

# Kernel Learning with a Million Kernels

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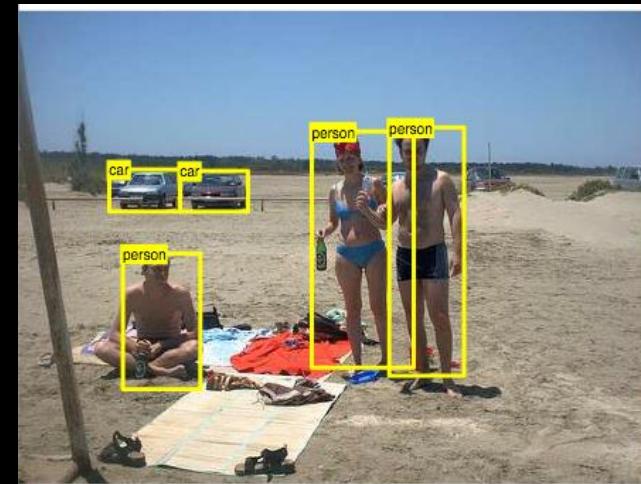
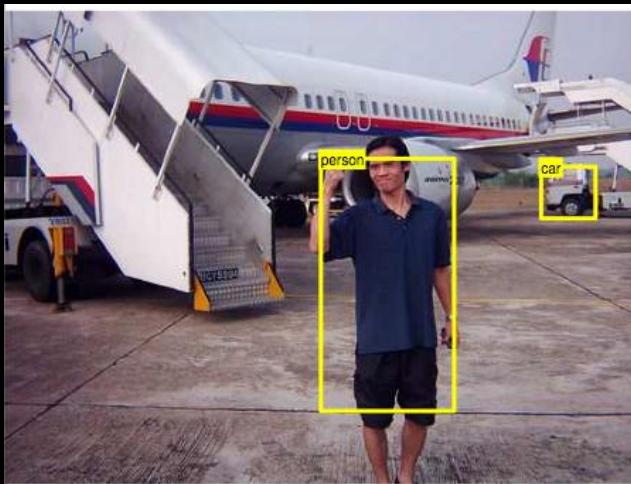
# Kernel Learning

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- The objective in kernel learning is to jointly learn both SVM and kernel parameters from training data.
- Kernel parameterizations
  - Linear :  $K = \sum_i d_i K_i$
  - Non-linear :  $K = \prod_i K_i = \prod_i e^{-d_i D_i}$
- Regularizers
  - Sparse  $l_1$
  - Sparse and non-sparse  $l_{p>1}$
  - Log determinant

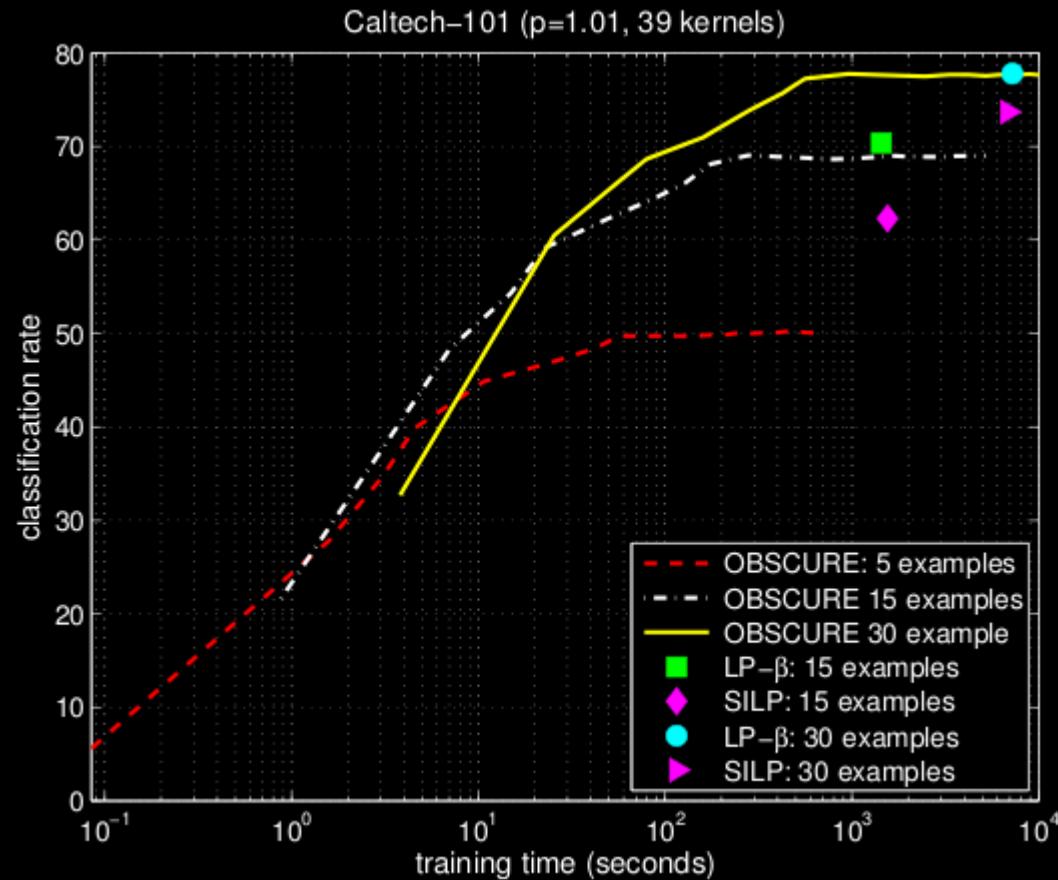
# Kernel Learning for Object Detection

- Vedaldi, Gulshan, Varma and Zisserman ICCV 2009



# Kernel Learning for Object Recognition

- Orabona, Jie and Caputo CVPR 2010



# Kernel Learning for Feature Selection

- Varma and Babu ICML 2009

FERET Gender Identification Data Set

# Feat	AdaBoost	Baluja <i>et al.</i> [IJCV 2007]	OWL-QN [ICML 2007]	LP-SVM [COA 2004]	SSVM QCQP [ICML 2007]	BAHSIC [ICML 2007]	Linear MKL	Non-Linear MKL
10	76.3 ± 0.9	79.5 ± 1.9	71.6 ± 1.4	84.9 ± 1.9	79.5 ± 2.6	81.2 ± 3.2	80.8 ± 0.2	<b>88.7 ± 0.8</b>
20	-	82.6 ± 0.6	80.5 ± 3.3	87.6 ± 0.5	85.6 ± 0.7	86.5 ± 1.3	83.8 ± 0.7	<b>93.2 ± 0.9</b>
30	-	83.4 ± 0.3	84.8 ± 0.4	89.3 ± 1.1	88.6 ± 0.2	89.4 ± 2.4	86.3 ± 1.6	<b>95.1 ± 0.5</b>
50	-	86.9 ± 1.0	88.8 ± 0.4	90.6 ± 0.6	89.5 ± 0.2	91.0 ± 1.3	89.4 ± 0.9	<b>95.5 ± 0.7</b>
80	-	88.9 ± 0.6	90.4 ± 0.2	-	90.6 ± 1.1	92.4 ± 1.4	90.5 ± 0.2	-
100	-	89.5 ± 0.2	90.6 ± 0.3	-	90.5 ± 0.2	94.1 ± 1.3	91.3 ± 1.3	-
150	-	91.3 ± 0.5	90.3 ± 0.8	-	90.7 ± 0.2	94.5 ± 0.7	-	-
252	-	93.1 ± 0.5	-	-	90.8 ± 0.0	94.3 ± 0.1	-	-
76.3(12.6)		-	91 (221.3)	91 (58.3)	90.8 (252)	-	91.6(146.3)	<b>95.5 (69.6)</b>

# The GMKL Primal Formulation

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$$\begin{aligned} P = \text{Min}_{\mathbf{w}, b, \mathbf{d}} \quad & \frac{1}{2} \mathbf{w}^t \mathbf{w} + C \sum_i L(\mathbf{w}^t \Phi_{\mathbf{d}}(\mathbf{x}_i) + b, y_i) + r(\mathbf{d}) \\ \text{s. t.} \quad & \mathbf{d} \in D \end{aligned}$$

- $K_{\mathbf{d}}(\mathbf{x}_i, \mathbf{x}_j) = \Phi_{\mathbf{d}}^t(\mathbf{x}_i) \Phi_{\mathbf{d}}(\mathbf{x}_j) > 0 \quad \forall \mathbf{d} \in D$
- $\nabla_{\mathbf{d}} K$  and  $\nabla_{\mathbf{d}} r$  exist and are continuous

# The GMKL Primal Formulation

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- The GMKL primal formulation for binary classification.

$$\begin{aligned} P = \text{Min}_{\mathbf{w}, b, \mathbf{d}, \xi} \quad & \frac{1}{2} \mathbf{w}^t \mathbf{w} + C \sum_i \xi_i + r(\mathbf{d}) \\ \text{s. t.} \quad & y_i (\mathbf{w}^t \Phi_{\mathbf{d}}(\mathbf{x}_i) + b) \geq 1 - \xi_i \\ & \xi_i \geq 0 \text{ & } \mathbf{d} \in D \end{aligned}$$

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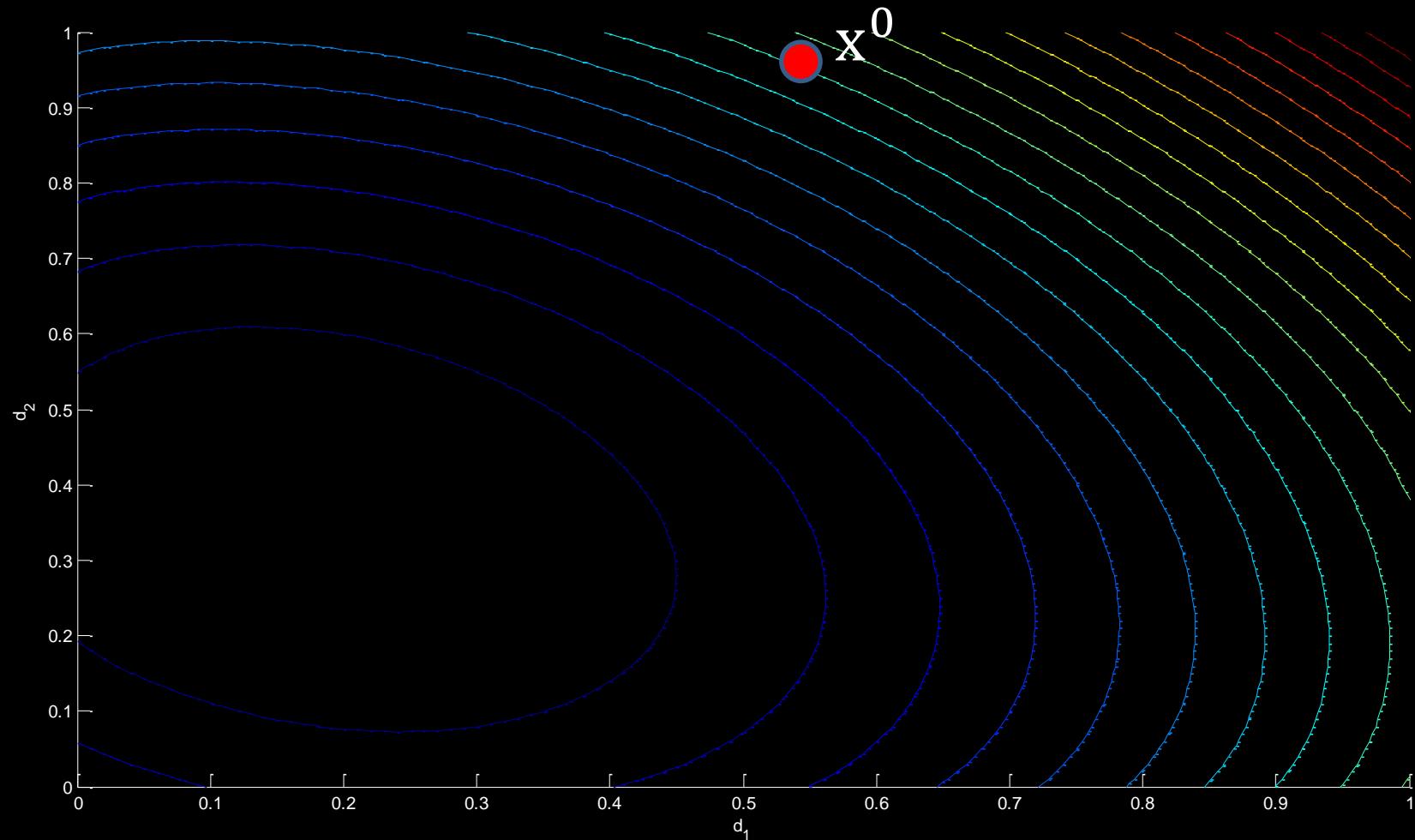
$$\begin{aligned} P = \text{Min}_{\mathbf{w}, b, \mathbf{d}, \xi} \quad & \frac{1}{2} \mathbf{w}^t \mathbf{w} + C \sum_i \xi_i + r(\mathbf{d}) \\ \text{s. t.} \quad & y_i (\mathbf{w}^t \Phi_{\mathbf{d}}(\mathbf{x}_i) + b) \geq 1 - \xi_i \\ & \xi_i \geq 0 \text{ & } \mathbf{d} \in D \end{aligned}$$

- Intermediate Dual

$$\begin{aligned} D = \text{Min}_{\mathbf{d}} \text{Max}_{\alpha} \quad & \mathbf{1}^t \alpha - \frac{1}{2} \alpha^t \mathbf{Y} \mathbf{K}_{\mathbf{d}} \mathbf{Y} \alpha + r(\mathbf{d}) \\ \text{s. t.} \quad & \mathbf{1}^t \mathbf{Y} \alpha = 0 \\ & 0 \leq \alpha \leq C \text{ & } \mathbf{d} \in D \end{aligned}$$

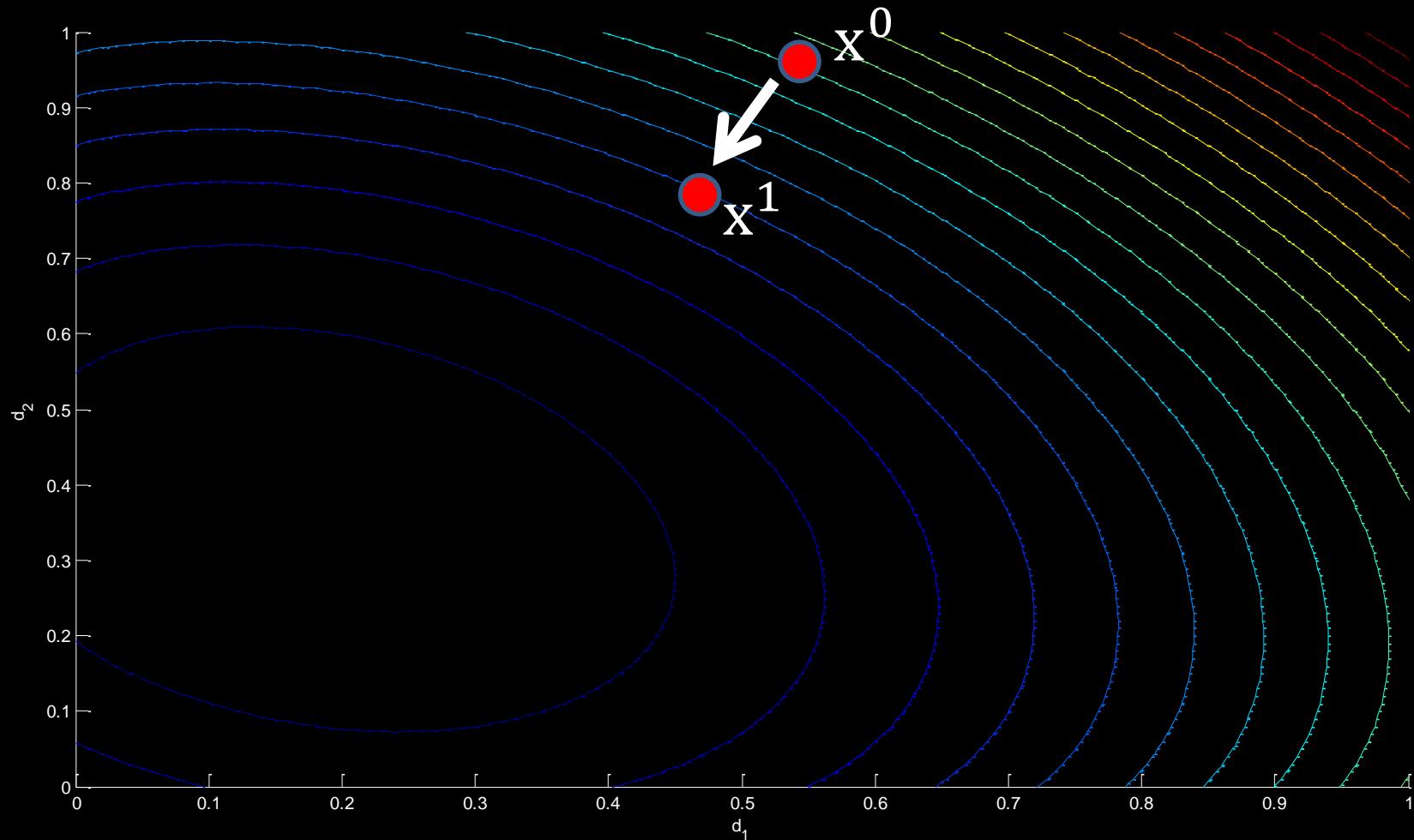
# Projected Gradient Descent

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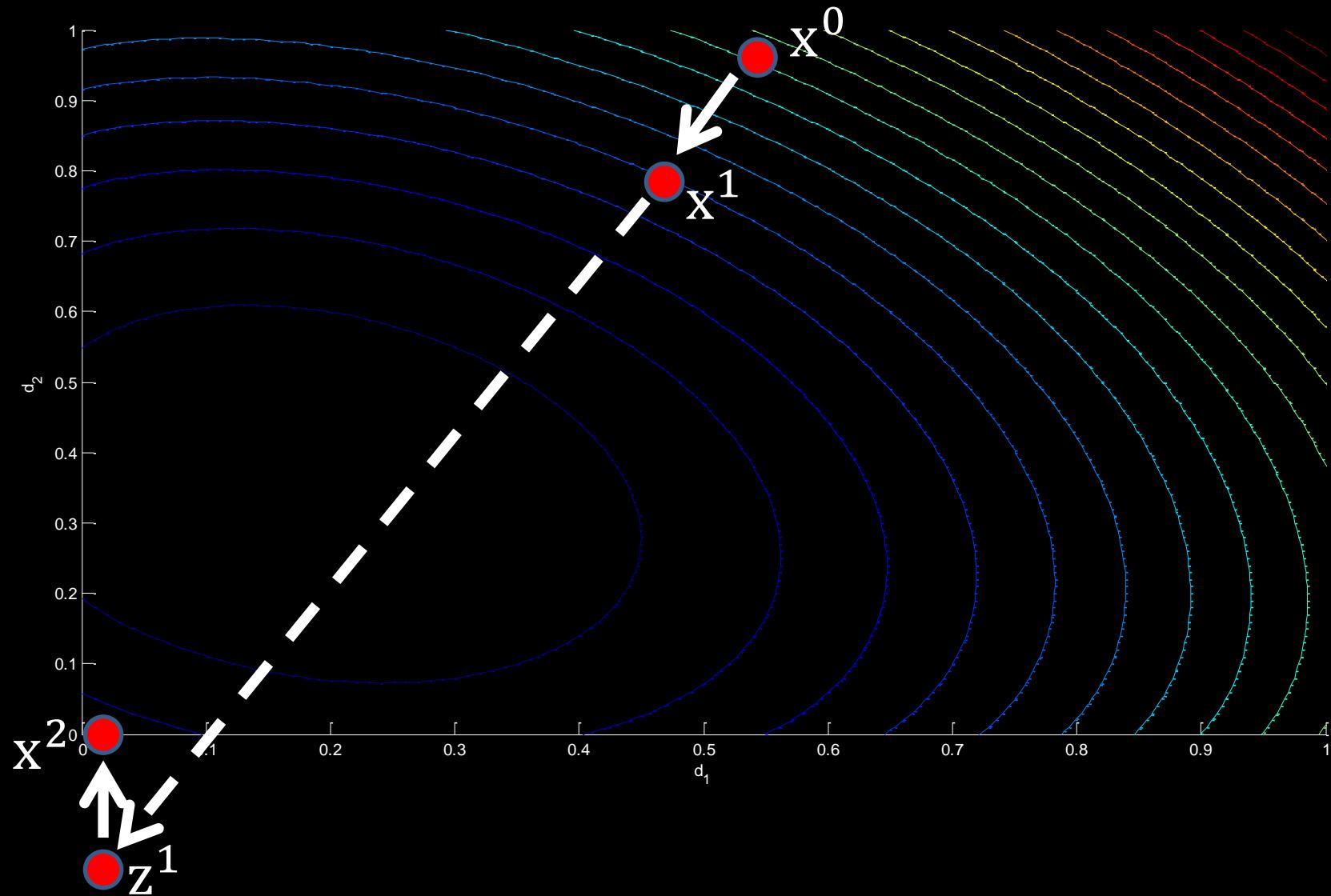


# Projected Gradient Descent

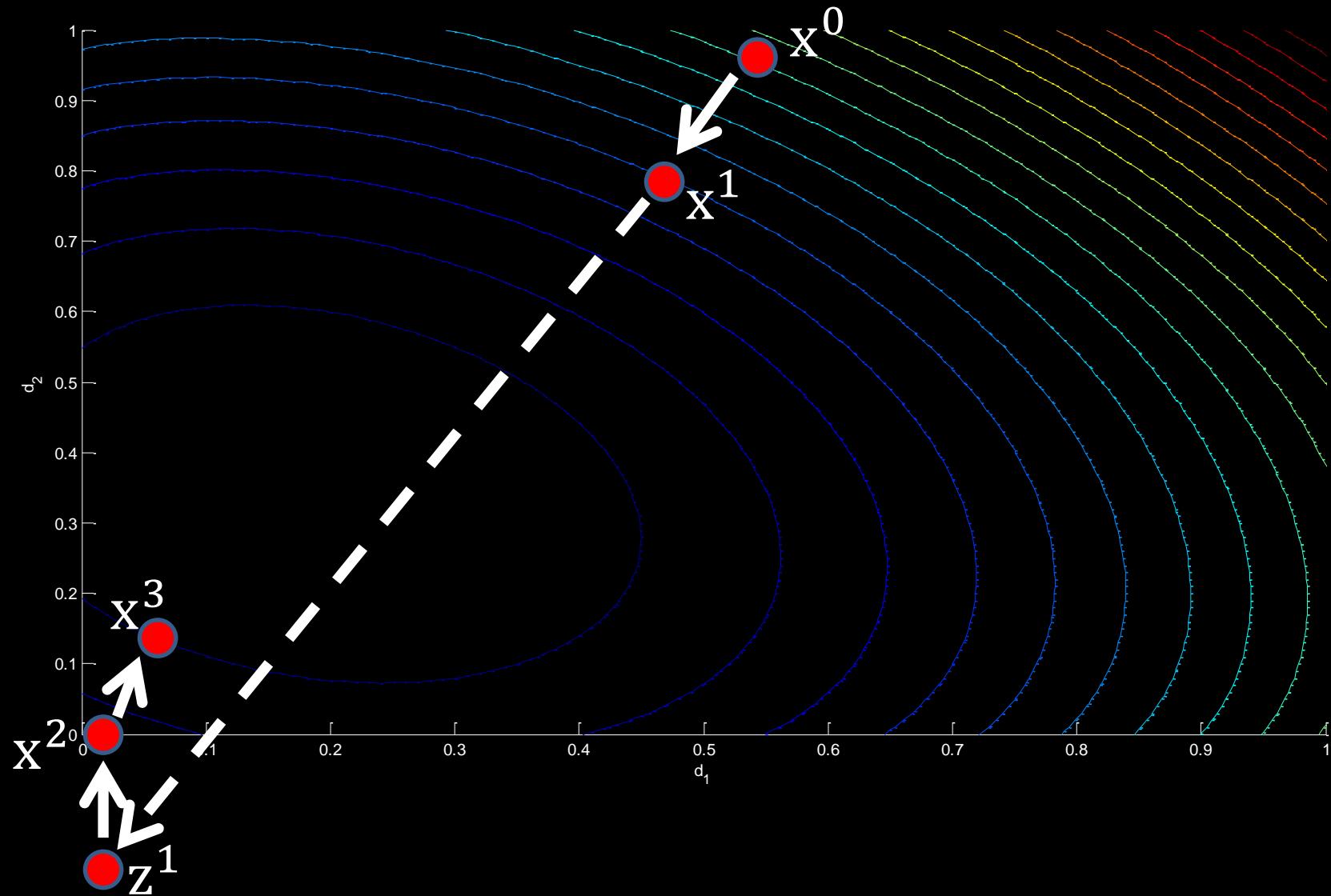
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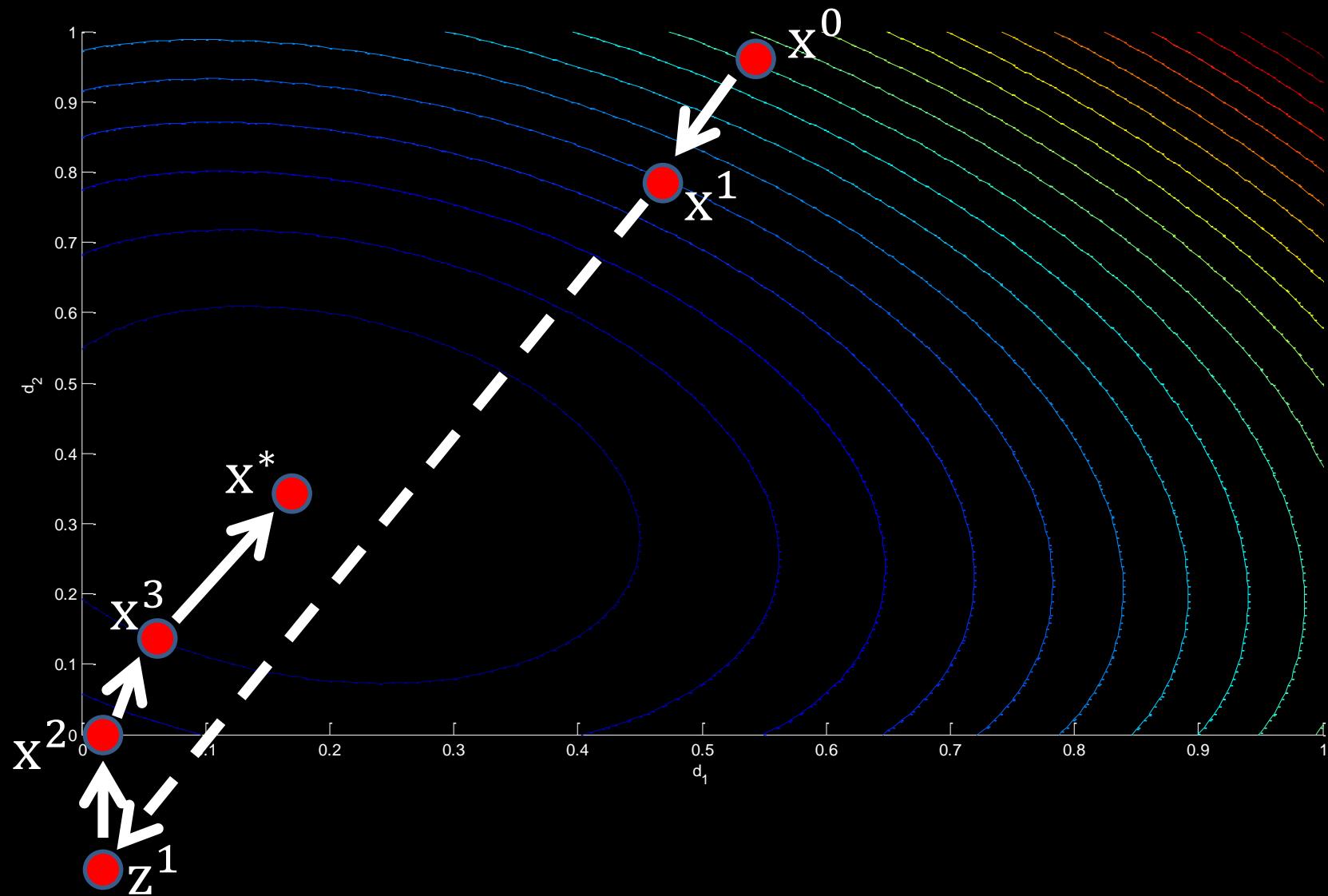
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# PGD Limitations

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- PGD requires many function and gradient evaluations as
  - No step size information is available.
  - The Armijo rule might reject many step size proposals.
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- Solving SVMs to high precision to obtain accurate function and gradient values is very expensive.

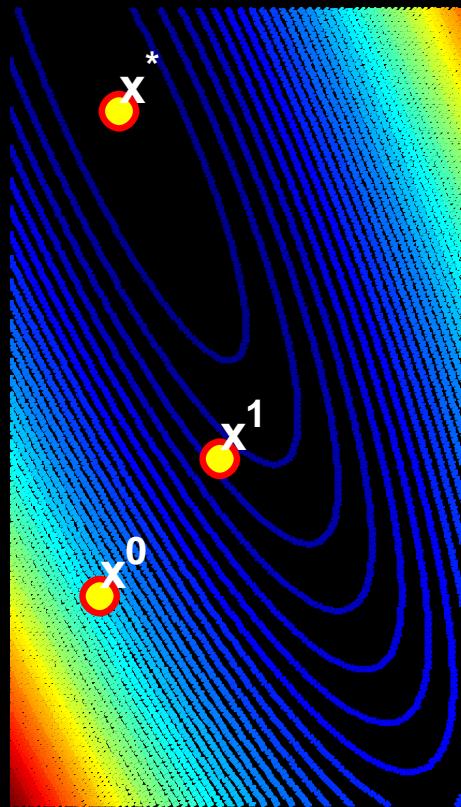
# PGD Limitations

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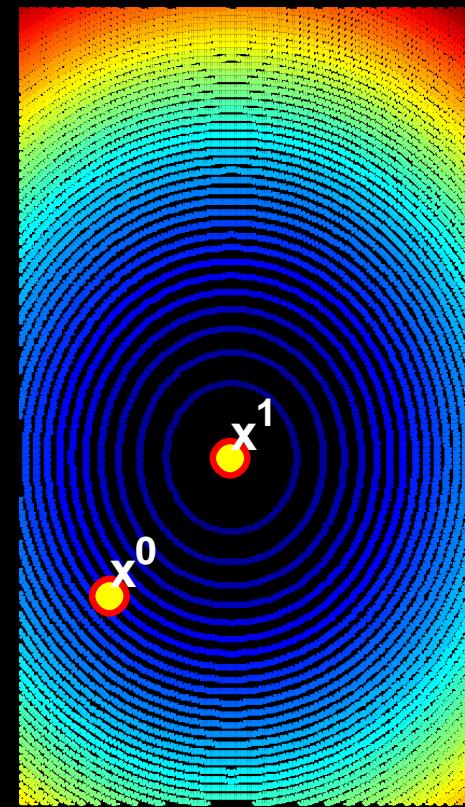
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  - The Armijo rule might reject many step size proposals.
  - Inaccurate gradient values can lead to many tiny steps.
- Noisy function and gradient values can cause PGD to converge to points far away from the optimum.
- Solving SVMs to high precision to obtain accurate function and gradient values is very expensive.
- Repeated projection onto the feasible set might also be expensive.

# SPG Solution – Spectral Step Length

- Quadratic approximation :  $\frac{1}{2}\lambda^{-1}\mathbf{x}^t\mathbf{x} + \mathbf{c}^t\mathbf{x} + d$
- Spectral step length :  $\lambda_{SPG} = \frac{\langle \mathbf{x}^n - \mathbf{x}^{n-1}, \mathbf{x}^n - \mathbf{x}^{n-1} \rangle}{\langle \mathbf{x}^n - \mathbf{x}^{n-1}, \nabla f(\mathbf{x}^n) - \nabla f(\mathbf{x}^{n-1}) \rangle}$



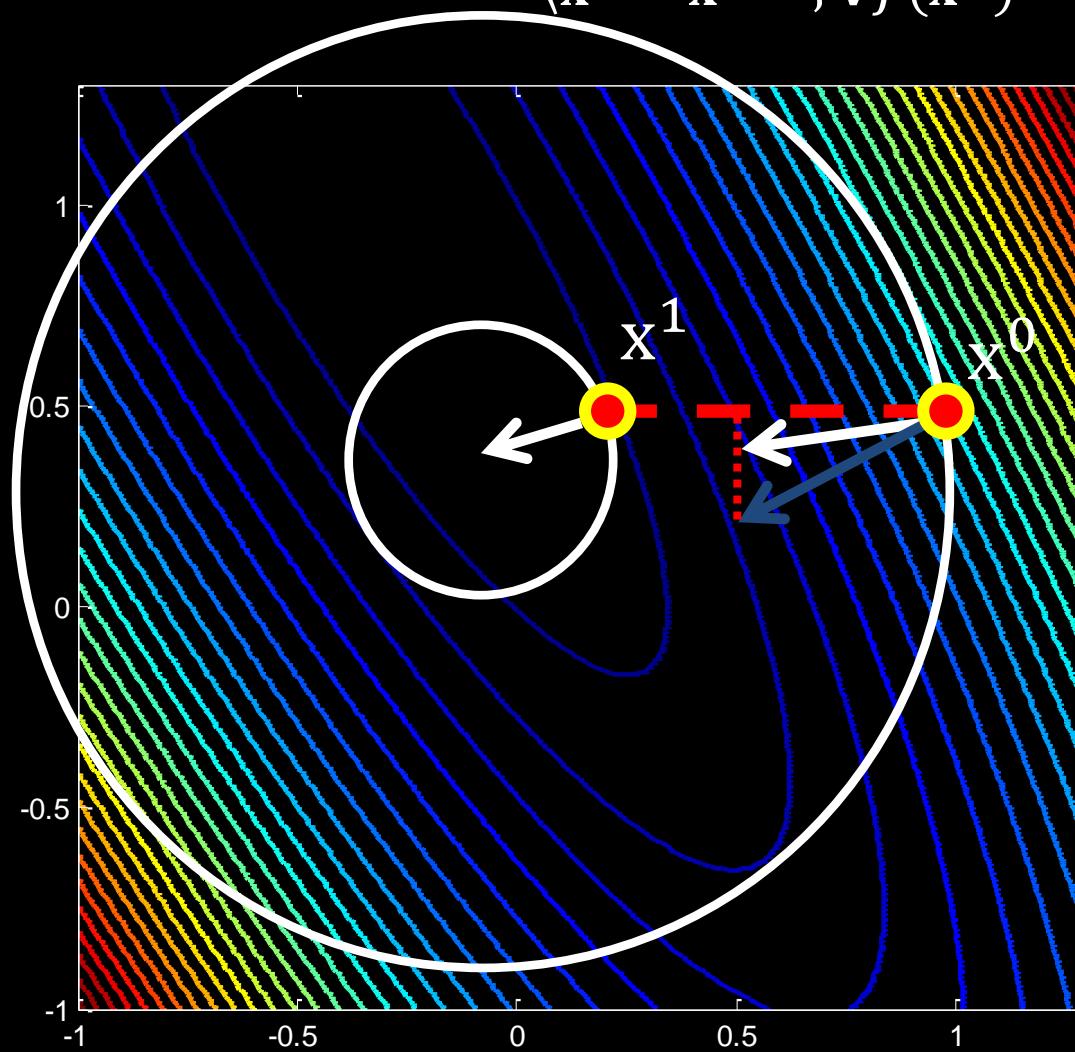
Original Function



Approximation

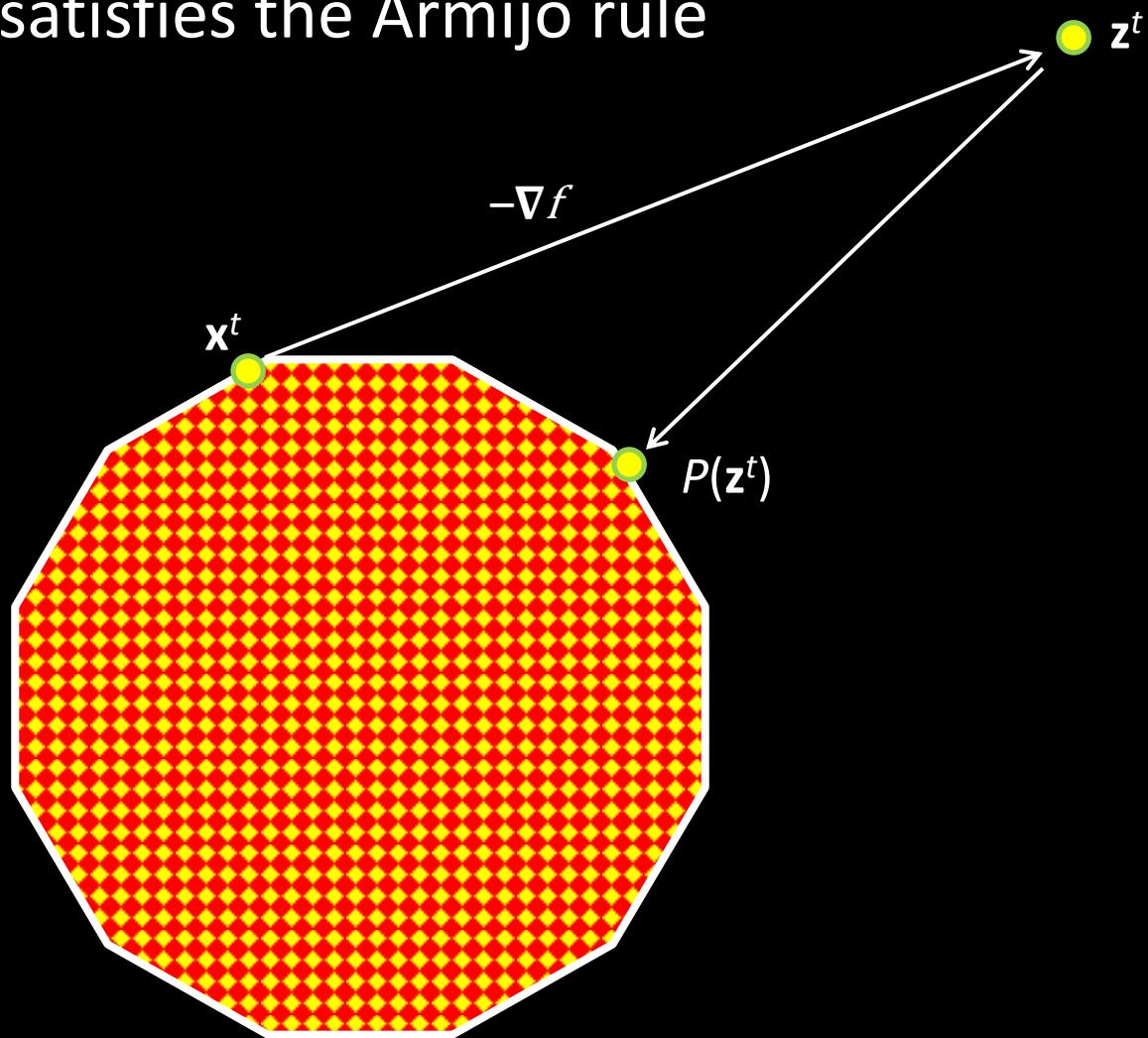
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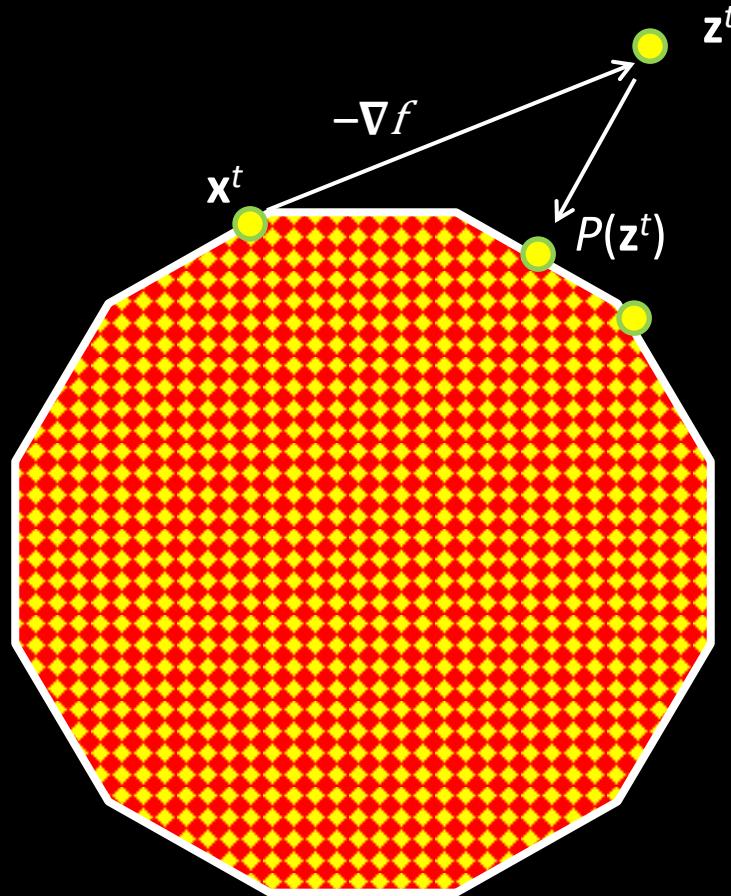
# PGD Limitations – Repeated Projections

- Accept  $P(\mathbf{z}^t)$  if it satisfies the Armijo rule



# PGD Limitations – Repeated Projections

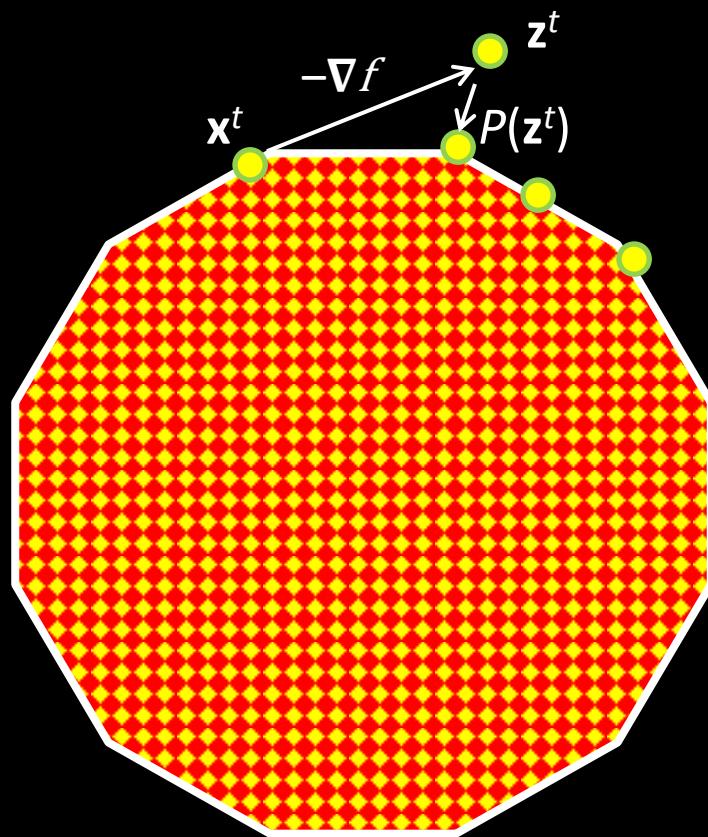
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# PGD Limitations – Repeated Projections

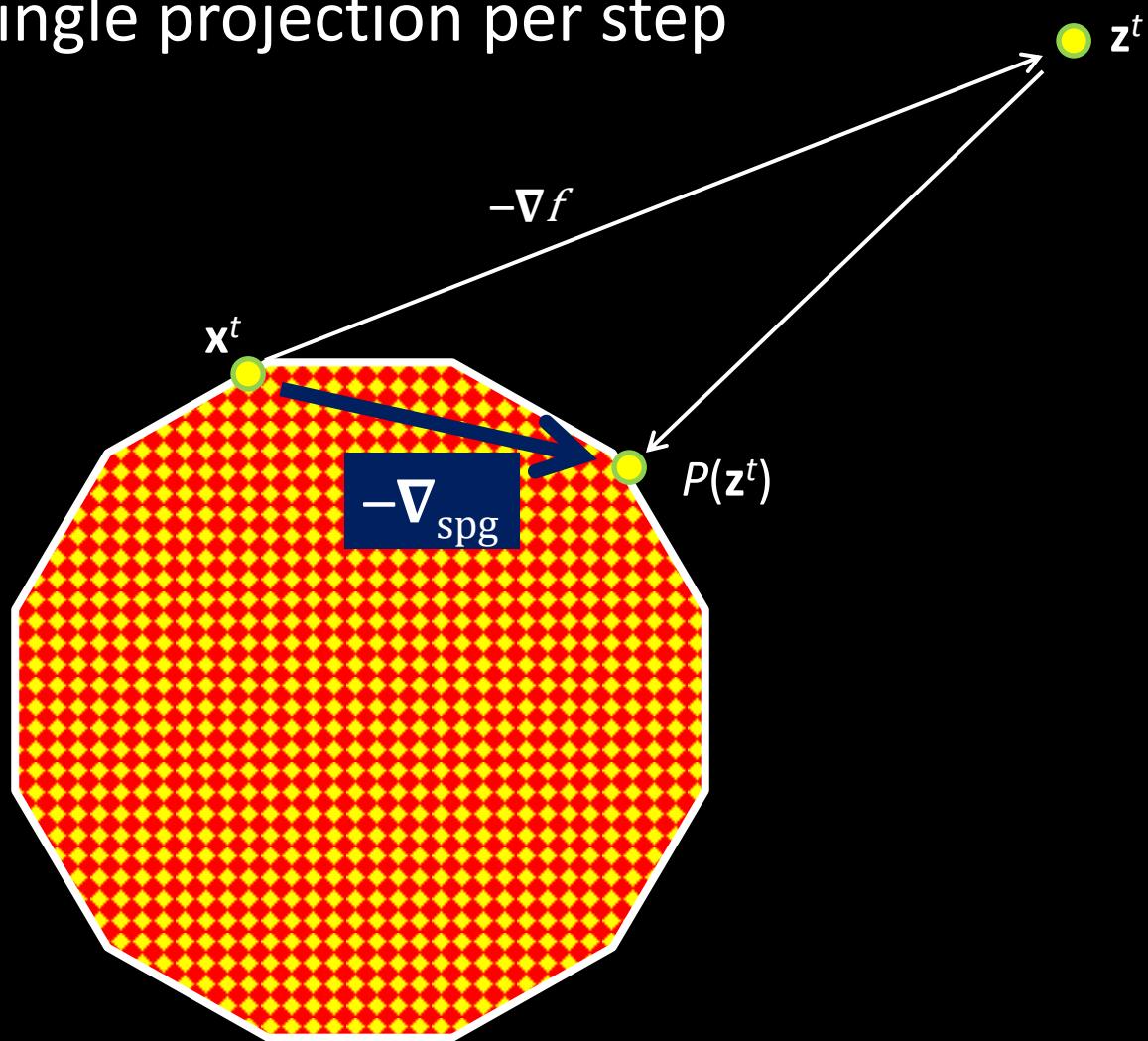
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- PGD might require many projections before accepting a point



# SPG Solution – Spectral Proj Gradient

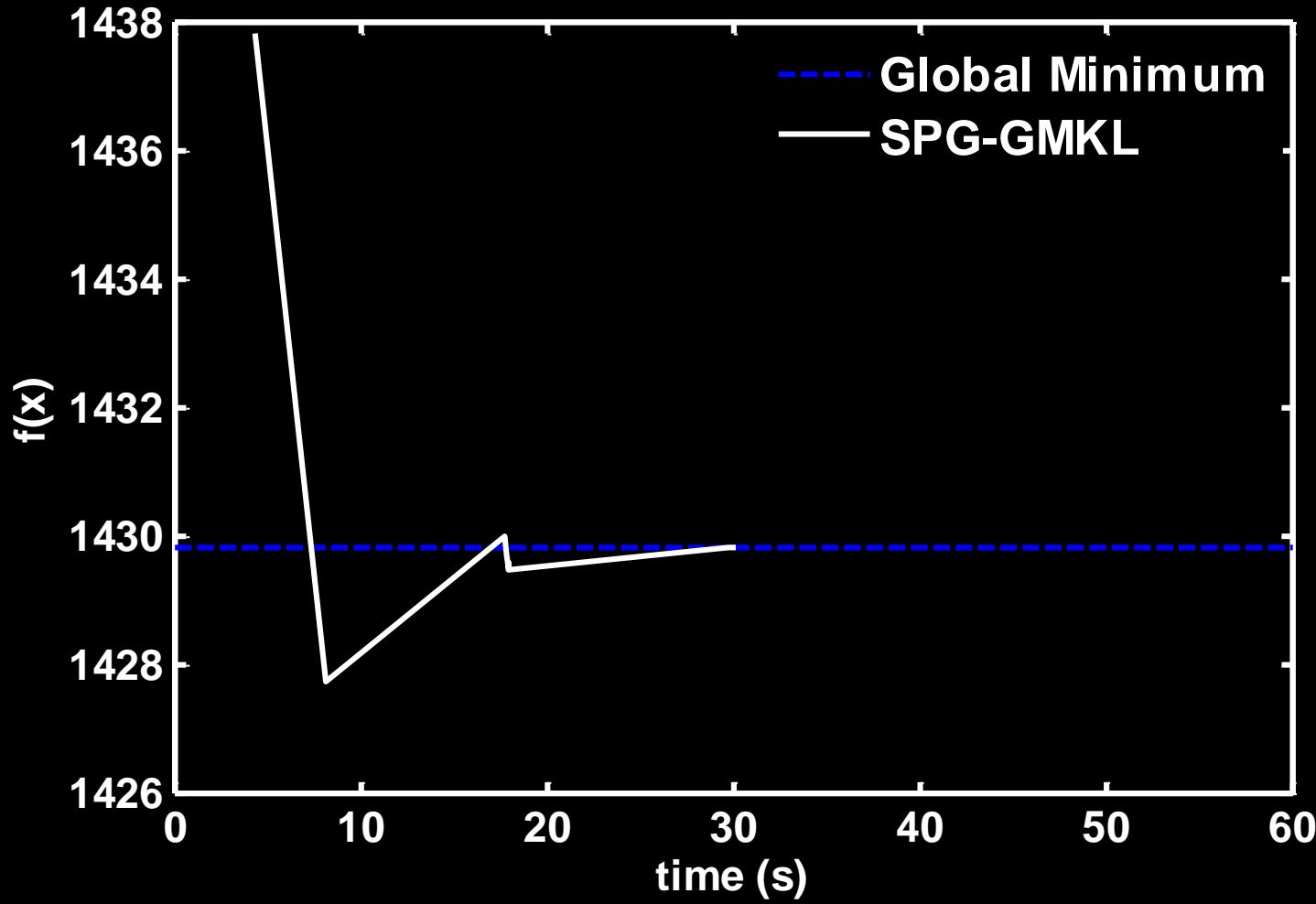
- SPG requires a single projection per step



# SPG Solution – Non-Monotone Rule

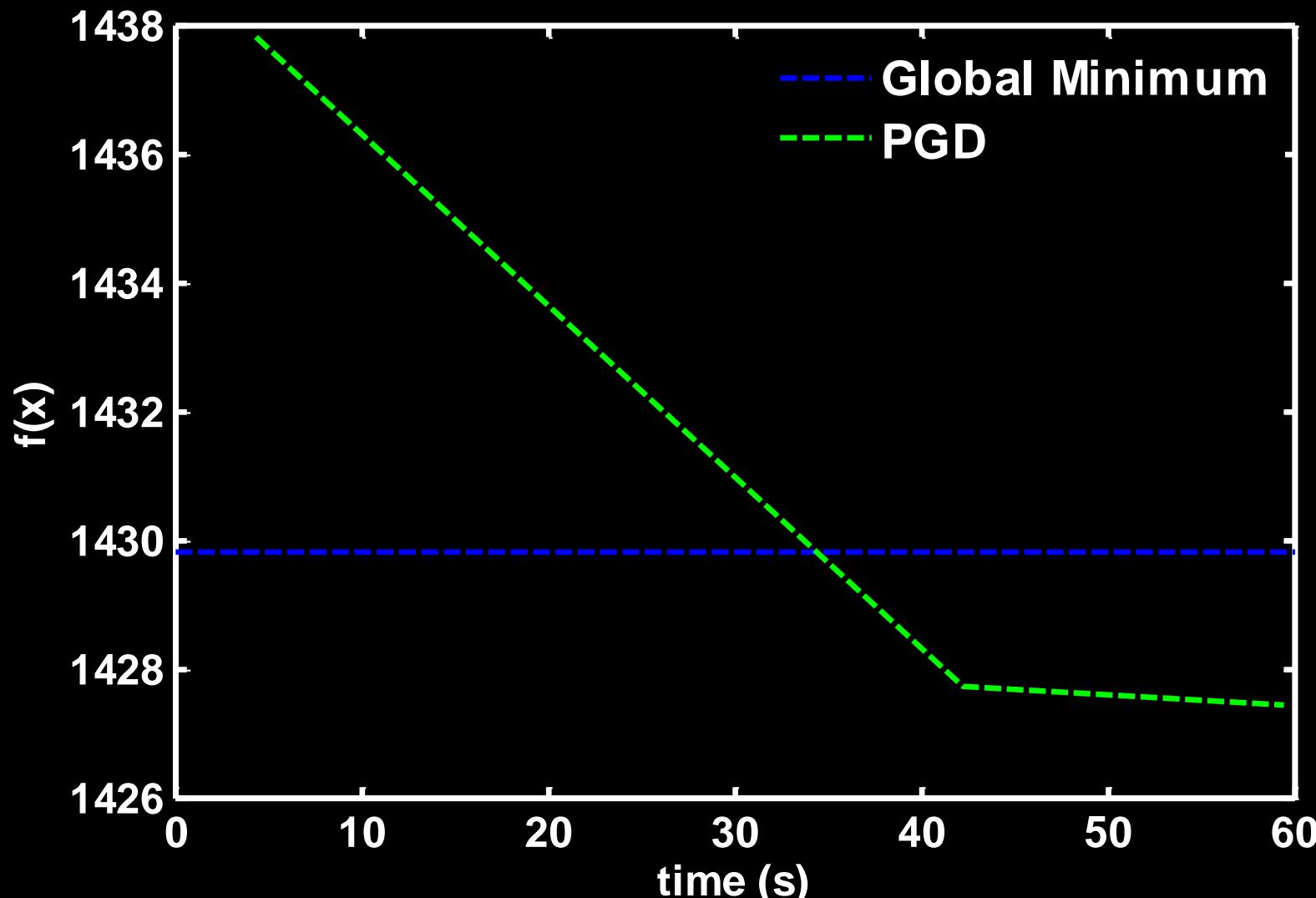
- Handling function and gradient noise.

- Non-monotone rule:  $f(x^t - s\nabla f(x^t)) \leq \max_{0 \leq j \leq M} f(x^{t-j}) - \gamma s |\nabla f(x^t)|_2^2$

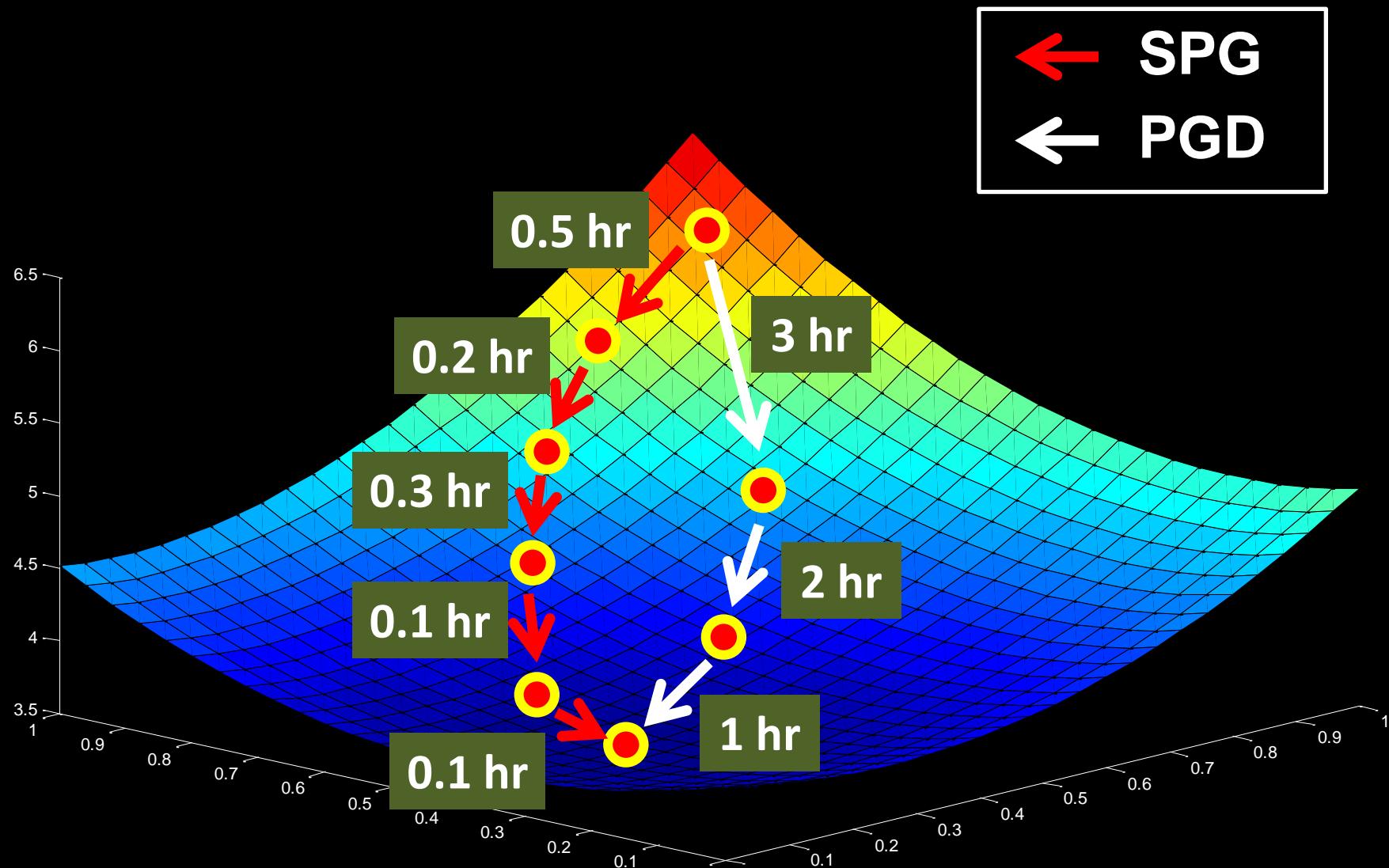


# PGD Limitations – Step Size Selection

- The Armijo rule might get stuck due to noisy function values



# SPG Solution – SVM Precision Tuning



# SPG Advantages

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- SPG requires fewer function and gradient evaluations due to
  - The 2<sup>nd</sup> order spectral step length estimation.
  - The non-monotone line search criterion.
- SPG is more robust to noisy function and gradient values due to the non-monotone line search criterion.
- SPG never needs to solve an SVM with high precision due to our precision tuning strategy.
- SPG needs to perform only a single projection per step.

# SPG Algorithm

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- 1:  $n \leftarrow 0$
- 2: Initialize  $\mathbf{d}^0$  randomly
- 3: **repeat**
- 4:    $\alpha^* \leftarrow \text{SolveSVM}(\mathbf{K}(\mathbf{d}^n), \epsilon)$
- 5:    $\lambda \leftarrow \text{SpectralStepLength}$
- 6:    $\mathbf{p}^n \leftarrow \mathbf{d}^n - \mathbf{P}(\mathbf{d}^n - \lambda \nabla W(\mathbf{d}^n, \alpha^*))$
- 7:    $s^n \leftarrow \text{Non-Monotone}$
- 8:    $\epsilon \leftarrow \text{TuneSMPrecision}$
- 9:    $\mathbf{d}^{n+1} \leftarrow \mathbf{d}^n - s^n \mathbf{p}^n$
- 10: **until** converged

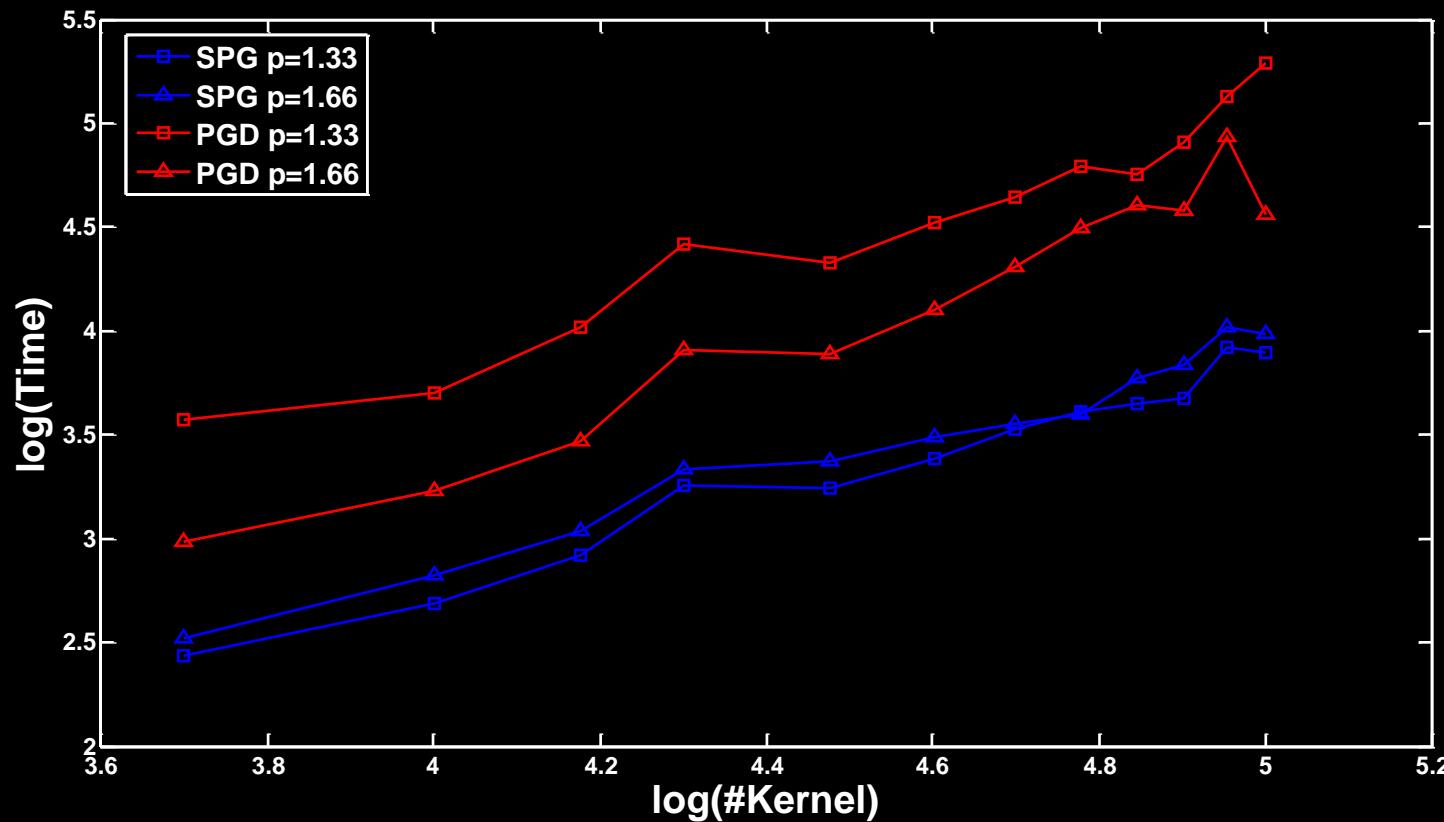
# Results on Large Scale Data Sets

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- Covertype: Sum of kernels subject to  $l_{1.33}$  regularization
  - Number of training points 581,012
  - Number of Kernels 5
  - SPG time taken 64.46 hrs
- SPG took 26 SVM evaluations
- First SVM evaluation took 44 hours
- Only 0.19% of SV were cached

# Results on Large Scale Data Sets

- Sonar: Sum of kernels subject to  $l_{1.33}$  regularization
  - Number of training points 208
  - Number of Kernels 1 Million
  - SPG time taken 105.62 hrs



# Results on Large Scale Data Sets

- Sum of kernels subject to  $l_{p \geq 1}$  regularization

Data Sets	# Train	# Kernels	$p=1$		$p=1.33$	
			PGD (hrs)	SPG (hrs)	PGD (hrs)	SPG (hrs)
Adult - 9	32,561	50	35.84	4.55	31.77	4.42
Cod - RNA	59,535	50	-	25.17	66.48	19.10
KDDCup04	50,000	50	-	40.10	-	42.20

# Results on Small Scale Data Sets

- Sum of kernels subject to  $l_1$  regularization

Data Sets	SimpleMKL (s)	Shogun (s)	PGD (s)	SPG (s)
Wpbc	$400 \pm 128.4$	$15 \pm 7.7$	$38 \pm 17.6$	$6 \pm 4.2$
Breast - Cancer	$676 \pm 356.4$	$12 \pm 1.2$	$57 \pm 85.1$	$5 \pm 0.6$
Australian	$383 \pm 33.5$	$1094 \pm 621.6$	$29 \pm 7.1$	$10 \pm 0.8$
Ionosphere	$1247 \pm 680.0$	$107 \pm 18.8$	$1392 \pm 824.2$	$39 \pm 6.8$
Sonar	$1468 \pm 1252.7$	$935 \pm 65.0$	–	$273 \pm 64.0$

# Results on Large Scale Data Sets

- Product of kernels subject to  $l_{p \geq 1}$  regularization

Data Sets	# Train	# Kernels	$p=1$		$p=1.33$	
			PGD (hrs)	SPG (hrs)	PGD (hrs)	SPG (hrs)
Letter	20,000	16	18.66	0.67	18.69	0.66
Poker	25,010	10	5.57	0.49	2.29	0.96
Adult - 8	22,696	42	–	1.73	–	3.42
Web - 7	24,692	43	–	0.88	–	1.33
RCV1	20,242	50	–	18.17	–	15.93
Cod - RNA	59,535	8	–	3.45	–	8.99

# Effect of Individual Components

- Sum of kernels subject to  $l_{1.1}$  regularization

Data Sets	PGD		PGD + N		PGD + S		PGD + N + S	
	Time (s)	# SVMs	Time (s)	# SVMs	Time (s)	# SVMs	Time (s)	# SVMs
Australian	$39.4 \pm 6.0$	3230	$32.7 \pm 3.6$	116	$317.0 \pm 49.1$	5980	$7.0 \pm 1.6$	621
Sonar	$785.5 \pm 471.1$	209461	$41.6 \pm 17.1$	3236	$40.2 \pm 24.6$	3806	$9.0 \pm 1.8$	2427
Breast - Cancer	$237.3 \pm 97.8$	109599	$42.2 \pm 4.1$	1187	$14.9 \pm 2.2$	3537	$8.6 \pm 2.2$	3006
Diabetes	$73.6 \pm 38.8$	29347	$26.3 \pm 9.5$	2966	$10.5 \pm 2.6$	1239	$4.1 \pm 0.5$	695
Wpbc	$44.4 \pm 11.6$	14376	$27.9 \pm 13.6$	9388	$2.9 \pm 0.8$	340	$1.2 \pm 0.4$	79

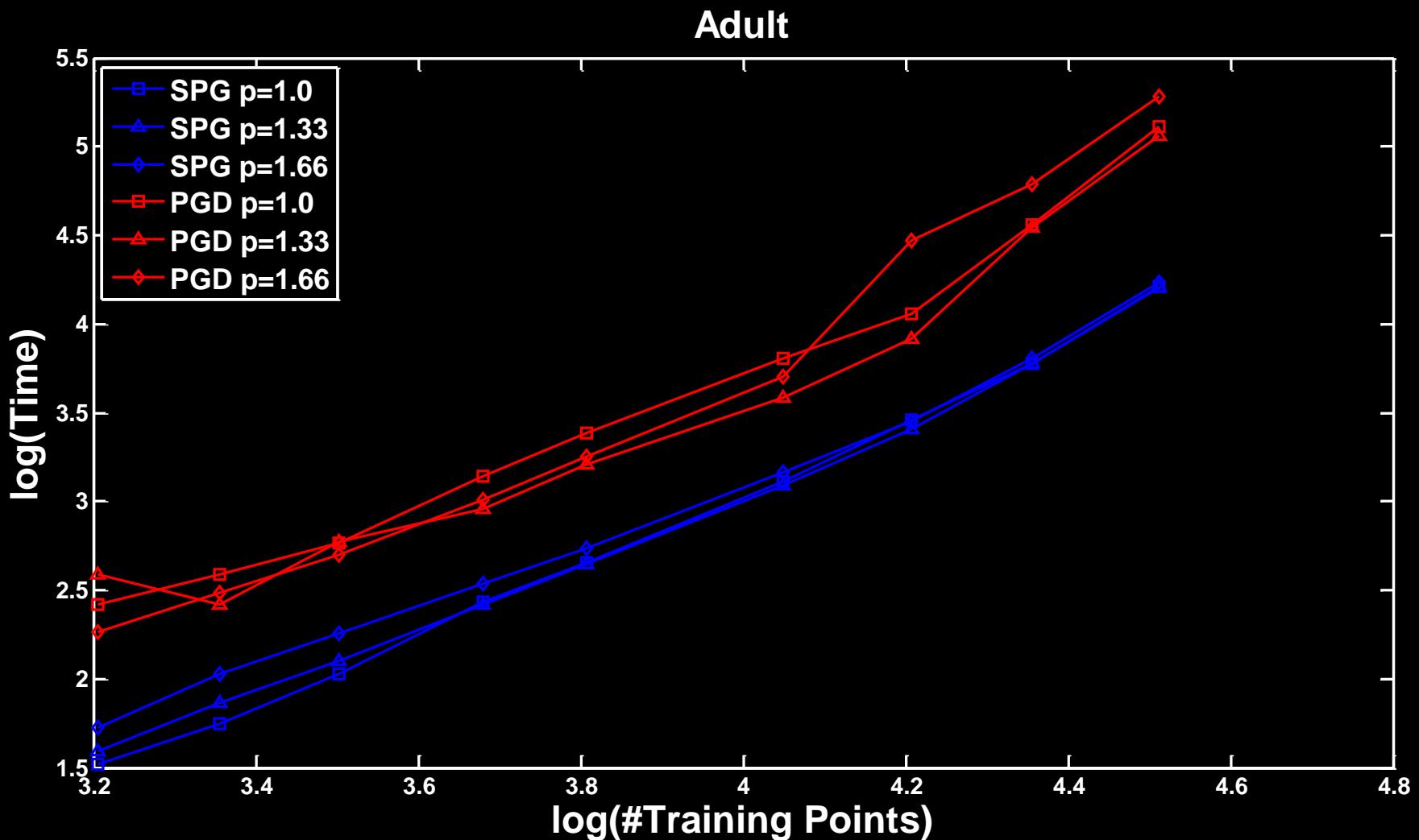
# SVM Precision Tuning

- Sum of kernels subject to  $l_{1.33}$  regularization

Data Sets	# Train	# Kernels	PGD (hrs)	PGD + N + S (hrs)	SPG (hrs)
Adult - 9	32,561	50	31.77	8.33	4.43
Web - 8	49,749	50	4.27	1.73	0.87
Sonar	208	100,000	53.91	3.35	2.19

# SPG Scaling Properties

- Scaling with the number of training points



# Conclusions

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- Developed a generic and efficient MKL optimizer.
- Experimented with four different MKL formulations and solved both small and large scale problems.
- Combining spectral step length and non-monotone rule gives best performance.
- Quasi Newton methods not suitable for MKL problems due to noisy gradient.

Code: <http://research.microsoft.com/en-us/um/people/manik/code/SPG-GMKL/download.html>

# Acknowledgements

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- Subhashis Banerjee (IITD)
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