

Application of the projective truncation and randomized singular value decomposition to a higher dimension.



[K.N. arXiv:2307.14191]

Katsumasa Nakayama (RIKEN)

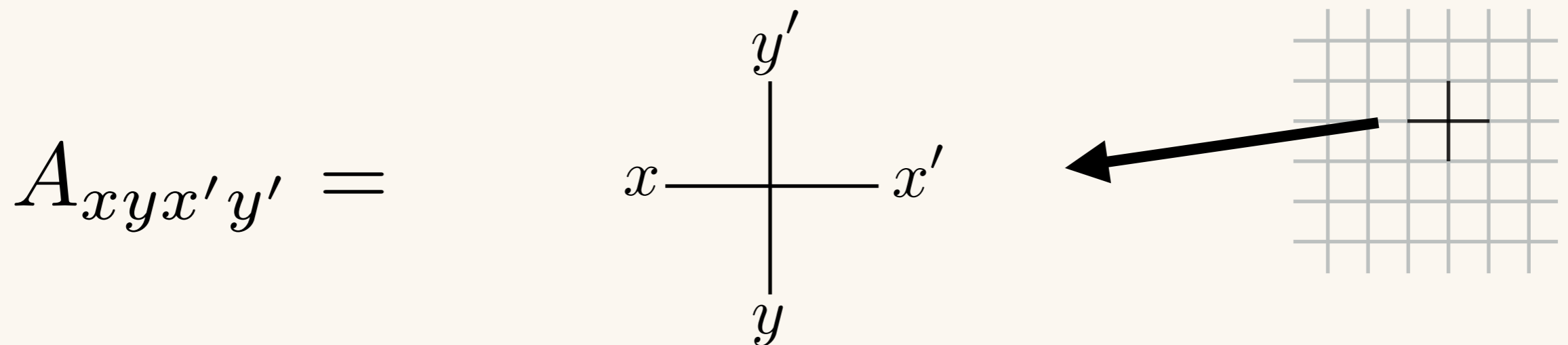
2023/07/31 @Online.

● Tensor renormalization group (TRG)

[M. Levin, C. P. Nave. arXiv:cond-mat/0611687]

- ◇ TRG calculate the physical quantity as trace of tensors.

$$Z = \text{Tr} \sum_{i \in \text{lattice}} A_{x_i y_i x'_i y'_i}$$



- Sign problem

- × High cost ($\text{dim} \geq 3$)

- Another representation

- △ Systematic error

How can we take whole contraction approximately?

→ Singular value decomposition (Frobenius norm)

● Motivation: The origin of the systematic error

◇ HOTRG [Z.Y. Xie, J. Chen, et al. arXiv:1201.1144]

Systematic error: isometry,

d : dimension

Cost: $O(D^{4d-1})$

D : truncated bond size

◇ Anisotropic TRG(ATRG)

[D. Adachi, T.Okubo, S. Todo. arXiv:1906.02007]

Systematic error: (isometry), decomposition, R-SVD

Cost: $O(D^{2d+1})$

◇ TriadTRG (TTRG) [D. Kadoh, K.N. arXiv:1912.02414]

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→ How about HOTRG with Randomized-SVD?

→ Can we reduce the systematic error from decomposition?

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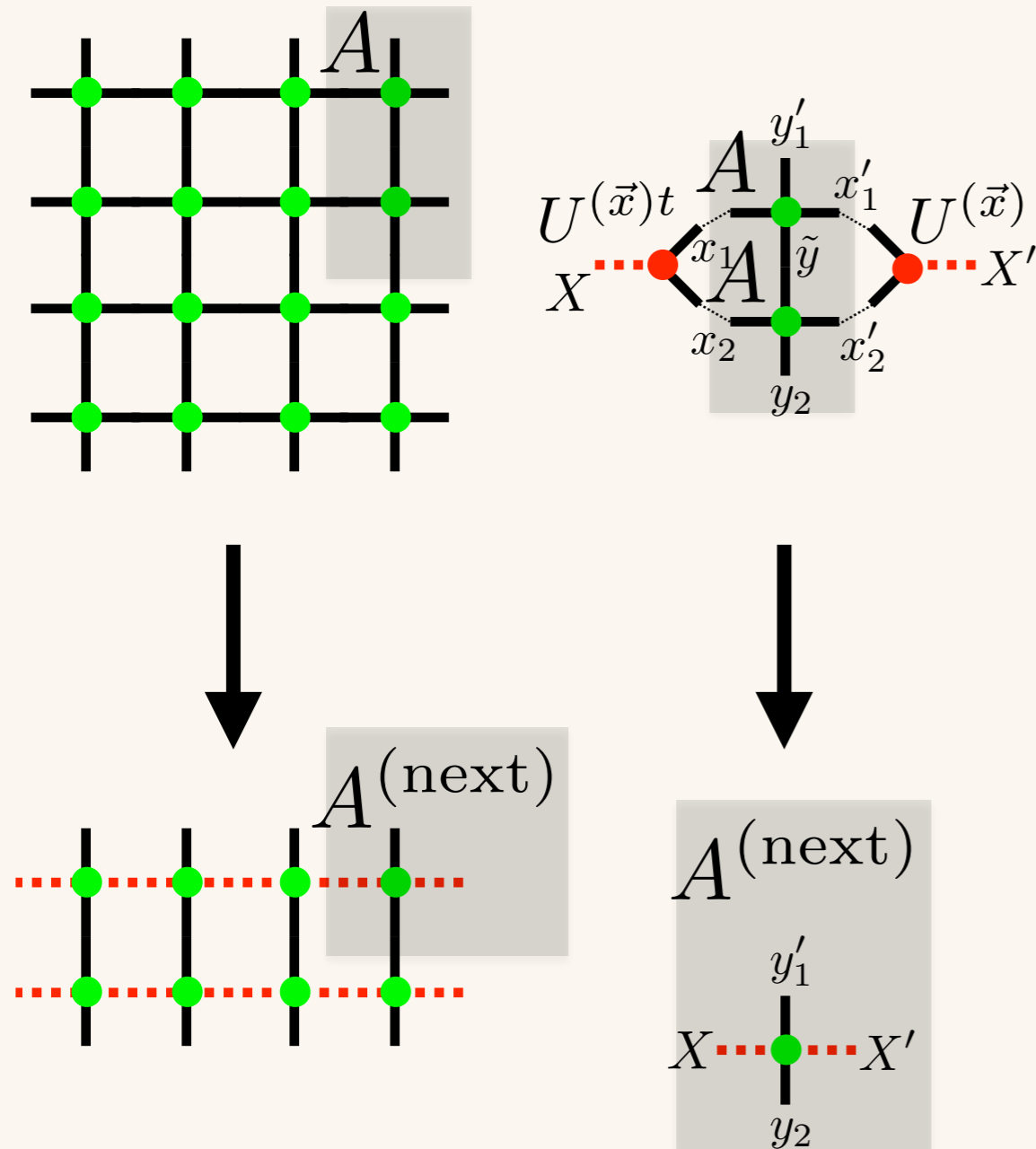
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HOTRG with randomized SVD

Higher-Order TRG (HOTRG)

[Z.Y. Xie, J. Chen, et al. arXiv:1201.1144]

◇ Contraction by projection operator U (isometry)



$$\Gamma^{(AA)} = AA \rightarrow A^{(\text{next})}$$

$\rightarrow U(\vec{x})$ is made by SVD of $\Gamma\Gamma^t$

$$[\Gamma\Gamma^t]_{[x_1 x_2][x_1^t x_2^t]} = \sum_{k=1}^{D^2} U_{[x_1 x_2]k}^{(x)} \lambda_k U_{[x_1^t x_2^t]k}^{(x)}$$

SVD Cutoff: $D^2 \rightarrow D$
 Cost: $O(D^6)$

$$U^t A A U = A^{(\text{next})}$$

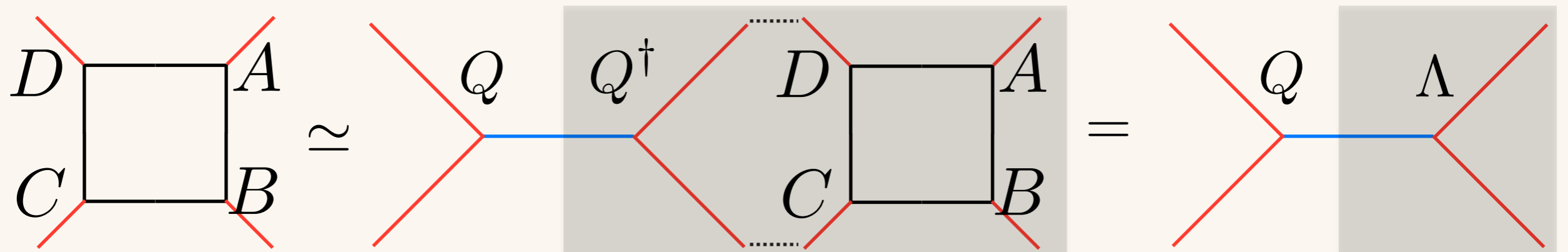
Contraction Cost: $O(D^7)$

● Randomized-SVD

[N. Halko, et al. arXiv:0909.4061]

[S. Morita, et al. arXiv:1712.01458]

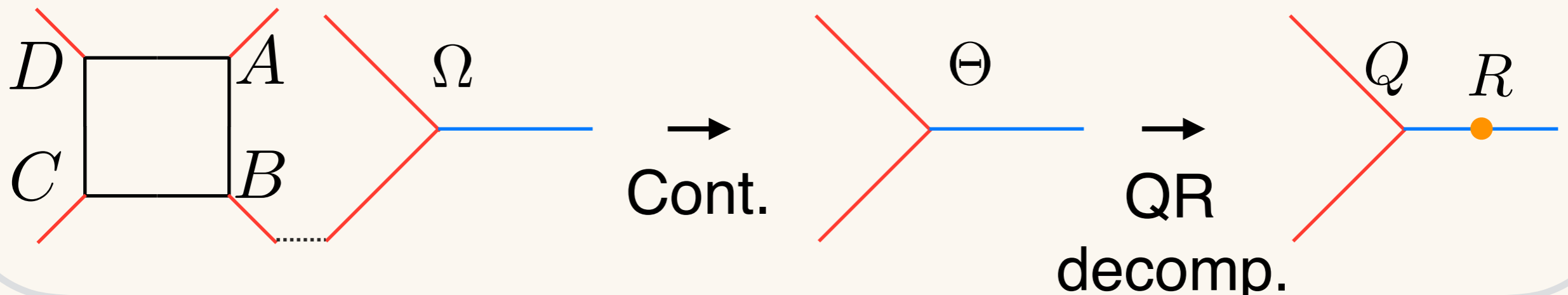
◇ Approximated contraction by orthogonal matrix Q



◇ SVD of $\Lambda \equiv Q^\dagger ABCD$.

◇ To prepare Q , we use randomized method & QR decomp.

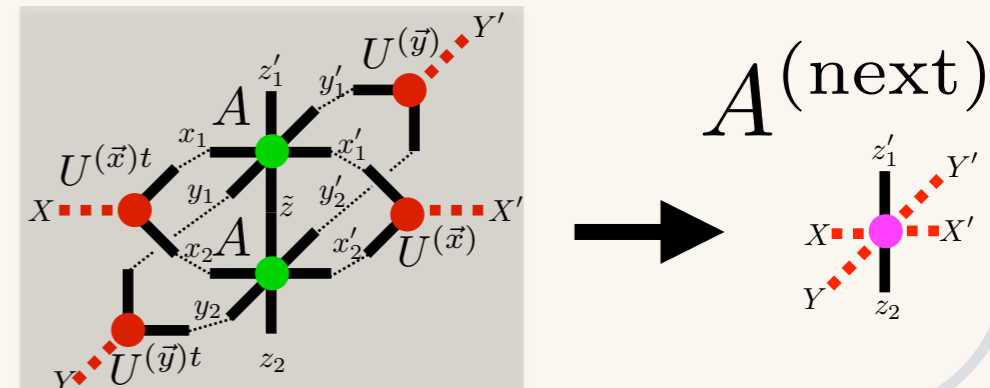
◇ Contraction with the random tensor Ω



● HOTRG [Z.Y. Xie, J. Chen, et al. arXiv:1201.1144]

◇ Contraction with U

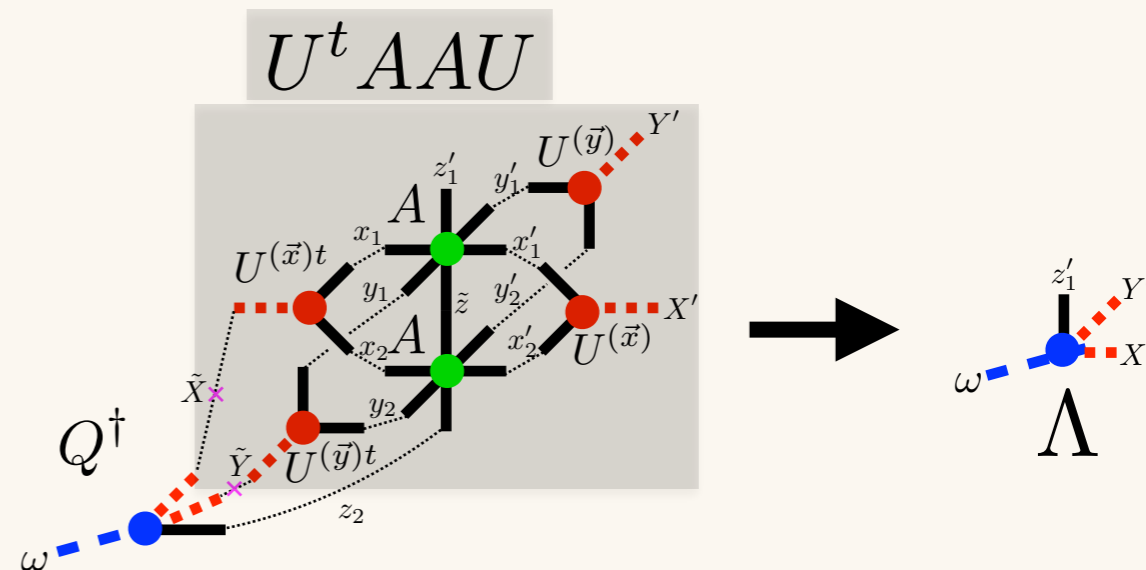
$$U(\vec{y})^t U(\vec{x})^t AAU(\vec{x})U(\vec{y}) \rightarrow A^{(\text{next})}$$



● HOTRG with R-SVD [K.N. arXiv:2307.14191]

◇ Contraction with U, Q

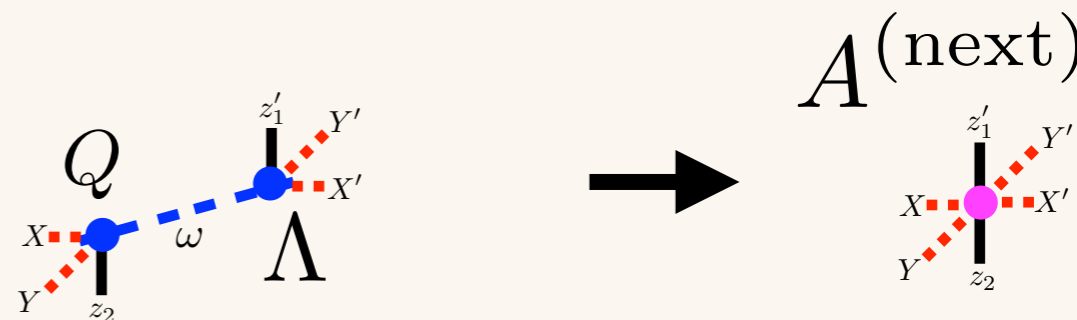
$$QQ^\dagger U(\vec{y})^t U(\vec{x})^t AAU(\vec{x})U(\vec{y}) \simeq A^{(\text{next})}$$



◇ Cost reduction

$$O(D^{4d-1}) \rightarrow O(D^{3d})$$

→ Further cost reduction?

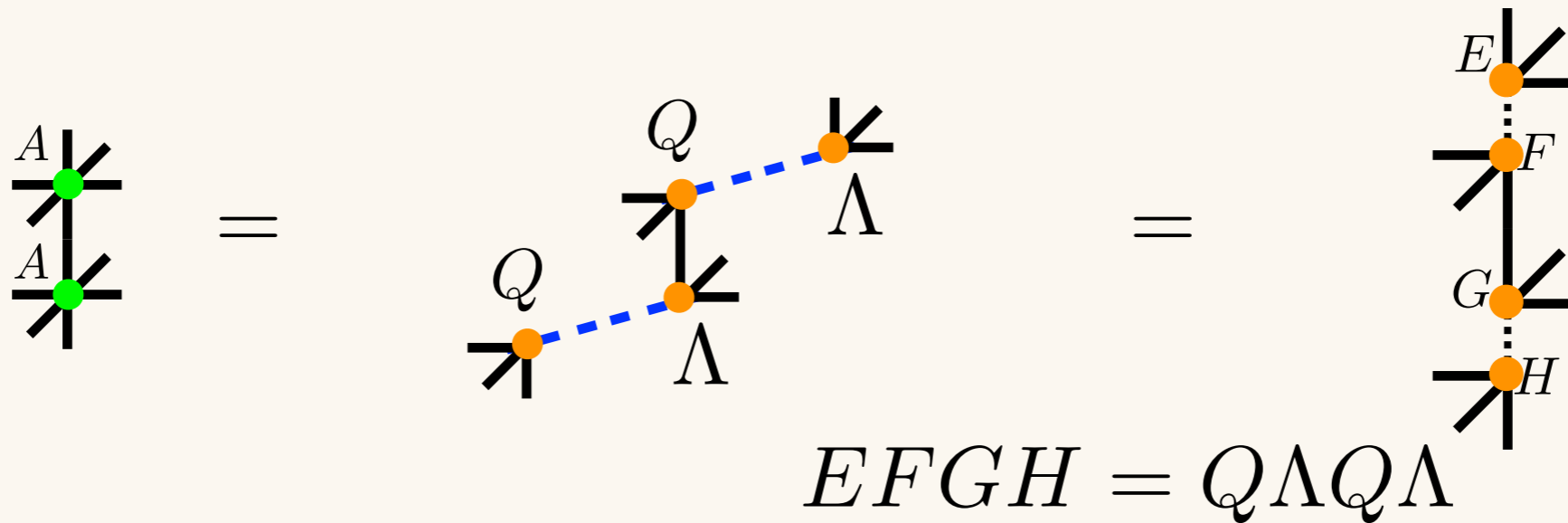


Cost and systematic error reduction

Minimally-decomposed TRG(MDTRG)

[K.N. arXiv:2307.14191]

→ We already have tensor of order $d+1$ rep. of Q and Λ .

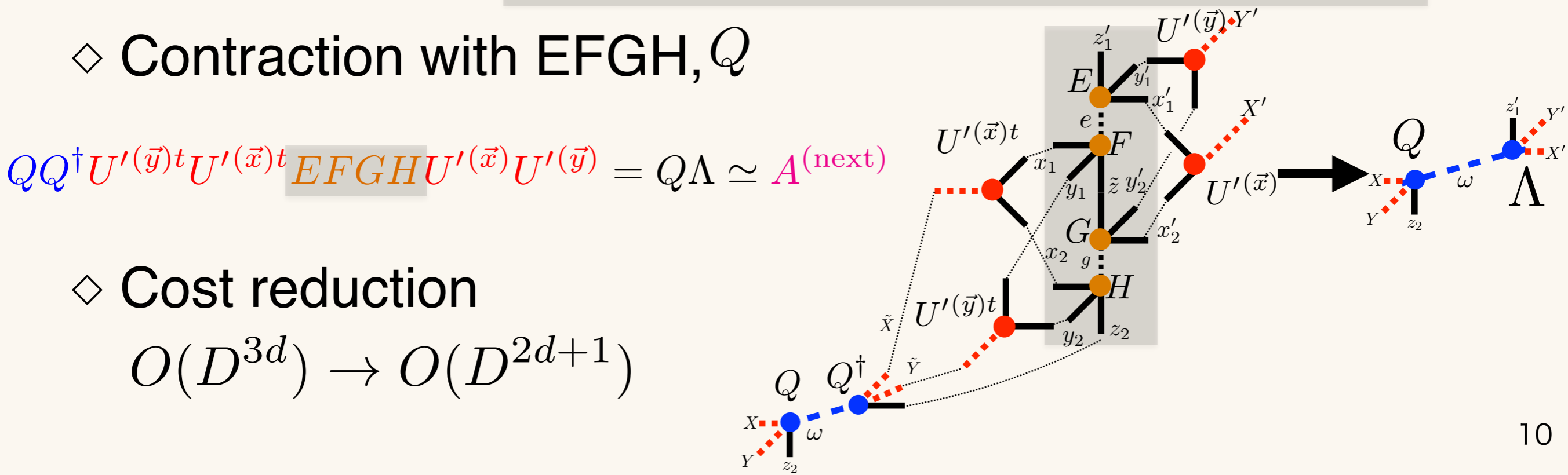


A: Order $2d$

E, F, G, H: Order $d+1$

Q, Λ : Order $d+1$

◇ Contraction with $EFGH, Q$

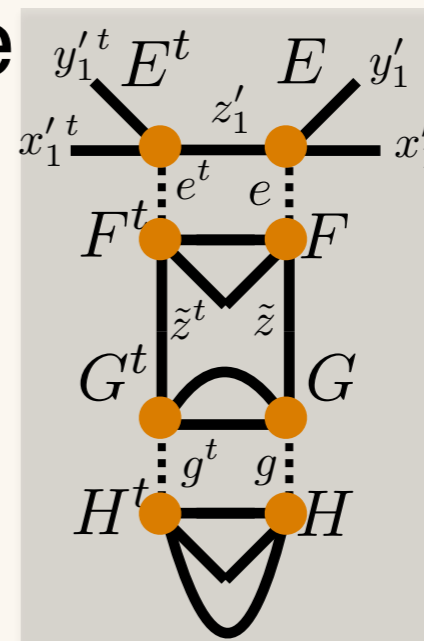
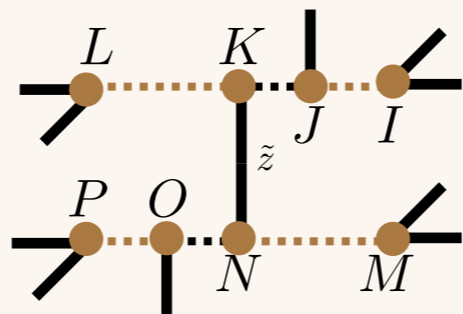
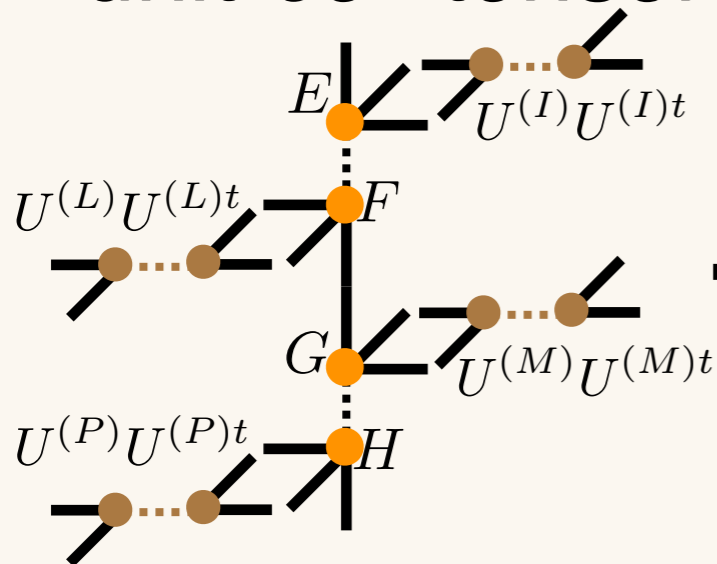


◇ Cost reduction

$$O(D^{3d}) \rightarrow O(D^{2d+1})$$

● Minimally-decomposed TRG on triad rep.

- ◇ We introduce triad rep. by SVD of the unit-cell tensor $\Gamma^{(EFGH)} = EFGH$.



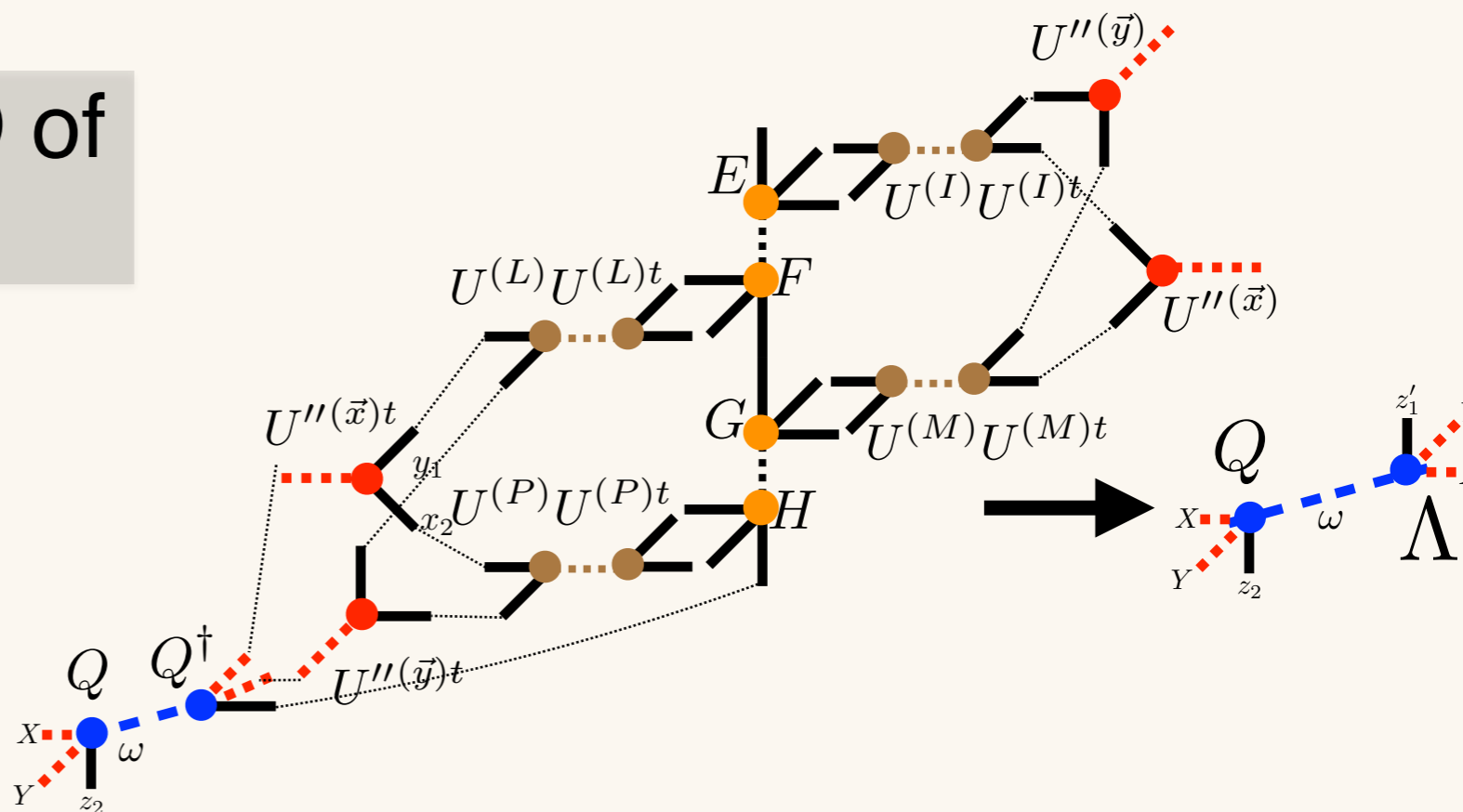
- ◇ We do NOT use SVD of E, F, G and H.

$$E = U^{(0)} s^{(0)} V^{(0)}$$

- ◇ We use SVD of $\Gamma\Gamma^t$.

- ◇ Cost reduction

$$O(D^{2d+1}) \rightarrow O(D^{d+3})$$



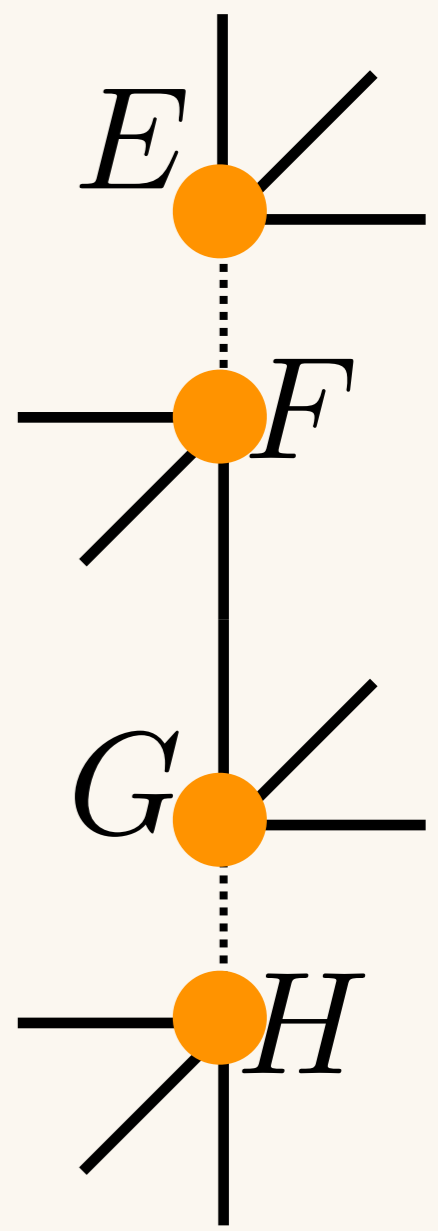
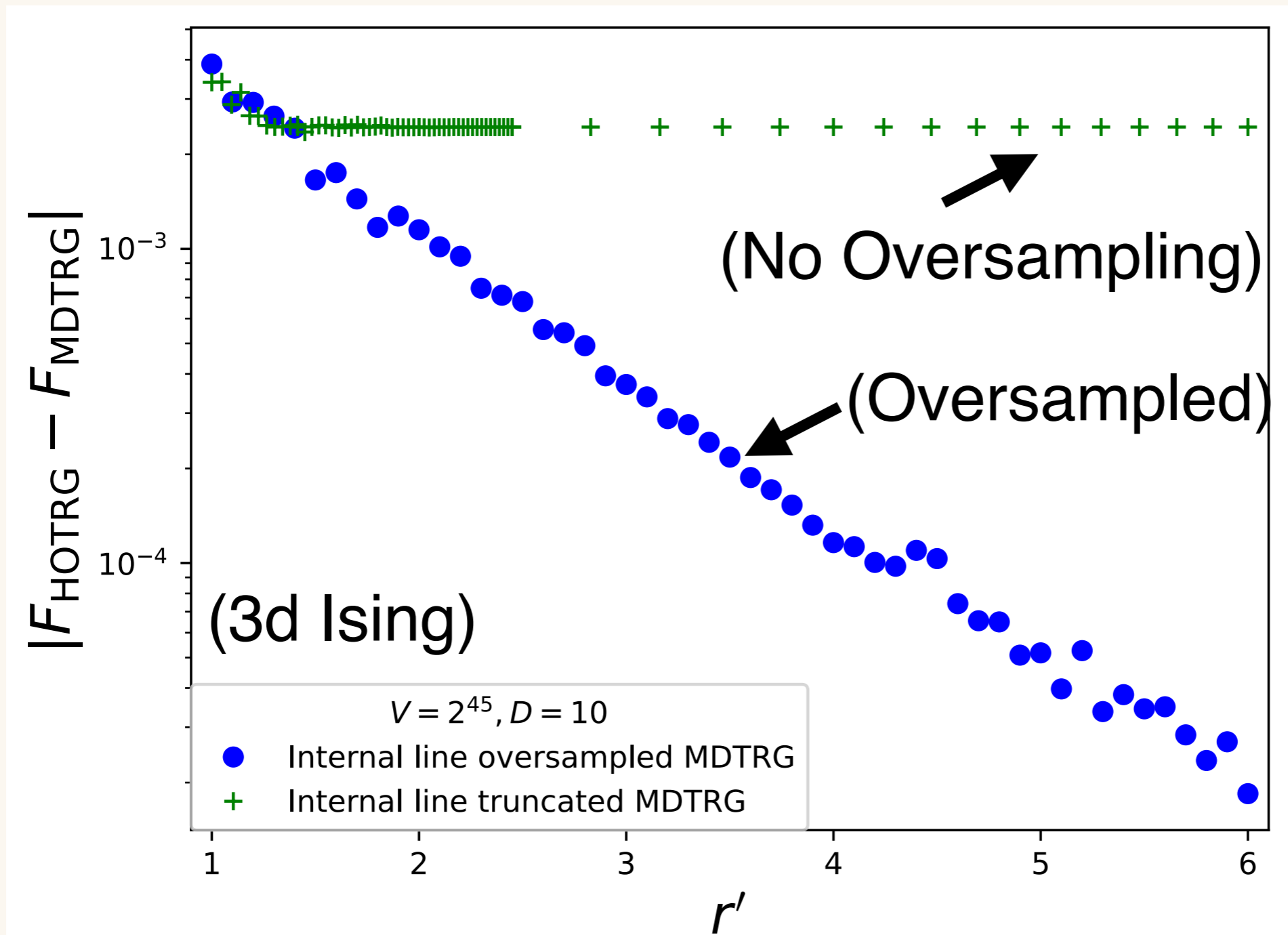
$$QQ^\dagger U''(\vec{y})^t U''(\vec{x})^t U^{(LP)} U^{(LP)t} EFGHU^{(IM)} U^{(IM)t} U''(\vec{x}) U''(\vec{y}) = Q\Lambda \simeq A^{(\text{next})} \quad 11$$

● HOTRG with R-SVD

	with R-SVD	w/o R-SVD	unit-cell order
◇ HOTRG	$O(D^{3d})$	$O(D^{4d-1})$	≠ $2d$
◇ ATRG	$O(D^{2d+1})$	$O(D^{3d})$	≠ $2d$
◇ MDTRG	$O(D^{2d+1})$	$O(D^{3d})$	≠ $d + 1$
◇ TTRG	$O(D^{d+3})$	$O(D^{d+4})$	≠ 3
◇ Triad-MDTRG	$O(D^{d+3})$	$O(D^{3d})$	≠ $d + 1$

→ Comparable cost. We will check the systematic error.

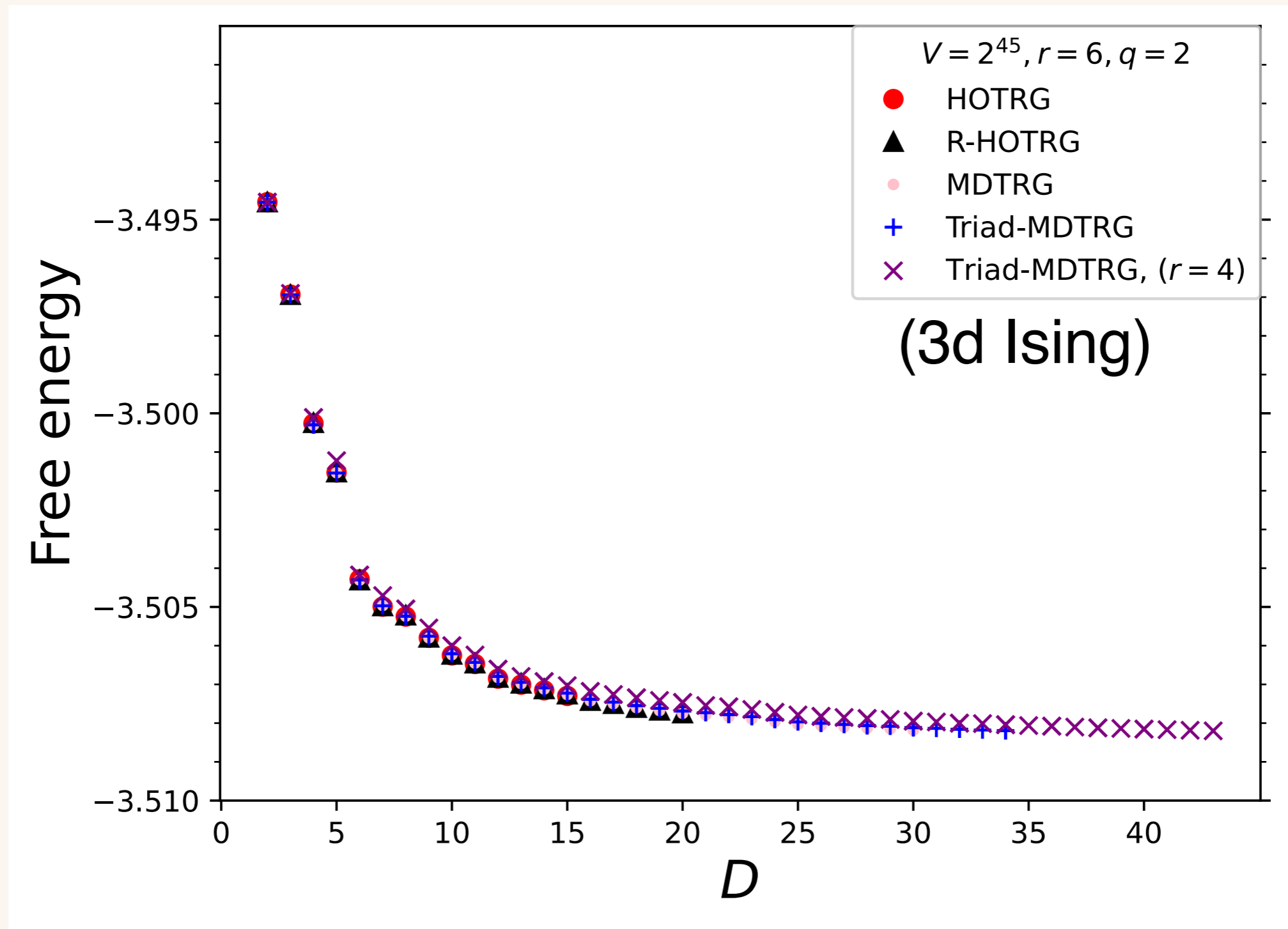
Internal line oversampling



($r = \text{Const.}$)

→ Internal lines (dotted lines) have to be oversampled ($D \rightarrow rD$) to reduce the systematic error from R-SVD.

Free energy density of 3d-Ising model

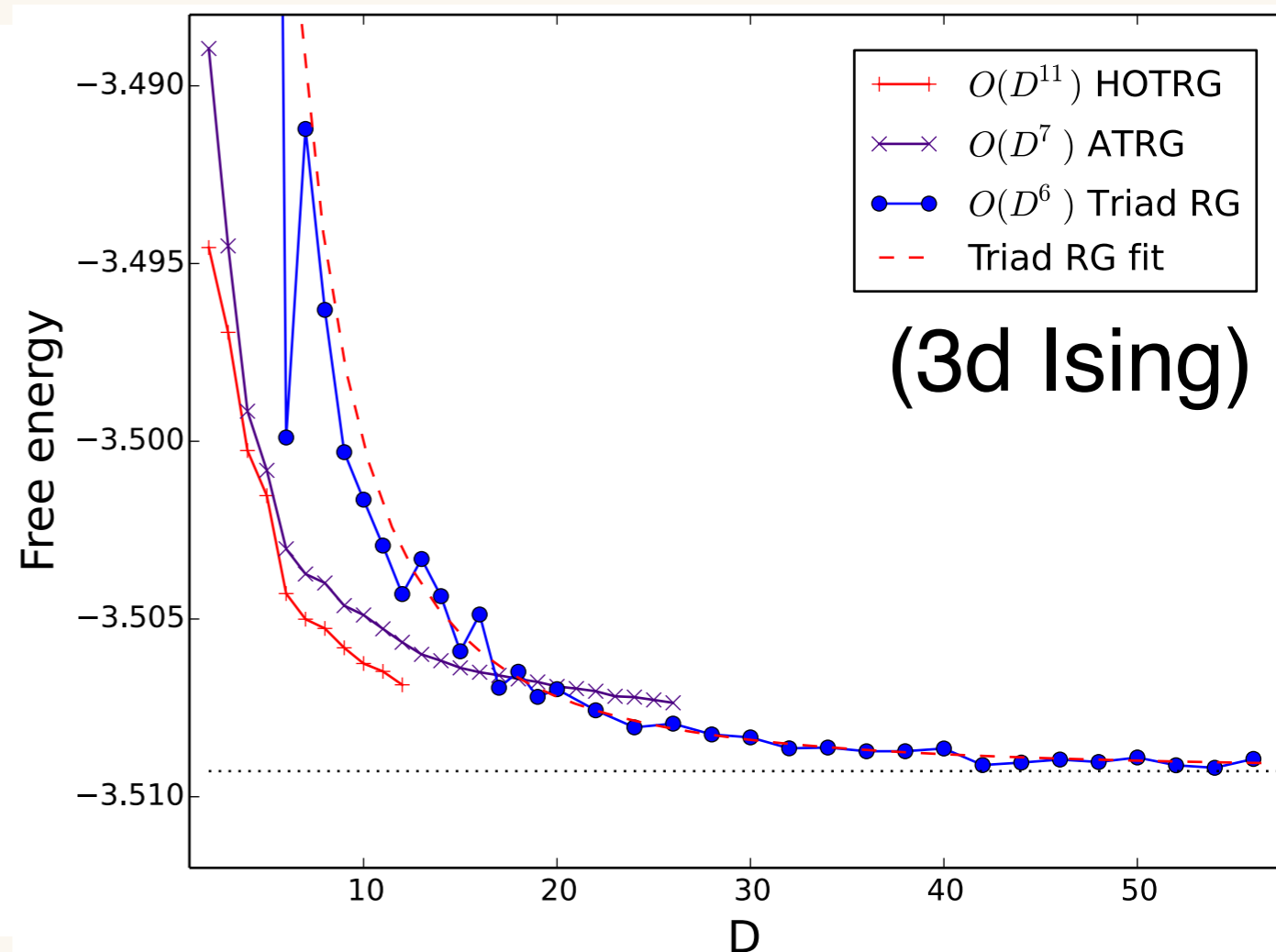
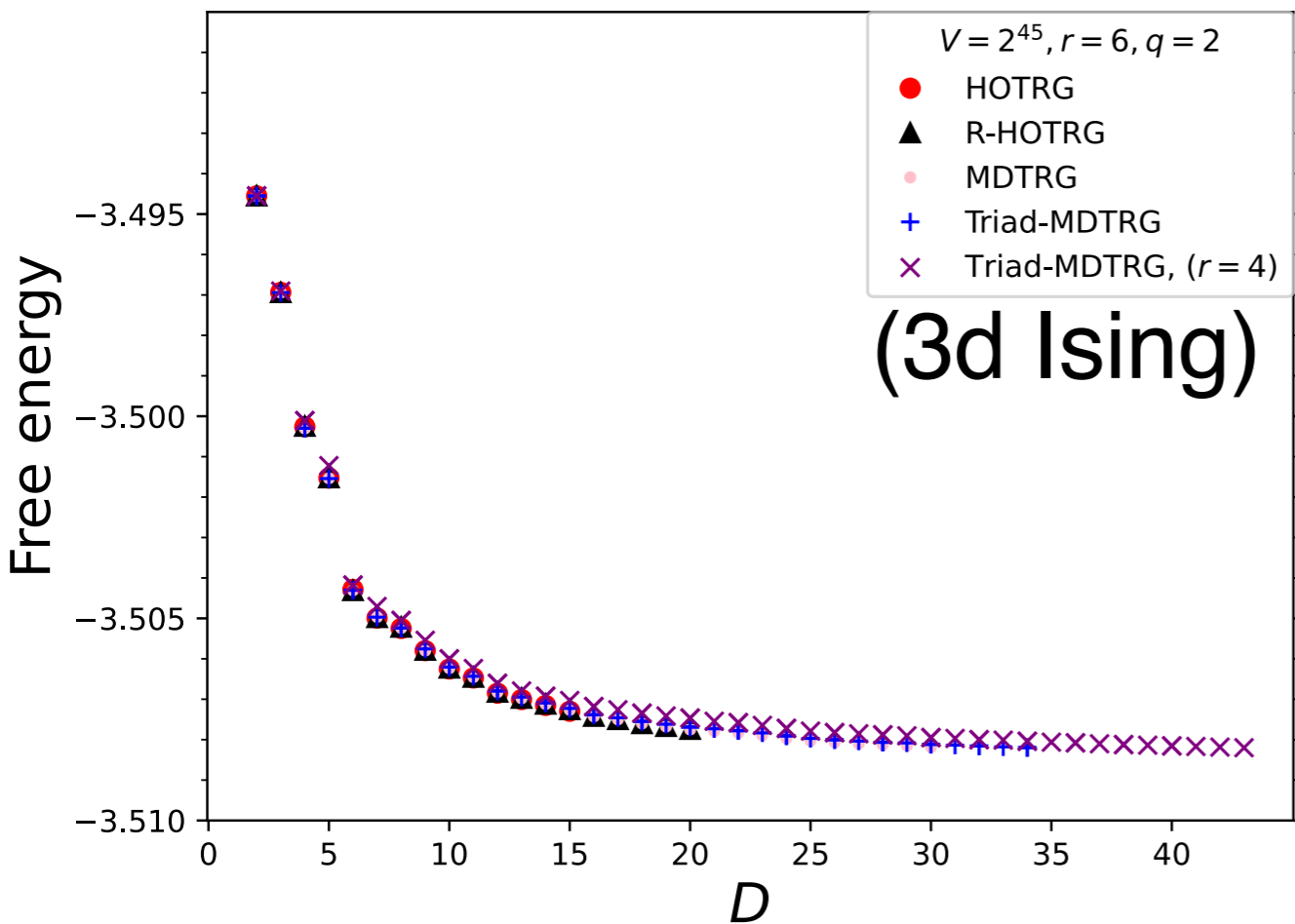


→ R-HOTRG, MDTRG, Triad-MDTRG results are consistent with HOTRG (additional systematic error is not dominant).

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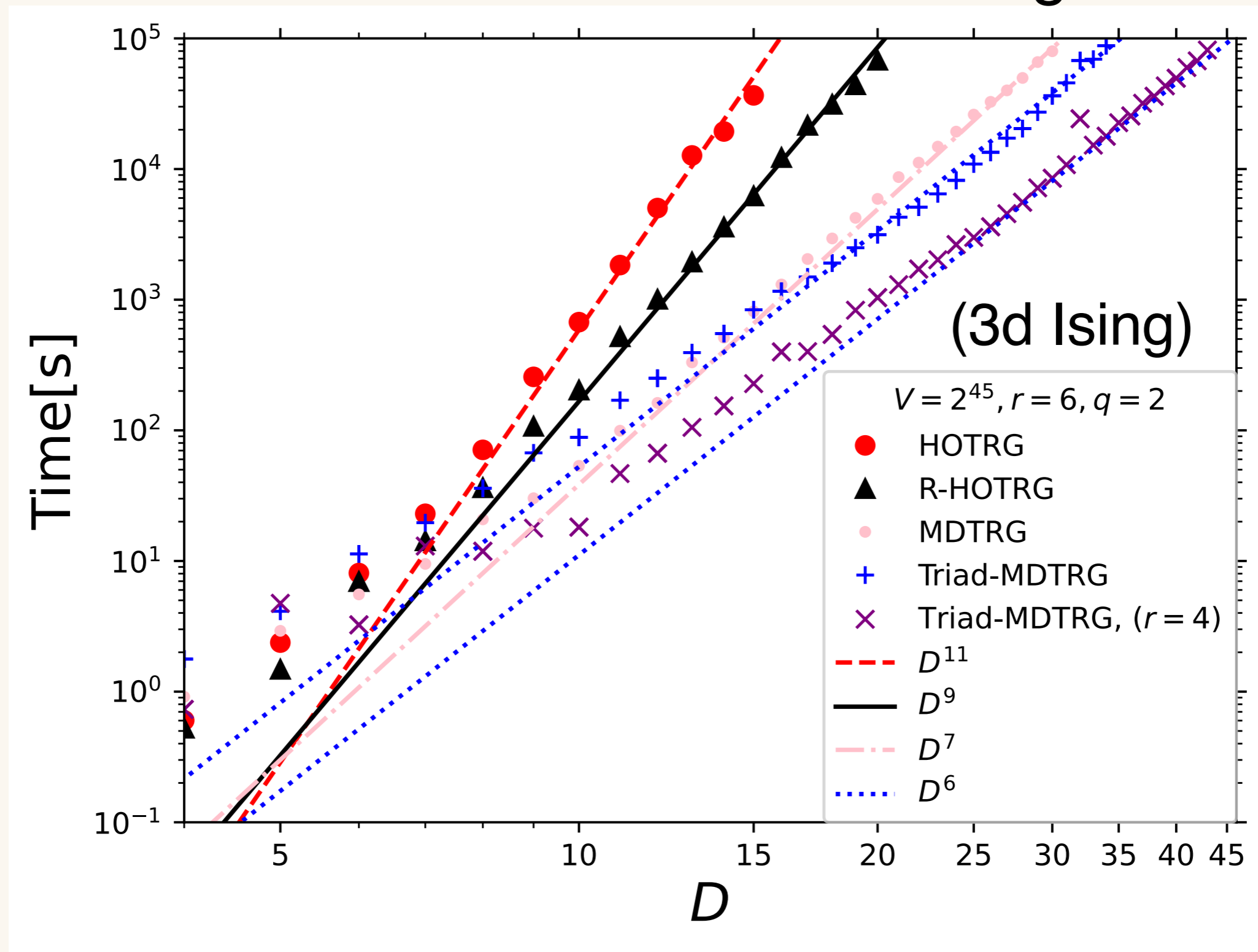
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● Truncated bond dimension D scalings



→ The scalings of D are confirmed.

● Summary

◇ How about HOTRG with Randomized-SVD?

	with R-SVD	w/o R-SVD	unit-cell order
◇ HOTRG	$O(D^{3d})$	$O(D^{4d-1})$	$\nrightarrow 2d$
◇ MDTRG	$O(D^{2d+1})$	$O(D^{3d})$	$\nrightarrow d+1$
◇ Triad-MDTRG	$O(D^{d+3})$	$O(D^{3d})$	$\nrightarrow d+1$

◇ Can we reduce the systematic error from decomposition?

→ R-HOTRG, MDTRG, and Triad-MDTRG produce consistent result with the HOTRG. The dominant systematic error is the truncation of the isometry.

Key ideas:

internal line oversampling, Isometry for unit-cell tensor.