(lets do it large scale with shogun)

Sören Sonnenburg

Shogun Toolbox Data Days 2013, Juli 12, Berlin

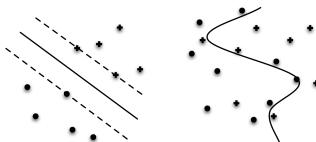


- SVMs
- 2 Large Scale SVMs
- 3 Applications
- 4 MKL
- MKL Extensions

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Classification

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



- Linear Classifier $f(\mathbf{x}) = \operatorname{sign}(\mathbf{w} \cdot \mathbf{x} + b)$
- Kernel Machine (e.g. Support Vector Machine), learn weighting $\alpha \in \mathbb{R}^N$ on training examples in kernel feature space $\Phi(\mathbf{x})$ $f(\mathbf{x}) = \operatorname{sign}\left(\sum_{i=1}^{N} y_i \alpha_i \mathbf{k}(\mathbf{x}, \mathbf{x}_i) + b\right),$

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

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Classification using Support Vector Machines

Via kernel: non-linear discrimination in input space

• learn weighting $\alpha \in \mathbb{R}^N$ on training examples $(\mathbf{x}_i, y_i)_{i=1}^N$ in kernel feature space $\Phi(\mathbf{x})$

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

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

SVMs rock!

- Kernels flexible!
- In many applications SVMs define the state-of-the-art!

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Support Vector Machines in Shogun I

Implementations supporting Kernels

- LibSVM, LibSVMOneClass, MulticlassLibSVM, LibSVR (sequential minimal optimization)
- GNPPSVM (ℓ₂-SVM)
- GPBTSVM (chunking algorithm based SVM)
- GMNPSVM (true multi-class SVM)
- LaRank (true multi-class SVM)
- MPDSVM (minimal primal-dual SVM)
- SVMLight, SVMLightOneClass, SVRLight (chunking based, thread-parallel non-GPL)
- ScatterSVM (multi-class approximation)

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Kernels

Popular Kernels

- Gaussian, Polynomial, String Kernels
- Custom Kernel (bring your own)

Kernel Normalizers

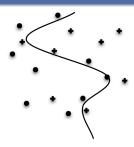
- SqrtDiagKernelNormalizer (points on hypersphere with R=1)
- ZeroMeanCenterKernelNormalizer (centered in feature space)
- AvgDiagKernelNormalizer, more http://bit.ly/133aG2V

Most comprehensive Kernel/SVM Framework

- Over 60 kernels available
- http://bit.ly/15zlCWY
- normalizers can be attached to kernels



Kernel based Support Vector Machines



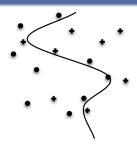
• SVMs learn weights $\alpha \in \mathbb{R}^m$ over training examples in kernel feature space $\Phi: \mathbf{x} \mapsto \mathbb{R}^n$

• Decision function
$$f(\mathbf{x}) = \operatorname{sign} \left(\sum_{i=1}^{m} y_i \alpha_i \mathbf{k}(\mathbf{x}, \mathbf{x}_i) + b \right)$$
, with kernel $k(\mathbf{x}, \mathbf{x}') = \langle \Phi(\mathbf{x}), \Phi(\mathbf{x}') \rangle$

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- But not large scale!

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Kernel based Support Vector Machines are a Dead End

The Curse of Support Vectors

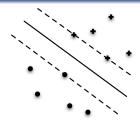
To compute output on all m examples $\mathbf{x}_1, \dots, \mathbf{x}_m$:

$$\forall j = 1, \ldots, m: \sum_{i=1}^{m_s} \alpha_i y_i \, \mathsf{k}(\mathbf{x}_i, \mathbf{x}_j) + b$$

Computational effort:

- All $\mathcal{O}(m_s mT)$, (T time to compute the kernel)
- Effort Scales linearly with $m_s = \mathcal{O}(m) := \#\mathsf{SVs}$
- ⇒ SVM's in bigO are not faster than standard k-NN.
- ⇒ Kernel Machines are just not large-scale!

What about Linear SVMs?

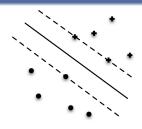


- Linear Support Vector Machines learn weights $\mathbf{w} \in \mathbb{R}^n$
- Decision function $f(x) = \langle w, x \rangle + b$

Recent Progress in Linear SVM solvers

- SGD (Bottou 2007), SGD-QN (Bordes et al., 2009)
- SVM^{perf} (Joachims 2006), liblinear (Fan et al. 2008)
- BMRM (Teo et.al. 2007), OCAS (Franc, Sonnenburg 2009)
- \Rightarrow Linear training Effort $\mathcal{O}(m)$
- \Rightarrow Computing Outputs Linear Effort $\mathcal{O}(nm)$

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- \Rightarrow Computing Outputs Linear Effort $\mathcal{O}(nm)$
- ... already linear time but just linear

Support Vector Machines in Shogun II

Linear SVMs

- OnlineLibLinear, LibLinear, MulticlassLibLinear (dual coordinate descent)
- NewtonSVM (newton primal svm)
- OnlineSVMSGD, SVMSGD (stochastic gradient descent)
- SGDQN (stochastic gradient descent with quasi-newton steps)
- SVMLin
- SVMOcas, MulticlassOCAS (cutting plane/bundle-method)

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0000000000 Linear SVMs

Shogun's computational framework for linear SVMs

AIM: Development of a large scale learning framework for SVMs

"Algorithm [linear SVM solver] improvements do not improve the order of test error convergence. They can simply improve constant factors and therefore compete evenly with the implementation improvements. Time spent refining the implementation is time well spent."

from: Bordes, Bottou, Gallinari: SQD-QN: Careful Quasi-Newton Stochastic Gradient Descent, JMLR 2009.

Towards a computational framework for linear SVMs

Linear SVM solvers like liblinear, SGD, BMRM, OCAS only require two operations to access data:

- (i) dot product between feature vector and the vector $\boldsymbol{w}\colon$
 - $r \leftarrow \langle \mathbf{x}, \mathbf{w} \rangle$

DOT

- (ii) multiplication with scalar $\alpha\in\Re$ and addition to vector $\mathbf{v}\in\Re^n$
 - $\mathbf{v} \leftarrow \alpha \mathbf{x} + \mathbf{v}$

ADD

COFFIN

COFFIN really is just two simple ideas:

On demand compute...

- Features $\Phi(\mathbf{x})$ (only non-zero dims)
 - Non-Linearity Possible
 - Examples: Low Degree Polynomial Kernel, Spectrum Kernel Weighted Degree Kernel
 - On-the-fly (de)compression
- Virtual Examples
 - Incorporating Invariances possible
 - Examples: Image translation, rotation, etc.

Needs efficient data structure for w!

- ..., dense, sorted array, trees, hashes
- fast only when $|\Phi_{\neq 0}(\mathbf{z})| \sim dim(\mathbf{z})$

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Data Structures

Effort of **ADD** and **DOT** for **z** and memory requirement of **w**.

	Dense	Sorted Array	Tree
Add	$\mathcal{O}(\Phi_{\neq 0}(\mathbf{z}))$	$\mathcal{O}(\mathbf{w} _{ eq 0}) + \Phi_{ eq 0}(\mathbf{z}))$	$\mathcal{O}(\Phi_{ eq 0}(\mathbf{z}))$
			to $\mathcal{O}(K \Phi_{ eq 0}(\mathbf{z}))$
Dot	$\mathcal{O}(\Phi_{\neq 0}(\mathbf{z}))$	$\mathcal{O}(\mathbf{w} _{\neq 0}) + \Phi_{\neq 0}(\mathbf{z}))$	$\mathcal{O}(\Phi_{\neq 0}(\mathbf{z}))$
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Mem	$\mathcal{O}(n)$	$\mathcal{O}(\sum_{i=1}^{m} \Phi_{\neq 0}(\mathbf{z}_i))$	$\mathcal{O}(\sum_{i=1}^m \Phi_{\neq 0}(\mathbf{z}_i))$

Sparse data structures have huge overhead!

Hashing to the Rescue (Shi et al (2009))

- Always use dense w with "compressed index"
- Hash function $h(J) \mapsto 1, \dots, 2^{\gamma}$,
- $(\widehat{\Phi}(\mathbf{z}))_j = \sum_{i \in J: h(i)=j} (\Phi(\mathbf{z}))_i$

Data Structures

Effort of **ADD** and **DOT** for **z** and memory requirement of **w**.

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COFFIN is implemented in Shogun's DotFeatures

DotFeatures

- Dense-,SparseFeatures
- PolyFeatures, BinnedDotFeatures
- CombinedDotFeatures
- ExplicitSpecFeatures, ImplicitWeightedSpecFeatures
- HashedDocDotFeatures

You can

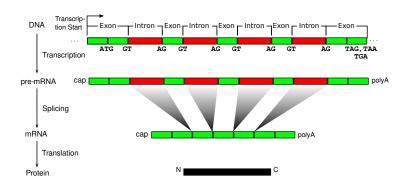
- Stack together features (dense, sparse,...)
- Provides abstraction to data representation

Use it!

Implemented for all linear SVMs



Splice Site Predictions



Application to Human Acceptor Splice Site Prediction

Splice Site Prediction

Discriminate true signal positions against all other positions

 \approx 150 nucleotides window around dimer

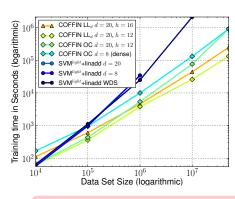
- True sites: fixed window around a true site
- Decoy sites: all other consensus sites

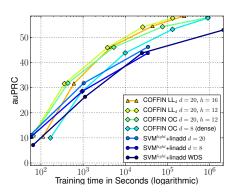
```
AAACAAATAAGTAACTAATCTTTTAGGAAGAACGTTTCAACCATTTTGAG
AAGATTAAAAAAAAACAAATTTTTAGCATTACAGATATAATAATCTAATT
CACTCCCCAAATCAACGATATTTTAGTTCACTAACACACTCCGTCTGTGCC
TTAATTTCACTTCCACATACTTCCAGATCATCAATCTCCAAAACCAACAC
TTGTTTTAATATTCAATTTTTTTACAGTAAGTTGCCAATTCAATGTTCCAC
TACCTAATTATGAAATTAAAATTCAGTGTGCTGATGGAAACGGAGAAGTC
```

- 50 million training examples
- COFFIN with kernels: weighted spectrum and weighted degree (explicit and hashed representation) $\approx 200,000,000$ dims



Applications



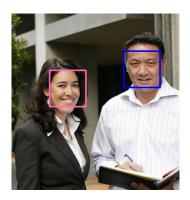


It's fast and works!

- Factor 47 faster on $10 \cdot 10^6$ examples than linadd
- New state-of-the-art results auPRC 58.57% vs. 53.01%

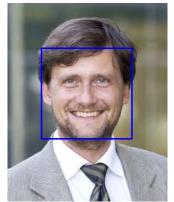
Gender Classification

Distinguish Females from Males solely based on Faces



- learn COFFIN on labelled faces
- virtual examples: translation, rotation, scale
- train \approx 5 million sample (that would require 50GB) on Vojtechs notebook

Gender Classification I



It's fast and works - again!

- auROC 95.44%
- (vs. auROC 89.57% without VE)

Klaus-Robert Müller is a male!

Gender Classification II



Gender Classification III



Gender Classification IV



Summary Support Vector Machines

Kernel SVMs

- Allows non-linear and complex models
- Applicable to mid-size datasets (1e6)
- General and often state-of-the art detectors

Linear SVMs

- Allows non-linearity with COFFIN/DotFeatures
- Applicable to huge datasets
- General and often state-of-the art detectors

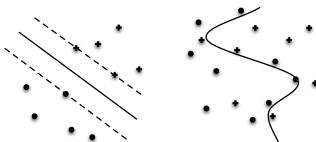
Datasets, Scripts, Efficient implementation

- Data and Scripts http://sonnenburgs.de/soeren/coffin
- Implementation http://www.shogun-toolbox.org
- More machine learning software http://mloss.org



Multiple Kernel Learning

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



- Linear Classifier $f(\mathbf{x}) = \operatorname{sign}(\mathbf{w} \cdot \mathbf{x} + b)$
- Kernel Machine (e.g. Support Vector Machine), learn weighting $\alpha \in \mathbb{R}^N$ on training examples in kernel feature space $\Phi(\mathbf{x})$ $f(\mathbf{x}) = \operatorname{sign}\left(\sum_{i=1}^N y_i \alpha_i \mathbf{k}(\mathbf{x}, \mathbf{x}_i) + b\right),$

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

Single Kernel

SVMs

- 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 $\Phi(\mathbf{x})$

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via kernel: non-linear discrimination in input space

Classification using Kernel Machines II

Multiple Kernels

SVMs

• 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:

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

Learn weighting over training examples lpha and kernels eta

Combining Heterogeneous Data

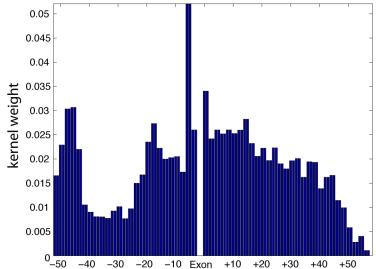
 Consider data from different domains: e.g Bioinformatics features: DNA-strings, binding energies, conservation, structure,...

AAACAAATAAGTAACTAATCTTTTAGGAAGAACGTTTCAACCATTTTGAG AAGATTAAAAAAAAACAAATTTTT<mark>AGCATTACAGATATAAT</mark>AATCTAATT CACTCCCCAAATCAACGATATTTTAGTTCACTAACACATCCGTCTGTGCC TTAATTTCACTTCCACATACTTCCAGATCATCAATCTCCAAAACCAACAC TTGTTTTAATATTCAATTTTTTACAGTAAGTTGCCAATTCAATGTTCCAC TACCTAATTATGAAATTAAAATTC<mark>AGTGTGCTG</mark>ATGGAAACGGAGAGTC

$$\begin{aligned} \mathbf{k}(\mathbf{x}, \mathbf{x}') &= \\ \beta_1 \, \mathbf{k}_{dna}(\mathbf{x}_{dna}, \mathbf{x}'_{dna}) + \beta_2 \, \mathbf{k}_{nrg}(\mathbf{x}_{nrg}, \mathbf{x}'_{nrg}) + \beta_3 \, \mathbf{k}_{3d}(\mathbf{x}_{3d}, \mathbf{x}'_{3d}) + \cdots \end{aligned}$$

Interpretability

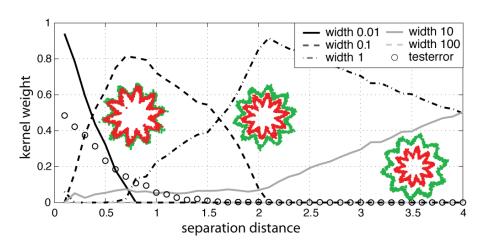
• Bioinformatics: One weight per position in sequence



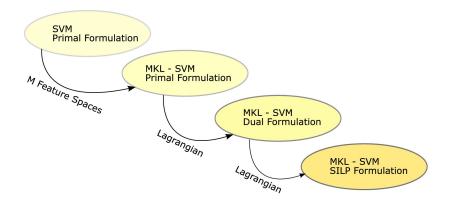


When is Multiple Kernel Learning useful?

Automated Model Selection



Derivation



For details see Sonnenburg, Rätsch, Schäfer, Schölkopf 2006

SVM Primal Formulation

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

MKL Primal Formulation

min
$$\frac{1}{2} \left(\sum_{j=1}^{M} \beta_{j} \| \mathbf{w}_{j} \|_{2} \right)^{2} + C \sum_{i=1}^{N} \xi_{n}$$
w.r.t.
$$\mathbf{w} = (\mathbf{w}_{1}, \dots, \mathbf{w}_{M}), \mathbf{w}_{j} \in \mathbb{R}^{D_{j}}, \quad \forall j = 1 \dots M$$

$$\beta \in \mathbb{R}_{+}^{M}, \ \boldsymbol{\xi} \in \mathbb{R}_{+}^{N}, \ b \in \mathbb{R}$$
s.t.
$$y_{i} \left(\sum_{j=1}^{M} \beta_{j} \mathbf{w}_{j}^{\mathsf{T}} \Phi_{j}(\mathbf{x}_{i}) + b \right) \geq 1 - \xi_{i}, \ \forall i = 1, \dots, N$$

$$\sum_{i=1}^{M} \beta_{j} = 1$$

Properties: equivalent to SVM for M=1; solution sparse in "blocks"; each block *i* corresponds to one kernel



MKL

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SVMs

MKL Dual Formulation

Bach, Lanckriet, Jordan 2004:

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

Properties: equivalent to SVM for M=1

Deriving the Semi-Infinite Linear Program

SVMs

The Semi-Infinite Linear Program I

$$\max \qquad \theta$$
w.r.t.
$$\theta \in \mathbb{R}, \beta \in \mathbb{R}_{+}^{M} \text{ with } \sum_{j=1}^{M} \beta_{j} = 1$$
s.t.
$$\sum_{j=1}^{M} \beta_{j} \left(\frac{1}{2} \sum_{r=1}^{N} \sum_{s=1}^{N} \alpha_{r} \alpha_{s} y_{r} y_{s} K_{j}(\mathbf{x}_{r}, \mathbf{x}_{s}) - \sum_{i=1}^{N} \alpha_{i} \right) \geq \theta$$

$$=: S_{j}(\alpha)$$
for all α with $0 \leq \alpha \leq C$ and $\sum_{i=1}^{N} y_{i} \alpha_{i} = 0$

Properties:

- ullet linear, optimize over a convex combination of $oldsymbol{eta}$
- infinitely many constraints, one for each $0 \le \alpha \le C$
- can use standard SVM to identify most violated constraints

SVMs

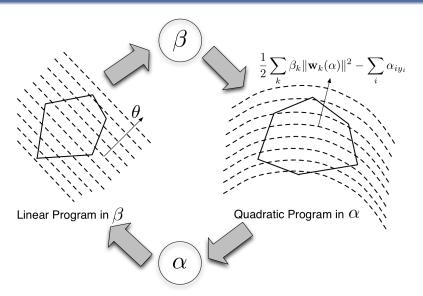
The Semi-Infinite Linear Program II

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

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



SVMs

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

$$\sum_{i=1}^{M} \beta_{j} \left(\frac{1}{2} S_{j}(\boldsymbol{\alpha}) - \sum_{i=1}^{N} \alpha_{i} \right) = \frac{1}{2} \sum_{r=1}^{N} \sum_{s=1}^{N} \alpha_{r} \alpha_{s} y_{r} y_{s} \sum_{i=1}^{M} \beta_{j} k_{j}(\mathbf{x}_{r}, \mathbf{x}_{s}) - \sum_{i=1}^{N} \alpha_{i},$$

- solved by taking set of most violated constraints into account
- most violated constraints given by SVM solution for fixed eta



Regression

Primal Formulation:

min
$$\frac{1}{2} \left(\sum_{j=1}^{M} \beta_{j} \| \mathbf{w}_{j} \| \right)^{2} + C \sum_{i=1}^{N} (\xi_{i} + \xi_{i}^{*})$$
w.r.t.
$$\mathbf{w} = (\mathbf{w}_{1}, \dots, \mathbf{w}_{M}), \mathbf{w}_{j} \in \mathbb{R}^{D_{j}}, \quad \forall j = 1 \dots M$$

$$\beta \in \mathbb{R}_{+}^{M}, \ \boldsymbol{\xi} \in \mathbb{R}^{N}, \ \boldsymbol{\xi}^{*} \in \mathbb{R}_{+}^{N}, \ b \in \mathbb{R}$$
s.t.
$$\sum_{j=1}^{M} \beta_{j} \mathbf{w}_{j}^{\mathsf{T}} \Phi_{j}(\mathbf{x}_{i}) + b \leq y_{i} + \varepsilon + \xi_{i}, \ \forall i = 1 \dots N$$

$$\sum_{j=1}^{M} \beta_{j} \mathbf{w}_{j}^{\mathsf{T}} \Phi_{j}(\mathbf{x}_{i}) + b \geq y_{i} - \varepsilon - \xi_{i}^{*}, \ \forall i = 1 \dots N$$

$$\sum_{j=1}^{M} \beta_{j} \mathbf{w}_{j}^{\mathsf{T}} \Phi_{j}(\mathbf{x}_{i}) + b \geq y_{i} - \varepsilon - \xi_{i}^{*}, \ \forall i = 1 \dots N$$

One Class

Primal Formulation:

$$\begin{aligned} & \min & \quad \frac{1}{2} \left(\sum_{j=1}^{M} \beta_{j} \left\| \mathbf{w}_{j} \right\|_{2} \right)^{2} + C \sum_{i=1}^{N} \xi_{i} - \rho \\ & \text{w.r.t.} & \quad \mathbf{w} = \left(\mathbf{w}_{1}, \dots, \mathbf{w}_{M} \right), \, \mathbf{w}_{j} \in \mathbb{R}^{D_{j}}, \quad \forall j = 1 \dots M \\ & \quad \beta \in \mathbb{R}_{+}^{M}, \, \, \boldsymbol{\xi} \in \mathbb{R}_{+}^{N} \\ & \text{s.t.} & \quad y_{i} \left(\sum_{j=1}^{M} \beta_{j} \mathbf{w}_{j}^{\mathsf{T}} \Phi_{j}(\mathbf{x}_{i}) \right) \geq \rho - \xi_{i}, \forall i = 1, \dots, N \\ & \quad \sum_{j=1}^{M} \beta_{j} = 1 \end{aligned}$$

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



SVMs

Multiclass

Primal Formulation (Zien, Ong 2007):

min
$$\frac{1}{2} \left(\sum_{j=1}^{M} \beta_{j} \|\mathbf{w}_{j}\|_{2} \right)^{2} + C \sum_{i=1}^{N} \xi_{n}$$
w.r.t.
$$\mathbf{w} = (\mathbf{w}_{1}, \dots, \mathbf{w}_{M}), \mathbf{w}_{j} \in \mathbb{R}^{k_{j}}, \quad \forall j = 1 \dots M$$

$$\boldsymbol{\beta} \in \mathbb{R}_{+}^{M}, \mathbf{s} \in \mathbb{R}^{N \times c}, \boldsymbol{\xi} \in \mathbb{R}_{+}^{N}, b \in \mathbb{R}$$
s.t.
$$\xi_{i} = \max_{u \neq y_{i}} s_{iu}, s_{iu} \geq 0,$$

$$\sum_{j=1}^{M} \beta_{j} \mathbf{w}_{j}^{\mathsf{T}} \left(\Phi_{j}(\mathbf{x}_{i}, y_{i}) - \Phi_{j}(\mathbf{x}_{i}, u) \right) + b_{y_{i}} - b_{u} \geq 1 - s_{iu},$$

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

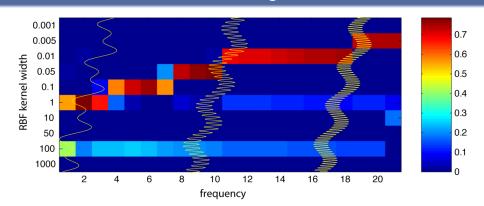
Further Extension

Extension

- \bullet ℓ_p norm MKL
- faster algorithms
- convergence bounds
- and many more
 - :

Automated Model Selection - Regression

Applications

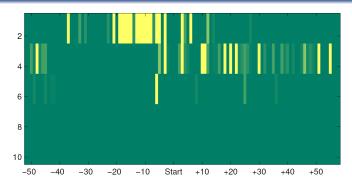


- $f(x) = \sin(ax) + \sin(bx) + cx$ for varying a
- Support Vector Regression with 10 RBF-Kernels of different width

Knowledge discovery

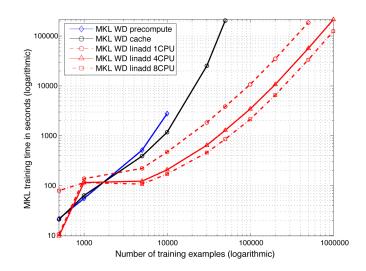
Feature Extraction

Applications



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

Speed



Summary and Outlook

MKL learns convex combination of kernels

- ⇒ allows (to some extend) 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.