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Approximate Manifold Regularization: Scalable Algorithm and Generalization Analysis

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1. Introduction

2. Algorithm Design

3. Theoretical Analysis

4. Experiments

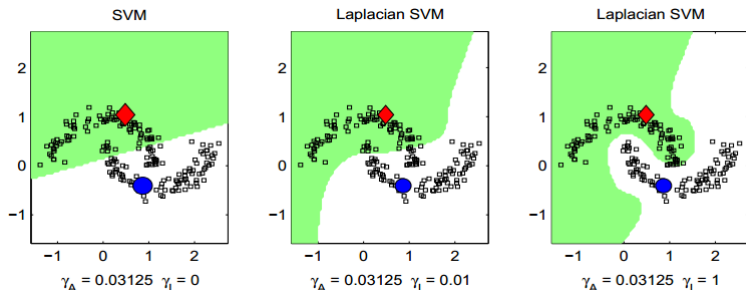
5. Conclusion

What is Manifold Regularization?

Consider common semi-supervised setting that a training dataset with n instances but only a few points m are labeled, where $m \ll n$.

$$\hat{f}_\lambda = \arg \min_{f \in \mathcal{H}} \sum_{i=1}^m \ell(y_i, f(\mathbf{x}_i)) + \lambda_A \|f\|_{\mathcal{H}}^2 + \lambda_I \mathbf{f}^T \mathbf{L} \mathbf{f}.$$

where \mathbf{L} is graph Laplacian by $\mathbf{L} = \mathbf{D} - \mathbf{W}$, $\mathbf{f} = [f(\mathbf{x}_1), \dots, f(\mathbf{x}_n)]^T$, $\mathbf{W} \in \mathbb{R}^{n \times n}$ measures similarities between all points and \mathbf{D} is a diagonal matrix $\mathbf{D}_{ii} = \sum_{j=1}^n W_{ij}$. [Belkin et al., 2006]



Scalability issues of LapRLS

1 Consider kernel ridge regression (KRR) with manifold regularization

- Representer Theorem $f(\mathbf{x}) = \sum_{i=1}^n \alpha_i K(\mathbf{x}_i, \mathbf{x})$
- The squared loss $\ell(y_i, f(x_i)) = (y_i - f(x_i))^2$
- Also called as Laplacian Regularized Least Squares (LapRLS)

2 LapRLS with closed-form solution

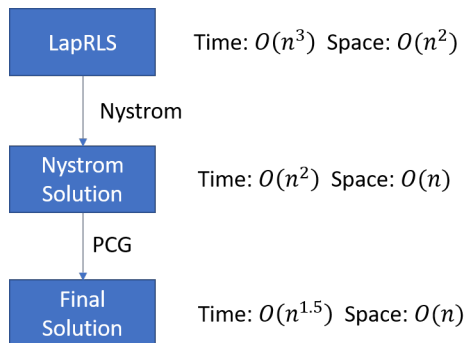
$$\hat{\alpha} = (\mathbf{JK} + \lambda_A \mathbf{I} + \lambda_I \mathbf{LK})^{-1} \mathbf{y}_n,$$

3 Scalability issues

- Space complexity: $\mathcal{O}(n^2)$.
e.g. Storing kernel matrix needs 18.6 GB when $n = 50,000$ while 74.5 GB when $n = 100,000$.
- Time complexity: $\mathcal{O}(n^3)$.

Unfeasible to deal with large scale semi-supervised tasks!!!

1 Core Idea : LapRLS + Nyström + PCG.



2 Contributions:

- Scalable Algorithm : $O(n)$ space and $O(n^{1.5})$ time.
- Theoretical Guarantee: Excess risk bounds with convergence rate $O(\frac{1}{\sqrt{m}})$.

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- 1 Uniform subsampling over the training set (n points $\rightarrow s$ Nyström centers)

$$\mathcal{H}_s = \{f \in \mathcal{H} | f = \sum_{i=1}^s \alpha_i K(\mathbf{x}_i, \cdot), \boldsymbol{\alpha} \in \mathbb{R}^s\},$$

- 2 Nyström LapRLS with a closed-form solution:

$$\hat{f}_\lambda^s(\mathbf{x}) = \sum_{i=1}^s \alpha_i K(\mathbf{x}_i, \mathbf{x}), \quad \text{with}$$
$$\boldsymbol{\alpha} = \underbrace{(\mathbf{K}_{ms}^T \mathbf{K}_{ms} + \lambda_A \mathbf{K}_{ss} + \lambda_I \mathbf{K}_{ns}^T \mathbf{L} \mathbf{K}_{ns})}^{\mathbf{H}} \underbrace{\mathbf{K}_{ms}^T \mathbf{y}}_{\mathbf{z}},$$

where \mathbf{H}^\dagger denotes the Moore-Penrose pseudoinverse and $\mathbf{H} \in \mathbb{R}^{s \times s}$.

- 3 Computation of \mathbf{H} needs $\mathcal{O}(ns^2)$. Consider iterative method: conjugate gradient (CG) to solve linear systems.

$$\mathbf{H}\boldsymbol{\alpha} = \mathbf{z}.$$

Preconditioned Conjugate Gradient (PCG)

- 1 Convergence properties of CG methods are determined by the condition number $\kappa(\mathbf{H})$: the larger $\kappa(\mathbf{H})$ is, the slower the improvement.

$$\mathbf{H}\boldsymbol{\alpha} = \mathbf{z}.$$

In most cases, $\kappa(\mathbf{H})$ is large (ill-conditioned), thus convergence is slow.

- 2 Preconditioning to reduce the condition number $\kappa(\mathbf{P}^{-1}\mathbf{H})$

$$\mathbf{P}^{-1}\mathbf{H}\boldsymbol{\alpha} = \mathbf{P}^{-1}\mathbf{z}.$$

The more similar \mathbf{H} and \mathbf{P} are, the smaller the condition number.
We provide two preconditioners to approximate \mathbf{H}

- $m \leq \sqrt{n}$

$$\mathbf{P} = \mathbf{K}_{ms}^T \mathbf{K}_{ms} + \lambda_A \mathbf{K}_{ss} + \frac{\lambda_I n^2}{s^2} \mathbf{K}_{ss} \mathbf{L}_{ss} \mathbf{K}_{ss}.$$

- $m > \sqrt{n}$

$$\mathbf{P} = \frac{m}{s} \mathbf{K}_{ss}^T \mathbf{K}_{ss} + \lambda_A \mathbf{K}_{ss} + \frac{\lambda_I n^2}{s^2} \mathbf{K}_{ss} \mathbf{L}_{ss} \mathbf{K}_{ss}.$$

Better scalability

- 1 Avoid matrix-matrix multiplications
For each iteration of PCG, we need to calculate

$$\mathbf{H}\mathbf{p}_t = (\mathbf{K}_{ms}^T \mathbf{K}_{ms} + \lambda_A \mathbf{K}_{ss} + \lambda_I \mathbf{K}_{ns}^T \mathbf{L} \mathbf{K}_{ns}) \mathbf{p}_t$$

where $\mathbf{p}_t \in \mathbb{R}^s$. If we figure out \mathbf{H} , it needs $\mathcal{O}(ns^2)$ times to perform matrix-matrix multiplications.

While a series of matrix-vector multiplications only need $\mathcal{O}(ns)$ time

$$\mathbf{H}\mathbf{p}_t = \mathbf{K}_{ms}^T (\mathbf{K}_{ms} \mathbf{p}_t) + \lambda_A \mathbf{K}_{ss} \mathbf{p}_t + \lambda_I \mathbf{K}_{ns}^T (\mathbf{L} (\mathbf{K}_{ns} \mathbf{p}_t)).$$

- 2 Block matrix multiplications
 - Kernel matrix \mathbf{K}_{ns} : $\mathcal{O}(ns)$ space
 - Decompose into $s \times s$ size block matrix multiplications: $\mathcal{O}(s^2)$ space
- 3 Space complexity: $\mathcal{O}(s^2)$. Time complexity: $\mathcal{O}(nst + s^3t)$.
 $\mathcal{O}(s^3)$ is due to the computation of $\mathbf{P}^{-1} \mathbf{r}_t$ in each iteration, where $\mathbf{r}_t \in \mathbb{R}^s$.

How many Nyström centers s and iterations t are needed?

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Theorem (Simple version)

Under common assumptions and

$$s \geq \mathcal{O}(\sqrt{n}) \quad \text{and} \quad t \geq \mathcal{O}(\log m)$$

then the following excess risk bound holds with high probability,

$$\mathcal{E}(\hat{f}_{\lambda,t}^s) - \mathcal{E}(f_{\mathcal{H}}) \leq \mathcal{O}\left(\frac{1}{\sqrt{m}}\right).$$

Technical challenges:

- Multi-penalty regularization. [[Rastogi and Sampath, 2017](#)]
- Integral operator for Nyström methods. [[Rudi et al., 2015](#)]
- Convergence of PCG. [[Rudi et al., 2017](#)]

The complexity:

- Space complexity: $\mathcal{O}(s^2) = \mathcal{O}(n)$.
- Time complexity: $\mathcal{O}(nst + s^3t) = \mathcal{O}(n\sqrt{n})$.

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Estimators	Time	Space
RLS-Direct	$\mathcal{O}(m^3)$	$\mathcal{O}(m^2)$
LapRLS-Direct	$\mathcal{O}(n^3)$	$\mathcal{O}(n^2)$
LapRLS-CG	$\mathcal{O}(n^{2.5})$	$\mathcal{O}(n^2)$
LapRLS-PCG	$\mathcal{O}(n^2)$	$\mathcal{O}(n^2)$
Nyström-Direct	$\mathcal{O}(n^2)$	$\mathcal{O}(n)$
Nyström-CG	$\mathcal{O}(n^{1.75})$	$\mathcal{O}(n)$
Nyström-PCG	$\mathcal{O}(n^{1.5})$	$\mathcal{O}(n)$

Table 1: Summary of time complexity and space complexity in terms of various methods. Here, we omit logarithmic terms.

RMSE, Iteration and Runing time

dataset	sample size	RLS-CG	LapRLS-CG	LapRLS-PCG	Nyström-CG	Nyström-PCG
madelon	2000	1.036±0.009	0.990±0.007	0.990±0.007	0.991±0.009	0.991±0.009
space_ga	3107	1.251±0.004	1.210±0.004	1.210±0.004	1.210±0.004	1.210±0.004
abalone	4177	4.55±0.2×10 ³	4.17±0.1×10³	4.17±0.1×10³	4.18±0.1×10 ³	4.18±0.1×10 ³
phishing	11055	0.426±0.049	0.294±0.005	0.273±0.007	0.295±0.005	0.275±0.008
a8a	22696	0.702±0.002	0.664±0.002	0.664±0.002	0.664±0.002	0.664±0.002
w7a	24692	0.291±0.002	0.283±0.002	0.283±0.002	0.284±0.002	0.284±0.002
a9a	32561	0.698±0.005	0.664±0.000	0.664±0.002	0.664±0.000	0.664±0.002
ijcnn1	49990	0.434±0.005	0.389±0.002	0.389±0.002	0.393±0.001	0.463±0.001
cod-rna	59535	0.686±0.002	/	/	0.614±0.001	0.614±0.001
connect-4	67757	0.781±0.015	/	/	0.739±0.002	0.739±0.002
skin_nonskin	245057	3.119±0.023	/	/	2.620±0.043	2.620±0.043
YearPrediction	463715	0.198±0.001	/	/	0.187±0.001	0.187±0.001

	RLS-CG		LapRLS-CG		LapRLS-PCG		Nyström-CG		Nyström-PCG	
	iter	time	iter	time	iter	time	iter	time	iter	time
madelon	32	0.003	13	0.029	6	0.032	12	0.043	1	0.006
space_ga	11	0.004	23	1.220	5	0.569	23	0.113	2	0.016
abalone	64	0.053	98	26.50	4	0.903	94	0.363	2	0.067
phishing	74	0.031	300	24.20	56	8.210	300	2.470	3	0.045
a8a	100	0.068	50	189.1	3	20.98	50	44.71	1	4.370
w7a	13	0.072	32	143.2	2	9.683	213	107.7	1	2.252
a9a	300	0.529	64	1699	3	30.30	65	70.40	1	4.034
ijcnn1	242	8.204	57	2154	9	72.41	53	108.8	5	4.186
cod-rna	96	7.178	/	/	/	/	55	134.6	7	8.154
connect-4	103	11.07	/	/	/	/	154	186.5	10	4.220
skin_nonskin	43	91.39	/	/	/	/	65	1490	3	40.05
YearPrediction	37	236.5	/	/	/	/	94	2479	2	116.1

We randomly select 10% samples ($m = 0.1n$) as labeled data and 10% samples ($s = 0.1n$) as Nyström centers.

Average RMSE for different labeled data proportion

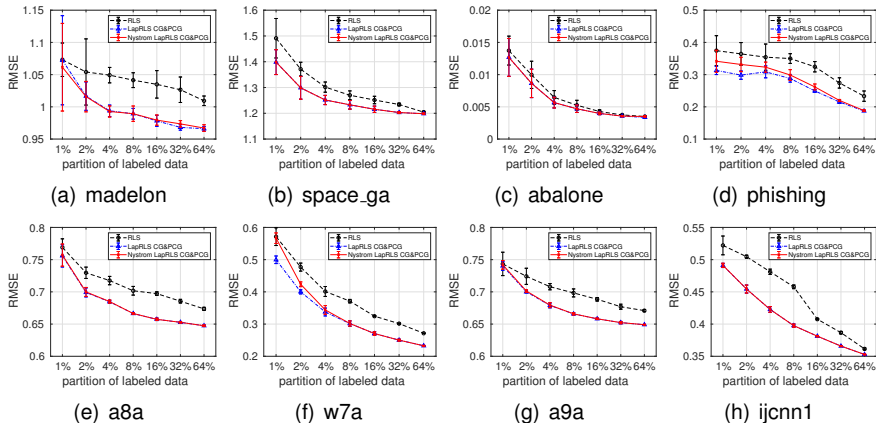


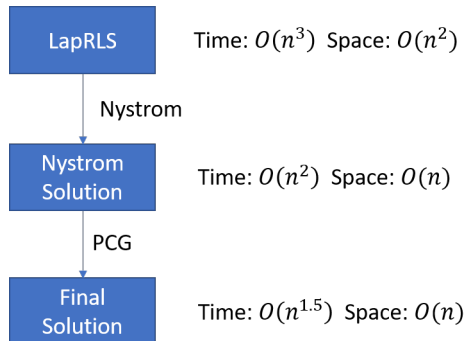
Figure 1: Average RMSE for different labeled data proportion.

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Conclusion

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References

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Any Questions?