error Adjusted Adaptive Lasso
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Depends R (>= 2.9.1), lars
Description Standard error adjusted adaptive lasso (SEA-lasso) is a version of the adaptive lasso, which incorporates OLS standard error to the L1 penalty weight. This method is intended for variable selection under linear regression settings (n > p). This new weight assignment strategy is especially useful when the collinearity of the design matrix is a concern.
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R topics documented:

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sealasso Standard Error Adjusted Adaptive Lasso

Description
This provides the solution path of some variants of the adaptive lasso (e.g. SEA-lasso, NSEA-lasso, OLS-adaptive lasso), and the corresponding optimal model selected by BIC criterion.
Usage

sealasso(x, y, method = c("nsealasso", "sealasso", "olsalasso", "lasso"))

Arguments

x                  The model matrix.
y                  The response.
method             One of "nsealasso", "sealasso", "olsalasso" and "lasso", which represent NSEA-lasso, SEA-lasso, OLS-adaptive lasso and the lasso, respectively. The default is "nsealasso".

Details

SEA-lasso and NSEA-lasso (Qian and Yang, 2010) are two versions of the adaptive lasso. They may be used for variable selection, especially in cases where condition index of the scaled model matrix is large (e.g. > 10) and collinearity is a concern. This function provides condition index, solution path and the suggested optimal model based on BIC. The estimated coefficients are also given for transition points of the path.

Value

method             The method used.
condition.index    Condition index of the scaled model matrix.
path               Solution path and corresponding BIC values at transition points.
beta               The estimated coefficients at transition points of solution path.
optim.beta         The estimated coefficients of the optimal model based on BIC criterion.

Author(s)

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References


Examples

# use the diabetes dataset from "lars" package
data(diabetes)
x <- diabetes$x
y <- diabetes$y
sealasso(x, y)

# with quadratic terms
x2 <- cbind(diabetes$x1, diabetes$x2)
object <- sealasso(x2, y, "sealasso")
**summary.sealasso**

The `summary.sealasso` function provides a summary method for `sealasso` objects. It prints out the method used, condition index and the optimal model selected by BIC.

### Usage

```r
## S3 method for class 'sealasso'
summary(object, ...)  
```

### Arguments

- `object`: a `sealasso` object
- `...`: other arguments to be passed to `summary`

### Details

A summary is provided for a `sealasso` object. To provide a more succinct output, only the method used, condition index and the selected optimal model are printed out.

### Value

- `method`: The method used. One of NSEA-lasso, SEA-lasso, OLS-adaptive lasso and Lasso.
- `condition.index`: Condition index of the scaled model matrix.
- `optim.beta`: The estimated coefficients of the optimal model based on BIC criterion.

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### References


### See Also

`sealasso`
Examples

```r
# use the diabetes dataset from "lars" package
data(diabetes)
y <- diabetes$y
# with quadratic terms
x2 <- cbind(diabetes$x1, diabetes$x2)
object <- sealasso(x2, y, "sealasso")
summary(object)
```
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