

# Kernel-Based Learning & Multivariate Modeling

## MIRI Master

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Use the R language with the package `kernlab`. You have to download the package, install it (once) and load it into R (every time you use it):

```
> install.packages('kernlab')  
> library(kernlab)
```

Notes:

1. For hyper-parameter selection (e.g.,  $C$ ,  $\varepsilon$ , ...), you should use standard resampling methods, like cross-validation.
2. Many times it is helpful to standardize the data prior to doing anything. I suggest to do it as a preprocess, using the `scale` function in R) –then you have to “deactivate” automatic scaling in the `ksvm` method by doing `ksvm(..., scaled=c(), ...)`.

### Problem 1 The SVM for regression in action

Use the `ksvm` method to perform SVM regression on some data set of your choice (as typical examples, use the Concrete Compressive Strength data set<sup>1</sup> or the Yacht Hydrodynamics data set<sup>2</sup>).

If you wish, you can also use standard multivariate regression methods, like (ridge) regression (`lm.ridge` in R); you can also use kernel (ridge) regression for a comparison against linear methods.

The SVM regression is found in the `ksvm` method (be sure to specify `eps-svr` for the 'type' parameter). Draw conclusions on your results.

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### Problem 2 The SVM for classification in action

Use the `ksvm` method to perform SVM classification on some data set of your choice (as typical examples, use the Ionosphere data set<sup>3</sup> or the Breast Cancer Wisconsin (Diagnostic) data set<sup>4</sup>).

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<sup>1</sup><http://archive.ics.uci.edu/ml/datasets/Concrete+Compressive+Strength>

<sup>2</sup><http://archive.ics.uci.edu/ml/datasets/Yacht+Hydrodynamics>

<sup>3</sup><http://archive.ics.uci.edu/ml/datasets/Ionosphere>

<sup>4</sup>[http://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+\(Diagnostic\)](http://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+(Diagnostic))

As less typical (and more challenging) examples, you can use the Pen-Based Recognition of Handwritten Digits data set<sup>5</sup> or the Splice (DNA sequences) data set<sup>6</sup>.

If you wish, you can also use standard multivariate classification methods, like discriminant analysis (`lda/qda` methods in `R`); you can also use logistic regression (`glm` method in `R`, for two classes) or multinomial regression (`multinom` (`nnet`) method, for more than two) for a comparison against linear methods. Draw conclusions on your results.

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<sup>5</sup><http://archive.ics.uci.edu/ml/datasets/Pen-Based+Recognition+of+Handwritten+Digits>

<sup>6</sup><http://www.cs.toronto.edu/~delve/data/splice/desc.html>