

Multiple Kernel Learning

(via Semi-Infinite Linear Programming)

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und Softwaretechnik

- 1 Introduction and Motivation
 - When is Multiple Kernel Learning useful?

- 2 Multiple Kernel Learning
 - Deriving the Semi-Infinite Linear Program
 - MKL SILP Algorithm

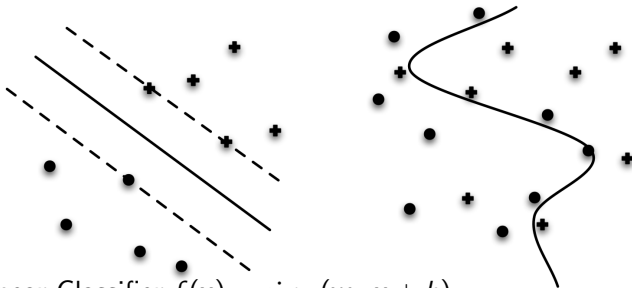
- 3 Applications, Extensions, Outlook
 - Extensions
 - Applications
 - Outlook

Outline

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Classification

Given training examples $(\mathbf{x}_i, y_i)_{i=1}^N \in (\mathcal{X}, \{-1, +1\})^N$



- Linear Classifier $f(\mathbf{x}) = \text{sign}(\mathbf{w} \cdot \mathbf{x} + b)$
- Kernel Machine (e.g. Support Vector Machine), learn weighting $\alpha \in \mathbb{R}^N$ on training examples in kernel feature space $\Phi(\mathbf{x})$

$$f(\mathbf{x}) = \text{sign} \left(\sum_{i=1}^N y_i \alpha_i k(\mathbf{x}, \mathbf{x}_i) + b \right),$$

where **Kernel** $k(\mathbf{x}, \mathbf{x}') = \Phi(\mathbf{x}) \cdot \Phi(\mathbf{x}')$

Classification using Kernel Machines I

Single Kernel

- Kernel Machine (e.g. Support Vector Machine)
 - learn weighting $\alpha \in \mathbb{R}^N$ on training examples $(\mathbf{x}_i, y_i)_{i=1}^N$ in kernel feature space $\Phi(\mathbf{x})$

$$f(\mathbf{x}) = \text{sign} \left(\sum_{i=1}^N y_i \alpha_i k(\mathbf{x}, \mathbf{x}_i) + b \right)$$

- where Kernel $k(\mathbf{x}, \mathbf{x}') = \Phi(\mathbf{x}) \cdot \Phi(\mathbf{x}')$
- still linear classifier $f(\mathbf{x}) = \text{sign}(\mathbf{w} \cdot \mathbf{x} + b)$ in kernel feature space, with weighting $\mathbf{w} = \sum_{i=1}^N y_i \alpha_i \Phi(\mathbf{x}_i)$ and examples $\mathbf{x} \mapsto \Phi(\mathbf{x})$

via **kernel: non-linear** discrimination in input space

Classification using Kernel Machines II

Multiple Kernels

- Linear combination of kernels $k(\mathbf{x}, \mathbf{x}') = \sum_{j=1}^M \beta_j k_j(\mathbf{x}, \mathbf{x}')$, $\beta_j \geq 0$. Learn α and β . Resulting classifier:

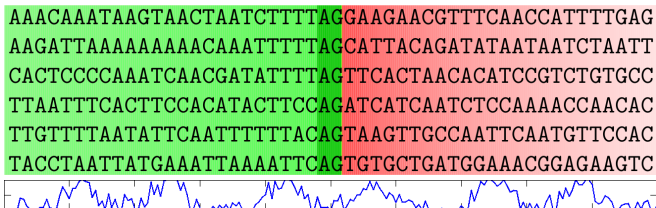
$$f(\mathbf{x}) = \text{sign} \left(\sum_{j=1}^M \beta_j \sum_{i=1}^N y_i \alpha_i k_j(\mathbf{x}, \mathbf{x}_i) + b \right)$$

Learn weighting over training examples α and kernels β

When is Multiple Kernel Learning useful?

Combining Heterogeneous Data

- Consider data from different domains: e.g. Bioinformatics features: DNA-strings, binding energies, conservation, structure,...



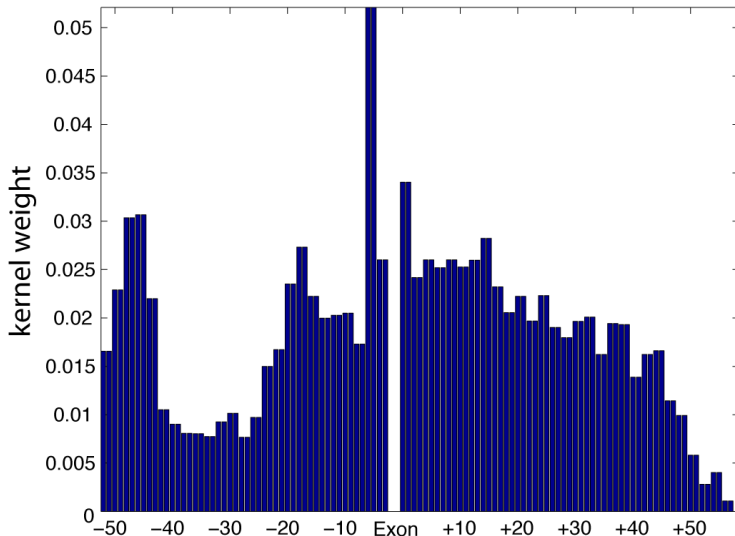
$$k(\mathbf{x}, \mathbf{x}') =$$

$$\beta_1 k_{dna}(\mathbf{x}_{dna}, \mathbf{x}'_{dna}) + \beta_2 k_{nrg}(\mathbf{x}_{nrg}, \mathbf{x}'_{nrg}) + \beta_3 k_{3d}(\mathbf{x}_{3d}, \mathbf{x}'_{3d}) + \dots$$

When is Multiple Kernel Learning useful?

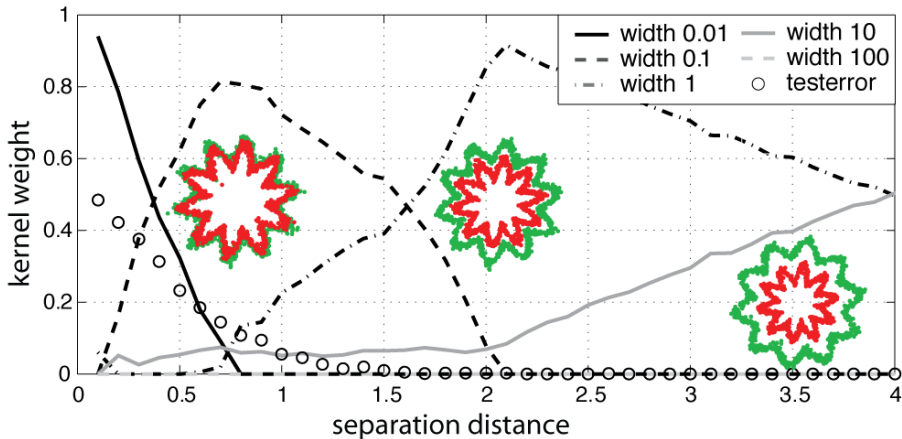
Interpretability

- Bioinformatics: One weight per position in sequence



When is Multiple Kernel Learning useful?

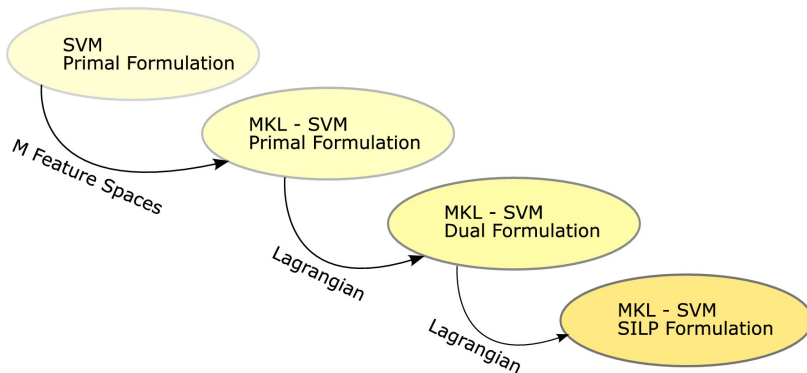
Automated Model Selection



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Derivation



For details see **Sonnenburg, Rätsch, Schäfer, Schölkopf 2006**

SVM Primal Formulation

$$\begin{aligned} \min \quad & \frac{1}{2} \|\mathbf{w}\|_2^2 + C \sum_{i=1}^N \xi_i \\ \text{w.r.t.} \quad & \mathbf{w} \in \mathbb{R}^D, \boldsymbol{\xi} \in \mathbb{R}_+^N, b \in \mathbb{R} \\ \text{s.t.} \quad & y_i \left(\mathbf{w}^\top \Phi(\mathbf{x}_i) + b \right) \geq 1 - \xi_i, \forall i = 1, \dots, N \end{aligned}$$

MKL Primal Formulation

$$\min \quad \frac{1}{2} \left(\sum_{j=1}^M \beta_j \|\mathbf{w}_j\|_2 \right)^2 + C \sum_{i=1}^N \xi_i$$

$$\text{w.r.t.} \quad \mathbf{w} = (\mathbf{w}_1, \dots, \mathbf{w}_M), \mathbf{w}_j \in \mathbb{R}^{D_j}, \quad \forall j = 1 \dots M$$

$$\beta \in \mathbb{R}_+^M, \xi \in \mathbb{R}_+^N, b \in \mathbb{R}$$

$$\text{s.t.} \quad y_i \left(\sum_{j=1}^M \beta_j \mathbf{w}_j^T \Phi_j(\mathbf{x}_i) + b \right) \geq 1 - \xi_i, \quad \forall i = 1, \dots, N$$

$$\sum_{j=1}^M \beta_j = 1$$

Properties: equivalent to SVM for $M = 1$; solution sparse in “blocks”; each block j corresponds to one kernel

MKL Dual Formulation

Bach, Lanckriet, Jordan 2004:

$$\begin{aligned} \min \quad & \gamma - \sum_{i=1}^N \alpha_i \\ \text{w.r.t.} \quad & \gamma \in \mathbb{R}, \boldsymbol{\alpha} \in \mathbb{R}^N \\ \text{s.t.} \quad & 0 \leq \boldsymbol{\alpha} \leq C, \sum_{i=1}^N \alpha_i y_i = 0 \\ & \frac{1}{2} \sum_{r=1}^N \sum_{s=1}^N \alpha_r \alpha_s y_r y_s K_j(\mathbf{x}_r, \mathbf{x}_s) - \gamma \leq 0, \forall j = 1, \dots, M \end{aligned}$$

Properties: equivalent to SVM for $M = 1$

The Semi-Infinite Linear Program I

$$\begin{aligned}
 \max \quad & \theta \\
 \text{w.r.t.} \quad & \theta \in \mathbb{R}, \beta \in \mathbb{R}_+^M \text{ with } \sum_{j=1}^M \beta_j = 1 \\
 \text{s.t.} \quad & \sum_{j=1}^M \beta_j \left(\underbrace{\frac{1}{2} \sum_{r=1}^N \sum_{s=1}^N \alpha_r \alpha_s y_r y_s K_j(\mathbf{x}_r, \mathbf{x}_s)}_{=: S_j(\alpha)} - \sum_{i=1}^N \alpha_i \right) \geq \theta \\
 & \text{for all } \alpha \text{ with } 0 \leq \alpha \leq C \text{ and } \sum_{i=1}^N y_i \alpha_i = 0
 \end{aligned}$$

Properties:

- linear, optimize over a convex combination of β
- infinitely many constraints, one for each $0 \leq \alpha \leq C$
- can use standard SVM to identify most violated constraints

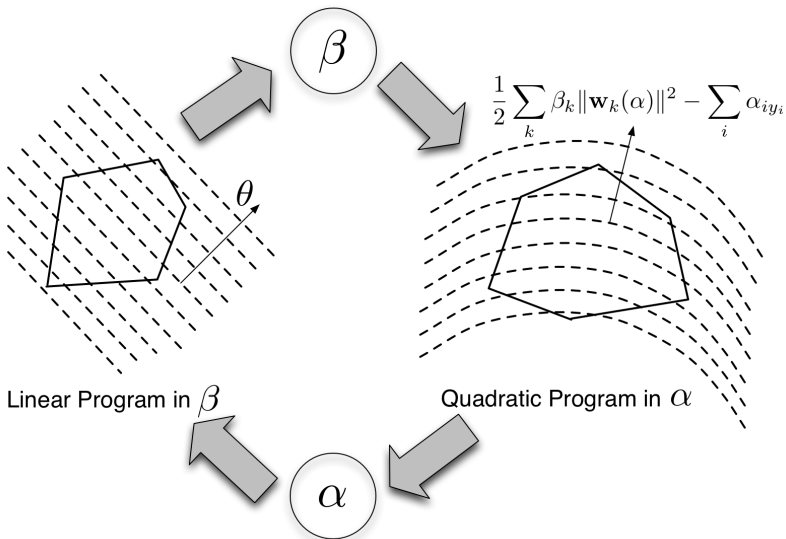
The Semi-Infinite Linear Program II

$$\begin{aligned} \max \quad & \theta \\ \text{w.r.t.} \quad & \theta \in \mathbb{R}, \boldsymbol{\beta} \in \mathbb{R}_+^M \text{ with } \sum_{j=1}^M \beta_j = 1 \\ \text{s.t.} \quad & \sum_{j=1}^M \beta_j \left(\frac{1}{2} S_j(\boldsymbol{\alpha}) - \sum_{i=1}^N \alpha_i \right) \geq \theta \\ & \text{for all } \boldsymbol{\alpha} \text{ with } 0 \leq \boldsymbol{\alpha} \leq C \text{ and } \sum_{i=1}^N y_i \alpha_i = 0 \end{aligned}$$

Solving the SILP:

- Column Generation
 - fast, but no known convergence rate
- Use Boosting like techniques: Arc-GV or AdaBoost*
 - known convergence rate $\mathcal{O}(\log(M)/\varepsilon^2)$
- Chunking like algorithm
 - consider suboptimal SVM solutions: empirically 3-5 times faster

Solving the SILP: Column Generation I



Solving the SILP: Column Generation II

$$\begin{array}{ll}
 \max & \theta \\
 \text{w.r.t.} & \theta \in \mathbb{R}, \boldsymbol{\beta} \in \mathbb{R}_+^M \text{ with } \sum_{j=1}^M \beta_j = 1 \\
 \text{s.t.} & \sum_{j=1}^M \beta_j \left(\frac{1}{2} S_j(\boldsymbol{\alpha}) - \sum_{i=1}^N \alpha_i \right) \geq \theta \\
 & \text{for all } \boldsymbol{\alpha} \text{ with } 0 \leq \boldsymbol{\alpha} \leq C \text{ and } \sum_{i=1}^N y_i \alpha_i = 0
 \end{array}$$

- iteratively find most violated constraints, solve linear program with current constraints, ..., till convergence to the global optimum

$$\sum_{j=1}^M \beta_j \left(\frac{1}{2} S_j(\boldsymbol{\alpha}) - \sum_{i=1}^N \alpha_i \right) = \frac{1}{2} \sum_{r=1}^N \sum_{s=1}^N \alpha_r \alpha_s y_r y_s \sum_{j=1}^M \beta_j k_j(\mathbf{x}_r, \mathbf{x}_s) - \sum_{i=1}^N \alpha_i,$$

- solved by taking set of most violated constraints into account
- most violated constraints given by SVM solution for fixed $\boldsymbol{\beta}$

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Regression

Primal Formulation:

$$\min \quad \frac{1}{2} \left(\sum_{j=1}^M \beta_j \|\mathbf{w}_j\| \right)^2 + C \sum_{i=1}^N (\xi_i + \xi_i^*)$$

$$\text{w.r.t.} \quad \mathbf{w} = (\mathbf{w}_1, \dots, \mathbf{w}_M), \mathbf{w}_j \in \mathbb{R}^{D_j}, \quad \forall j = 1 \dots M$$

$$\beta \in \mathbb{R}_+^M, \xi \in \mathbb{R}^N, \xi^* \in \mathbb{R}_+^N, b \in \mathbb{R}$$

$$\text{s.t.} \quad \sum_{j=1}^M \beta_j \mathbf{w}_j^T \Phi_j(\mathbf{x}_i) + b \leq y_i + \varepsilon + \xi_i, \quad \forall i = 1 \dots N$$

$$\sum_{j=1}^M \beta_j \mathbf{w}_j^T \Phi_j(\mathbf{x}_i) + b \geq y_i - \varepsilon - \xi_i^*, \quad \forall i = 1 \dots N$$

$$\sum_{j=1}^M \beta_j = 1$$

One Class

Primal Formulation:

$$\min \quad \frac{1}{2} \left(\sum_{j=1}^M \beta_j \|\mathbf{w}_j\|_2 \right)^2 + C \sum_{i=1}^N \xi_i - \rho$$

$$\text{w.r.t.} \quad \mathbf{w} = (\mathbf{w}_1, \dots, \mathbf{w}_M), \mathbf{w}_j \in \mathbb{R}^{D_j}, \quad \forall j = 1 \dots M$$

$$\beta \in \mathbb{R}_+^M, \quad \boldsymbol{\xi} \in \mathbb{R}_+^N$$

$$\text{s.t.} \quad y_i \left(\sum_{j=1}^M \beta_j \mathbf{w}_j^\top \Phi_j(\mathbf{x}_i) \right) \geq \rho - \xi_i, \forall i = 1, \dots, N$$

$$\sum_{j=1}^M \beta_j = 1$$

Generalized for arbitrary strictly convex differentiable loss functions (Sonnenburg, Rätsch, Schäfer, Schölkopf 2006)

Multiclass

Primal Formulation (Zien, Ong 2007):

$$\min \quad \frac{1}{2} \left(\sum_{j=1}^M \beta_j \|\mathbf{w}_j\|_2 \right)^2 + C \sum_{i=1}^N \xi_n$$

$$\text{w.r.t.} \quad \mathbf{w} = (\mathbf{w}_1, \dots, \mathbf{w}_M), \mathbf{w}_j \in \mathbb{R}^{k_j}, \quad \forall j = 1 \dots M$$

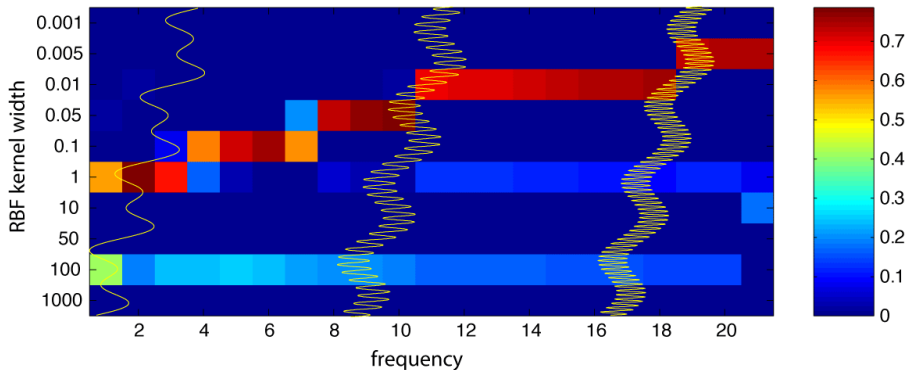
$$\beta \in \mathbb{R}_+^M, \mathbf{s} \in \mathbb{R}^{N \times c}, \boldsymbol{\xi} \in \mathbb{R}_+^N, \mathbf{b} \in \mathbb{R}$$

$$\text{s.t.} \quad \xi_i = \max_{u \neq y_i} s_{iu}, \quad s_{iu} \geq 0,$$

$$\sum_{j=1}^M \beta_j \mathbf{w}_j^T (\Phi_j(\mathbf{x}_i, y_i) - \Phi_j(\mathbf{x}_i, u)) + b_{y_i} - b_u \geq 1 - s_{iu},$$

$$\forall i = 1 \dots N, \forall u = 1 \dots c$$

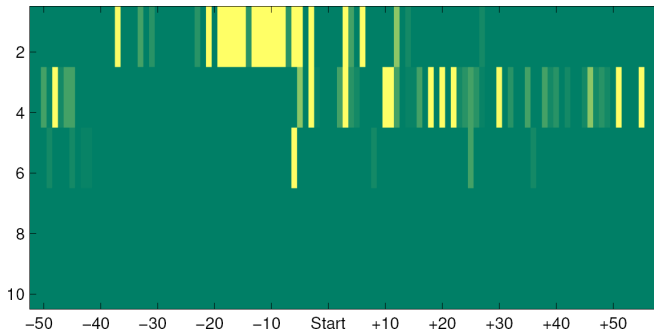
Automated Model Selection - Regression



- $f(x) = \sin(ax) + \sin(bx) + cx$ for varying a
- Support Vector Regression with 10 RBF-Kernels of different width

Knowledge discovery

Feature Extraction



- Support Vector Classification on Bioinformatics problem, distinguish “splice sites” from “fake sites” (aligned DNA sequences)
- One weight β_j per position and per sub-sequence length
- Displayed: Learned weights of 500 kernels

Summary and Outlook

MKL learns convex combination of kernels

- ⇒ allows (to some extent) for automated model selection
- ⇒ allows for interpreting SVM result
- ⇒ matches prior knowledge on real-world bioinformatics problem

- **Simple:** iterative wrapper algorithm around single kernel SVM
- **General:** same technique applicable to a wide range of problems (1-class, 2-class, Multiclass, Regression, ...)
- **Fast:** suitable for large scale problems ($> 100,000$ examples)

Download free source <http://www.shogun-toolbox.org>.