# Introduction to Machine Learning NPFL 054

http://ufal.mff.cuni.cz/course/npf1054

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- library(e1071), but there are also other libraries (kernlab, shogun ...)
- training: function svm()
- prediction: function predict()
- svm() can work in both classification and regression mode
- if target attribute (response variable) is categorical (factor) the engine switches to classification

### model = svm(formula, data=, kernel=, cost=, cross=, ...)

- ?svm
- kernel defines the kernel used in training and prediction. The options are: linear, polynomial, radial basis and sigmoid, default = radial
- cost cost of constraint violation (default: 1)
- cross optional, with the value k the k-fold cross-validation is performed

Kernel name	Formula	Learning parameters and their default values		
linear	$\mathbf{x}_i \cdot \mathbf{x}_j$			
polynomial	$(\gamma \mathbf{x}_i \cdot \mathbf{x}_j + c)^d$	$\gamma$ , gamma=1/(data dimension) c, coef0=0 d, degree=3		
radial	$\exp(-\gamma(  \mathbf{x}_i - \mathbf{x}_j  ^2))$	$\gamma$ , gamma=1		
sigmoid	$ anh(\gamma \mathbf{x}_i \cdot \mathbf{x}_j + c)$	$\gamma$ , gamma=1/(data dimension) c, coef0=0		

- polynomial kernel
  - smaller degree can generalize better
  - higher degree can fit (only) training data better
- radial basis
  - very robust
  - you should try and use it when polynomial kernel is weak to fit your data

## SVM Parameter tuning with tune.svm

- SVM is a more complicated method in comparison with the previous and usually requires parameter tuning!
- parameter tuning can take a very long time on big data, use a reasonably smaller part is often recommended

```
> model.tune= tune.svm(class ~ ., data=train.small,
                       kernel = "radial",
                       gamma = c(0.001, 0.005, 0.01, 0.015, 0.02),
                       cost = c(0.5, 1, 5, 10))
> model.tune
Parameter tuning of 'svm':
 sampling method: 10-fold cross validation
 best parameters:
gamma cost
 0.01 1
 best performance: 0.739
```

### K-fold cross-validation

#### parameter cross

 class.weights parameter
 In case of asymmetric class sizes you may want to avoid possibly overproportional influence of bigger classes. Weights may be specified in a vector with named components, like
 m <- svm(x, y, class.weights = c(A = 0.3, B = 0.7))</li>

- Note that SVMs may be very sensible to the proper choice of parameters, so always check a range of parameter combinations, at least on a reasonable subset of your data.
- Be careful with large datasets as training times may increase rather fast.
- C-classification with the RBF kernel (default) can often be a good choice because of its good general performance and the few number of parameters (only two: cost and gamma).
- When you use C-classification with the RBF kernel: try small and large values for cost first, then decide which are better for the data by cross-validation, and finally try several gamma values for the better cost.

target	True	False	False	class			F1
class	Positive	Positive	Negative	weight	Precision	Recall	score
C <sub>1</sub>	$TP_1$	$FP_1$	$FN_1$	$W_1$	$P_1$	$R_1$	$F_1$
C <sub>2</sub>	$TP_2$	$FP_2$	$FN_2$	W2	$P_2$	$R_2$	$F_2$
 C	 TP <i>k</i>	 FP,	 FN⊭		 P,	 P.	 E.
$C_k$	IFk	Γ <b>Γ</b> k	r ink	$W_k$	гk	$R_k$	$F_k$

- class weight w<sub>i</sub> is the relative frequency of C<sub>i</sub> class in the data
- macro-averaged F1 score =  $\sum_{i=1}^{k} F_i/k$
- weighted-averaged F1 score =  $\sum_{i=1}^{k} w_i F_i / k$

- In general, if you are working with an imbalanced dataset where all classes are equally important, using the macro average would be a good choice.
- If you have an imbalanced dataset but want to assign greater contribution to classes with more examples in the dataset, then the weighted average is preferred.