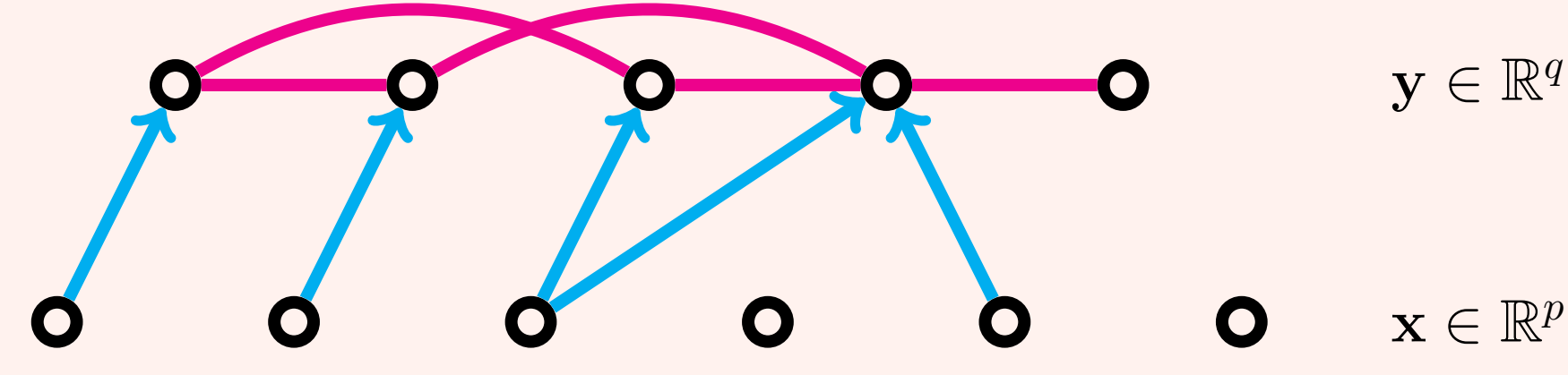


Motivation

Sparse Conditional Gaussian Graphical Model

$$p(\mathbf{y}|\mathbf{x}; \Lambda, \Theta) = \exp\{-\mathbf{y}^T \Lambda \mathbf{y} - 2\mathbf{x}^T \Theta \mathbf{y}\} / Z(\mathbf{x})$$

$$Z(\mathbf{x}) = (2\pi)^{q/2} |\Lambda|^{-1} \exp(\mathbf{x}^T \Theta \Lambda^{-1} \Theta^T \mathbf{x})$$



Sparse Estimation

Given empirical covariances: $\mathbf{S}_{xx} \in \mathbb{R}^{p \times p}$, $\mathbf{S}_{xy} \in \mathbb{R}^{p \times q}$, $\mathbf{S}_{yy} \in \mathbb{R}^{q \times q}$

$$\min_{\Lambda > 0, \Theta} f(\Lambda, \Theta) = g(\Lambda, \Theta) + h(\Lambda, \Theta)$$

$$g(\Lambda, \Theta) = -\log |\Lambda| + \text{tr}(\mathbf{S}_{yy} \Lambda + 2\mathbf{S}_{xy}^T \Theta + \Lambda^{-1} \Theta^T \mathbf{S}_{xx} \Theta)$$

$$h(\Lambda, \Theta) = \lambda_\Lambda \|\Lambda\|_1 + \lambda_\Theta \|\Theta\|_1$$

Convex but **difficult** problem due to **last term**

Previous Optimization Algorithms

- OWL-QN [1]
- FISTA [2]
- **Newton Coordinate Descent [3]**
 - Second order approximation minimized over active set
 - Proximal Newton subproblem solved via coordinate descent
 - Step size found via backtracking

Second order approximation on both Λ and Θ

$$\bar{g}_{\Lambda, \Theta}(\Delta_\Lambda, \Delta_\Theta) = \text{vec}(\nabla g(\Lambda, \Theta))^T \text{vec}([\Delta_\Lambda \ \Delta_\Theta]) + \frac{1}{2} \text{vec}([\Delta_\Lambda \ \Delta_\Theta])^T \nabla^2 g(\Lambda, \Theta) \text{vec}([\Delta_\Lambda \ \Delta_\Theta])$$

Scalability Problems

Time: Genomic dataset with $p = 34k$, $q = 10k$: **> 50 hours**
Memory: Requires $O(pq+q^2)$ memory: **>100 Gb** when $p+q = 80k$

$$\nabla g(\Lambda, \Theta) = [\mathbf{S}_{yy} - \Sigma - \Psi \quad 2\mathbf{S}_{xy} + 2\Gamma]$$

$$\nabla^2 g(\Lambda, \Theta) = \begin{bmatrix} \Sigma \otimes (\Sigma + 2\Psi) & -2\Sigma \otimes \Gamma^T \\ -2\Sigma \otimes \Gamma & 2\Sigma \otimes \mathbf{S}_{xx} \end{bmatrix}$$

$$\Sigma = \Lambda^{-1}$$

$$\Gamma = \mathbf{S}_{xx} \Theta \Lambda^{-1}$$

$$\Psi = \Lambda^{-1} \Theta^T \mathbf{S}_{xx} \Theta \Lambda^{-1}$$

Alternating Newton Coordinate Descent

Update Λ given fixed Θ :

- Solve for Newton direction via CD
- $\bar{g}_{\Lambda, \Theta}(\Delta_\Lambda) = \text{vec}(\nabla_\Lambda g(\Lambda, \Theta))^T \text{vec}(\Delta_\Lambda) + \frac{1}{2} \text{vec}(\Delta_\Lambda)^T \nabla_\Lambda^2 g(\Lambda, \Theta) \text{vec}(\Delta_\Lambda)$
- Run backtracking line search

faster computation

Update Θ given fixed Λ :

- Solve *Lasso* problem directly via CD
- $g_\Lambda(\Theta) = \text{tr}(2\mathbf{S}_{xy}^T \Theta + \Lambda^{-1} \Theta^T \mathbf{S}_{xx} \Theta)$

Second order approximation only on Λ

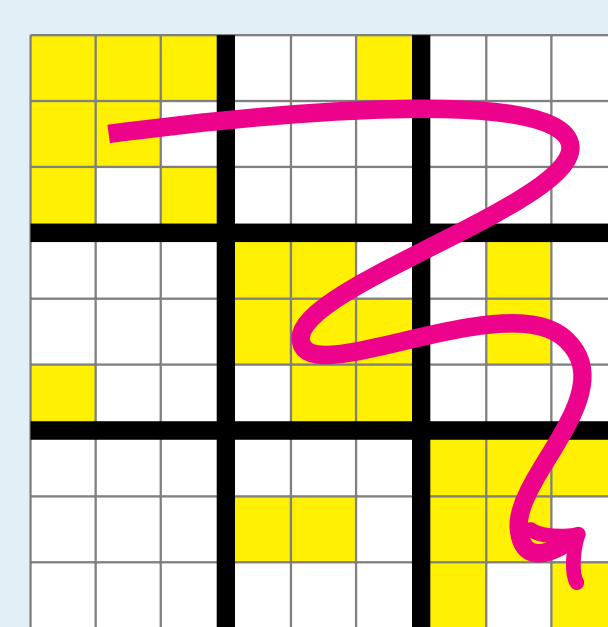
- ✓ Eliminate Γ
- ✓ Reduce CD time complexity
- ✓ Backtrack only for Λ – faster early convergence

Alternating Newton Block Coordinate Descent

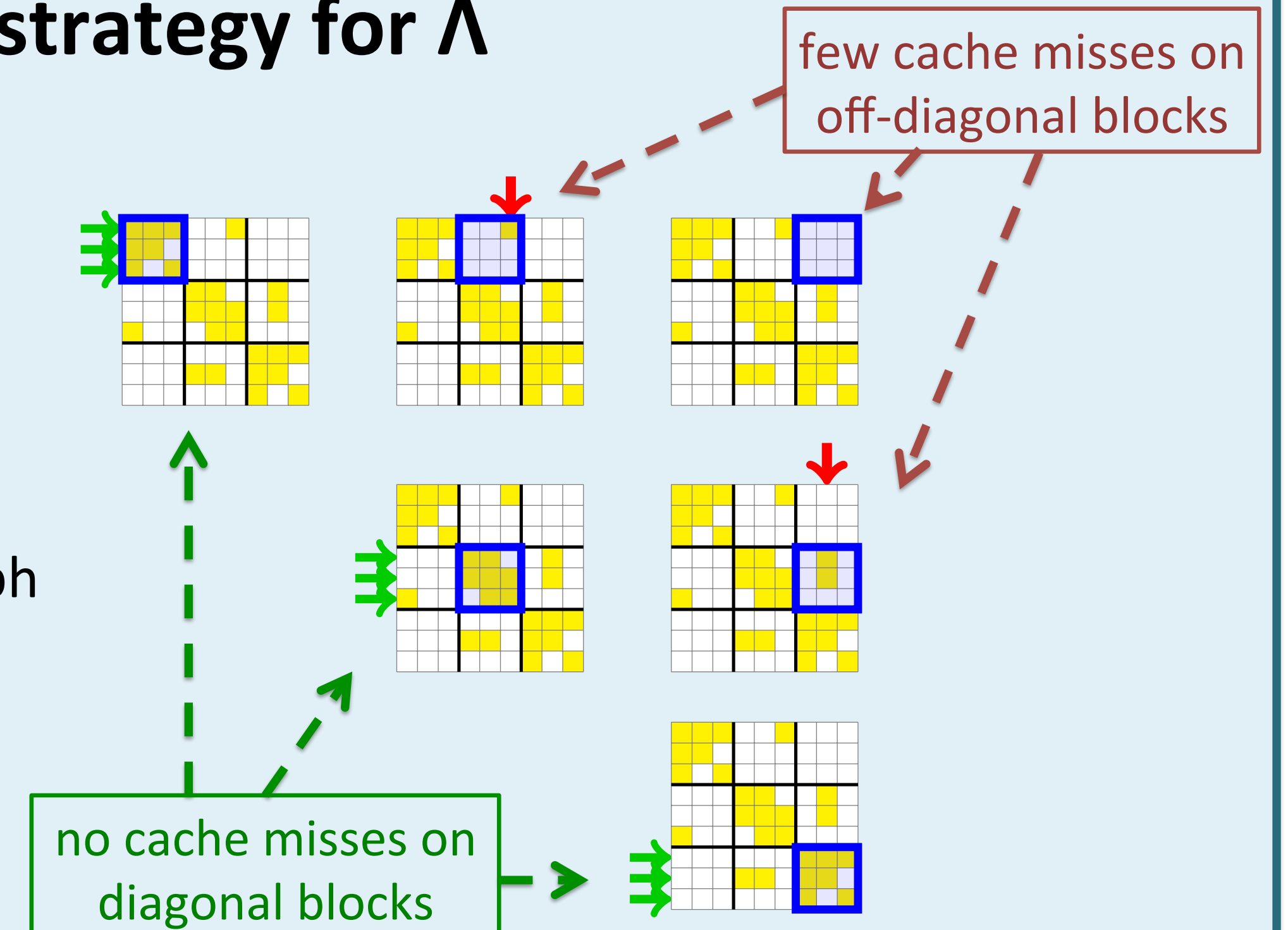
- ☑ Pre-compute all matrices: **lots of memory, fast**
- ☑ Compute as needed: **little memory, slow (many cache misses)**
- ☑ Block coordinate descent: **fast as possible given available memory (few cache misses)**

no memory restriction

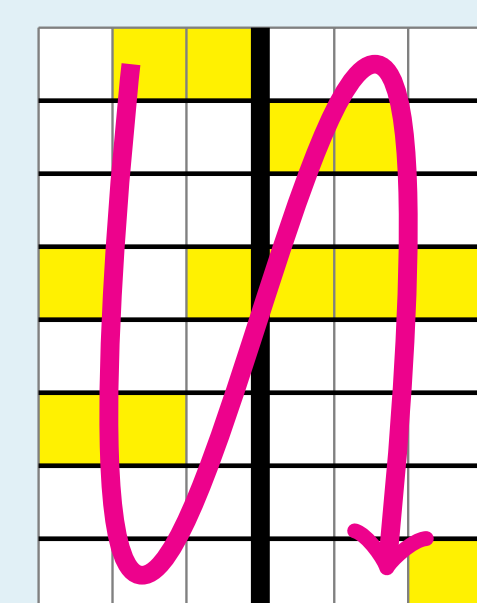
Block-wise strategy for Λ



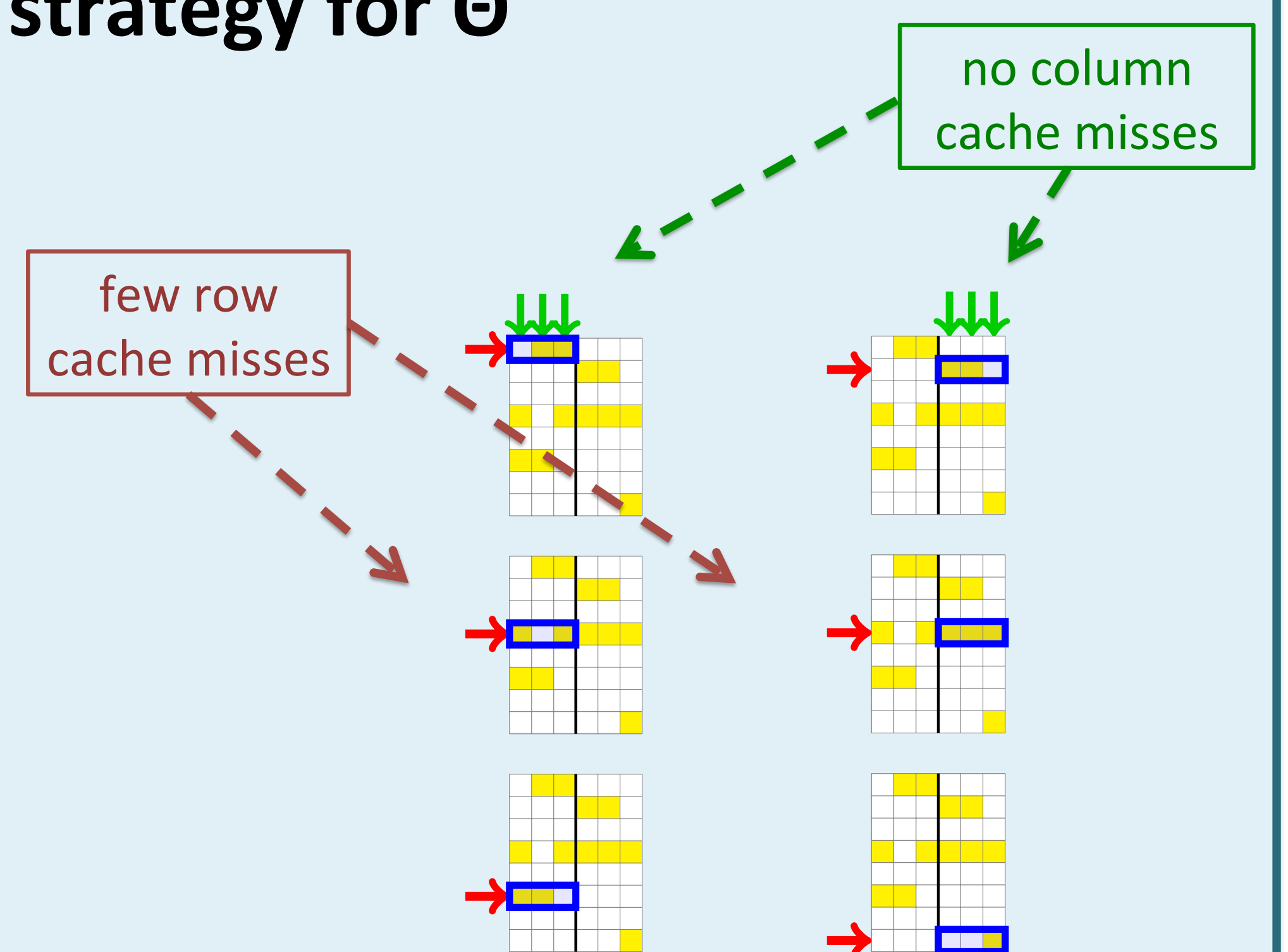
- Share computations across groups of rows
- Choose blocks via graph clustering on Λ



Block-wise strategy for Θ

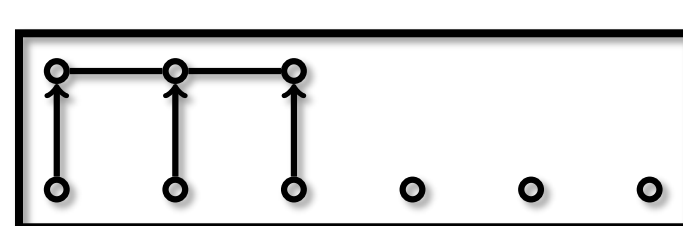


- Share computations across groups of columns
- Choose blocks via graph clustering on $\Theta^T \Theta$
- Exploit row-wise sparsity

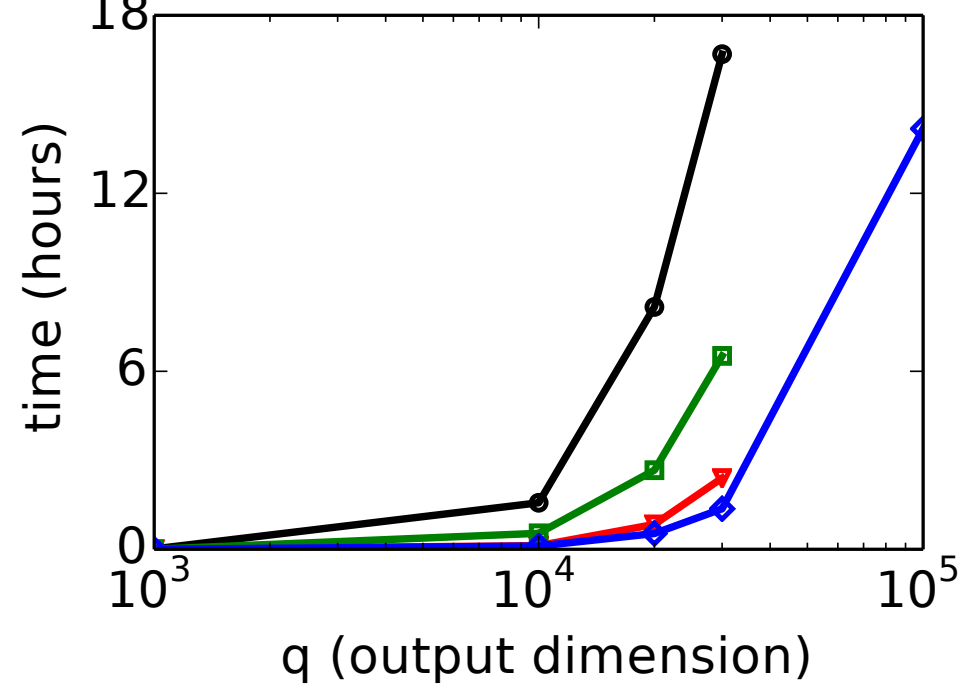


Simulation Results

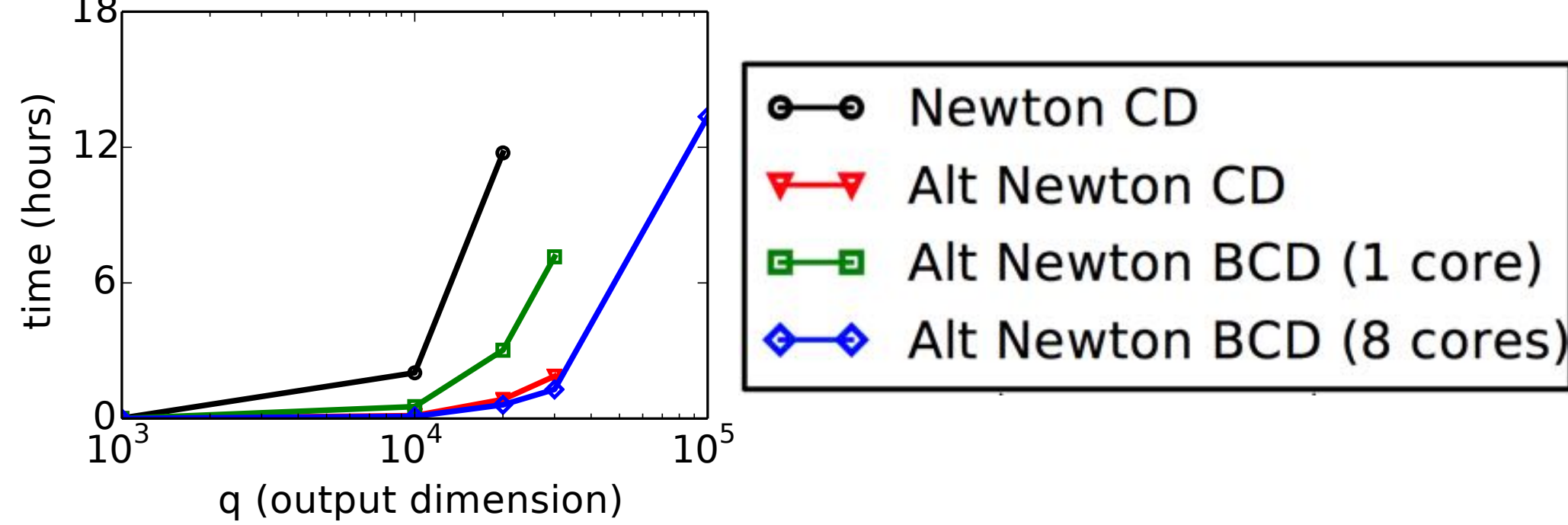
Linear chain graphs



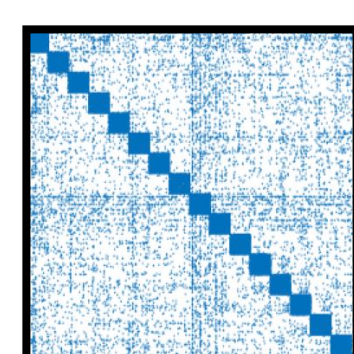
Scaling q (#outputs=q)



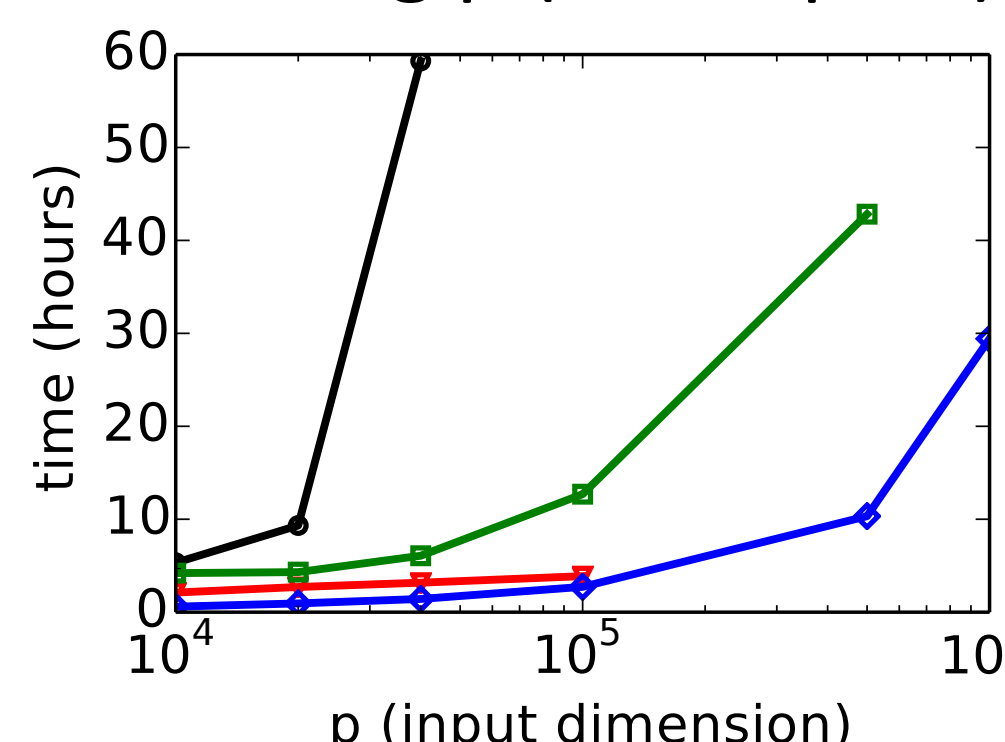
Scaling q (#outputs=2q)



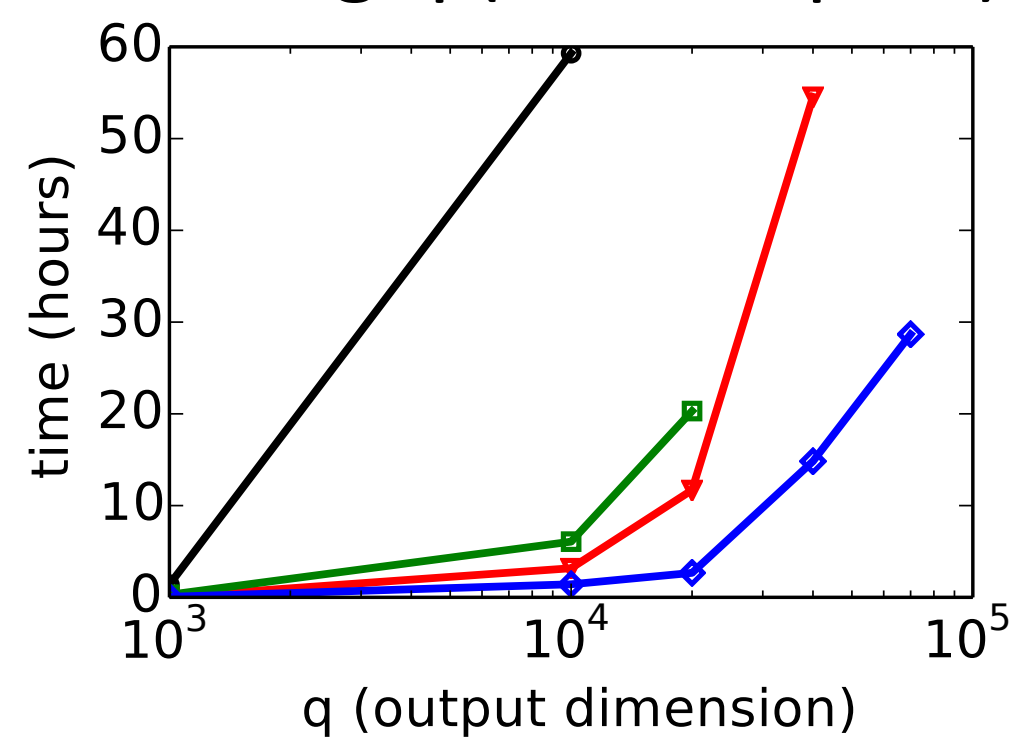
Random graphs with clustering



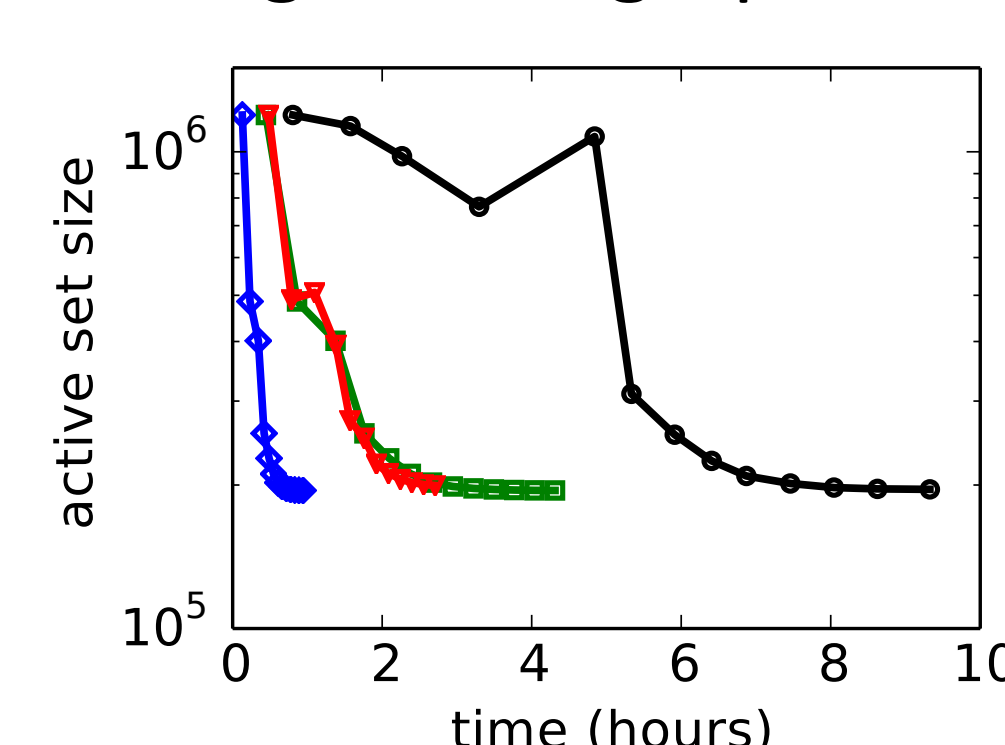
Scaling p (# of inputs)



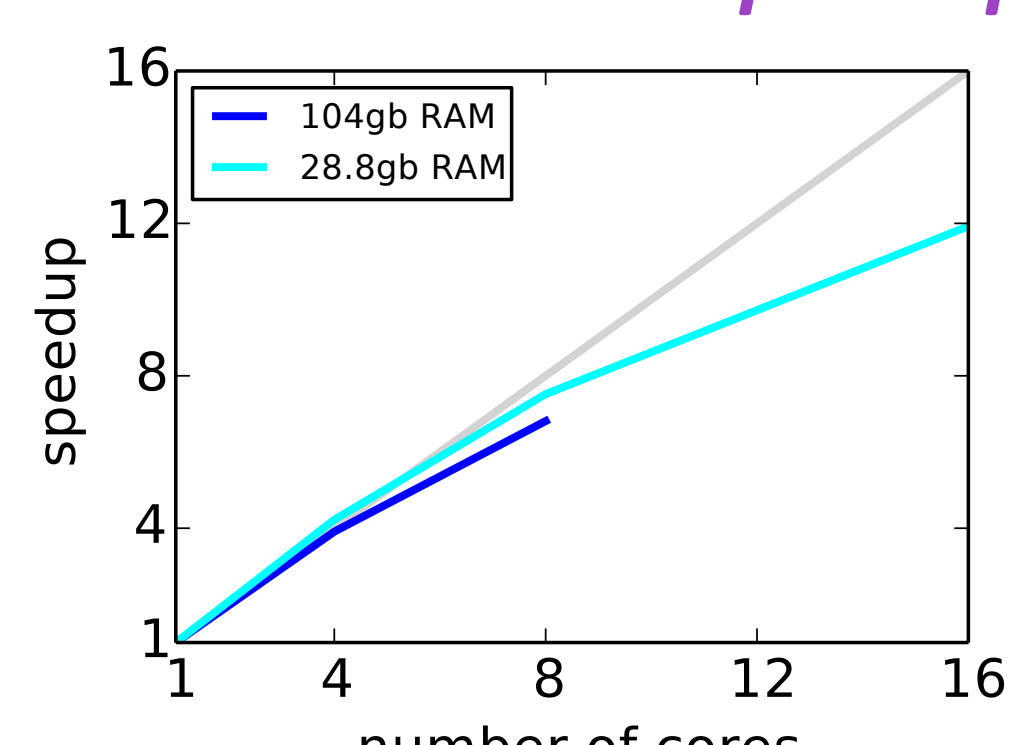
Scaling q (# of outputs)



Convergence in graph structure



Parallelization speedup

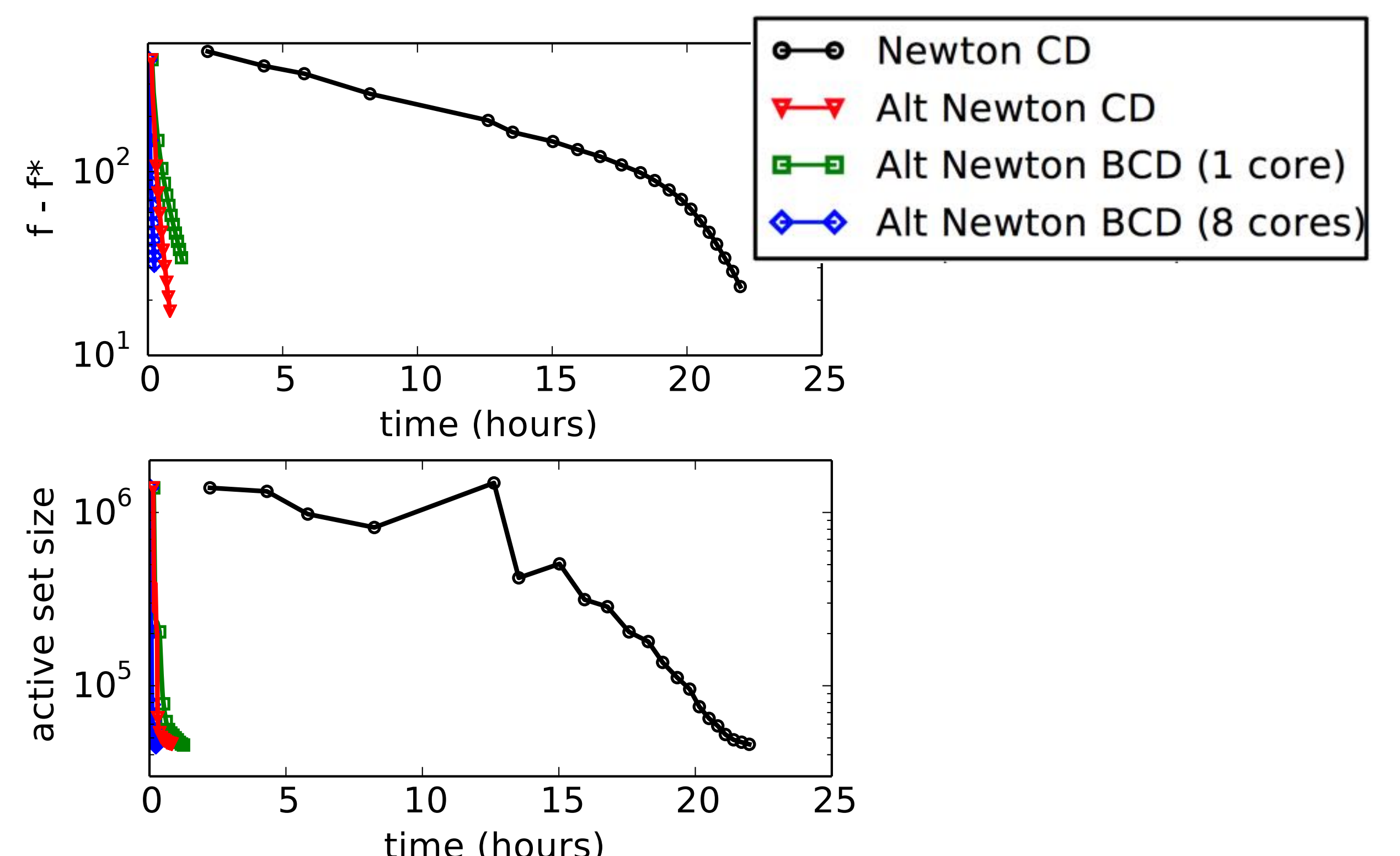


Genomic Data Analysis

- Genotypes and gene expression levels for 171 individuals with asthma
- Contains 442,440 SNPs and 10,256 genes with variance > 0.01
- Regularization parameters chosen to learn graphs with 10q edges

p	q	Newton CD	Alt Newton CD	Alt Newton BCD
34,249	3,268	22.0	0.51	0.24
34,249	10,256	> 50	2.4	2.3
442,440	3,268	*	*	11

Results for dataset with 34,249 SNPs from chromosome 1 and 3,268 genes:



References

- [1] Sohn & Kim. Joint estimation of structured sparsity and output structure in multiple-output regression via inverse-covariance regularization. AISTATS 2012.
- [2] Yuan & Zhang. Partial gaussian graphical model estimation. IEEE Transactions on Information Theory. 2014.
- [3] Wytock & Koltar. Sparse Gaussian conditional random fields: algorithms, theory, and application to energy forecasting. ICML 2013.
- [4] Hsieh et al. BIG & QUIC: Sparse inverse covariance estimation for a million variables. NIPS 2013.