

1. Original formulation of DMRG

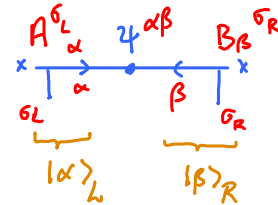
[White1992], [White1993], [Schollwöck2011, Sec 2.2]

Goal: finding ground state of infinite chain

Infinite-size DMRG (iDMRG)

Diagonalize small system (e.g. 2 sites), write ground state in the form

$$|\psi\rangle = \sum_{\alpha\beta} |\beta\rangle_R |\alpha\rangle_L \psi^{\alpha\beta}$$



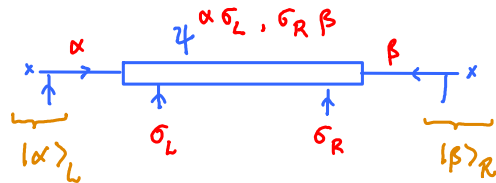
(1)

'Block L' describes left part of system, with basis  $\{|\alpha\rangle_L\}$

'Block R' describes right part of system, with basis  $\{|\beta\rangle_R\}$

Now add two sites between blocks L and R, and seek new ground state of  $H_{L\cdots R}$  of the form

$$|\psi\rangle = |\beta\rangle_R |\sigma_R\rangle |\sigma_L\rangle |\alpha\rangle_L \psi^{\alpha\sigma_L, \sigma_R\beta}$$



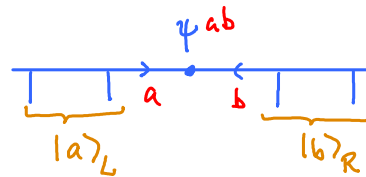
(2)

by minimizing (Lanzcos) 
$$\frac{\langle\psi|\hat{H}_{L\cdots R}|\psi\rangle}{\langle\psi|\psi\rangle}$$

Bond dimension has grown from  $D\times D$  for  $\psi^{\alpha\beta}$  to  $Dd\times Dd$  for  $\psi^{\alpha\sigma_L, \sigma_R\beta}$ , so truncation is needed. Split enlarged system in the middle, and call left side (new) block L, right side (new) block R.

Write ground state in the form

$$|\psi\rangle = |b\rangle_R |a\rangle_L \psi^{a,b}$$



(3)

with composite indices  $a = (\alpha, \sigma_L), b = (\beta, \sigma_R)$

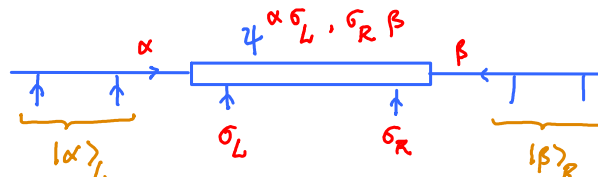
of dimension  $D_a = D_\alpha d$ . White's truncation prescription: compute reduced DM of  $L$ ,

$$P_{L_0} = \text{Tr}_R |\psi\rangle\langle\psi| = |a'\rangle_L \underbrace{\psi^{a'b'} \bar{\psi}_{b'a}}_{(P_{L_0})^{a'a}} \langle a| = \sum_c |\tilde{c}\rangle_L \rho_c \langle\tilde{c}| \quad \text{diagonalize} \quad (4)$$

Construct truncated basis for block L, using the  $D$  eigenvectors  $|\tilde{c}\rangle_L$  with the largest eigenvalues  $\rho_c$ . Rename:  $|\alpha\rangle_L^{\text{new}} := |\tilde{c}\rangle_L$ . here truncation happens

Ditto for block R.

Then iterate, : add two more sites, etc.



(5)

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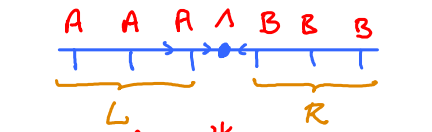
Remark: we established early on (see MPS-II.2) that the eigenvalues  $\rho_c$  of reduced density matrix of L, are obtained by SVD of  $\psi^{ab}$

$$a \text{---} \psi \text{---} b = a \text{---} U \text{---} S \text{---} V^T \text{---} b \Rightarrow \rho_c = (S^c_c)^2$$

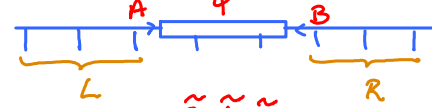
So retaining the  $\rho_L$ -eigenstates with largest  $\rho_c$  is equivalent to just doing SVD-truncation on  $\psi^{ab}$ .

Modern formulation

Start with MPS in bond-canonical form:

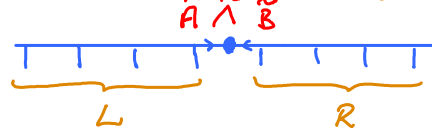


Add two central sites and find ground state (Lanczos):



Do SVD to split chain into two larger blocks, and truncate:

$$\psi = U S V^T = \tilde{A} \tilde{\Lambda} \tilde{B}$$

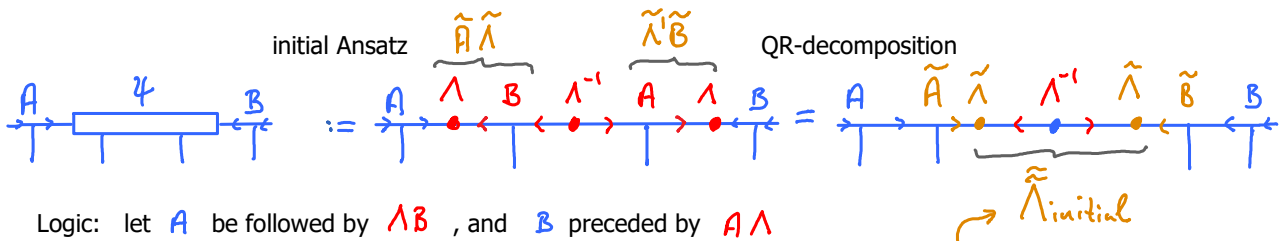


Iterate: make chain longer and longer, until ground state energy per site converges.

'iDMRG state prediction'

[McCulloch2008], [Schollwöck2011, Sec. 10.1]

To speed up Lanczos search for ground state, construct initial guess for  $\psi$  from previous data:



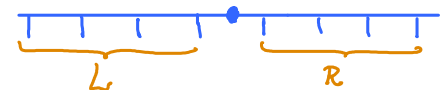
Logic: let A be followed by  $\Lambda B$ , and B preceded by  $A \Lambda$  to reverse arrows between B and A, use  $\Lambda^{-1}$

use this to initialize Lanczos

This leads to 'dramatic speedup' of iDMRG.

Finite-size DMRG

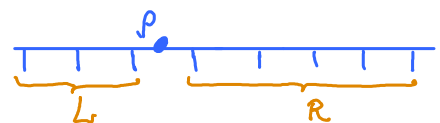
Grow chain to some length N using infinite-size DMRG algorithm.



Then reduce L, enlarge R, optimize  $\psi$ :



Diagonalize  $\rho_L$ , truncate.



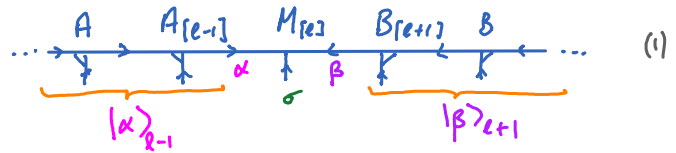
Iterate: sweep back and forth until convergence.

This is conceptually identical to variational optimization with two-site update.

Single-site DMRG is also possible  $\leftrightarrow$  variational single-site update.

Problem: when exploiting quantum numbers, 1-site DMRG performs poorly, because it does not explore subspaces with different quantum numbers. An early remedy for this is 2-site DMRG, but that is computationally much more expensive than 1-site DMRG. Subsequent suggestions for 1-site DMRG with symmetries are 'density matrix perturbation' [White2005], the 'center matrix wave function formalism' [McCulloch2007], and, most recently, 'subspace expansion' [Hubig2015], which performs best.

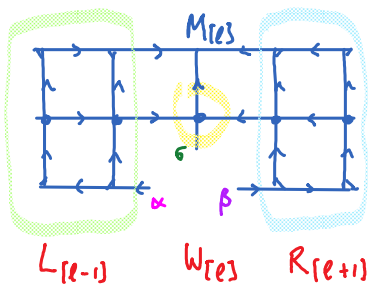
Reminder of 1-site DMRG, in site-canonical representation:



Local basis:  $|\alpha, \sigma, \beta\rangle := |\beta\rangle_{e+1} |\sigma\rangle |\alpha\rangle_{e-1}$

Minimize energy with constraint of fixed normalization, 1 site at a time:

$$\frac{\partial}{\partial B_{e1}^\dagger} \left[ \langle \psi | \hat{H} | \psi \rangle - \lambda \langle \psi | \psi \rangle \right] = 0 \quad (2)$$



$$- \lambda \left[ \text{Diagram of } M[e] \text{ tensor} \right] = \alpha \rightarrow \left[ \text{Diagram of } M[e] \text{ tensor} \right] \beta \quad (3)$$

close zipper

$$H_{[e]}^{a'} M_{[e]}^a = \lambda M_{[e]}^{a'} \quad a = (\alpha, \sigma, \beta)$$

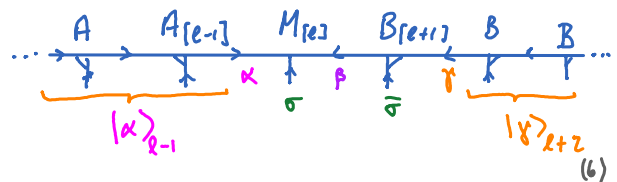
with  $H_{[e]} = L_{[e-1]} W_{[e]} R_{[e+1]} = \left[ \text{Diagram of } H_{[e]} \text{ tensor} \right] \quad (4)$

Solve for 'eigenvector' with lowest eigenvalue, say  $\tilde{M}_{[e]}$ , then do SVD on it to move to next site:

$$D \left[ \text{Diagram of } \tilde{M}_{[e]} \text{ tensor} \right] B_{[e+1]} \stackrel{\text{SVD}}{=} D \left[ \text{Diagram of } (U) (S) (V^\dagger) B_{[e+1]} \right] = D \left[ \text{Diagram of } \tilde{A}_{[e]} M_{[e+1]} \right] \quad (5)$$

Important: dimensions of  $\tilde{M}_{[e]}$  are fixed, hence truncation is neither needed nor possible!

Reminder of 2-site DMRG, in site-canonical representation:



Local basis:  $|\alpha, \sigma, \bar{\sigma}, \gamma\rangle := |\gamma\rangle_{e+2} |\bar{\sigma}\rangle |\sigma\rangle |\alpha\rangle_{e-1}$

Minimize energy two sites at a time:

$$H_{[e, e+1]}^{a'} \left( M_{[e]} B_{[e+1]} \right)^a = \lambda \left( M_{[e]} B_{[e+1]} \right)^{a'} \quad a = (\alpha, \sigma, \bar{\sigma}, \gamma)$$

$$\left[ \text{Diagram of } M_{[e]} B_{[e+1]} \text{ tensor} \right] = \alpha \rightarrow \left[ \text{Diagram of } M_{[e]} B_{[e+1]} \text{ tensor} \right] \gamma \quad (7)$$

Solve for 'eigenvector' with lowest eigenvalue,  $(\tilde{M}B)$ , then do SVD and truncate (!) to move to next site:

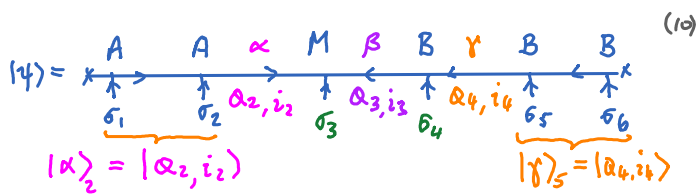
$$\alpha, \sigma \left[ \text{Diagram of } \tilde{M}B \text{ tensor} \right] \bar{\sigma}, \beta \stackrel{\text{SVD}}{=} \alpha, \sigma \left[ \text{Diagram of } (U) (S) (V^\dagger) \right] \bar{\sigma}, \beta \stackrel{\text{truncate}}{\approx} \alpha, \sigma \left[ \text{Diagram of } (\tilde{S}) (V^\dagger) \right] \bar{\sigma}, \beta := \alpha \left[ \text{Diagram of } \tilde{A}_{[e]} M_{[e+1]} \right] \beta \quad (8)$$

Problem of single-site optimization: it is constrained to a variational space defined by outgoing state spaces  $|\beta\rangle_{L+1} |\alpha\rangle_{L-1}$ . If the ranges of quantum numbers  $Q_\alpha$  and  $Q_\beta$  for these spaces are too small to accurately represent the ground state, single-site DMRG has no way to enlarge them.

Two-site optimization does not have this problem: the action of H on two sites enlarges bond dimension in between, adding the full range of quantum numbers needed on that bond. If a certain quantum number was missing on that bond before the action of H, but appears afterwards with non-negligible weight, it will survive after SVD and truncation. Hence: two-site optimization can add missing quantum numbers, if needed.

But this comes at a cost: effective two-site Hamiltonian has dimension  $D^2 d^2 \times D^2 d^2$ .  
By contrast, effective one-site Hamiltonian has dimension  $D^2 d \times D^2 d$ . (9)

Example: 6-site chain of S=1/2 spins, exploiting Sz quantum numbers:  $\sigma_e \in \{+1, -1\}$



spin conservation:  $-(Q_2 + \sigma_3) = Q_3 = \sigma_4 + Q_4$  (11)

$$\begin{aligned}
 |\psi\rangle &= \sum_{\{\sigma\}} |\bar{\sigma}\rangle (A^{\sigma_1} A^{\sigma_2})^\alpha M_{[3]}^{\alpha \sigma_3 \beta} B_{[4]}^{\sigma_4 \gamma} (B^{\sigma_5} B^{\sigma_6})^\gamma \\
 &= |\gamma\rangle_5 |\sigma_4\rangle |\sigma_3\rangle |\alpha\rangle_2 M_{[3]}^{\alpha \sigma_3 \beta} B_{[4]}^{\sigma_4 \gamma} \\
 &= |\alpha, i_4\rangle |\sigma_3\rangle |\alpha, i_2\rangle M_{[3]}^{\alpha, i_2, \sigma_3, \alpha, i_3} B_{[4]}^{\sigma_4, \alpha, i_4}
 \end{aligned}$$

(12)

[see SYM-I.3]

$M_{[3]}  \alpha_3, i_3\rangle$	$ \alpha_3, i_3\rangle$	$ -3, 1\rangle$	$ -1, 1\rangle$	$ -1, 2\rangle$	$ -1, 3\rangle$	$ +1, 1\rangle$	$ +1, 2\rangle$	$ +1, 3\rangle$	$ +3, 1\rangle$
$\langle \alpha_2, i_2   \langle \sigma_2  $	$\gamma \uparrow$	$\uparrow \uparrow$	$\uparrow \downarrow$	$\downarrow \uparrow$	$\downarrow \downarrow$	$\uparrow \uparrow$	$\uparrow \downarrow$	$\downarrow \uparrow$	$\downarrow \downarrow$
$\langle +2, 1   \langle +1  $	$\uparrow \uparrow$	$\square$							
$\langle +2, 1   \langle -1  $	$\uparrow \downarrow$		$\square$						
$\langle 0, 1   \langle +1  $	$\downarrow \uparrow$			$\square$					
$\langle 0, 1   \langle -1  $	$\downarrow \downarrow$				$\square$				
$\langle 0, 2   \langle +1  $	$\downarrow \uparrow$					$\square$			
$\langle 0, 2   \langle -1  $	$\downarrow \downarrow$						$\square$		
$\langle -2, 1   \langle +1  $	$\downarrow \uparrow$							$\square$	
$\langle -2, 1   \langle -1  $	$\downarrow \downarrow$								$\square$

In general, the product  $M_{[3]} B_{[4]}$  has block matrix product structure, where sum over bond index  $\beta = (\alpha_3, i_3)$  matches blocks with same  $Q_3$ :

$$M_{[3]} B_{[4]} = \begin{pmatrix} \square & & & \\ & \square & & \\ & & \square & \\ & & & \square \end{pmatrix} \begin{matrix} \alpha \sigma_3, \beta \\ \alpha_3 \\ -3 \\ -1 \\ +1 \\ +3 \end{matrix} \begin{pmatrix} \square & & & \\ & \square & & \\ & & \square & \\ & & & \square \end{pmatrix} \begin{matrix} \sigma_4 \gamma \\ \beta \end{matrix} = \begin{pmatrix} \square & & & \\ & \square & & \\ & & \square & \\ & & & \square \end{pmatrix} \begin{matrix} \alpha \sigma_3, \sigma_4 \gamma \\ \beta \end{matrix}$$

(13)

Bond dimensions:  $D_2 \cdot d_3 \times D_3 = 4 \cdot 2 \times 8$        $D_3 \times d_4 \cdot D_4 = 8 \times 2 \cdot 4$        $4 \cdot 2 \times 4 \cdot 2$

But in practice, we use truncated state spaces. Suppose  $D_{\max} = 4$ ,

with  $D_2 = 4, d_3 = 2, D_3 = 4$  and that random initial state does not contain  $Q_3 = \pm 3$

Then the corresponding two blocks are missing in both  $M_{[3]}$  and  $B_{[4]}$ :

Truncated:

$$M_{[3]} B_{[4]} = \begin{pmatrix} \text{---} & \text{---} \\ \text{---} & \text{---} \\ \text{---} & \text{---} \\ \text{---} & \text{---} \end{pmatrix} \cdot \begin{pmatrix} \text{---} & \text{---} \\ \text{---} & \text{---} \\ \text{---} & \text{---} \\ \text{---} & \text{---} \end{pmatrix} = \begin{pmatrix} \text{---} & \text{---} \\ \text{---} & \text{---} \\ \text{---} & \text{---} \\ \text{---} & \text{---} \end{pmatrix} \quad (14)$$

1-site optimization of truncated  $M_{[3]}$  will never find a good ground state if the latter has non-negligible contributions from missing blocks.

2-site optimization reinstates missing block: E.g.:

$$S_z: \begin{array}{cccccccc} & Q_2 & Q_3 & Q_4 & & & & \\ & 1 & 2 & -1 & -2 & -1 & & \\ \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow \\ +1 & +1 & -1 & +1 & -1 & -1 & & \\ \underbrace{\phantom{+1 +1}}_{|\alpha\rangle_2} & & \underbrace{\phantom{-1 +1}}_{\sigma_3 \sigma_4} & & \underbrace{\phantom{-1 -1}}_{|\gamma\rangle_5} & & & \\ \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow \\ & \hat{S}^+ & \hat{S}^- & & & & & \end{array} = \begin{array}{cccccccc} & Q_2 & Q_3 & Q_4 & & & & \\ & 1 & 2 & -3 & -2 & -1 & & \\ \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow \\ +1 & +1 & +1 & -1 & -1 & -1 & & \\ & & & & & & & \end{array} \quad (15)$$

action of spin-flip term on sites 3 and 4  $\rightarrow$  can increase range of  $Q_3$

### Subspace expansion

1-site scheme that can 'expand' the variational space to recover missing blocks!

Suppose site  $l$  has been optimized by 1-site DMRG, yielding  $M_{[l]}$ , defining 'current ground state'  $|\psi_0\rangle$

current approx. for ground state

Conceptual idea: if the variational space  $\text{span}\{|\psi_0\rangle\}$  is too small, expand it to  $\text{span}\{|\psi_0\rangle, \hat{H}|\psi_0\rangle\}$

all MPS with specified bond dimensions and quantum numbers

explores more quantum numbers (except if  $|\psi_0\rangle$  is an exact eigenstate)

Concretely: define  $|\tilde{\psi}\rangle = |\psi_0\rangle + c |\psi_\alpha\rangle$  and adjust  $c$  to further minimize the energy. (16)

$\leftarrow \sim \hat{H}|\psi_0\rangle$

$\leftarrow$  'mixing coefficient'

$$|\psi_0\rangle = \underbrace{|\psi\rangle_{l+2} |\bar{\sigma}\rangle |\sigma\rangle |\alpha\rangle_{l-1}}_{\text{shorthand } \rightarrow |\alpha, \sigma, \bar{\sigma}, \gamma\rangle} (M_{[l]})^{\alpha\sigma\beta} B_{[l+1]}^{\bar{\sigma}\gamma} = \begin{array}{cccccccc} A & A & M_{[l]} & B_{[l+1]} & & & & \\ \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow \\ \alpha & & \beta & & \bar{\sigma} & & \gamma & \end{array} \quad (17)$$

$$|\psi_\alpha\rangle = \hat{P} \hat{H} |\psi_0\rangle = \begin{array}{cccccccc} A & A & M_{[l]} & B_{[l+1]} & & & & \\ \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow \\ \alpha & & \beta & & \bar{\sigma} & & \gamma & \end{array} =: \begin{array}{cccccccc} A & A & \tilde{L}_{[l]} & \tilde{R}_{[l+1]} & B & B & & \\ \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow & \uparrow \\ \alpha & & \beta & & \bar{\sigma} & & \gamma & \end{array} \quad (18)$$

projector into local basis  $\sum |\alpha, \sigma, \bar{\sigma}, \gamma\rangle \langle \alpha, \sigma, \bar{\sigma}, \gamma|$

Crucial point:  $\tilde{\beta}$  enumerates all states possibly generated by action of  $H$  on sites  $l, l+1$ , hence its range can be larger than that of  $\beta$  in (17) for  $|\psi_0\rangle$ , possibly including new quantum numbers!

Thus, virtual state space on this bond has been 'expanded':

$$\dim(\tilde{\beta}) = \dim(\alpha) \cdot \dim(\sigma)$$

[for an example, see (15)]

$$\tilde{D} = D d$$

(19)

For 6-site chain discussed earlier:

for each fixed  $\mu$ :

$$\dots \alpha\sigma\beta$$

$$\dots \bar{\sigma}\gamma$$

$$\dots \tilde{\mu} \alpha\sigma\tilde{\beta}$$

$$\dots \bar{\sigma}\gamma$$

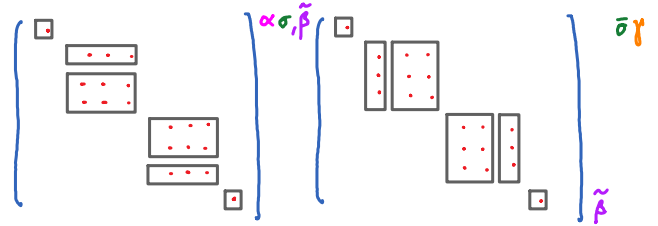
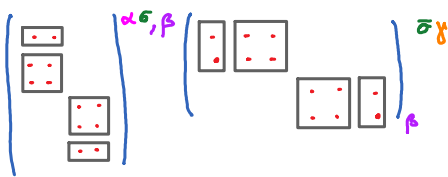
For 6-site chain discussed earlier:

for each fixed  $\mu$  :

$$(M_{[e]})^{\alpha\sigma\beta} \cdot (B_{[e+1]})_{\beta}^{\bar{\sigma}\gamma}$$

and

$$(\bar{L}_{[e]})^{\alpha\sigma\tilde{\beta}} \cdot (\bar{R}_{[e+1]\mu})_{\tilde{\beta}}^{\bar{\sigma}\gamma}$$



(20)

In (18), we identified coefficients tied to  $|\sigma\rangle|\kappa\rangle_{e-1}$  and to  $|\gamma\rangle|\bar{\sigma}\rangle$  :

$$\bar{L}_{[e]}^{\alpha\sigma\mu\tilde{\beta}} := \begin{array}{c} \xrightarrow{\alpha} \\ \begin{array}{|c|} \hline \xrightarrow{\sigma} \\ \hline \end{array} \\ \xleftarrow{\mu} \\ \xleftarrow{\tilde{\beta}} \end{array} \quad (21a)$$

$$\bar{R}_{[e+1]\mu}^{\bar{\sigma}\gamma} := \begin{array}{c} \begin{array}{|c|} \hline \xrightarrow{\tilde{\beta}} \\ \hline \end{array} \\ \begin{array}{|c|} \hline \xrightarrow{\bar{\sigma}} \\ \hline \end{array} \\ \xrightarrow{\mu} \\ \xrightarrow{\gamma} \end{array} \quad (21b)$$

$$= (L_{[e-1]} W_{[e]} M_{[e]})^{\alpha\sigma,\mu\tilde{\beta}} \quad (22a)$$

$$= (B_{[e+1]} W_{[e+1]} R_{[e+2]})_{\tilde{\beta}\mu}^{\bar{\sigma}\gamma} \quad (22b)$$

dimensions:  $(D \cdot d) \times (d_W \cdot \tilde{D})$

dimensions:  $(D \cdot d) \times (d_W \cdot \tilde{D})$  (23)

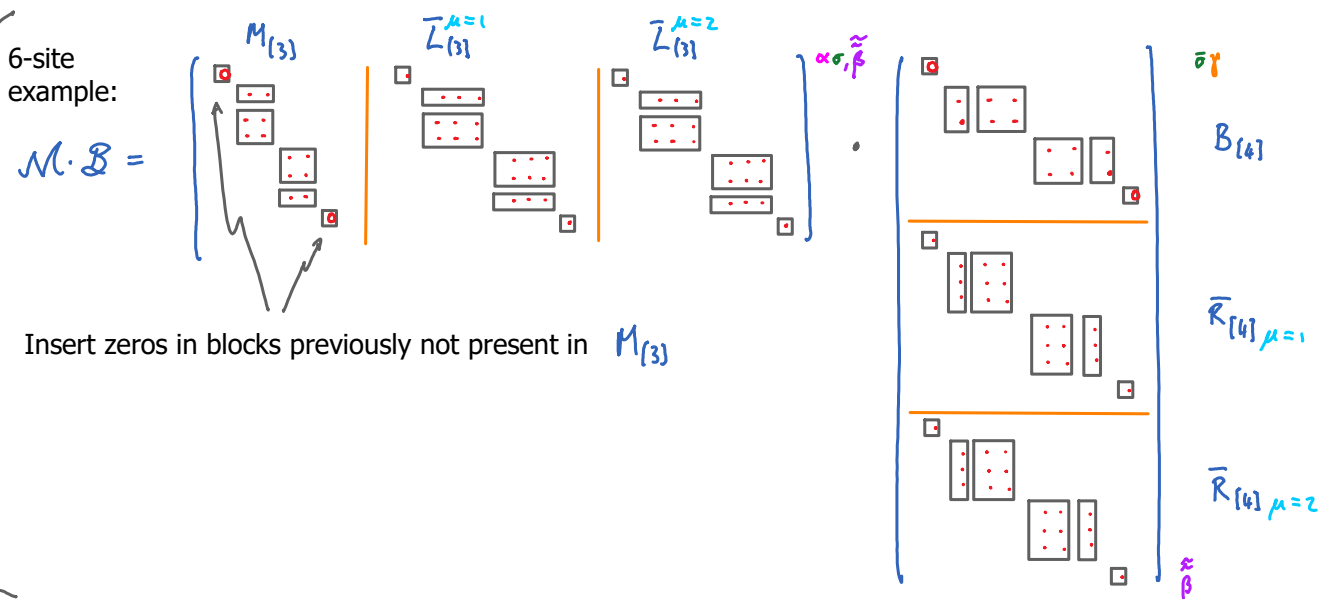
Now we add the two MPS in (16), by yet again expanding virtual state space and block-multiplying their tensors:

$$|\tilde{\Psi}\rangle = |\Psi_0\rangle + c|\Psi_a\rangle = |\alpha, \sigma, \bar{\sigma}, \gamma\rangle \left[ (M_{[e]})^{\alpha\sigma\beta} (B_{[e+1]})_{\beta}^{\bar{\sigma}\gamma} + c (\bar{L}_{[e]})^{\alpha\sigma\mu\tilde{\beta}} (\bar{R}_{[e+1]\mu})_{\tilde{\beta}\mu}^{\bar{\sigma}\gamma} \right] \quad (24)$$

$$\begin{pmatrix} M_{[e]} & c\bar{L}_{[e]} \end{pmatrix} \begin{matrix} \alpha\sigma \\ \tilde{\beta} \end{matrix} \cdot \begin{matrix} B_{[e+1]} \\ \bar{R}_{[e+1]\mu} \end{matrix} \begin{matrix} \bar{\sigma}\gamma \\ \tilde{\beta} \end{matrix} =: (\mathcal{M} \cdot \mathcal{B})^{\alpha\sigma\bar{\sigma}\gamma} \quad (25)$$

$(Dd) \times (Dd)$

$$\dim(\tilde{\beta}) = \dim(\beta) + \dim(\tilde{\beta}) \cdot \dim(\mu) \quad (26)$$



To deal with large-dimension bond  $\tilde{\beta}$ , use SVD on  $\mathcal{M} = u s v^t$  and truncate back to (!!)

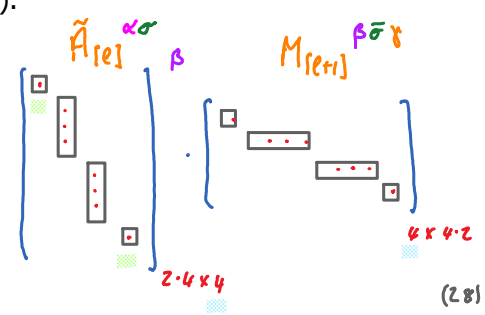
$$\begin{array}{c} \mathcal{M} \quad \mathcal{B} \\ \leftarrow \quad \leftarrow \\ \text{D} \quad \text{D} \\ \uparrow \quad \uparrow \\ d \quad d \\ \approx \tilde{\text{D}} \end{array} \xrightarrow{\text{SVD}} \begin{array}{c} u \quad s \quad v^t \quad \mathcal{B} \\ \leftarrow \quad \leftarrow \quad \leftarrow \quad \leftarrow \\ \text{D} \quad \text{D} \quad \text{D} \quad \text{D} \\ \uparrow \quad \uparrow \quad \uparrow \quad \uparrow \\ d \quad d \quad d \quad d \\ \approx \tilde{\text{D}} \end{array} \xrightarrow{\text{truncate}} \begin{array}{c} (u) \tilde{s} \quad v^t \quad \mathcal{B} \\ \leftarrow \quad \leftarrow \quad \leftarrow \quad \leftarrow \\ \text{D} \quad \text{D} \quad \text{D} \quad \text{D} \\ \uparrow \quad \uparrow \quad \uparrow \quad \uparrow \\ d \quad d \quad d \quad d \\ \approx \tilde{\text{D}} \end{array} = \begin{array}{c} \text{D} \\ \leftarrow \quad \leftarrow \\ \text{D} \quad \text{D} \\ \uparrow \quad \uparrow \\ d \quad d \\ \approx \tilde{\text{D}} \end{array} \begin{array}{c} \text{D} \\ \leftarrow \quad \leftarrow \\ \tilde{A}_{[l]} \quad M_{[l+1]} \\ \uparrow \quad \uparrow \\ \text{D} \quad \text{D} \\ \approx \tilde{\text{D}} \end{array} \quad (27)$$

Left-normalized  $\tilde{A}_{[l]}$  is final result of optimization of site  $l$ , and  $M_{[l+1]}$  initializes optimization of site  $l+1$ .

Crucial point: as mentioned earlier,  $\tilde{\beta}$  may include larger range of quantum numbers than  $\beta$ . If these contribute significant weight to  $|\psi_\alpha\rangle$ , then they will have significant weight in SVD of (27), hence will survive the truncation in (27), and thus be present in  $\tilde{A}_{[l]}$ .

In this way, subspace expansion succeeds in 'reinstating missing quantum numbers' (without ever needing to do Lanczos with huge effective 2-site Hamiltonian of 2-site DMRG!).

6-site example: if  $Q_3 = \pm 3$  carries significant weight, then SVD, followed by truncation with  $\dim(\beta) = 4$ , will reinstate those blocks:



Remark: [Hubig2015] choose  $\bar{R}_{[l+1]} = \circ$  [instead of (22b)], arguing that it only affects  $M_{[l+1]}$ , which will be updated anyway when optimizing site  $l+1$  via Lanczos ground state solver in next iteration. But using (21b) speeds up convergence for that Lanczos solver [Weichselbaum (private communication)].

Summary of subspace expansion algorithm 'strictly single-site DMRG (DMRG3S)'

Left-to-right sweep:

1. In site-canonical basis for site  $l$ , use 1-site DMRG to find optimized  $M_{[l]}$

with  $H_{[l]} M_{[l]} = E M_{[l]}$  (29)  
↙ new current energy estimate

2. Expand subspace,  $M_{[l]} B_{[l+1]} \rightarrow \mathcal{M}_{[l]} \cdot \mathcal{B}_{[l+1]} = \begin{pmatrix} M_{[l]} & c \bar{L}_{[l]} \end{pmatrix} \begin{pmatrix} B_{[l+1]} \\ \bar{R}_{[l+1]} \end{pmatrix} \approx \tilde{A}_{[l]} M_{[l+1]}$  (30)  
 do SVD + truncation:

3. Obtain 'post-expansion' energy estimate  $\tilde{E} =$  (31)  
 Back to 1. and repeat.

Right-to-left sweep is analogous, with

2. Expand subspace,  $A_{[l-1]} M_{[l]} \rightarrow \mathcal{A}_{[l-1]} \mathcal{M}_{[l]} = \begin{pmatrix} A_{[l-1]} & c \bar{L}_{[l-1]} \end{pmatrix} \begin{pmatrix} M_{[l]} \\ \bar{R}_{[l]} \end{pmatrix} \approx M_{[l-1]} \tilde{B}_{[l]}$  (32)  
 do SVD + truncation:

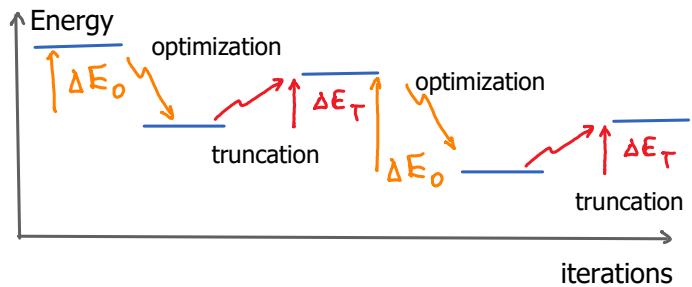
3. Obtain 'post-expansion' energy estimate

$$\tilde{E} = \begin{array}{c} \begin{array}{|c|c|c|} \hline \begin{array}{c} \leftarrow \\ \leftarrow \\ \leftarrow \end{array} \\ \hline \begin{array}{c} \leftarrow \\ \leftarrow \\ \leftarrow \end{array} \\ \hline \begin{array}{c} \leftarrow \\ \leftarrow \\ \leftarrow \end{array} \\ \hline \end{array} \\ \begin{array}{c} M_{[e-1]} \quad \tilde{B}_{[e]} \\ L_{[e-2]} \quad M_{[e-1]}^f \quad \tilde{\delta}_{[e]}^f \quad R_{[e+1]} \end{array} \end{array} \quad (33)$$

Choice of mixing coefficient  $c$  :

$\Delta E_0$  energy change from 1-site optimization, usually  $< 0$

$\Delta E_T$  energy change from truncation, can be either  $< 0$  or  $> 0$ .



Behavior of energy changes depends on  $c$  :

If  $c$  is too small, one can get stuck in local minimum, i.e.  $\Delta E_0 \approx 0$

If  $c$  is too large, truncation destroys changes made by previous local optimization, and  $0 < \Delta E_T \approx |\Delta E_0|$   
Then reduce  $c$  for next iteration on next site. (34)

If  $\Delta E_T$  is much smaller than  $|\Delta E_0|$  or even negative, then one can increase  $c$  for next iteration.

In practice, start with  $c = 1$  and aim for  $\Delta E_T \approx 0.3 |\Delta E_0|$  (for scheme of [Hubig2015], with  $\bar{R} = 0$ ) and change  $c$  by multiplying with some factor greater/smaller than 1 (rather than subtracting/adding). After several DMRG sweeps back and forth through the chain, the 'correct' quantum numbers will typically have been found. So, one can eventually take  $c \rightarrow 0$ , returning to pure 1-site DMRG.

Remark: there may be better ways to choose  $c$  : build the 2-dimensional Krylov space  $K = \{|\psi_0\rangle, |\psi_a\rangle\}$

and choose  $c$  such that  $|\tilde{\psi}\rangle = |\psi_0\rangle + c|\psi_a\rangle$  is the ground state of in this space. (35)

This corresponds to the first step of a Lanczos iteration, cf. DMRG-I.2, Eqs. (6-11).

Orthogonalize:  $b_1 |\psi_1\rangle := |\tilde{\psi}_1\rangle := \hat{P} \hat{H} |\psi_0\rangle - |\psi_0\rangle \langle \psi_0 | \hat{P} \hat{H} | \psi_0 \rangle$  (36)

$$= |\psi_a\rangle - a_0 |\psi_0\rangle \quad (37)$$

Computing  $|\tilde{\psi}_1\rangle$  involves addition of two MPS, can be done as in (24,25) [no need for SVD + truncation here], to arrive at an MPS with same bond dimensions as  $|\psi_0\rangle$ .

$$b_1 = \sqrt{\langle \tilde{\psi}_1 | \tilde{\psi}_1 \rangle} = \langle \psi_1 | H | \psi_0 \rangle, \quad a_1 = \langle \psi_1 | H | \psi_1 \rangle \quad (38)$$

Then diagonalize  $H_K = \begin{pmatrix} a_0 & b_1 \\ b_1 & a_1 \end{pmatrix}$ , its lower-energy eigenvector yields optimal  $c$ . (39)

Then use that to compute a new  $|\tilde{\psi}\rangle$ , and from that compute  $M-B = \tilde{A}_{[e]} M_{[e+1]}$  as described in (24,25,27). Explore whether this yields better convergence behavior!

How can one estimate / control error during extrapolations with increasing bond dimension? See [Hubig2018].