# Regularization Algorithms for Learning

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Alessandro Verri Regularization Algorithms for Learning

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- motivation
- setting
- elastic net regularization
  - iterative thresholding algorithms
  - error estimates and parameter choice
- applications

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- starting point of many learning schemes is a fixed data representation
- if prediction is the goal, black box solutions are acceptable
- in many problems the primary goal is the identification of the variables/measures relevant for prediction
- it is often the case that variables are dependent

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**model**:  $X \times Y$  is endowed with a probability distribution

$$p(y,x) = p(y|x)p(x)$$

input space:  $X \subset \mathbb{R}^d$ regression:  $Y \in [-M, M]$ classification:  $Y \in \{-1, 1\}$ 

the distribution *p* is fixed but unknown

**DATA**: we are given a set of examples, i.e.  $(x_1, y_1), \ldots, (x_n, y_n)$  sampled i.i.d. according to p(y, x)

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• regression function:  $f^*(x) = \mathbb{E}[y|x]$  minimizes

$$\mathcal{E}(f) = \mathbb{E}\left[|y - f(x)|^2\right]$$

• Bayes rule:  $f_b(x) = \text{sign}(f^*(x))$  minimizes  $\mathcal{R}(f) = P(yf(x) \le 0)$ 

• for any f

$$\mathcal{R}(f) - \mathcal{R}(f_b) \leq \sqrt{\mathcal{E}(f) - \mathcal{E}(f^*)}$$

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## Hypotheses Space and Dictionary

The search for a solution is restricted to some space of hypotheses  $\ensuremath{\mathcal{H}}$ 

- dictionary:  $\mathcal{D} = \{\varphi_{\gamma} : X \to \mathbb{R} \mid \gamma \in \Gamma\}$
- atoms or features:  $\varphi_{\gamma}$
- hypotheses space:  $\mathcal{H} = \{f | f = f_{\beta} = \sum_{\gamma \in \Gamma} \varphi_{\gamma} \beta_{\gamma} \}$
- The atoms are not linearly independent,
- the dictionary can be infinite dimensional,
- the atoms can be seen as measures (features) on the input objects in *X*,
- the solution is a weighted combination of the features  $f_{\beta}(x) = \sum_{\gamma \in \Gamma} \varphi_{\gamma}(x) \beta_{\gamma}$

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#### learning task

given  $(x_1, y_1), \ldots, (x_n, y_n)$  find an estimator  $f_n$  such that

 $f_n(x) \sim f^*(x)$ 

### An Important Distinction

- The problem of prediction/generalization is that of estimating f\*.
- The problem of **selection** is that of detecting a meaningful  $\beta^*$  (with  $f^* = \sum_{\gamma \in \Gamma} \beta^*_{\gamma} \varphi_{\gamma}$ ).

- learning can be seen as an ill-posed inverse problem and regularization is the theory of choice to restore well-posedness (Girosi and Poggio...)
- in the recent years the genova gang explored the connection between learning and inverse problems in a series of works covering theory, algorithms and applications.

a classical way to avoid overfitting: *penalized empirical risk minimization* 

$$eta_n^{\lambda} = \operatorname*{argmin}_{eta} \left( \frac{1}{n} \sum_{i=1}^n |y_i - f_{eta}(x_i)|^2 + \lambda pen(eta) \right)$$

different penalizations corresponds to different algorithms:

#### examples

- tikhonov/ridge regression:  $pen(\beta) = \sum_{\gamma \in \Gamma} \beta_{\gamma}^2$
- basis pursuit/lasso:  $pen(\beta) = \sum_{\gamma \in \Gamma} |\beta_{\gamma}|$

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we study the regularization scheme defined by

$$\beta_n^{\lambda} = \underset{\beta}{\operatorname{argmin}} \left( \frac{1}{n} \sum_{i=1}^n |y_i - f_{\beta}(x_i)|^2 + \lambda (\sum_{\gamma \in \Gamma} w_{\gamma} |\beta_{\gamma}| + \varepsilon \sum_{\gamma \in \Gamma} \beta_{\gamma}^2) \right),$$

(see Zou and Hastie'06)

Vector notation

$$\left\| \hat{Y} - \Phi_n \beta \right\|_n^2 + \lambda (\|\beta\|_{1,w} + \varepsilon \|\beta\|_2^2)$$

-  $\hat{Y} = (y_1, \dots, y_n)$ - $\Phi_n$  is  $n \times \Gamma$  (possible infinite dimensional matrix).

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$$pen(\beta) = \lambda(\|\beta\|_{1,w} + \varepsilon \|\beta\|_2)$$

- $\varepsilon > 0$  grouping effect in selection
- $\varepsilon > 0$  strictly convex approximation to basis pursuit
- more stable w.r.t. to noise in the measurements

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we (approximately) solve

$$\hat{\mathbf{Y}} = \Phi_n \beta$$

where  $\Phi$  is  $n \times \Gamma$ . We look for the minimal *pen*( $\beta$ ) solution.



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### questions:

$$\min\{\left\|\hat{\mathbf{Y}}-\Phi_{n\beta}\right\|_{n}^{2}+\lambda(\|\beta\|_{1,w}+\varepsilon\,\|\beta\|_{2}^{2})\}$$

Q1: statistical convergence of the algorithm

Q2: solution of the optimization problem

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### **(**) summability condition: $\Gamma$ denumerable and, for some $\kappa > 0$ ,

$$orall x \in X$$
  $\sum_{\gamma \in \Gamma} |arphi_{\gamma}(x)|^2 \leq \kappa.$ 

(2) there exists  $(\beta^*{}_{\gamma})_{\gamma\in\Gamma}$  such that

$$\sum_{\gamma\in\Gamma} w_{\gamma}|\beta_{\gamma}^{*}| < +\infty \qquad \text{and} \qquad f^{*}(x) = \sum_{\gamma\in\Gamma} \varphi_{\gamma}(x)\beta_{\gamma}^{*} \quad x\in\mathcal{X}.$$

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let  $\Phi_n$  be the  $n \times \Gamma_{\lambda}$  matrix (with transpose  $\Phi_n^T$ ). We can define an iteration  $\beta^{\ell}$  converging to  $\beta_n^{\lambda}$ 

let  $\beta^0 = 0$ , for  $\ell = 1, 2, ...$  $\beta^{\ell} = \frac{1}{C + \lambda \varepsilon} \mathbf{S}_{\lambda} \left( (CI - \Phi_n^T \Phi_n) \beta^{\ell-1} + \Phi_n^T \hat{Y} \right)$ 

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## thresholding function



$$\mathcal{S}_{\mu}\left(x
ight) = \left\{egin{array}{ccc} x - rac{\mu}{2} & ext{if} & x > rac{\mu}{2} \ 0 & ext{if} & |x| \leq rac{\mu}{2} \ x + rac{\mu}{2} & ext{if} & x < -rac{\mu}{2} \end{array}
ight.$$

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# generalized and distribution dependent solutions

### generalized solution

 $\beta^{\varepsilon}$  solves

$$\min\{\|\beta\|_{1,w} + \varepsilon \|\beta\|_2\}$$
  
s.t.  $f^* = \sum_{\gamma \in \Gamma} \varphi_{\gamma} \beta^*$ 

### elastic net distribution dependent solution

 $\beta^{\lambda}$  solves

$$\min\{\mathbb{E}\left[|f_{\beta}(x) - y|^{2}\right] + \lambda pen(\beta)\}$$

we also let  $f^{\lambda} = \sum_{\gamma \in \Gamma} \varphi_{\gamma} \beta_{\gamma}^{\lambda}$ 

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## error analysis

Error decomposition for fixed  $\lambda > 0$ 



 $\bullet$  under some assumptions on the noise with probability greater than  $1-4e^{-\delta}$ 

$$\left\|\beta_{n}^{\lambda}-\beta^{\lambda}\right\|_{2}\leq\left\|\beta^{\lambda}-\beta\right\|_{2}\left(\frac{C\sqrt{\delta}}{\sqrt{n\lambda}}\right)+\left(\frac{C\sqrt{\delta}}{\sqrt{n\lambda}}\right)$$

• The approximation error satisfies

$$\lim_{\lambda \to 0} \left\| \beta^{\lambda} - \beta^{\varepsilon} \right\|_{2} \to 0$$

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if we choose  $\lambda_n$  s.t.  $\lambda_n \sqrt{n} \to \infty$ , when  $n \to \infty$  then

$$\mathbb{E}\left[\left\|\beta_n^{\lambda_n}-\beta^{\varepsilon}\right\|_2\right]\to\mathbf{0},$$

moreover we also have

$$\mathbb{E}\left[\mathcal{E}(f_n^{\lambda_n}) - \mathcal{E}(f^*)\right] \to \mathbf{0}$$

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## related work

- many algorithms for sparse selection, (Mallat et al. Gilbert and Tropp OMP, Candes and Tao 06 - Dantzig estimator, Donoho et al. - Basis Pursuit, Fuguereido and Nowak -Projected Gradient, Freund and Shapiro- Boosting...)
- many theoretical results on L2 regularization (Smale and Zhou 05, Caponnetto and De Vito 05, Bauer et al. 06...).
- Recently many results for sparsity based scheme. Mostly in different settings - fixed design regression, signal processing, linear coding - (Donoho '85...Candes and Tao 05... Daubachies et al. 05...)
- fewer results in the context of learning (Barron et al. 06, Bunuea et al. 06, Koltchinskii 07...)

### (Destrero, De Mol, Odone, Verri 07)

face detection integrated in a monitoring system in our department



data:image size 20x20, 2000 + 2000 training, 1000 + 1000 validation, 2000 + 2000 test.

overcomplete dictionary of rectangular features capturing the local geometry of faces (Viola and Jones, 2004),



- features computed at at all locations, scales, aspect ratios: roughly 64000 per image.

- highly correlated features

## selected features



42 features extracted by a 2 stage selection scheme.

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## results



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## microarray data



in micro-array classification the number of samples is much smaller than the number of genes expressions

(Mosci, De Mol, Traskine, Verri 07) Algorithms were tested on three datasets:

- leukemia (patients 72 (38-34), genes 7129)
- lung cancer (patients 181 (91-90), genes 12533)
- prostate cancer (patients 102 (51-51), genes 12533)

$\lambda = 0.07$	test error	# of	intersection w.
	(test set	selected	genes selected
$\lambda arepsilon$	size: 90)	genes	for bigger $arepsilon$
0	0	22	100%
0.001	0	28	100%
0.0025	1	37	100%
0.005	1	54	100%
0.01	1	80	99%
0.1	1	247	96%
1	1	743	_

Previous: 8 genes with 91-98 % correct classification on test (Gordon et al. 02)

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### prostate cancer

$\lambda = 0.06$	test error	# of	intersection w.
	(test set	selected	genes selected
$\lambda arepsilon$	size: 51)	genes	for bigger $arepsilon$
0	5	19	95%
0.001	5	20	100%
0.0025	5	25	100%
0.005	4	31	97%
0.01	5	40	98%
0.1	6	85	94%
1	5	121	_

Previous: 5-8 genes with 82,9-95,7 % correct classification on test using ranking and K-NN (Singh et al 2002).

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- approximation properties and connections with work in signal processing
- design good (data-dependent?) dictionary

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