

Advanced Machine Learning & Perception

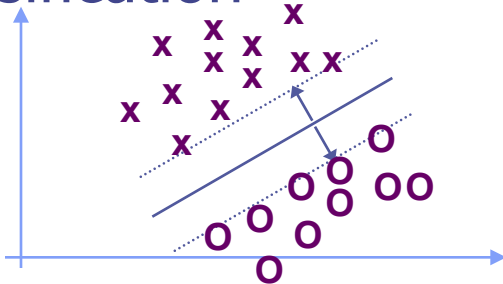
Instructor: Tony Jebara

Multi-Task Learning

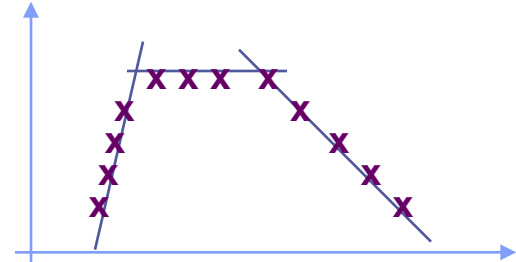
- Review MED SVM
- MED Feature Selection
- MED Kernel Selection
- Multi-Task MED
- Adaptive Pooling

SVM Extensions

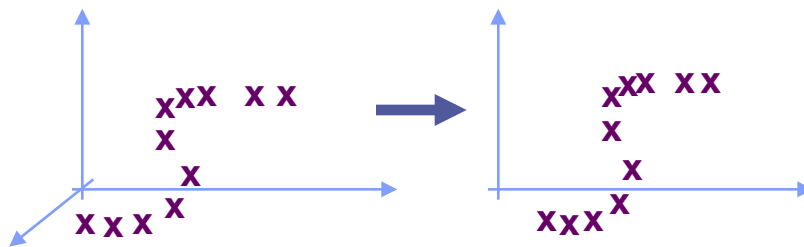
Classification



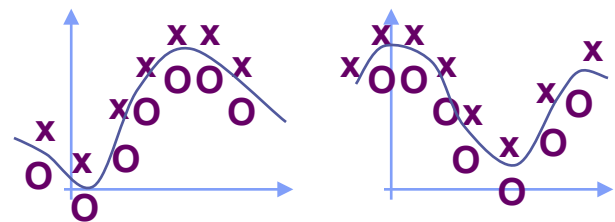
Regression



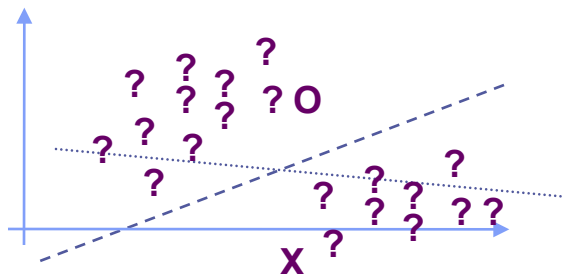
Feature/Kernel Selection



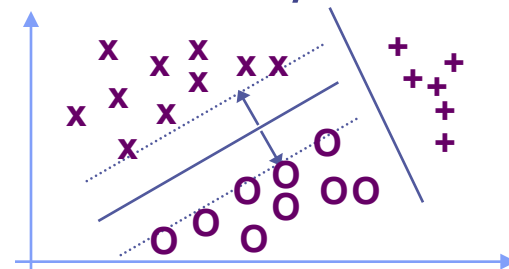
Meta/Multi-Task Learning



Transduction



Multi-Class / Structured



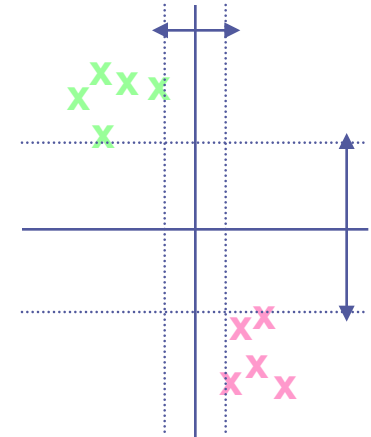
MED Support Vector Machine

- MED approach to find discriminant: $L(X; \Theta) = \theta^T X + b$
- Get $P(\theta)$:
$$\min_P KL(P \| P_0) + C \sum_t \xi_t$$

$$s.t. \int P(\Theta) y_t L(X_t; \Theta) d\Theta \geq 1 - \xi_t, \forall t$$
- Solution:
$$P(\Theta) = \frac{1}{Z(\lambda)} P_0(\Theta) \exp\left(\sum_t \lambda_t \left[y_t L(X_t; \Theta) - 1\right]\right)$$
- Partition:
$$Z(\lambda) = \int_{\Theta} P_0(\Theta) \exp\left(\sum_t \lambda_t \left[y_t L(X_t; \Theta) - 1\right]\right) d\Theta$$
- Objective:
$$J(\lambda) = \max_{\lambda} \sum_t \lambda_t - \frac{1}{2} \sum_{t, t'} \lambda_t \lambda_{t'} y_t y_{t'} (X_t^T X_{t'})$$

 (same as SVM)
$$s.t. 0 \leq \lambda_t \leq C, \sum_t \lambda_t y_t = 0$$
- Prediction:
$$\hat{y} = \operatorname{sgn} \int P(\Theta) L(X; \Theta) d\Theta = \operatorname{sgn} \sum_t y_t \lambda_t X_t^T X + b$$

MED Feature Selection



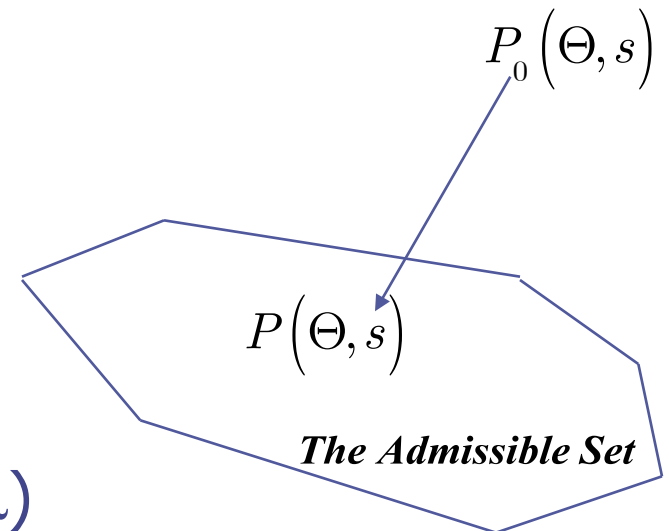
- Goal: pick 100 of 10000 features to get largest margin classifier (NP)
- Turn features on/off via binary switches $s_i \in \{0, 1\}$

- Discriminant is now $L(X; \Theta) = \sum_i s_i \theta_i X_i + b$

- Introduce a prior on switches:

$$P_{s,0}(s_i) = \rho^{s_i} (1 - \rho)^{1-s_i}$$

- This is a Bernoulli distribution where ρ controls a priori pruning level
- MED finds discriminative $P(\theta, s)$ close to prior by maximizing $J(\lambda) = -\log Z(\lambda)$

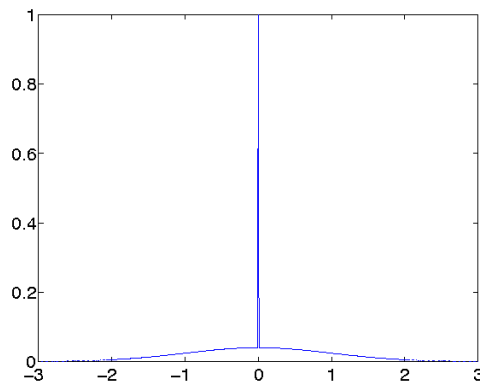


MED Feature Selection

- Discriminant function now is $L(X; \Theta) = \sum_{i=1}^D s_i \theta_i X_i + b$
- The model $\Theta = \{b, \theta_1, \dots, \theta_D, s_1, \dots, s_D\}$ contains binary $s_i \in \{0, 1\}$ parameters (with Bernoulli priors) to prune features

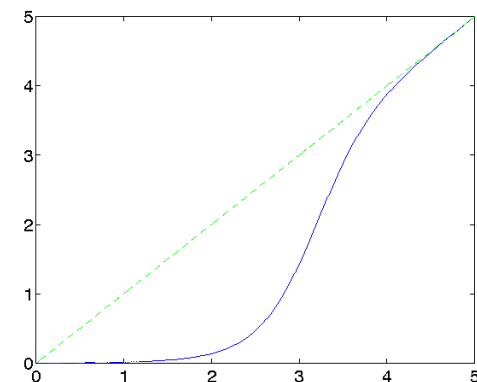
Prior: $P_0(\Theta) = P_0(b) P_0(\theta) P_0(s) = N(b | 0, \infty) N(\theta | 0, I) \prod_i P_0(s_i)$

Partition: $Z(\lambda) = \sum_{s_1=0}^1 \dots \sum_{s_D=0}^1 \int_b \int_{\theta} P_0(\Theta) \exp\left(\lambda_t \left[y_t L(X_t; \Theta) - 1\right]\right) d\theta db$



Prior on $s_i \theta_i$

Aggressive attenuation
of linear coefficients
at low values (rho=.01).

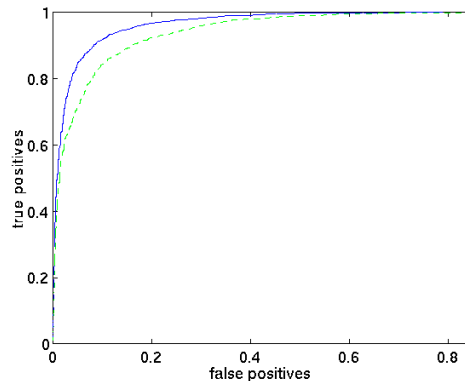


MED Feature Selection

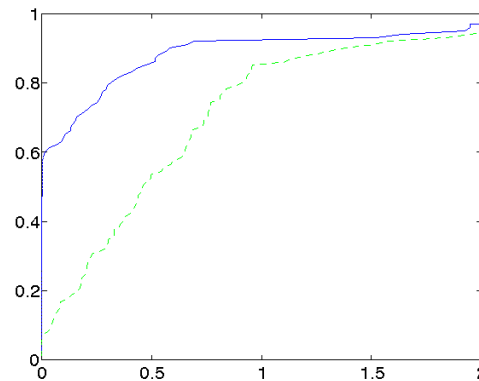
Objective is now:
$$J(\lambda) = \sum_t \lambda_t - \sum_{i=1}^D \log \left[1 - \rho + \rho e^{\frac{1}{2} \left(\sum_t \lambda_t y_t X_{t,i} \right)^2} \right]$$

s.t. $0 \leq \lambda_t \leq C, \quad \sum_t \lambda_t y_t = 0$

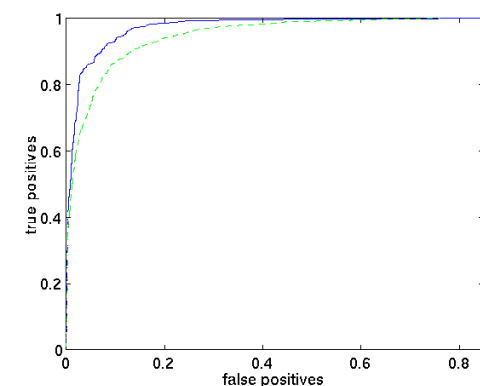
DNA Data: 2-class, 100 element binary vectors. Train/Test=500/4724



ROC of DNA Splice Site
100 Features
Original 25xGATC



CDF of Linear Coeffs
DNA Splice Site
100 Features



ROC DNA Splice Site
~5000 Features
Quadratic Kernel

Dashed line: $\rho = 0.99999$

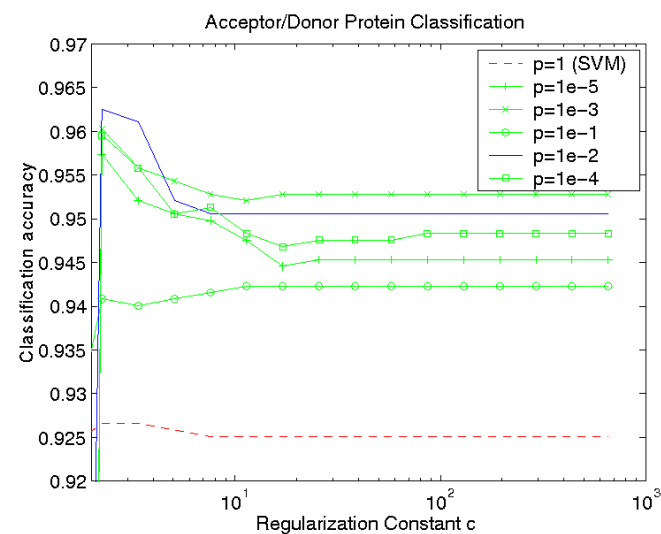
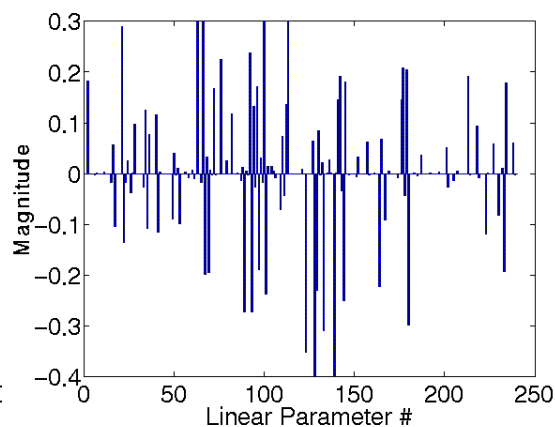
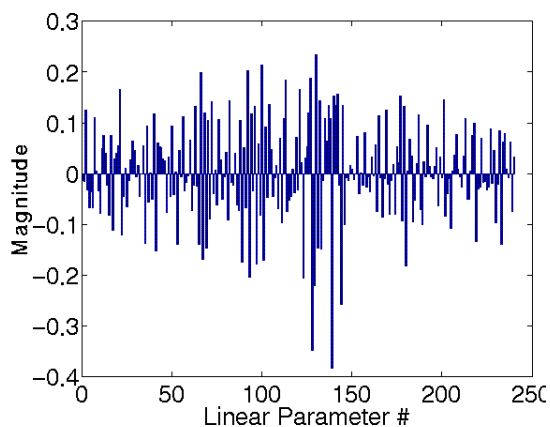
Solid line: $\rho = 0.00001$

MED Feature Selection

$$J(\lambda) = \sum_t \lambda_t - \sum_{i=1}^D \log \left[1 - \rho + \rho e^{\frac{1}{2} \left(\sum_t \lambda_t y_t X_{t,i} \right)^2} \right]$$

$$s.t. \quad 0 \leq \lambda_t \leq C, \quad \sum_t \lambda_t y_t = 0$$

Example: Intron-Exon Protein Classification:
UCI: 240 dims; 200 train, 1300 test



MED Feature Selection

- MED can also use switches in regression, objective is then:

$$J(\lambda) = \sum_t y_t (\lambda'_t - \lambda_t) - \epsilon \sum_t (\lambda'_t + \lambda_t) - \sum_i \log \left(1 - p_0 + p_0 e^{\frac{1}{2} \left[\sum_t (\lambda_t - \lambda'_t) X_{t,i} \right]^2} \right)$$

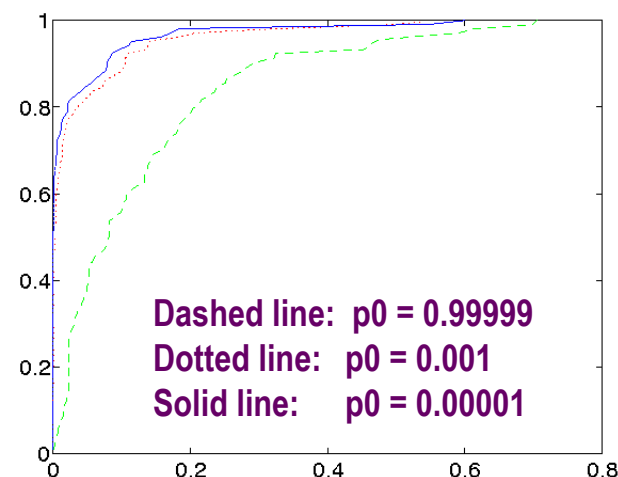
$$s.t. 0 \leq \lambda'_t, \lambda_t \leq C, \sum_t \lambda'_t - \lambda_t = 0$$

- Boston Housing Data: predict price from 13 scalars

Train/Test=481/25

Explicit Quadratic Kernel Expansion

Linear Model Estimator	Epsilon-Sensitive Linear Loss
Least-Squares	1.7584
MED p0 = 0.99999	1.7529
MED p0 = 0.1	1.6894
MED p0 = 0.001	1.5377
MED p0 = 0.00001	1.4808



- Cancer Data: predict expression from 67 other cancer levels
Train/Test = 50/3951

Linear Model Estimator	Epsilon-Sensitive Linear Loss
Least-Squares	3.609e+03
MED p0 = 0.00001	1.6734e+03

MED Kernel Selection

- Purpose: pick mixture of subset of D Kernel matrices to get largest margin classifier (i.e. learn the Gram matrix)
- Turn kernels on/off via binary switches $s_i \in \{0, 1\}$
- Switch Prior: Bernoulli distribution $P_{s,0}(s_i) = \rho^{s_i} (1 - \rho)^{1-s_i}$
- Discriminant uses D models with multiple nonlinear mappings of datum $L(X; \Theta) = \sum_i s_i \theta_i^T \Phi_i(X) + b$
- MED solution has analytic concave objective fn:

$$J(\lambda) = \sum_t \lambda_t - \sum_{i=1}^D \log \left[1 - \rho + \rho \exp \left(\frac{1}{2} \sum_{t=1}^T \sum_{t'=1}^T \lambda_t \lambda_{t'} y_t y_{t'} k_i(X_t, X_{t'}) \right) \right]$$

$$s.t. \quad 0 \leq \lambda_t \leq C, \quad \sum_t \lambda_t y_t = 0$$

Meta Feature Selection

- Given a series of classification tasks: $m \in [1..M]$
 map inputs to binary: $X_{tm} \rightarrow y_{tm} \quad \forall t \in [1..T_m]$
 using M discriminants with 1 feature selection vector:

$$L(X; s, \theta_m, b_m) = \sum_i s_i \theta_{m,i} X_i + b_m$$

Subject to MED classification constraints:

$$\int P(s, \theta_1, \dots, \theta_M, b_1, \dots, b_M) \left[y_{tm} \left(L(X_{tm}; s, \theta_m, b_m) - 1 \right) \right] d\Theta \geq 0, \quad \forall t \forall m$$

Solve by optimizing joint objective function for all Lagranges

$$J(\lambda) = \sum_{t,m} \lambda_{tm} - \sum_{i=1}^D \log \left(1 - \rho + \rho \exp \left(\frac{1}{2} \sum_{m=1}^M \left[\sum_{t=1}^{T_m} \lambda_{tm} y_{tm} X_{tm,i} \right]^2 \right) \right)$$

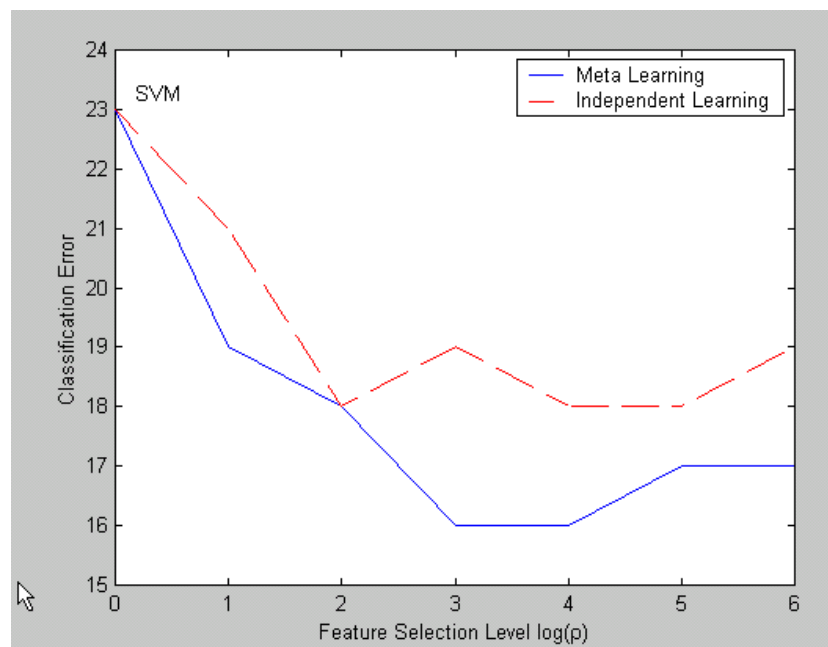
$$s.t. \quad 0 \leq \lambda_{tm} \leq C, \quad \sum_t \lambda_{tm} y_{tm} = 0 \quad \forall m$$

Meta Feature Selection Results

- Have many classification tasks with common feature selection. To ensure coupled tasks, turn multi-class data set into multiple 1 versus many tasks

UCI Dermatology Dataset: 200 trains, 166 tests, 33 features, 6 classes

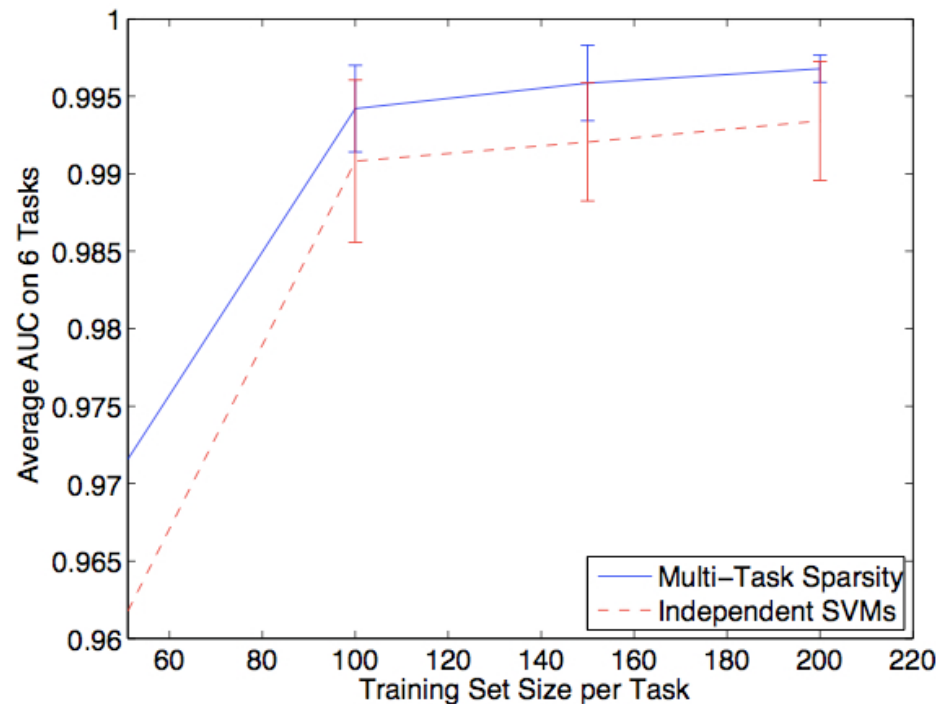
**Cross-validating over
Regularization Levels**



Meta Feature Selection Results

Can also cross validate over ρ (or $\alpha=(1-\rho)/\rho$) as well as C

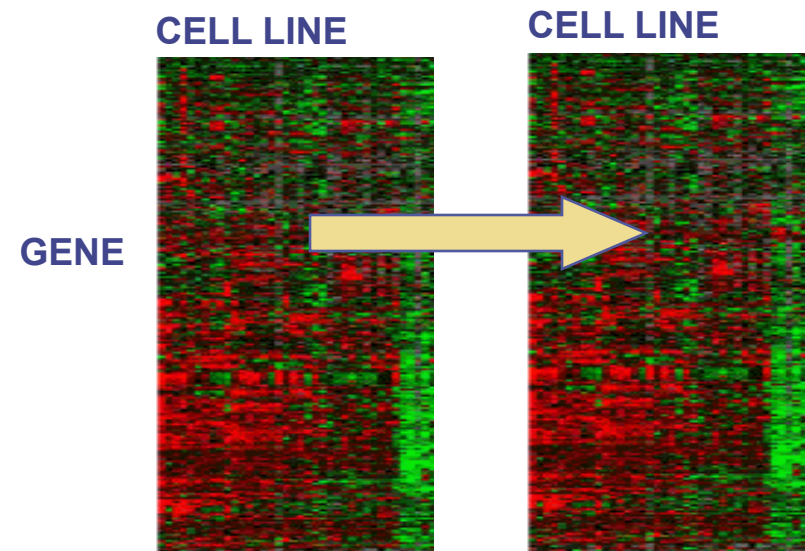
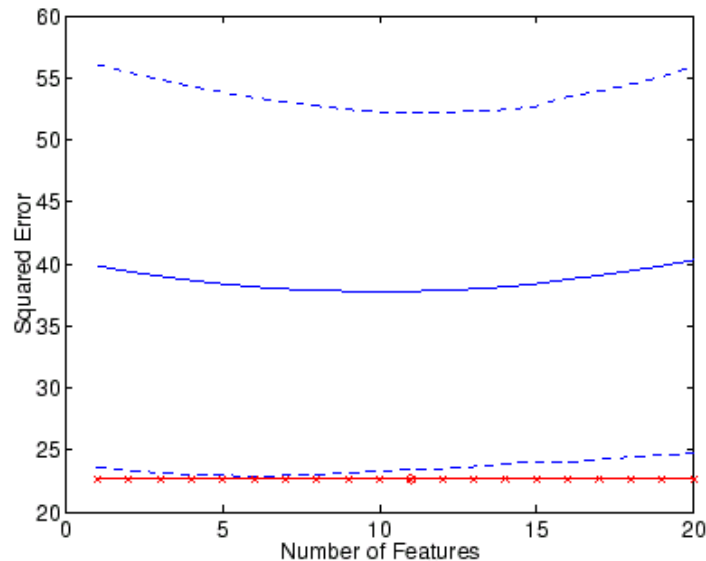
Example: UCI Dermatology dataset (6 tasks)



Meta Feature Select Regression

- Can also solve many regression tasks with one common feature selection

**D. Ross Cancer Data: 67 expression level features.
Use subset of 800 genes to predict all others
Compared with random feature selection**



Meta Kernel Selection

- Given many tasks with common (unknown) kernel matrix
- Use M discriminants with one feature selection vector:

$$L\left(X; s, \Theta_m, b_m\right) = \sum_i s_i \theta_{m,i}^T \Phi_i(X) + b_m$$

- Subject to MED classification constraints:

$$\int P\left(s, \Theta_1, \dots, \Theta_M, b_1, \dots, b_M\right) \left[y_{tm} L\left(X_{tm}; s, \Theta_m, b_m\right) - \gamma \right] ds db_m d\Theta_m \geq 0, \quad \forall t \forall m$$

optimize joint objective function over Lagrange multipliers

$$J(\lambda) = \sum_{t,m} \lambda_{tm} - \sum_{i=1}^D \log \left[1 - \rho + \rho \exp \left(\frac{1}{2} \sum_{m=1}^M \sum_{t=1}^{T_m} \sum_{t'=1}^{T_m} \lambda_{tm} \lambda_{tm'} y_{tm} y_{tm'} k_i(X_{tm}, X_{tm'}) \right) \right]$$

$$s.t. \quad 0 \leq \lambda_{tm} \leq C, \quad \sum_t \lambda_{tm} y_{tm} = 0 \quad \forall m$$

Meta Kernel Selection as QP

- The objective function is convex but not quite a QP

$$J(\lambda) = \sum_{t,m} \lambda_{tm} - \sum_{i=1}^D \log \left[1 - \rho + \rho \exp \left(\frac{1}{2} \sum_{m=1}^M \sum_{t=1}^{T_m} \sum_{t'=1}^{T_m} \lambda_{tm} \lambda_{tm'} y_{tm} y_{tm'} k_i(X_{tm}, X_{tm'}) \right) \right]$$

$$s.t. \quad 0 \leq \lambda_{tm} \leq C, \quad \sum_t \lambda_{tm} y_{tm} = 0 \quad \forall m$$

- Use a bound on each log term to make it quadratic in λ

$$-\log \left(\alpha + \exp \left(\frac{\mathbf{u}^T \mathbf{u}}{2} \right) \right) \geq -\log \left(\alpha + \exp \left(\frac{\mathbf{v}^T \mathbf{v}}{2} \right) \right) - \frac{\exp \left(\frac{\mathbf{v}^T \mathbf{v}}{2} \right)}{\alpha + \exp \left(\frac{\mathbf{v}^T \mathbf{v}}{2} \right)} \mathbf{v}^T (\mathbf{u} - \mathbf{v}) - \frac{1}{2} (\mathbf{u} - \mathbf{v})^T (\mathcal{G} \mathbf{v}^T \mathbf{v} + I) (\mathbf{u} - \mathbf{v})$$

$$\text{where } \mathcal{G} = \frac{\tanh \left(\frac{1}{2} \log \left(\alpha \exp \left(-\mathbf{v}^T \mathbf{v} / 2 \right) \right) \right)}{2 \log \left(\alpha \exp \left(-\mathbf{v}^T \mathbf{v} / 2 \right) \right)}$$

- As with EM, maximize the lower bound, update & repeat

- Converges in fewer steps than $\left\lceil \frac{\log(1/\epsilon)}{\log(\min(1 + 1/\alpha, 2))} \right\rceil$

Meta Kernel Selection as QP

- Code for learning the weights for $d=1\dots D$ kernels

Algorithm 1 Multitask SVM Learning

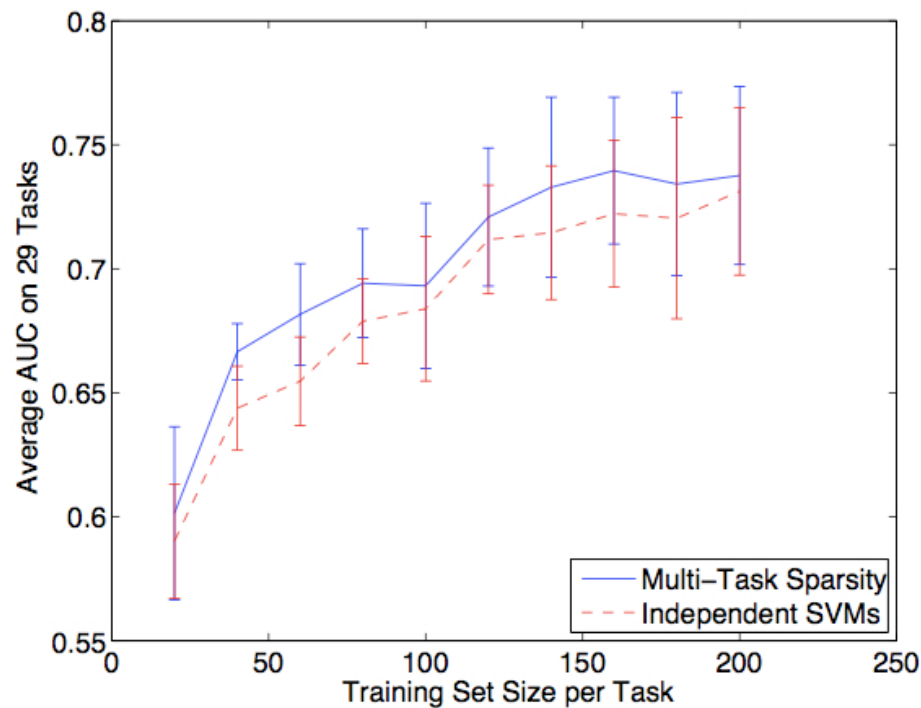
0	Input dataset $\mathcal{D}$, $C > 0$, $\alpha \geq 0$, $0 < \varpi < 1$ and kernels k_d for $d = 1, \dots, D$.
1	Initialize Lagrange multipliers to zero $\lambda = \mathbf{0}$.
2	Store $\tilde{\lambda} = \lambda$.
3	For $m = 1, \dots, M$ do:
3a	Set $g_d = \alpha \exp \left(-\frac{1}{2} \sum_{m=1}^M \sum_{t=1}^{T_m} \sum_{\tau=1}^{T_m} \lambda_{m,t} \lambda_{m,\tau} y_{m,t} y_{m,\tau} k_d(\mathbf{x}_{m,t}, \mathbf{x}_{m,\tau}) \right)$ for all d . Set $\mathcal{G}_d = \frac{\tanh(\frac{1}{2} \log(g_d))}{2 \log(g_d)}$ for all d . Set $\hat{s}(d) = \frac{1}{1+g_d}$ for all d . Set $\hat{y}_{m,t}(d) = \sum_{\tau=1}^{T_m} \lambda_{m,\tau} y_{m,\tau} k_d(\mathbf{x}_{m,t}, \mathbf{x}_{m,\tau})$ for all t and d .
3b	Update each of the λ_m vectors with the SVM QP: $\max_{\lambda_m} \sum_{t=1}^{T_m} \lambda_{m,t} - \sum_{t=1}^{T_m} \lambda_{m,t} y_{m,t} \sum_{d=1}^D \hat{s}(d) \hat{y}_{m,t}(d)$ $+ \sum_{t=1}^{T_m} \sum_{\tau=1}^{T_m} \lambda_{m,t} \tilde{\lambda}_{m,\tau} y_{m,t} y_{m,\tau} \sum_{d=1}^D (\mathcal{G}_d \hat{y}_{m,t}(d) \hat{y}_{m,\tau}(d) + k_d(\mathbf{x}_{m,t}, \mathbf{x}_{m,\tau}))$ $- \frac{1}{2} \sum_{t=1}^{T_m} \sum_{\tau=1}^{T_m} \lambda_{m,t} \lambda_{m,\tau} y_{m,t} y_{m,\tau} \sum_{d=1}^D (\mathcal{G}_d \hat{y}_{m,t}(d) \hat{y}_{m,\tau}(d) + k_d(\mathbf{x}_{m,t}, \mathbf{x}_{m,\tau}))$ s.t. $0 \leq \lambda_{m,t} \leq C \ \forall t = 1, \dots, T_m$ and $\sum_{t=1}^{T_m} y_{m,t} \lambda_{m,t} = 0$.
4	If $\ \lambda - \tilde{\lambda}\ > \varpi \ \lambda\ $ go to 2.
5	Output: $\hat{s}$ and λ .

- Final kernel to use in the SVMs: $k(X, X') = \sum_{d=1}^D \hat{S}(d) k_d(X, X')$

Meta Kernel Selection Results

Can also cross validate over ρ (or $\alpha=(1-\rho)/\rho$) as well as C

Example: Landmine dataset (29 tasks) with RBF kernels



Meta or Adaptive Pooling

- Another type of meta-learning is adaptive pooling
- Assume $m \in [1..M]$ datasets predicting binary labels
- Here, datasets are all labeled for the same task
- But, inputs are sampled from slightly different distributions
- E.g. Dataset 1: color face images labeled as male/female
Dataset 2: gray face images labeled as male/female
- Pooling: combine both datasets and learn one classifier

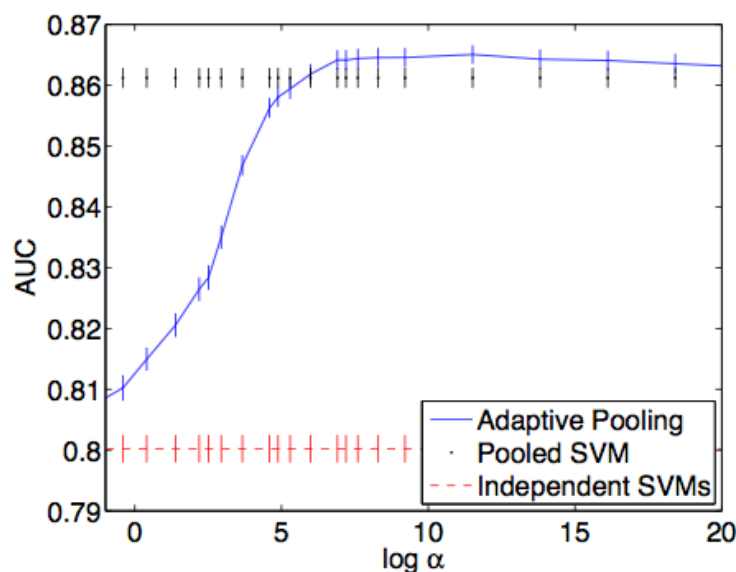
$$L_m(X) = \theta^T X + b$$
- Independent learning: learn a separate classifier for each

$$L_m(X) = \theta_m^T X + b_m$$
- Adaptive pooling: each classifier is a mix of the shared model and a specialized model

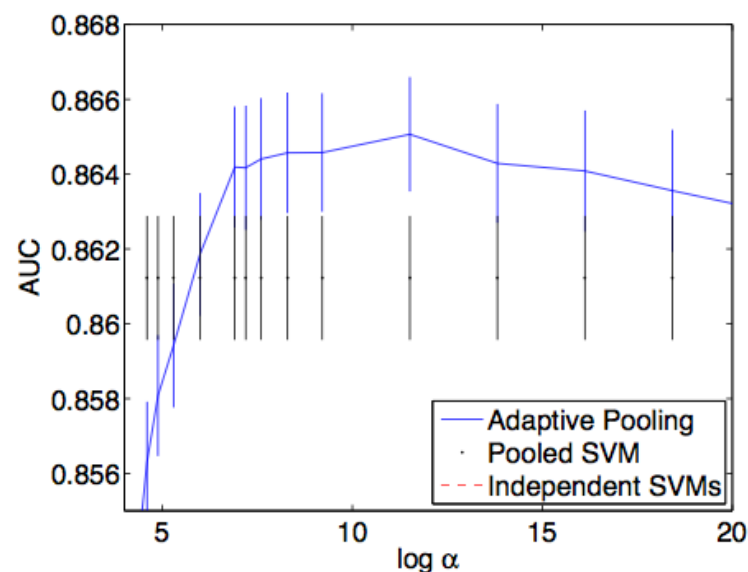
$$L_m(X) = s_m \left(\theta_m^T X + b_m \right) + \left(\theta^T X + b \right)$$
- Once again MED solution is straightforward...

Meta or Adaptive Pooling

- Compare to full pooling and independent learning



(a) Average AUC



(b) Average AUC zoomed in

$$J(\lambda) = \sum_{t,m} \lambda_{tm} - \sum_m \sum_{m'} \frac{1}{2} \sum_{t=1}^{T_m} \sum_{t'=1}^{T_{m'}} \lambda_{tm} \lambda_{t'm'} y_{tm} y_{t'm'} k(X_{tm}, X_{t'm'})$$

$$\sum_{m=1}^M \log \left[\alpha + \exp \left(\frac{1}{2} \sum_{t=1}^{T_m} \sum_{t'=1}^{T_{m'}} \lambda_{tm} \lambda_{t'm'} y_{tm} y_{t'm'} k_m(X_{tm}, X_{t'm'}) \right) \right] + M \log(\alpha + 1)$$

$$s.t. \ 0 \leq \lambda_{tm} \leq C, \ \sum_t \lambda_{tm} y_{tm} = 0 \quad \forall m$$