

Randomized higher-order tensor renormalization group for higher dimension.



[D. Kadoh, K.N. arXiv:1912.02414]
[K.N. arXiv:2307.14191]

Katsumasa Nakayama (RIKEN)
2023/07/31 @Online.

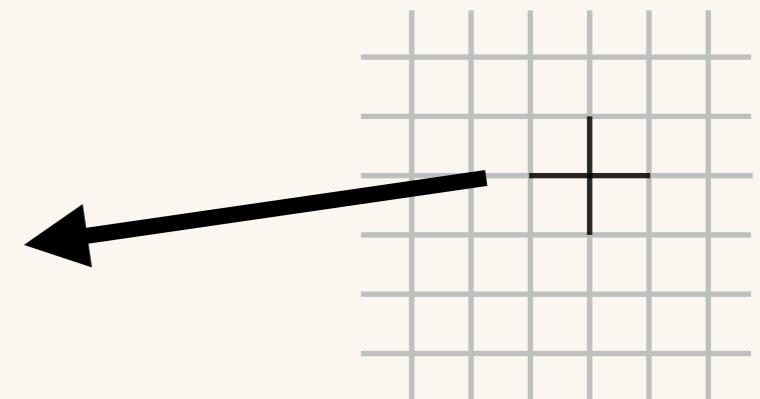
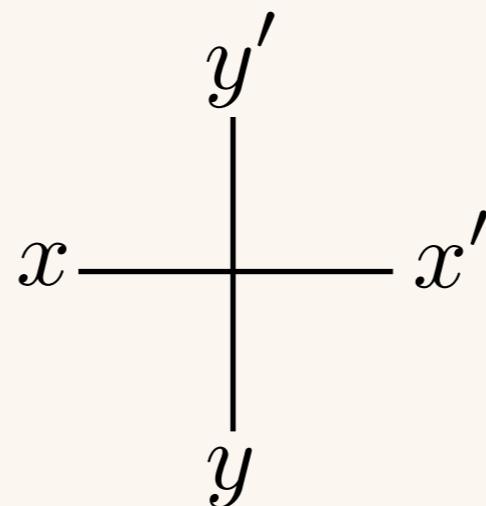
Tensor renormalization group (TRG)

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

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

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

$$A_{xyx'y'} =$$



- Sign problem
- Another representation

- ✗ High cost ($\dim \geq 3$)
- △ Systematic error

How can we take whole contraction approximately?

→ Singular value decomposition (Frobenius norm)

● Singular Value Decomposition (SVD)

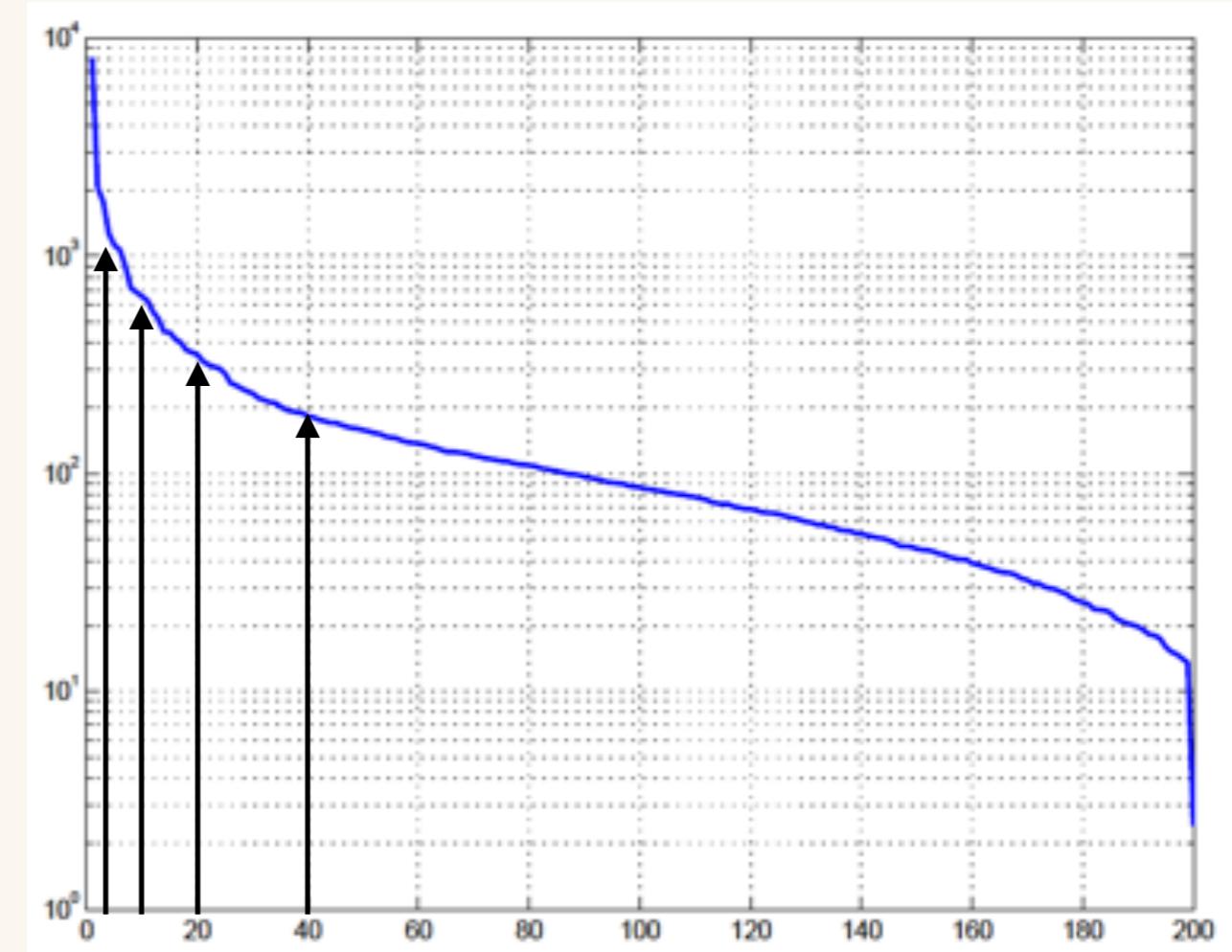
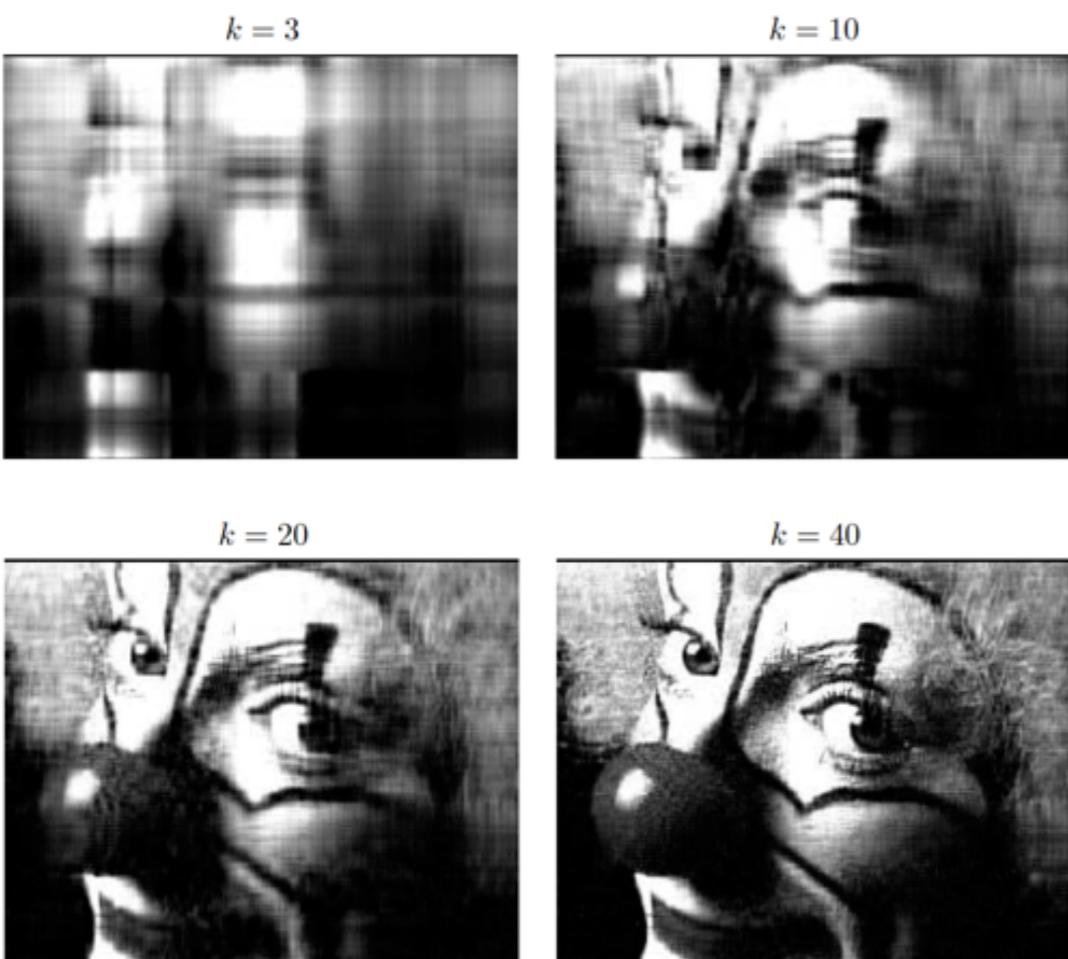
$$T_{abcd} = \sum_k^D A_{ab}{}^k \lambda^k B_{cd}{}^k$$

The diagram illustrates the Singular Value Decomposition (SVD) of a matrix T_{abcd} . The matrix T is decomposed into three components: $A_{ab}{}^k$, λ^k , and $B_{cd}{}^k$. The matrix A has dimensions $a \times k$, and the matrix B has dimensions $k \times c$. The singular value λ is represented by a red double-headed arrow between the two matrices, indicating its magnitude.

- ◊ Larger singular values λ have much “information” of T
→ (Frobenius norm)
- We can approximate the matrix by the cutoff of index k

$$\dim(k) = \dim(a)\dim(b) \rightarrow D$$

● SVD for a coarse graining (e.g. Image)



[<http://www.na.scitec.kobe-u.ac.jp/~yamamoto/lectures/cse-introduction2009/cse-introduction090512.PPT>]

● Motivation: The origin of the systematic error

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

Systematic error: isometry,

Cost:

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

d : dimension

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]

Systematic error: isometry, decomposition, R-SVD

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→ How about HOTRG with Randomized-SVD?

→ Can we reduce the systematic error from decomposition?

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Systematic error: isometry, decomposition, R-SVD

Cost: $O(D^{d+3})$

→ How about HOTRG with Randomized-SVD?

→ Can we reduce the systematic error from decomposition?

● HOTRG with R-SVD

	with R-SVD	w/o R-SVD
◇ HOTRG	?	$O(D^{4d-1})$
◇ ATRG	$O(D^{2d+1})$	$O(D^{3d})$
◇ TTRG	$O(D^{d+3})$	$O(D^{d+4})$

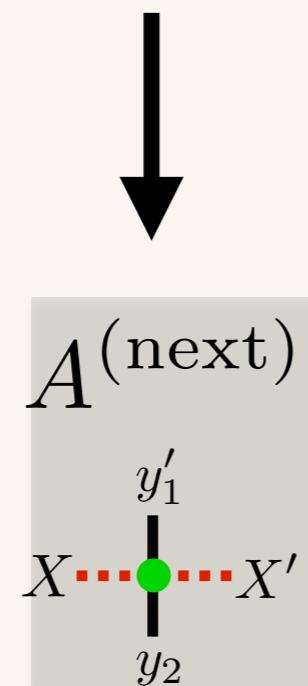
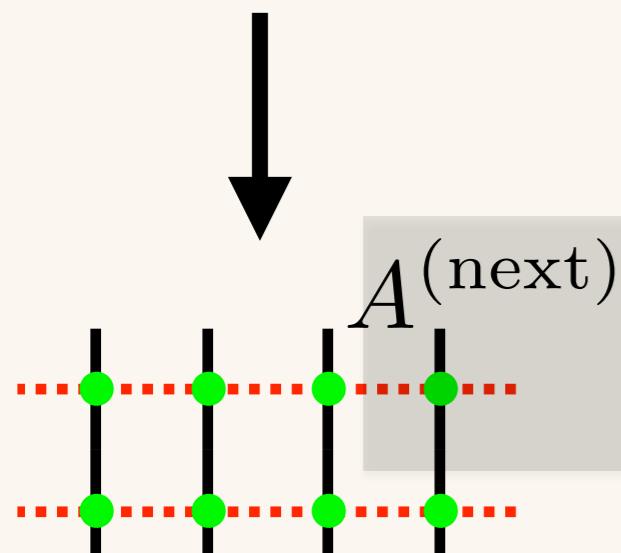
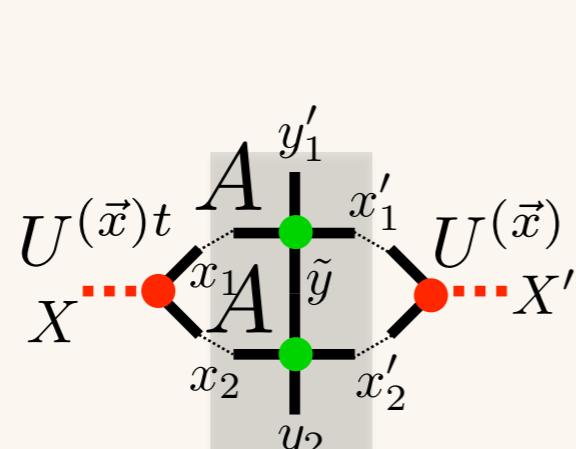
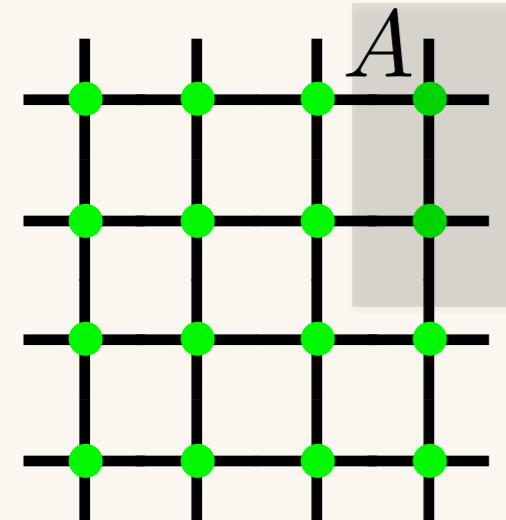
→ We propose HOTRG with R-SVD.

HOTRG with randomized SVD

● Higher-Order TRG (HOTRG)

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

◇ Contraction by projection operator U (isometry)



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

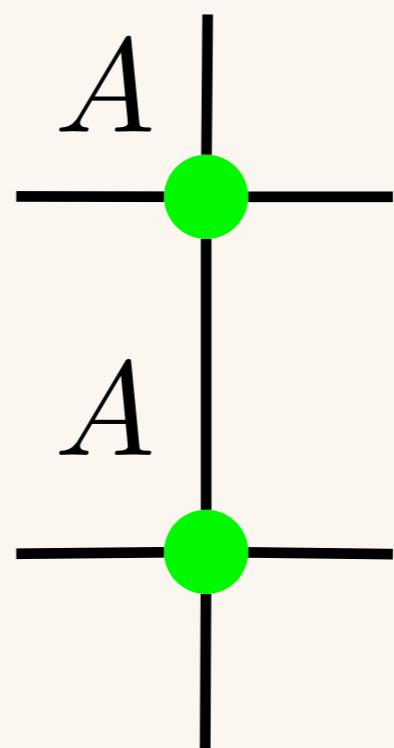
$\rightarrow U^{(\vec{x})}$ is made by SVD of $\Gamma\Gamma^t$

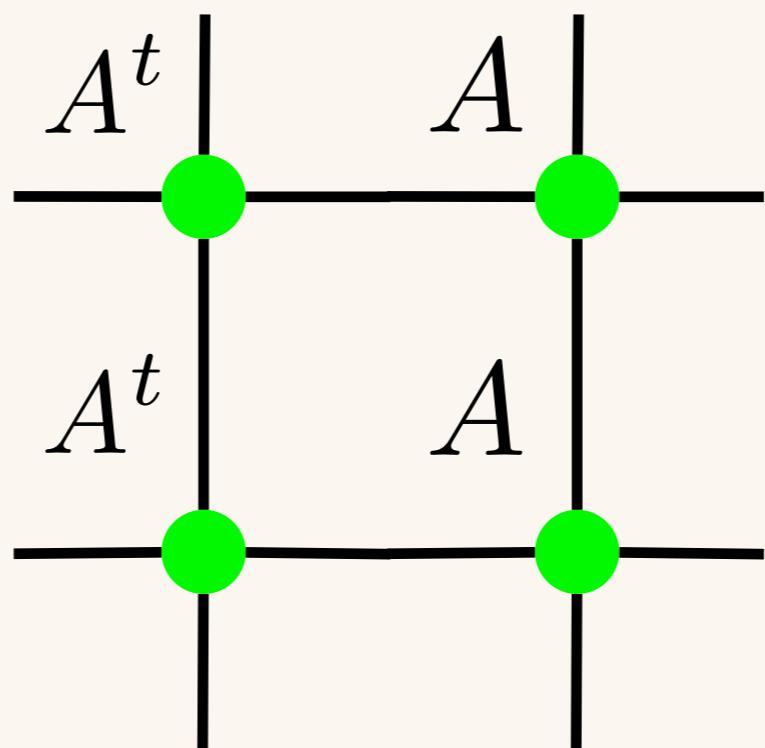
$$[\Gamma\Gamma^t]_{[x_1x_2][x_1^tx_2^t]} = \sum_{k=1}^{D^2} U_{[x_1x_2]k}^{(x)} \lambda_k U_{[x_1^tx_2^t]k}^{(x)}$$

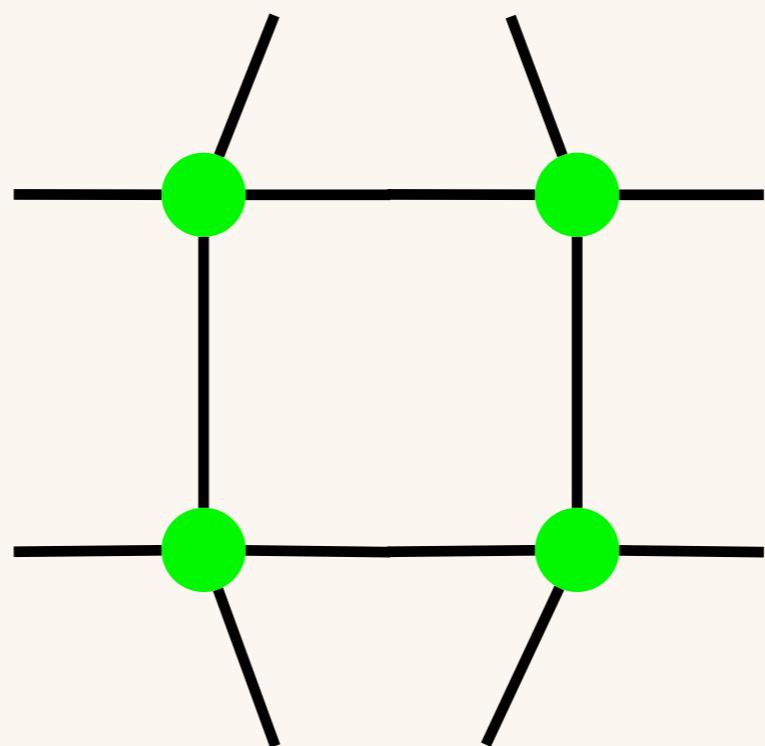
SVD Cutoff: $D^2 \rightarrow D$
Cost: $O(D^6)$

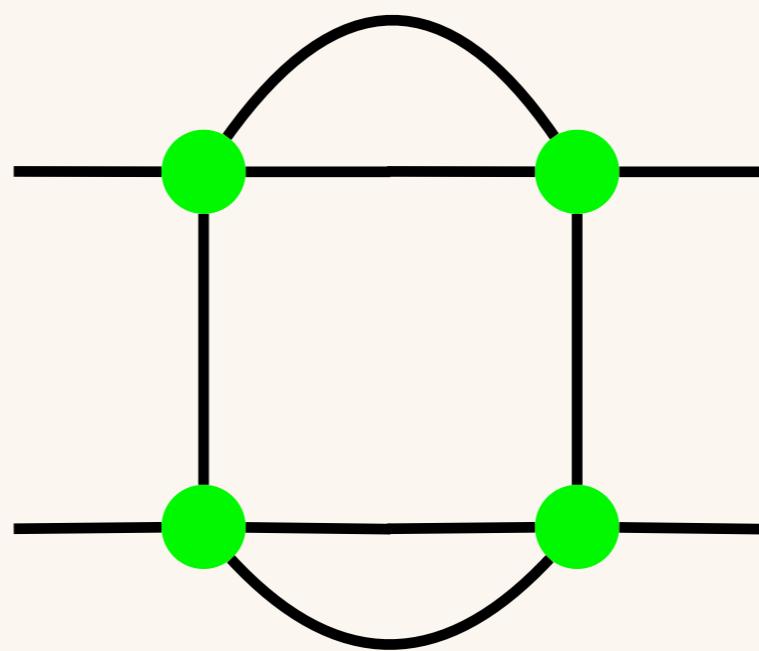
$$U^t A A U = A^{(\text{next})}$$

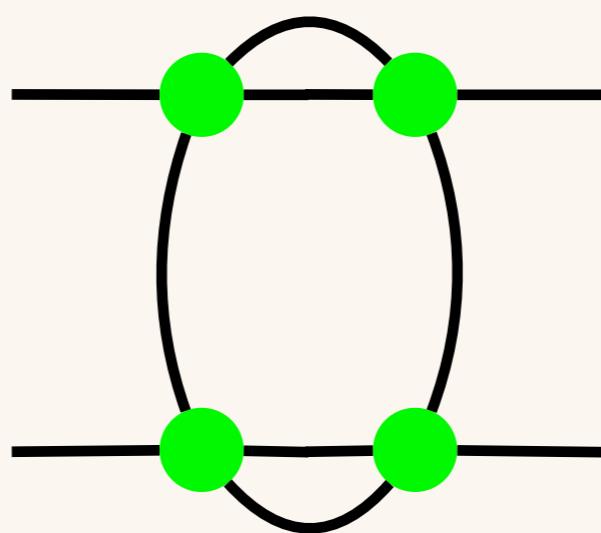
Contraction Cost: $O(D^7)$

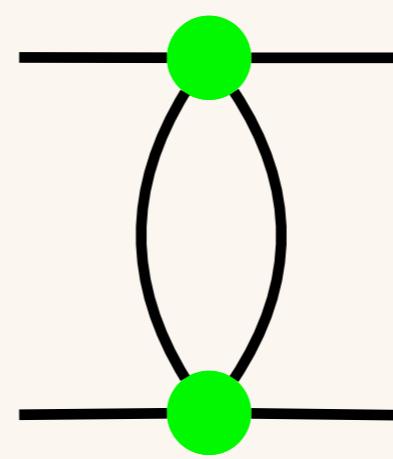


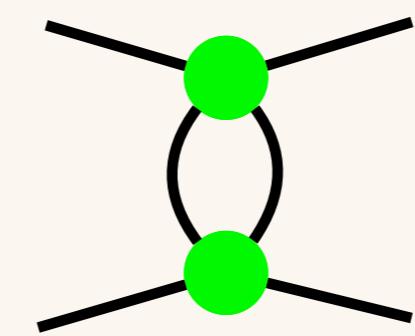


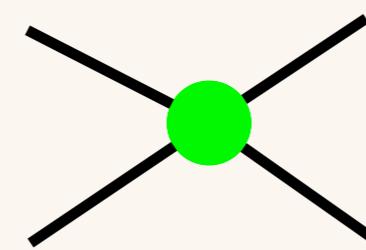




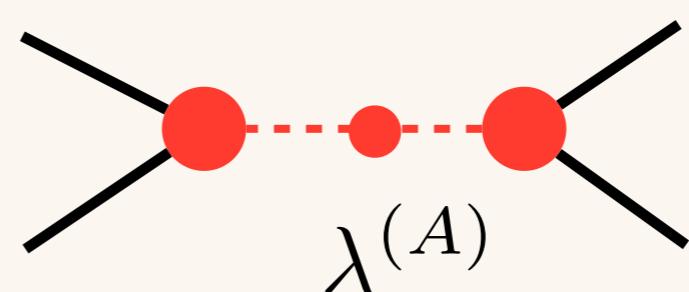


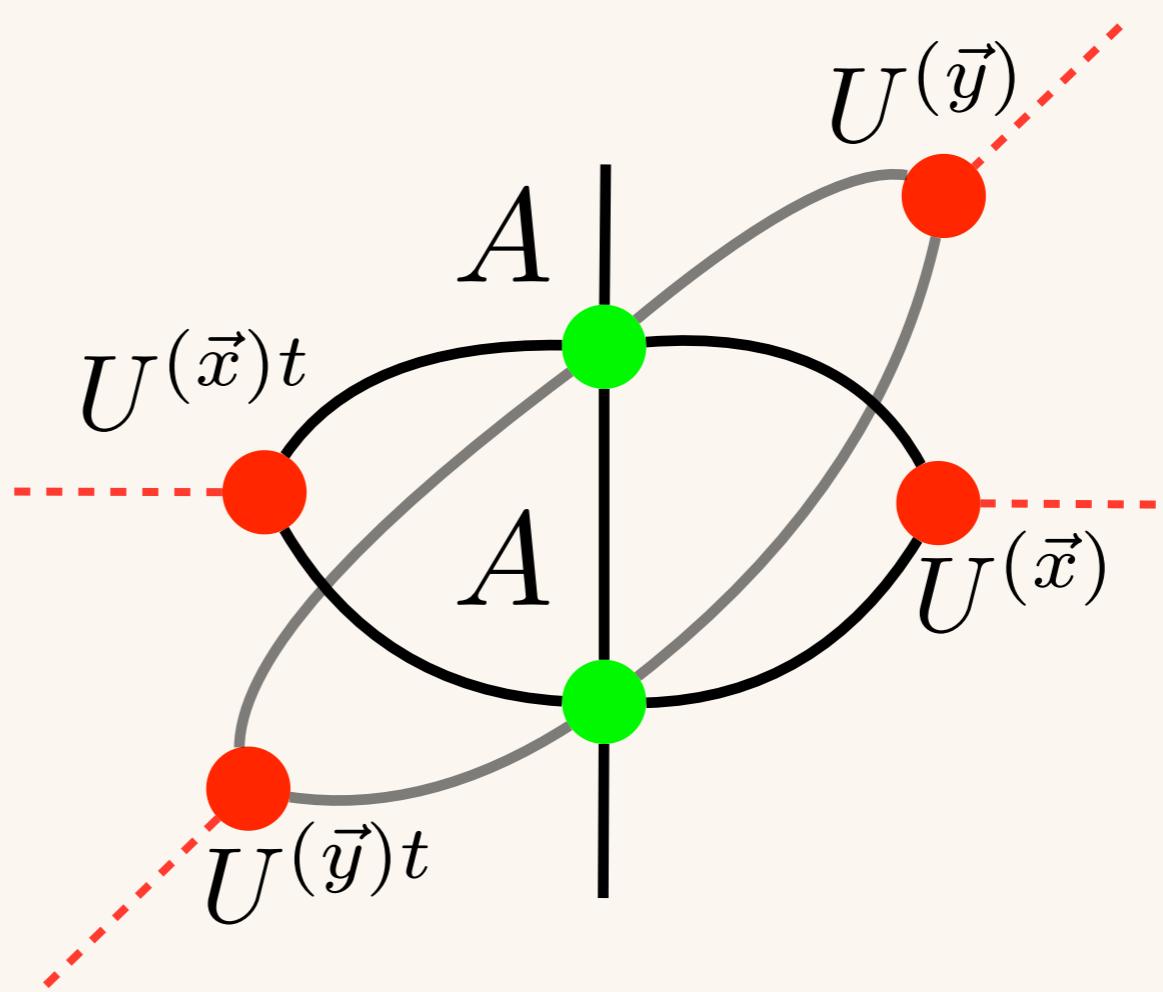


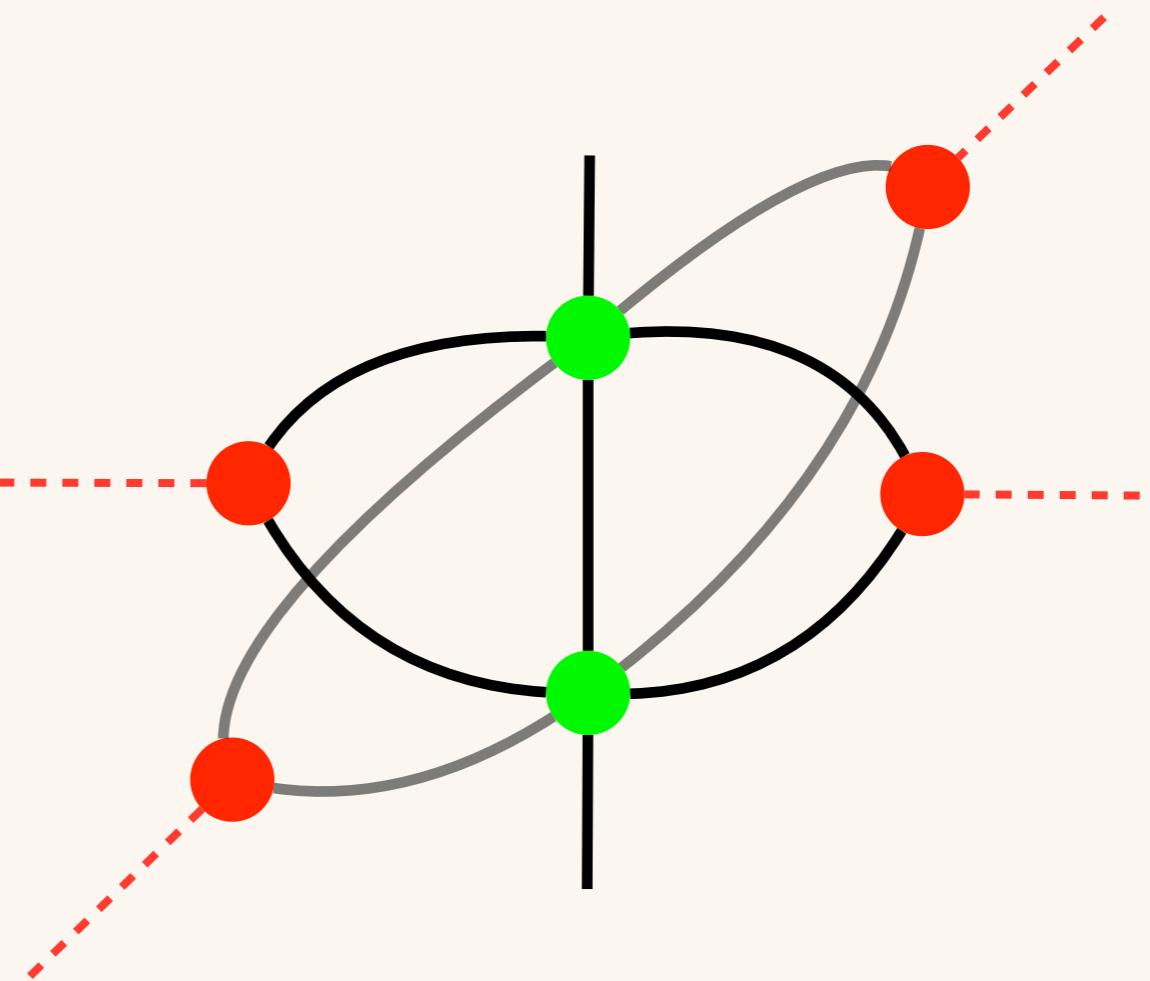


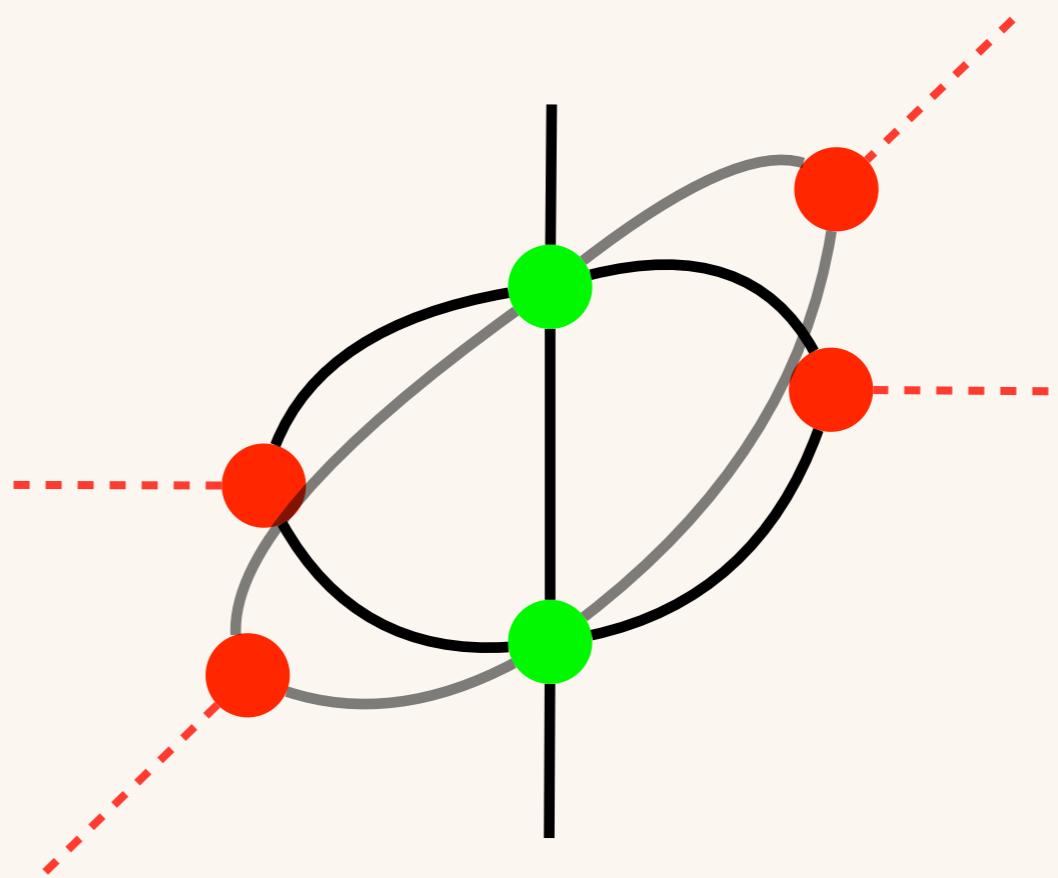


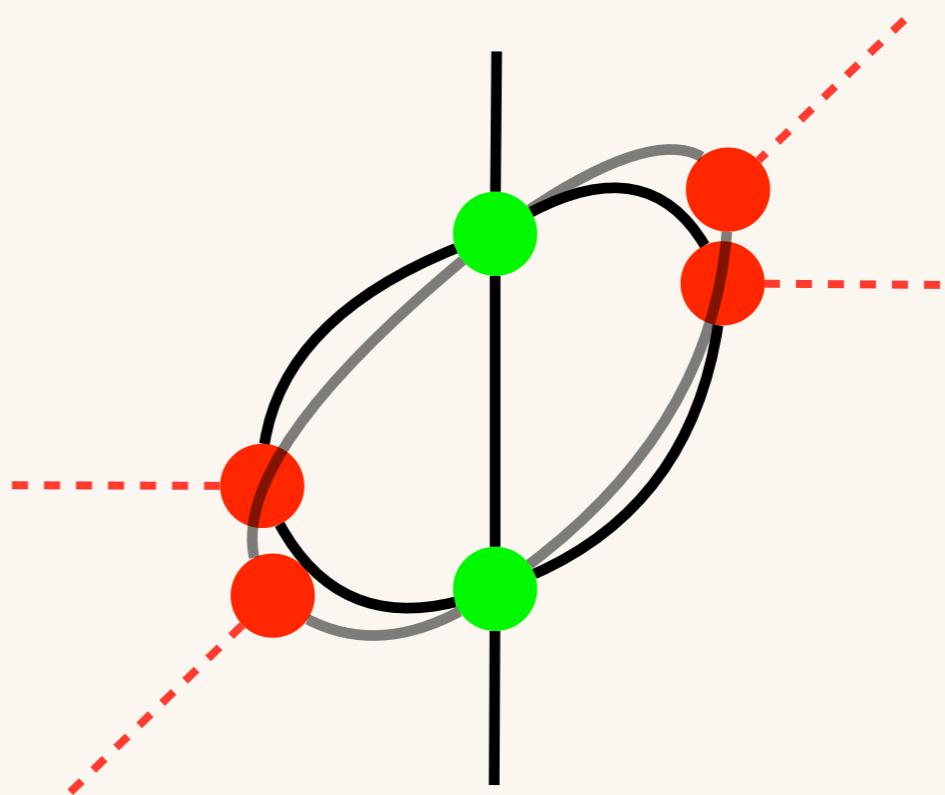
$$U(\vec{x})t \quad U(\vec{x})$$

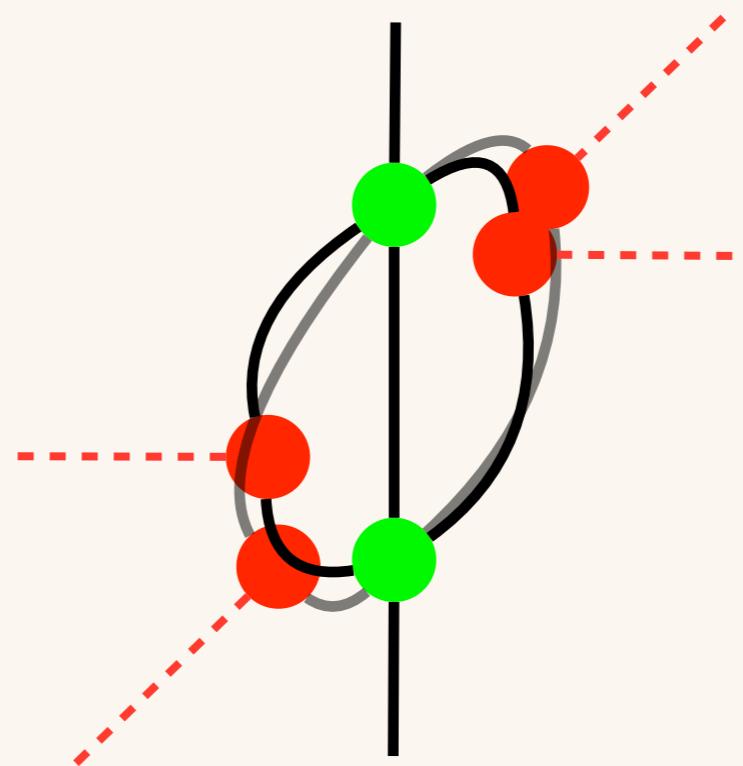


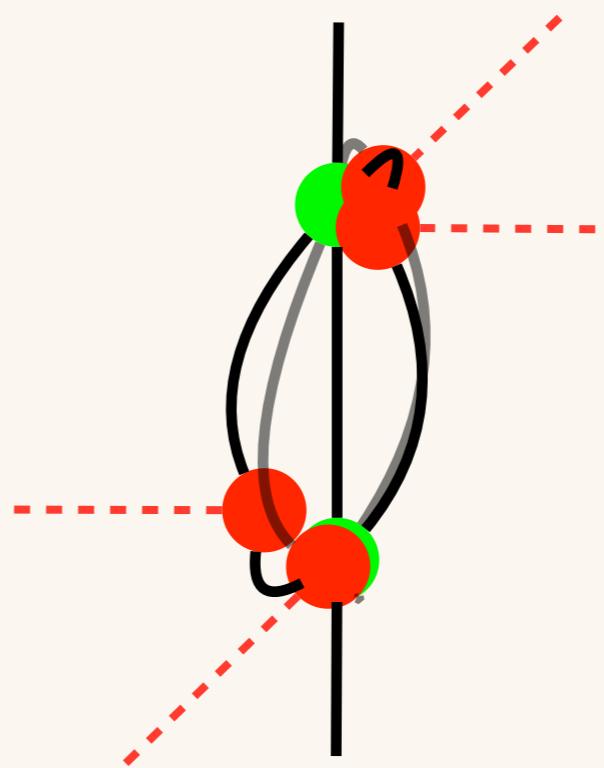


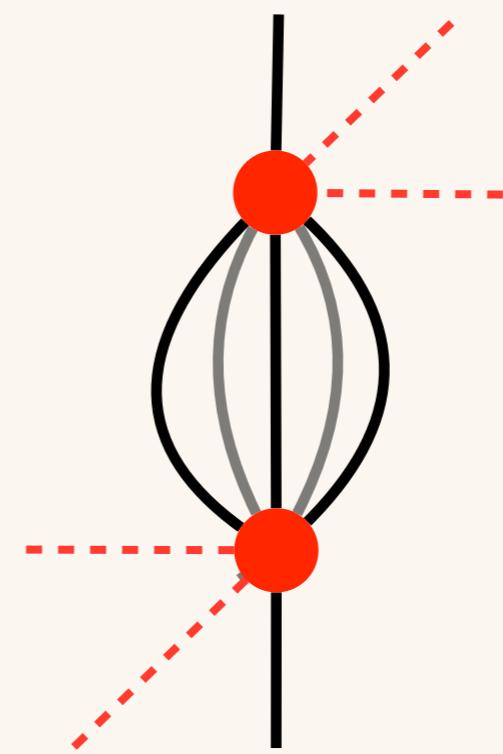


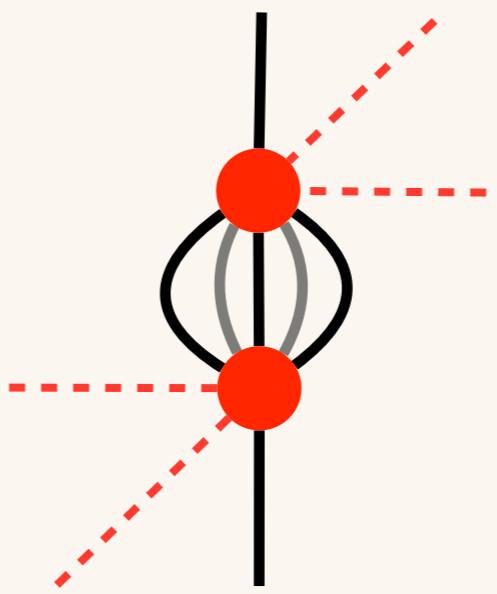




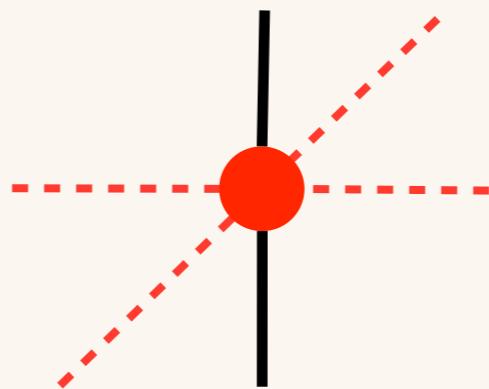








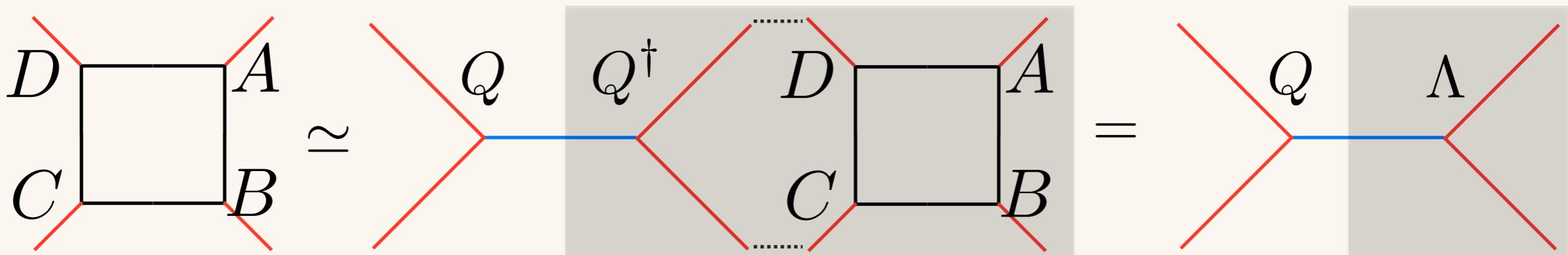
$A^{(\text{next})}$



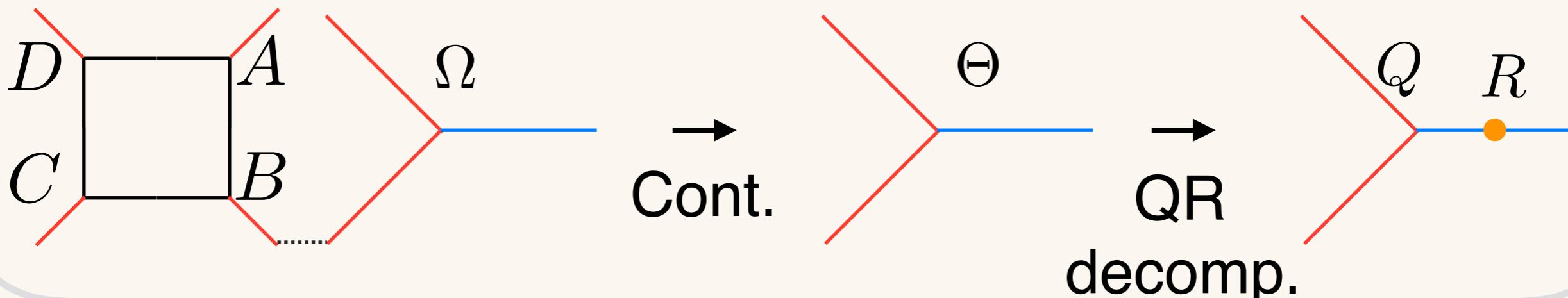
Randomized-SVD

[N. Halko, et al. arXiv:0909.4061]
 [S. Morita, et al. arXiv:1712.01458]

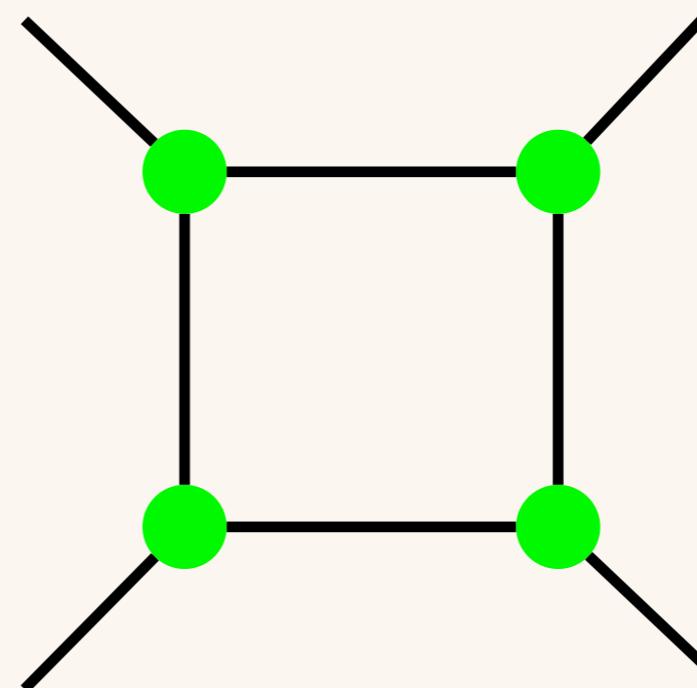
- ◊ Approximated contraction by orthogonal matrix Q



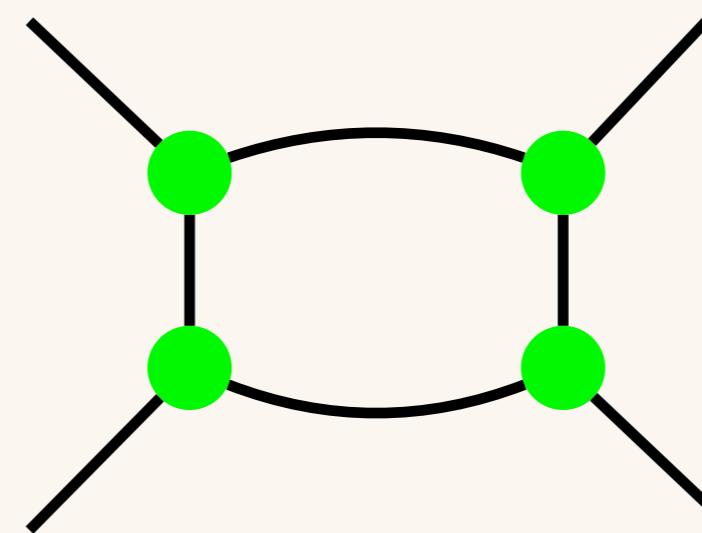
- ◊ SVD of $\Lambda \equiv Q^\dagger ABCD$.
- ◊ To prepare Q , we use randomized method & QR decomp.
- ◊ Contraction with the random tensor Ω



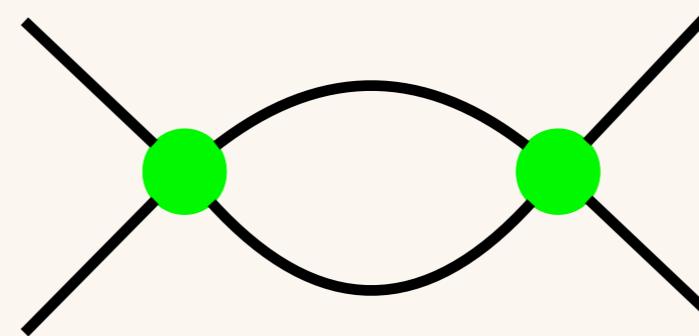
◇ Step 1: $O(D^5)$



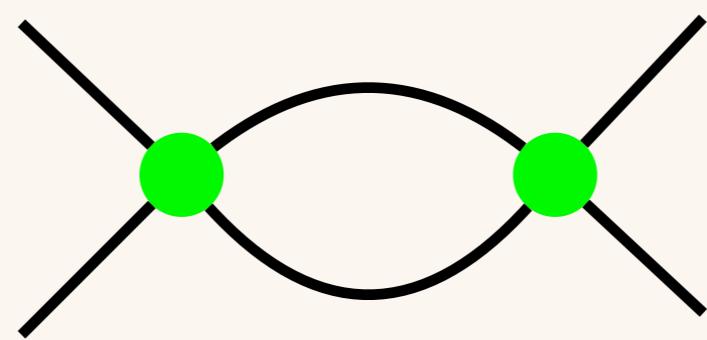
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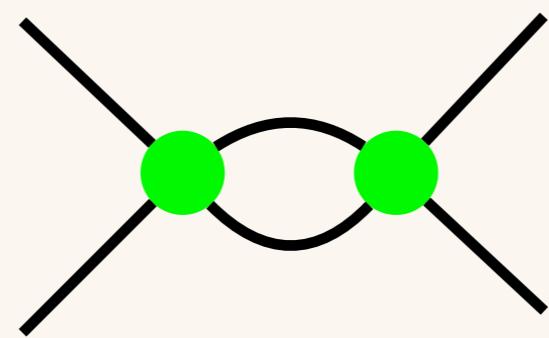
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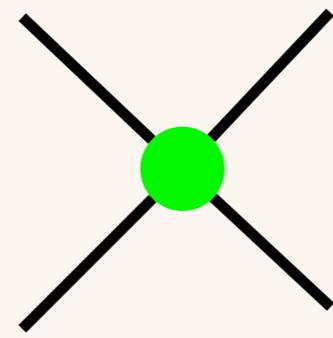
◇ Step 2: $O(D^6)$



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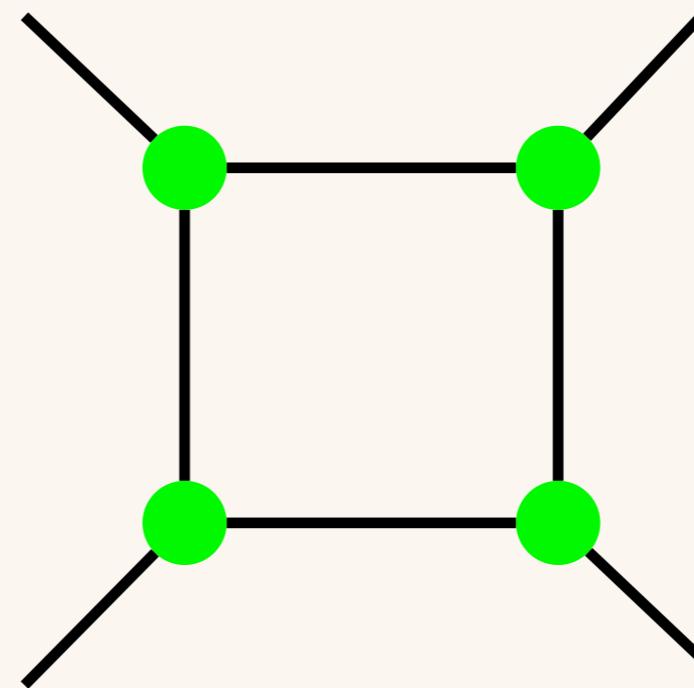
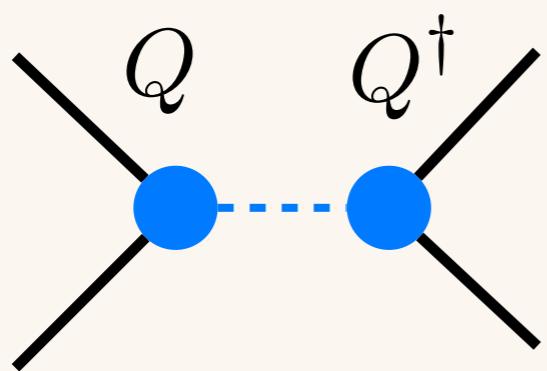


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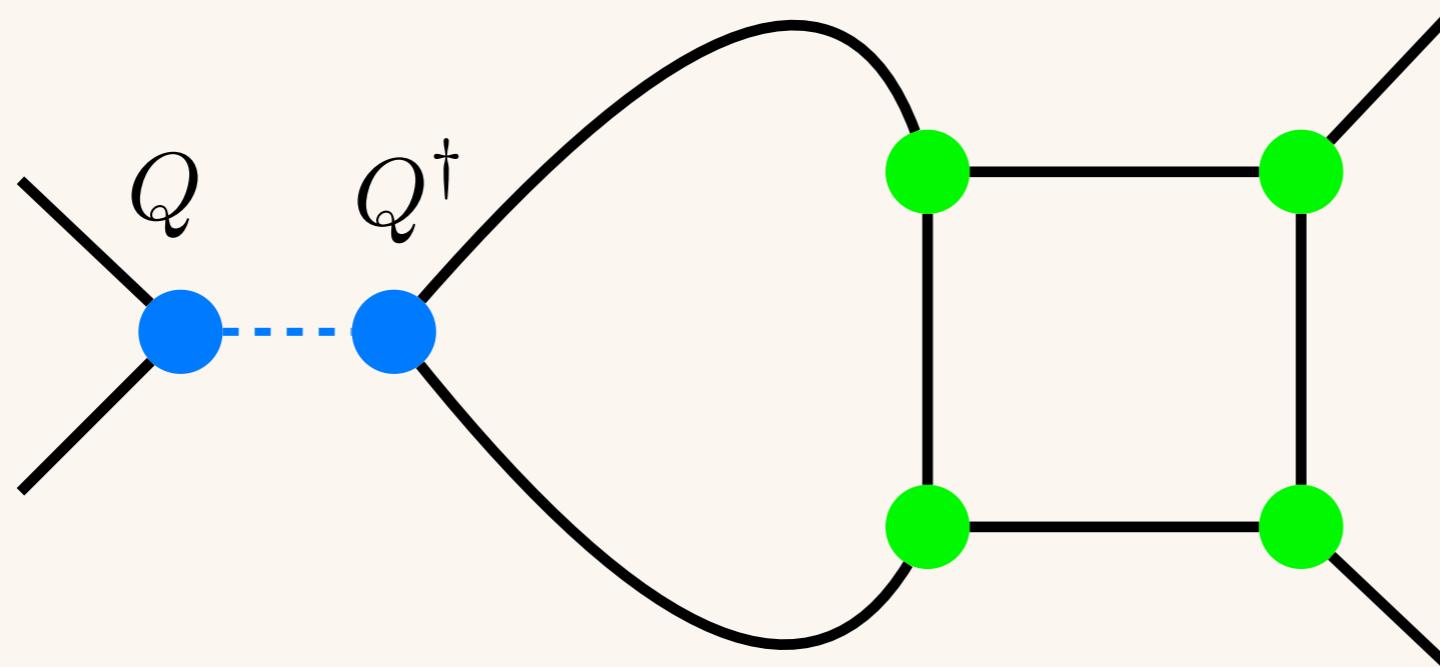


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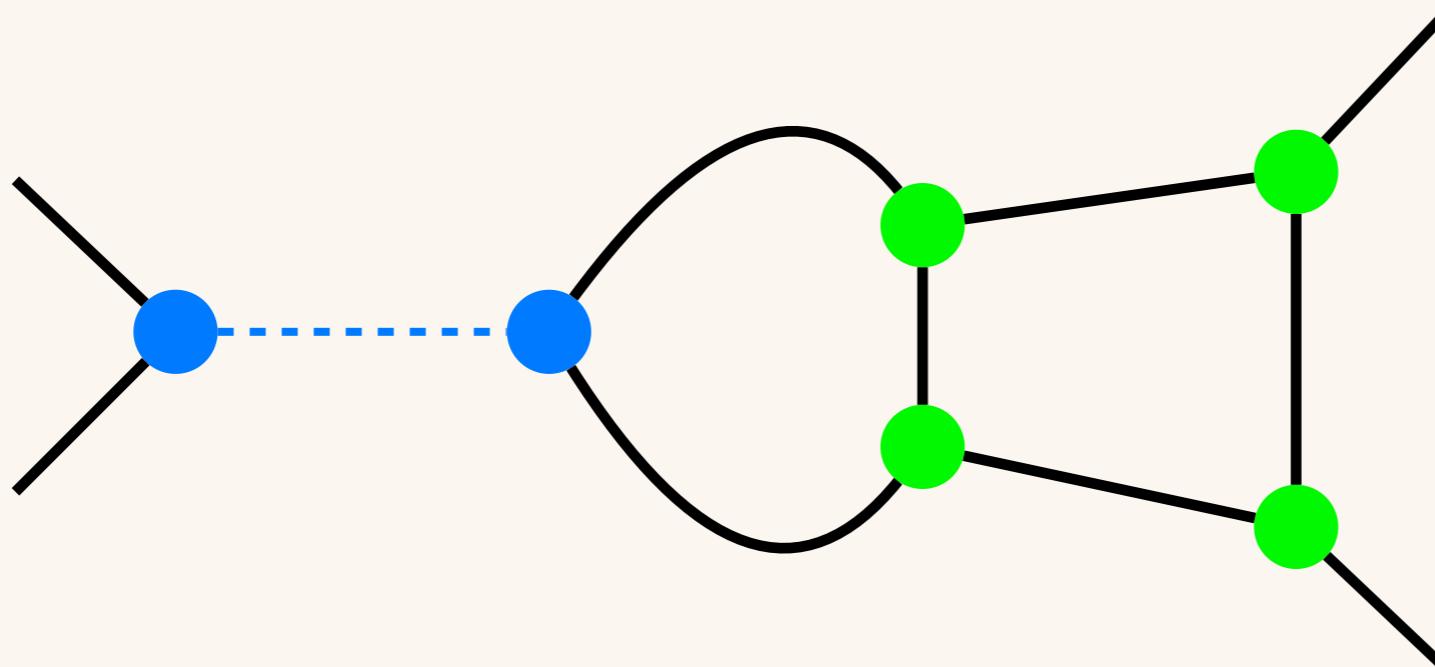
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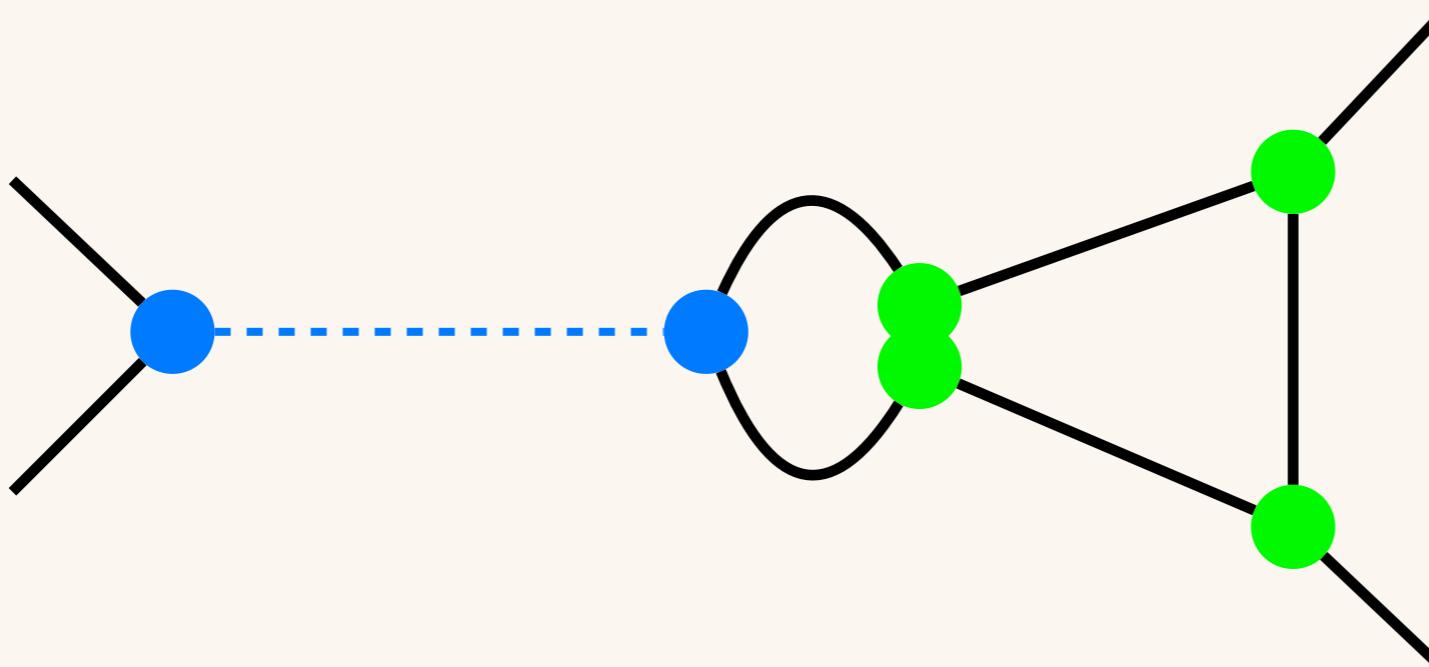
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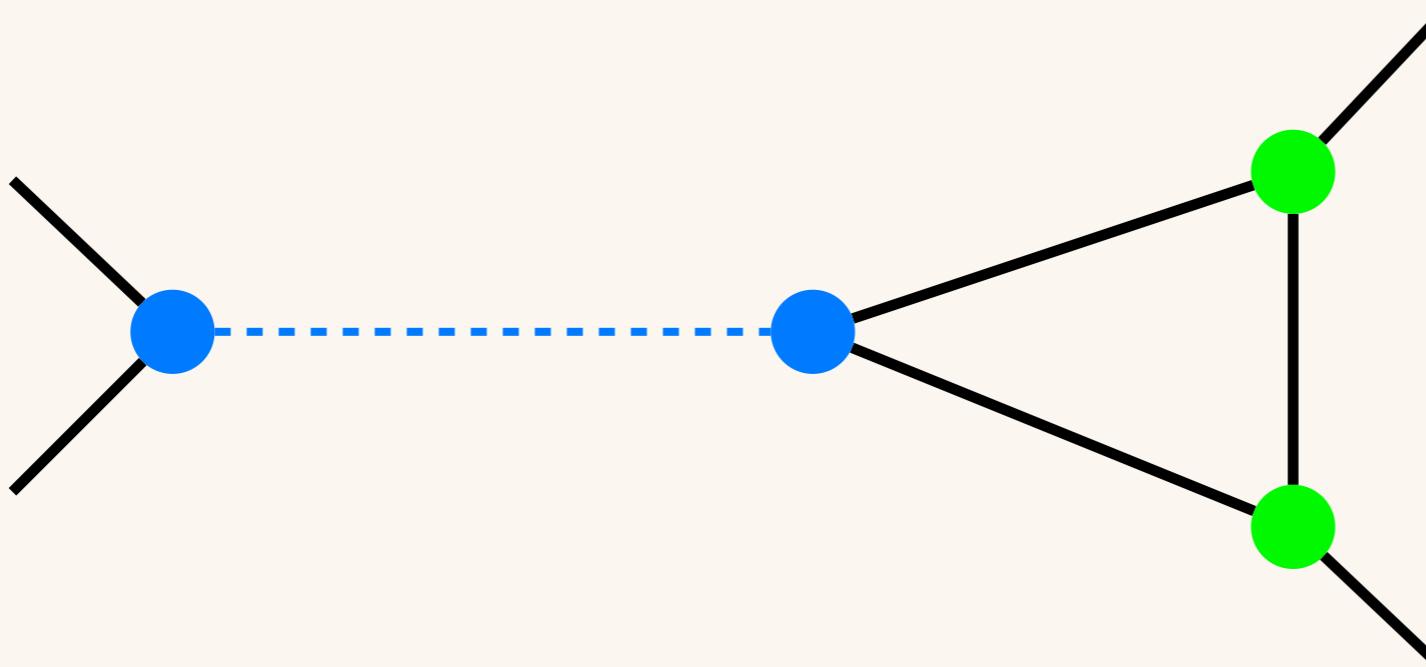
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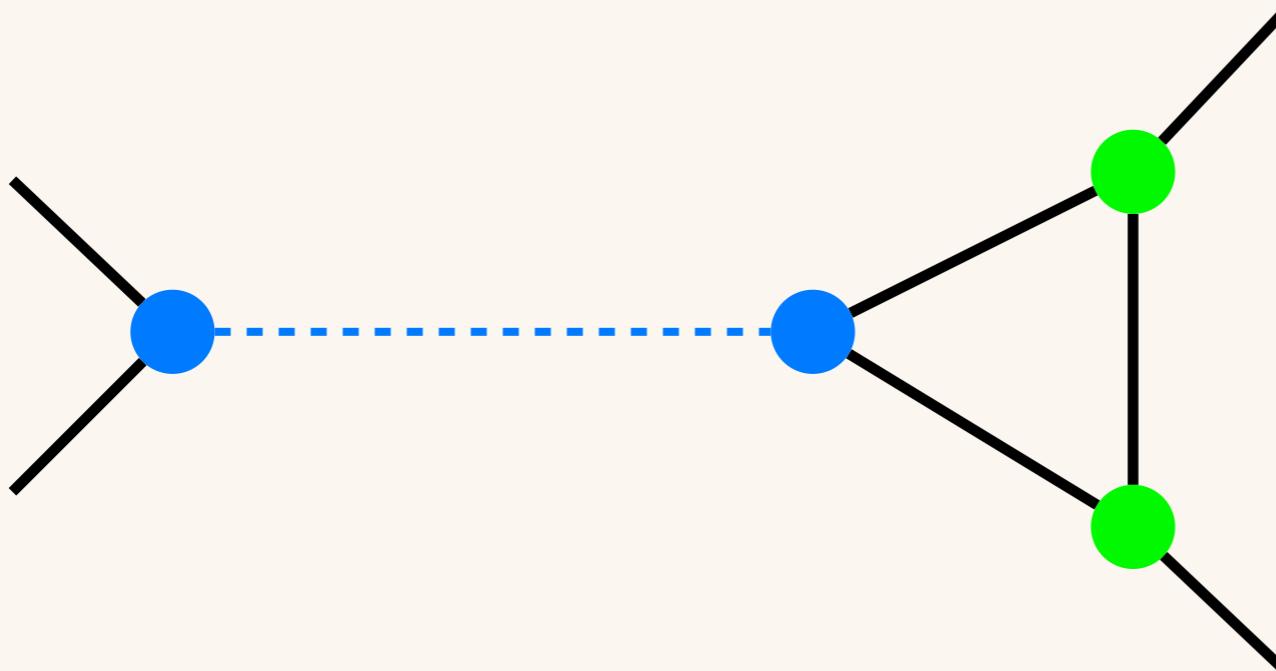
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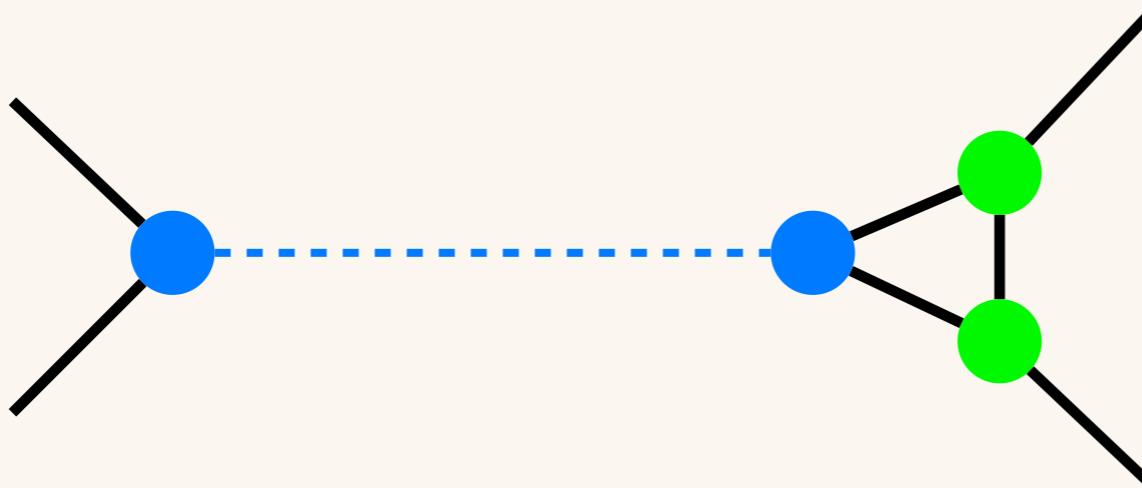
◇ Step 1: $O(D^5)$



◇ Step 2: $O(D^5)$



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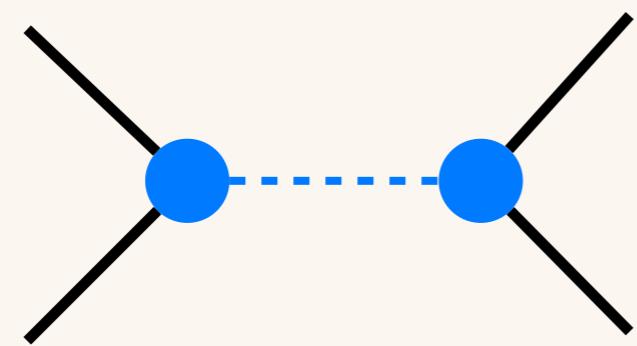
◇ Step 2: $O(D^5)$

◇ Step 3: $O(D^5)$

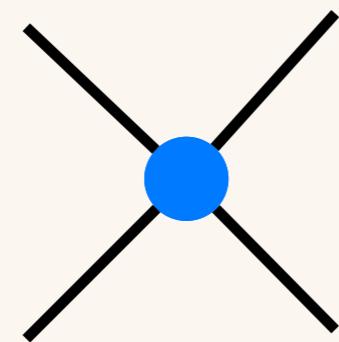


◇ Step 2: $O(D^5)$

◇ Step 3: $O(D^5)$



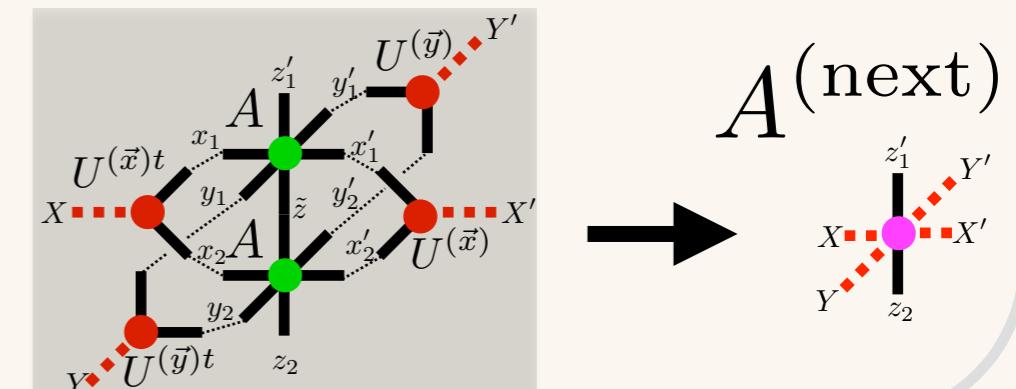
◇ Step 3: $O(D^5)$



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

◇ Contraction with U

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



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

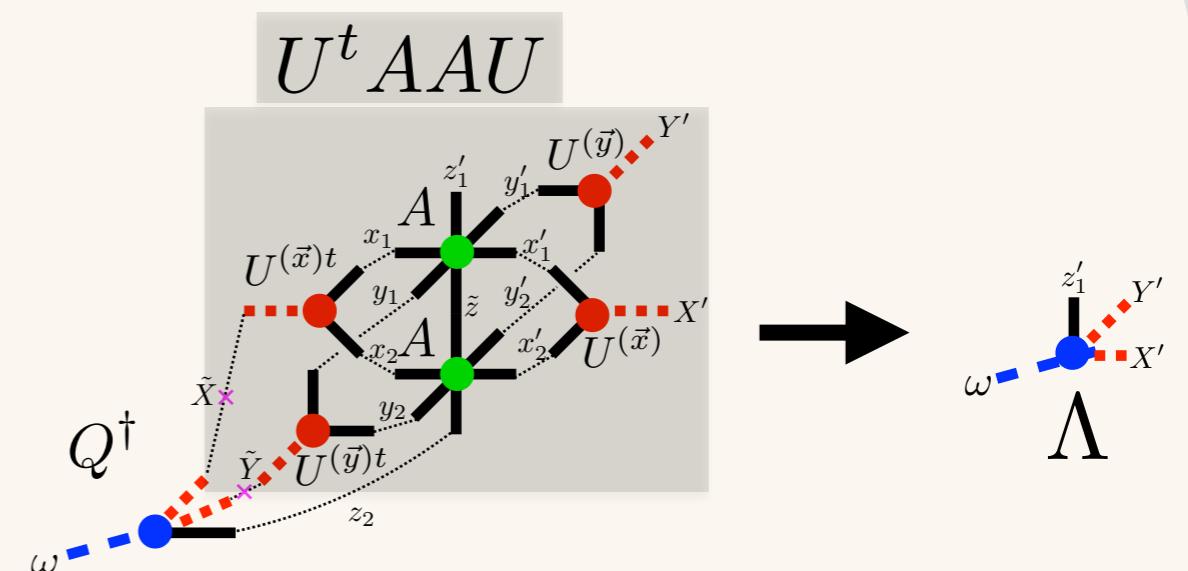
◇ Contraction with U, Q

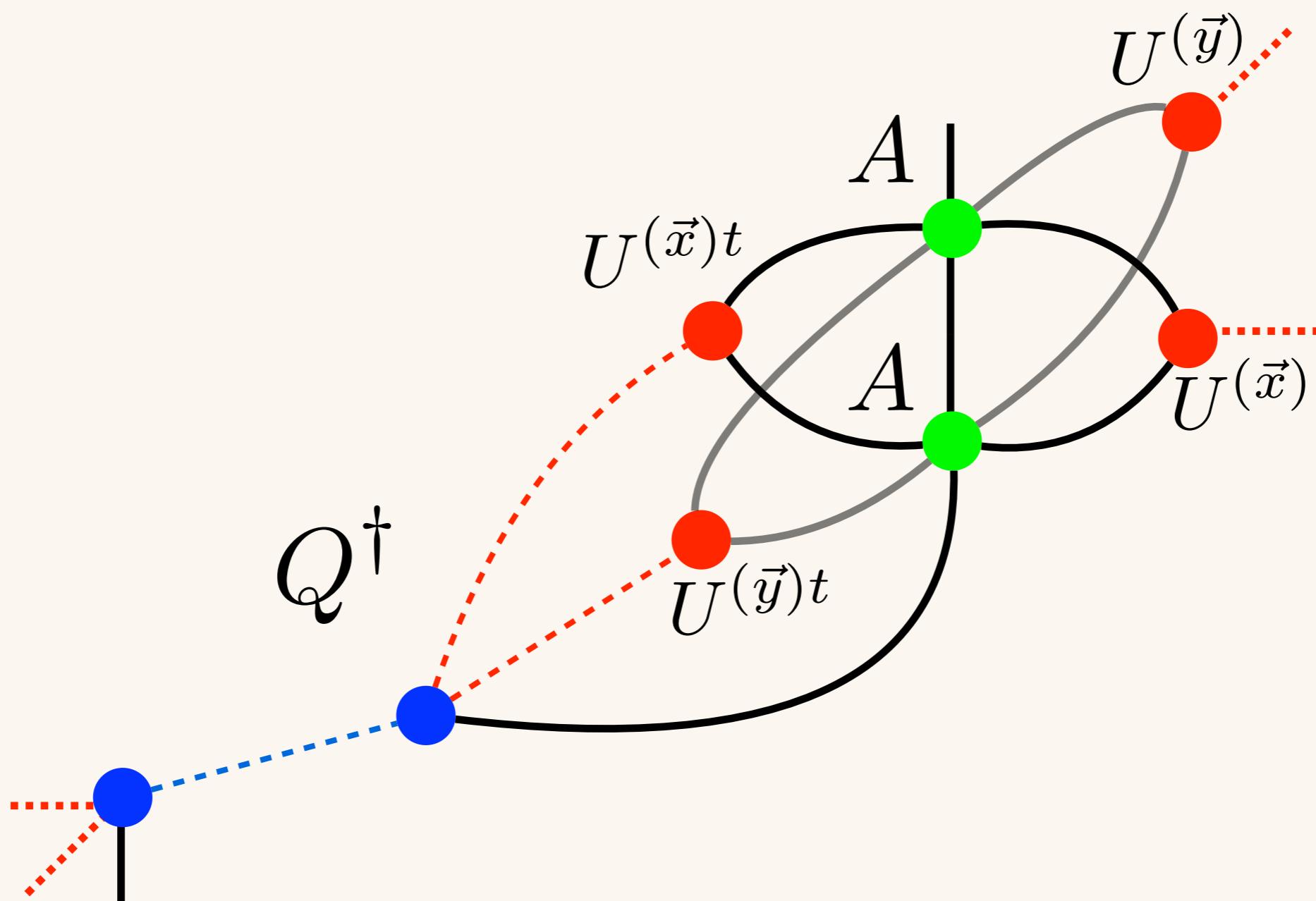
$$Q Q^\dagger U(\vec{y})^t U(\vec{x})^t A A U(\vec{x}) U(\vec{y}) \simeq A^{(\text{next})}$$

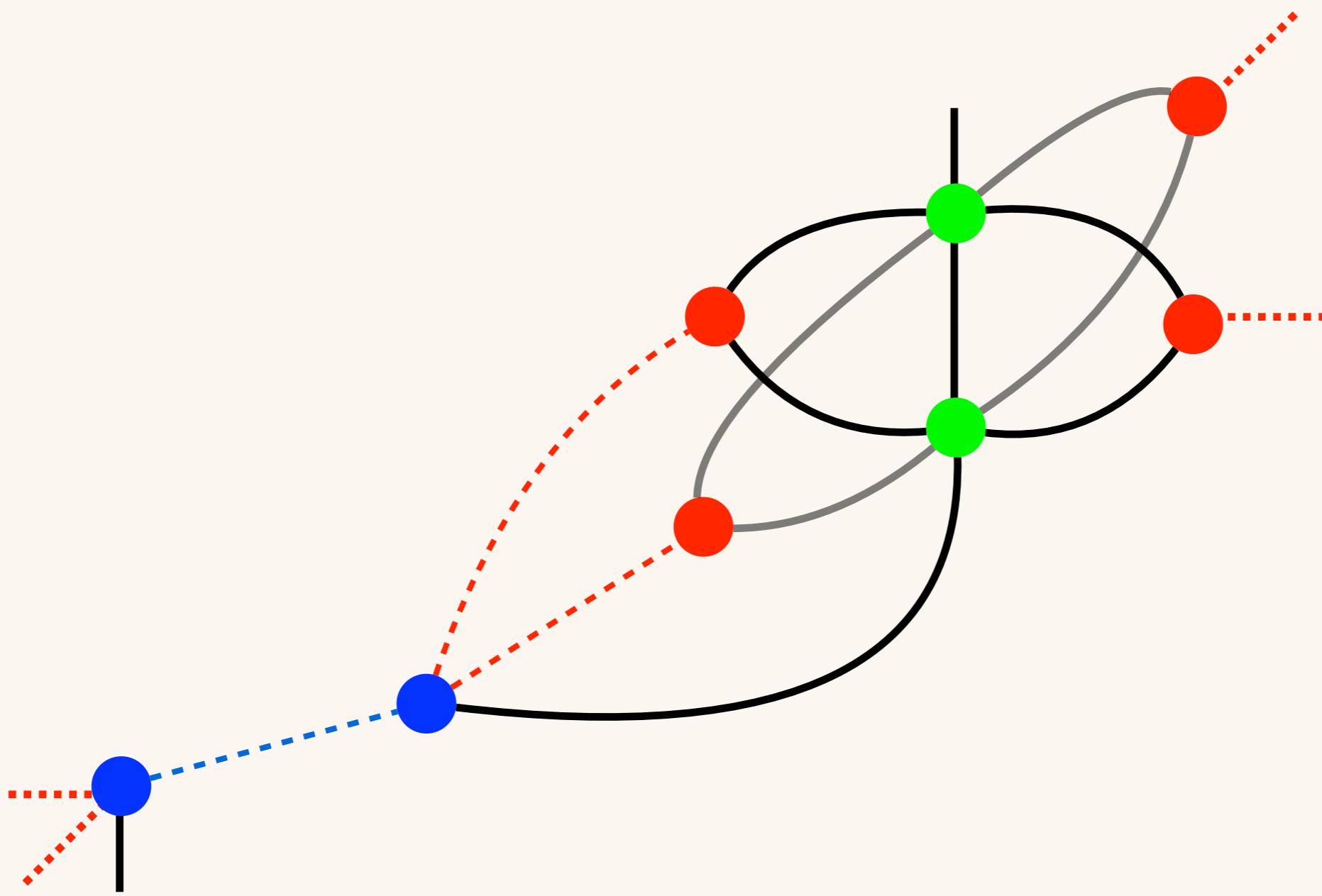
◇ Cost reduction

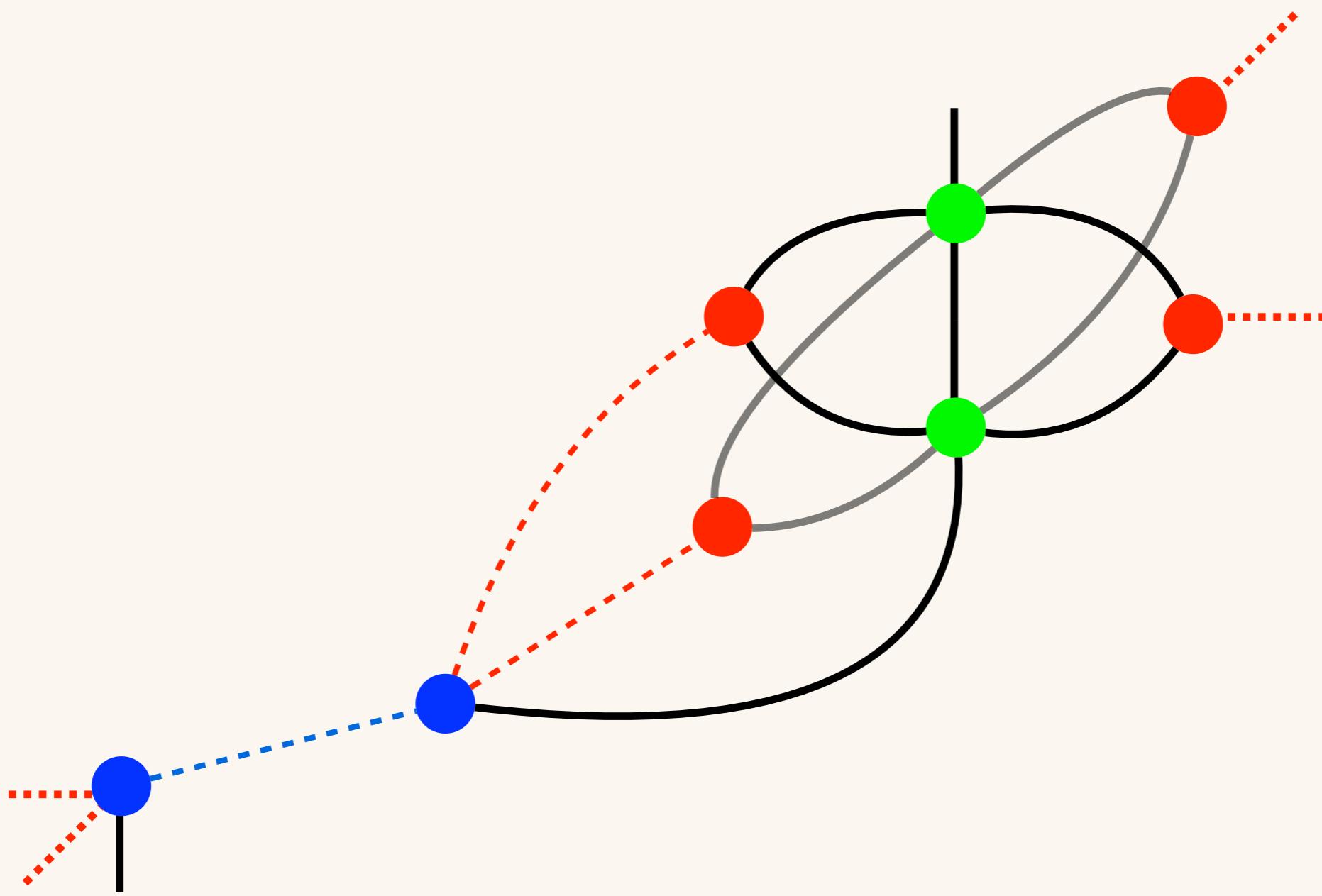
$$O(D^{4d-1}) \rightarrow O(D^{3d})$$

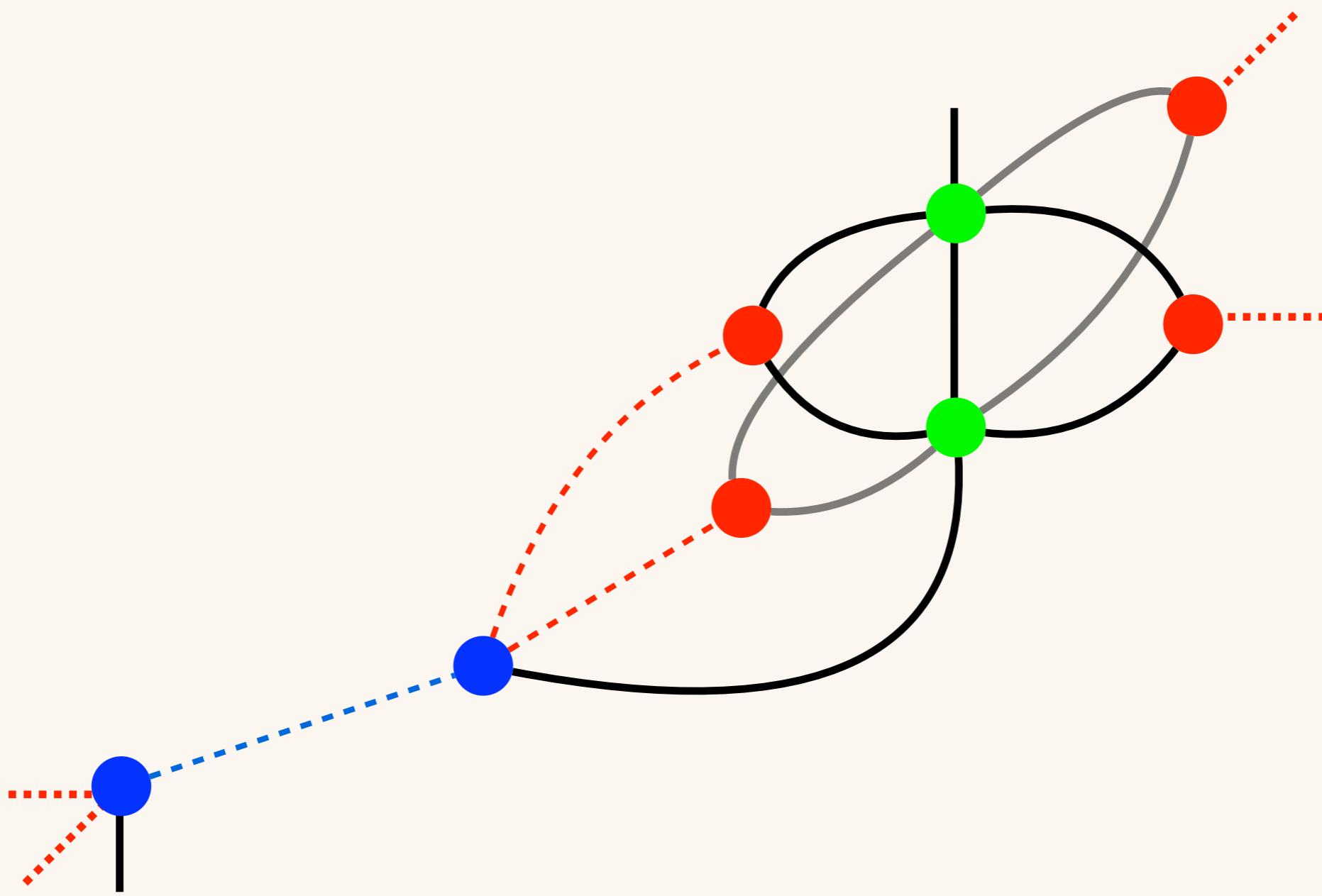
→ Further cost reduction?

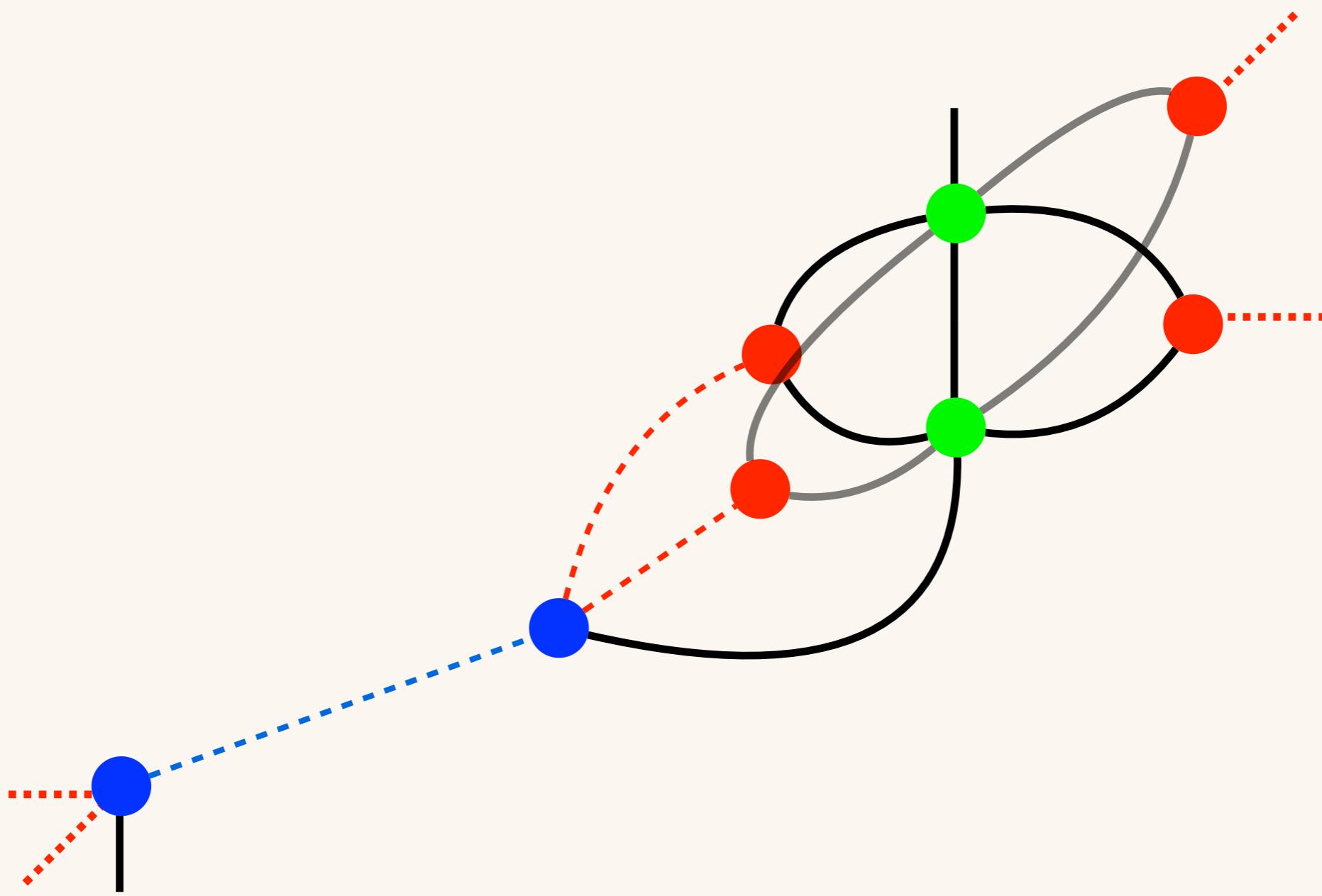


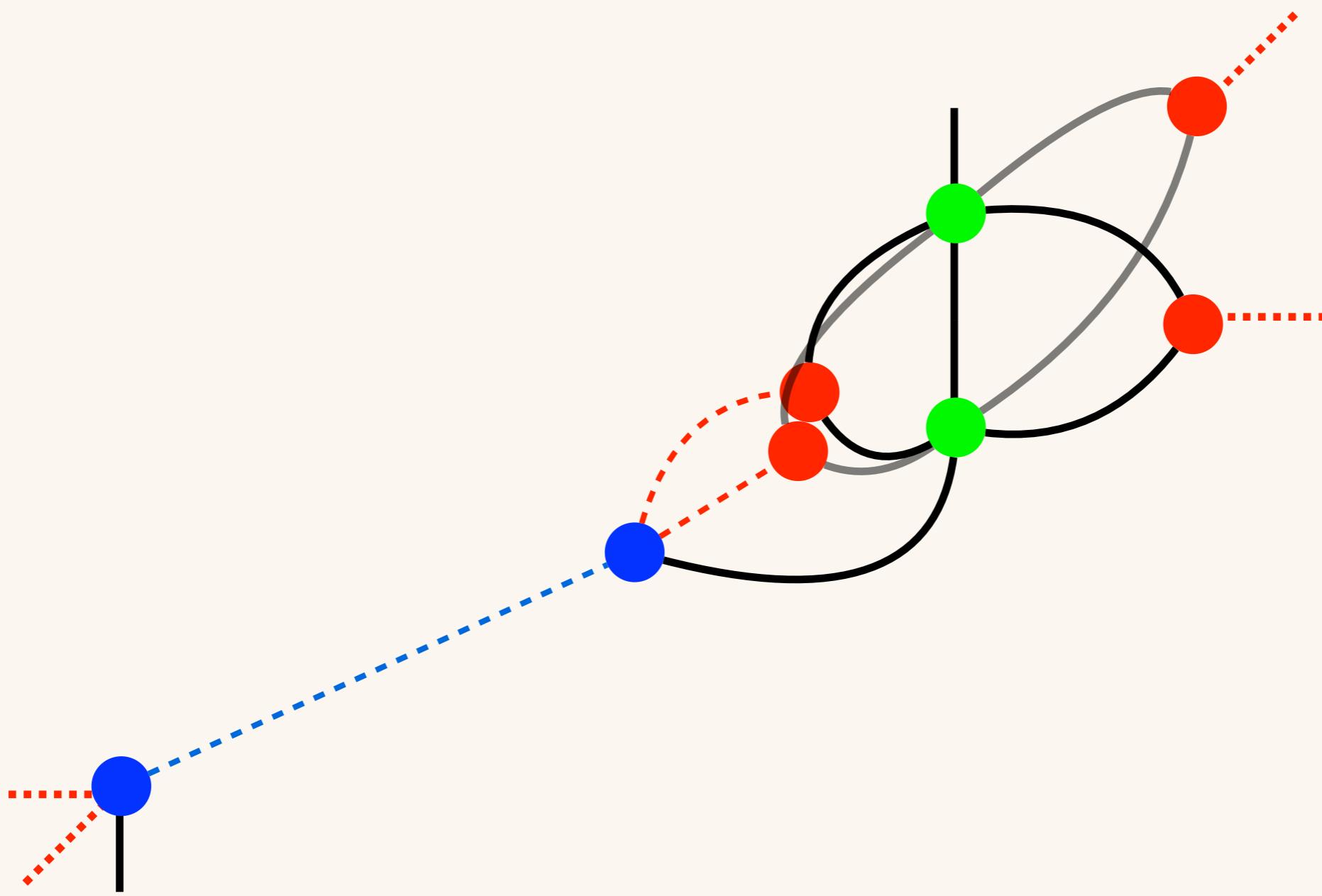


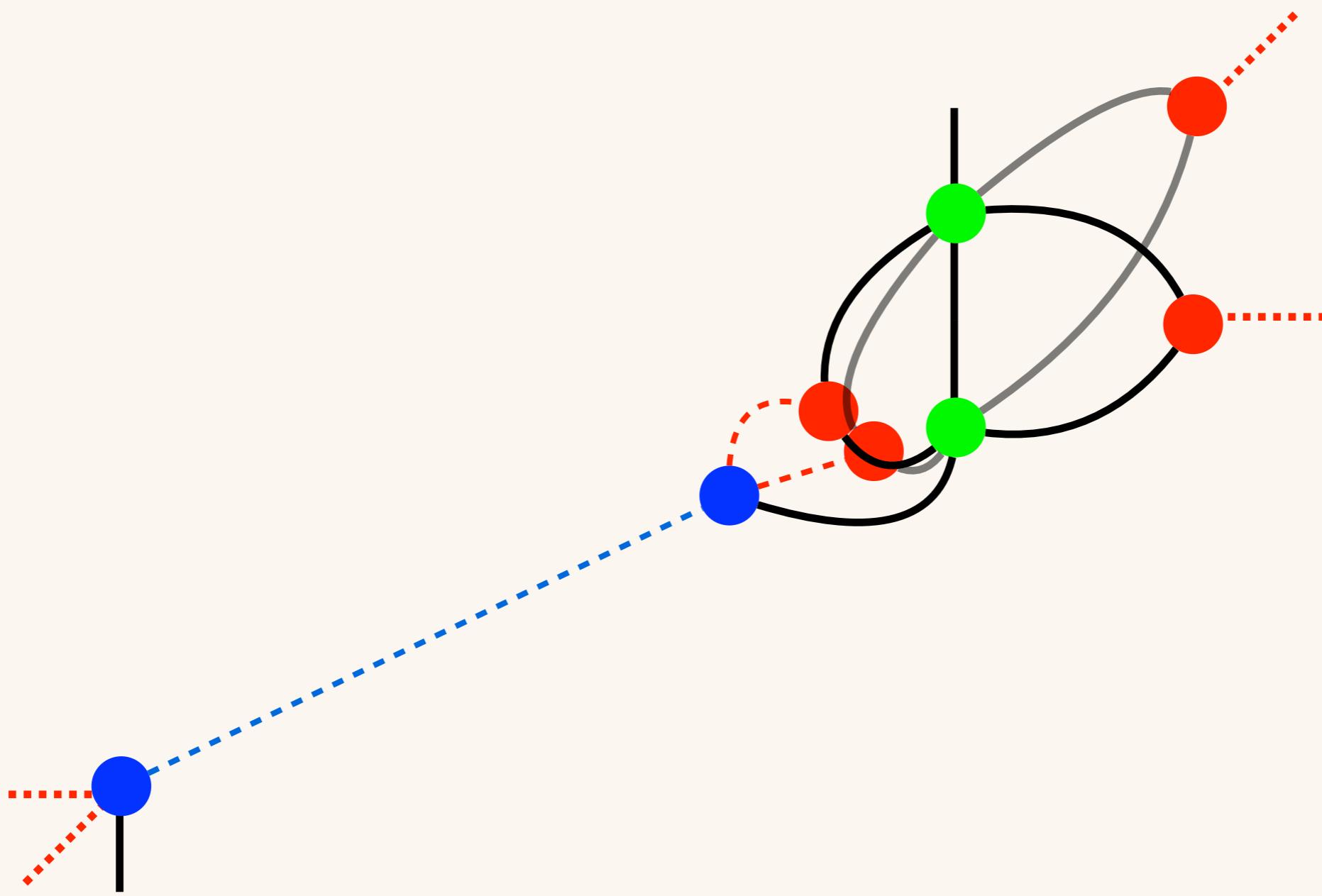


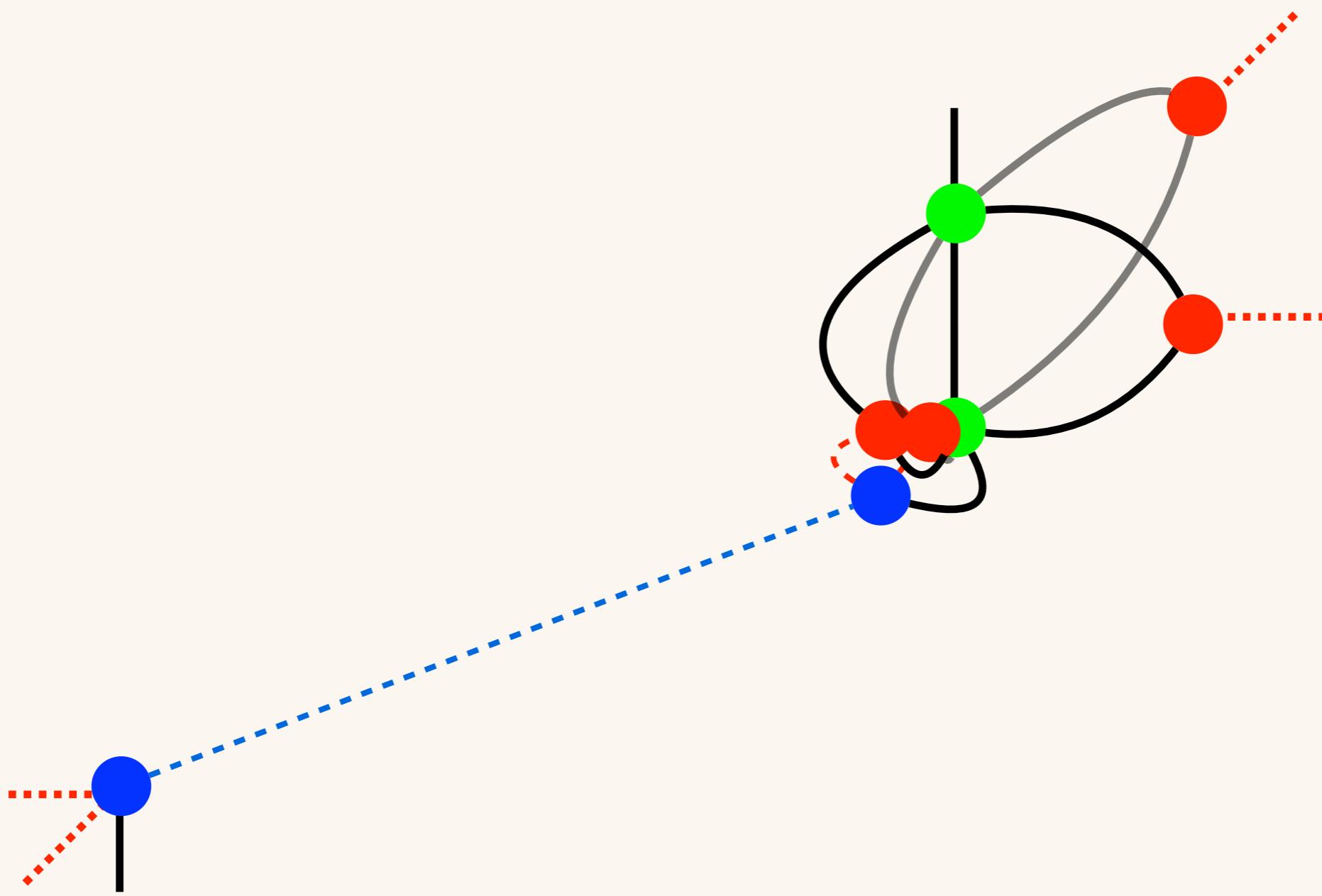


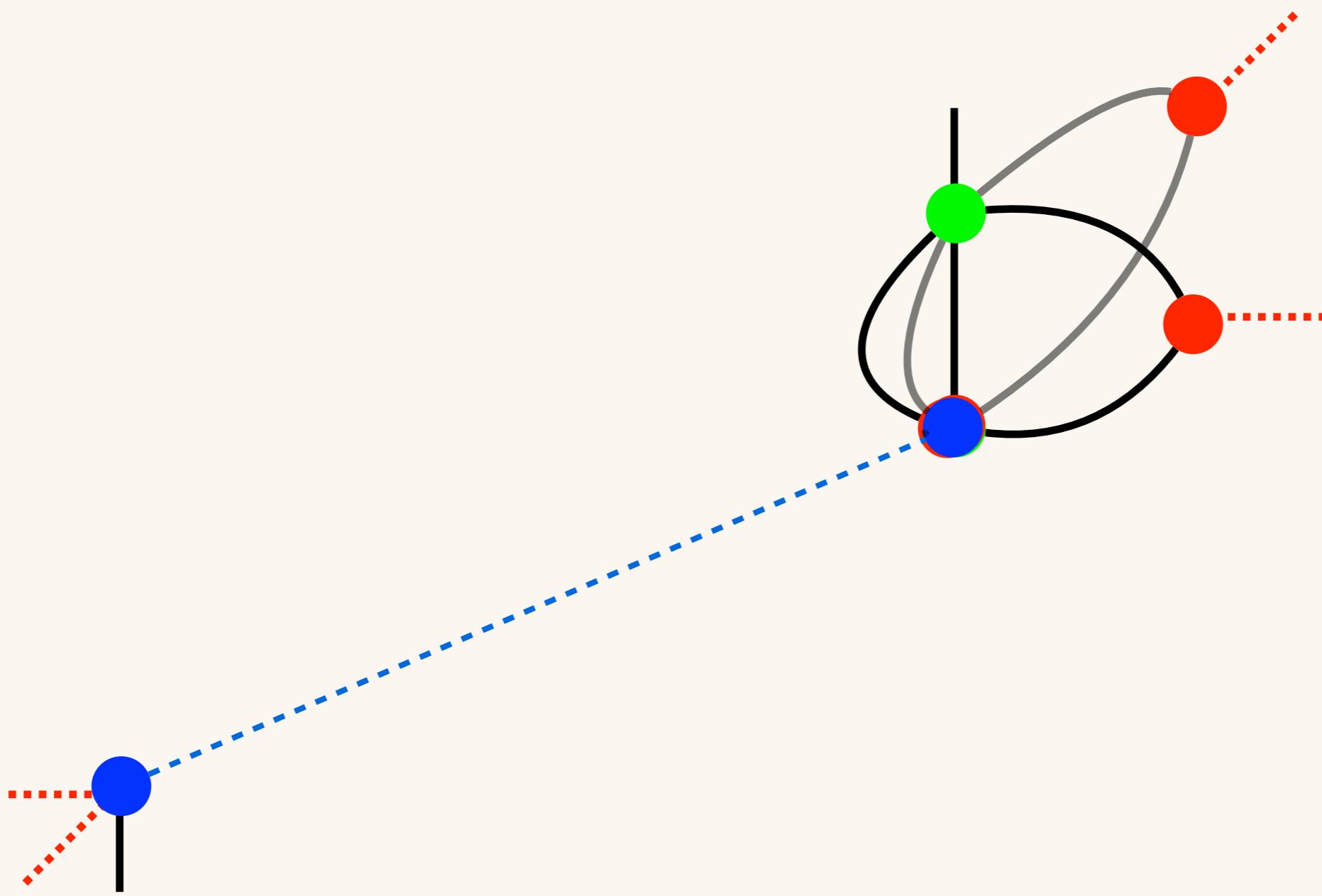


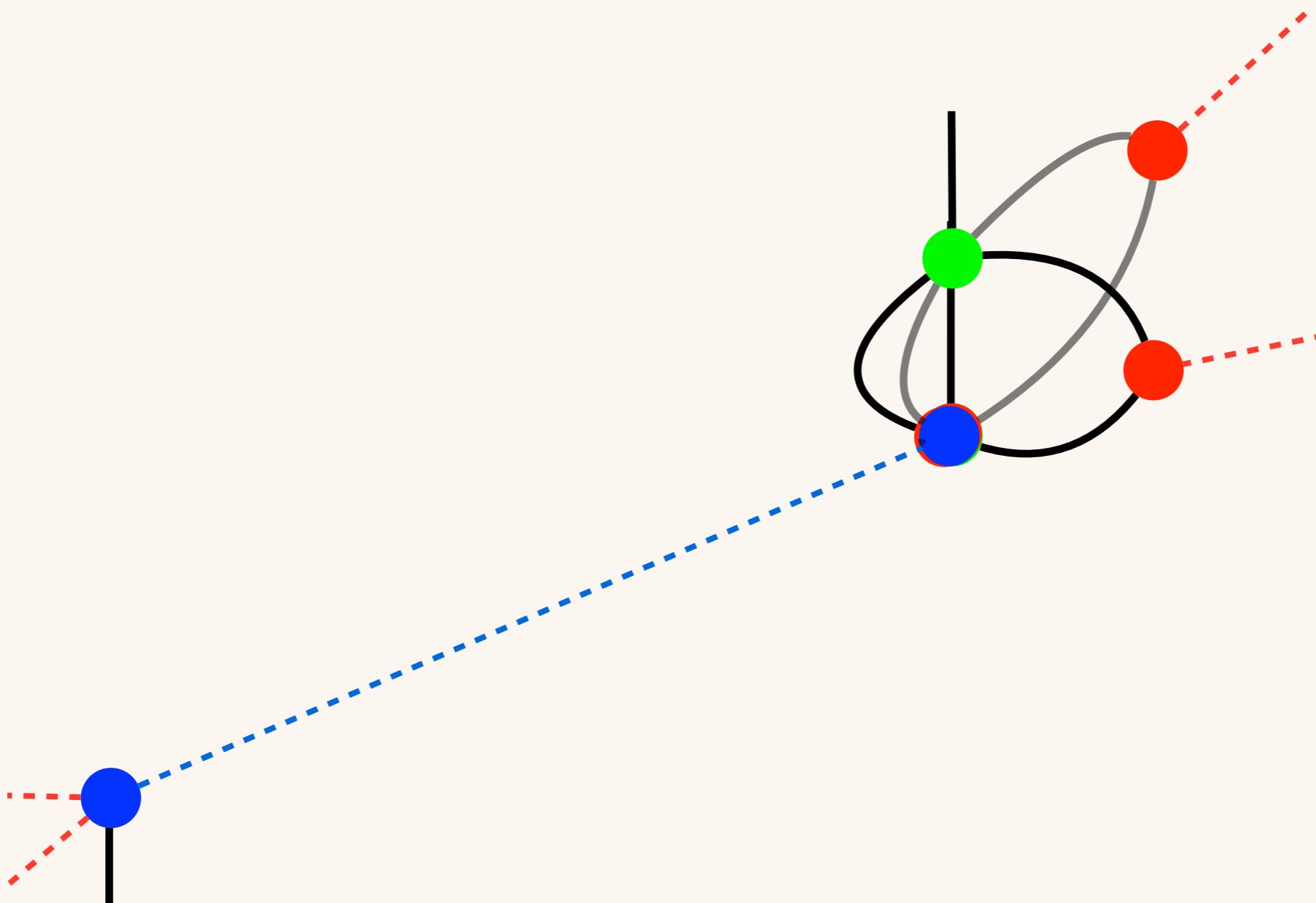


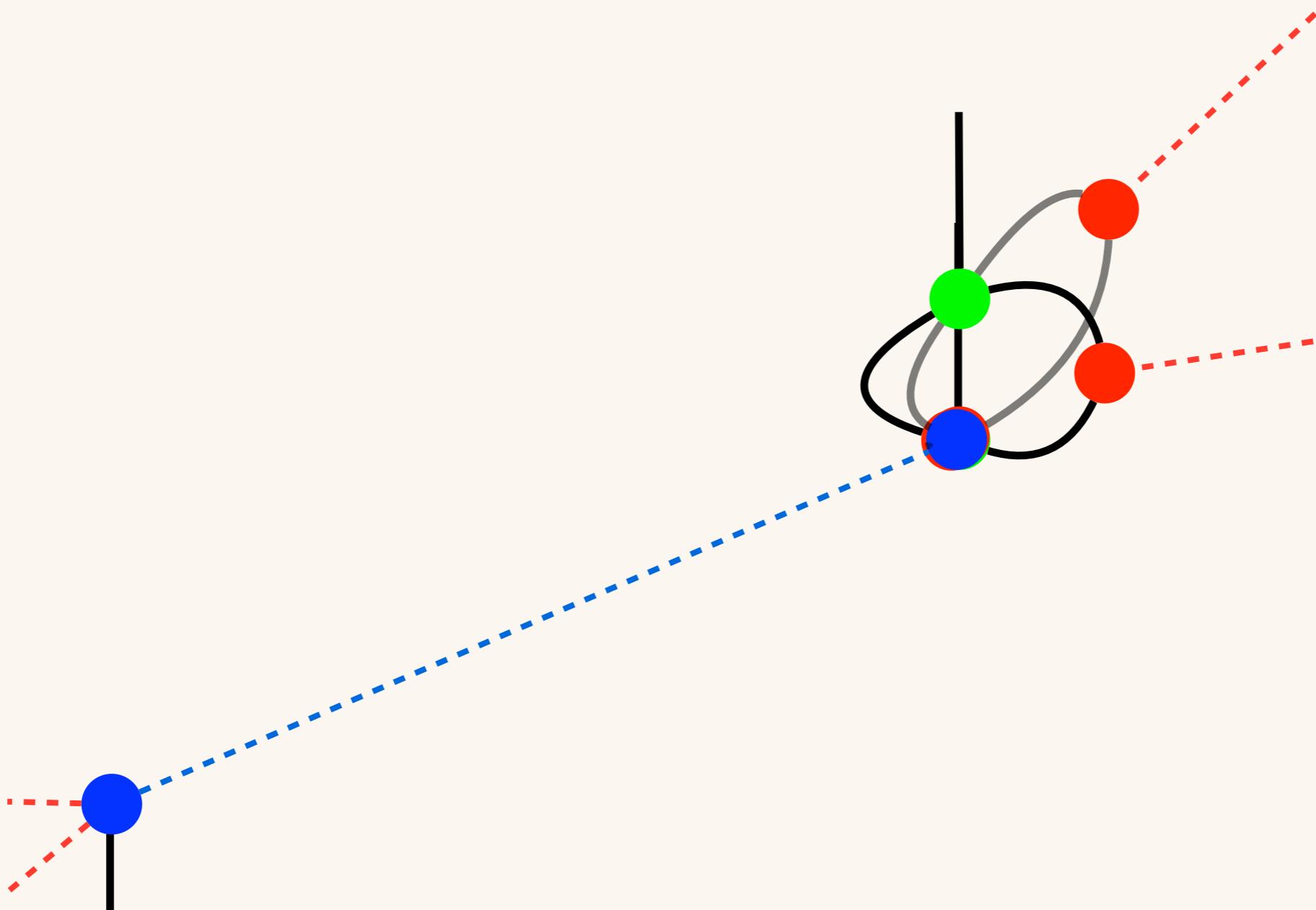


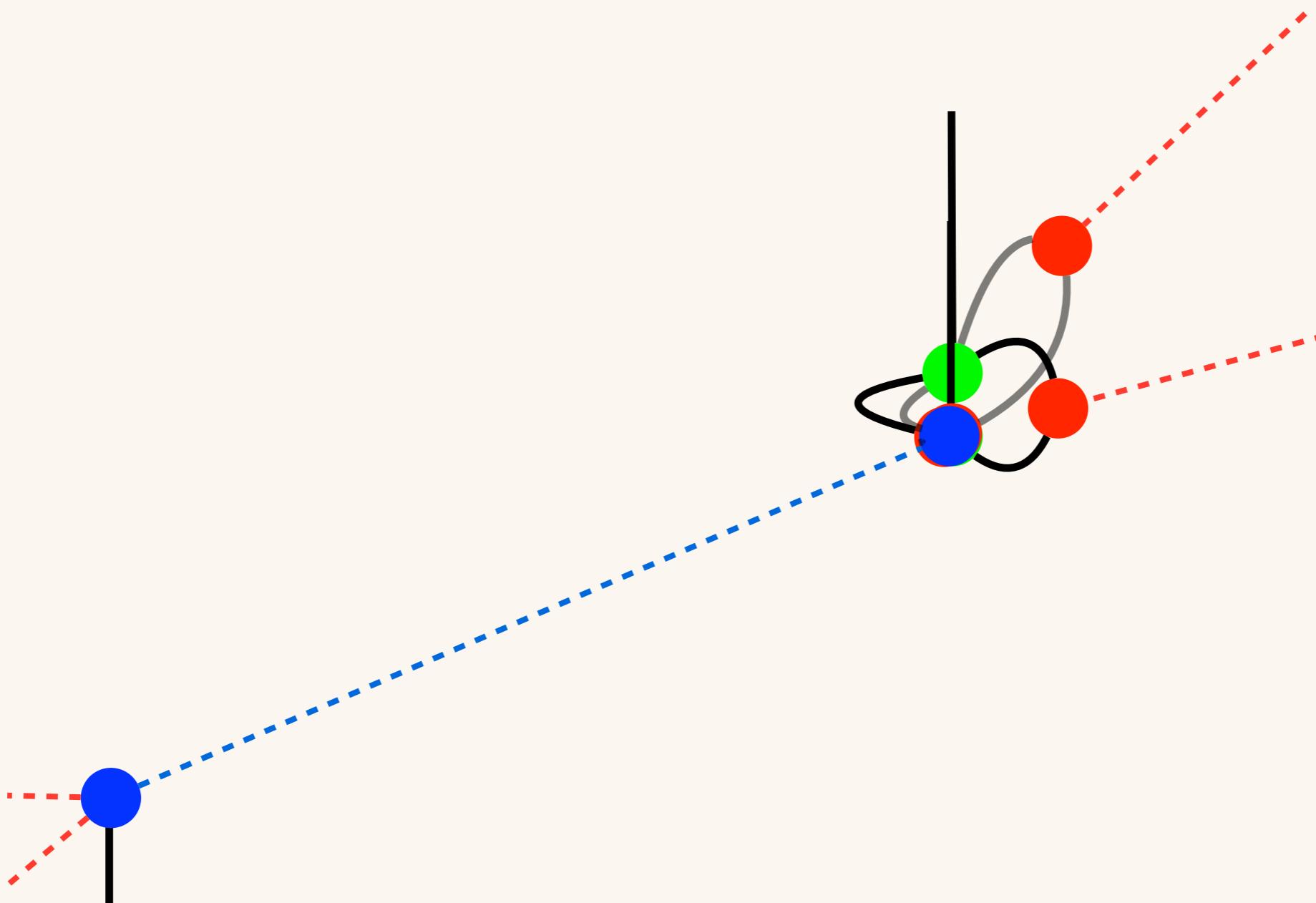


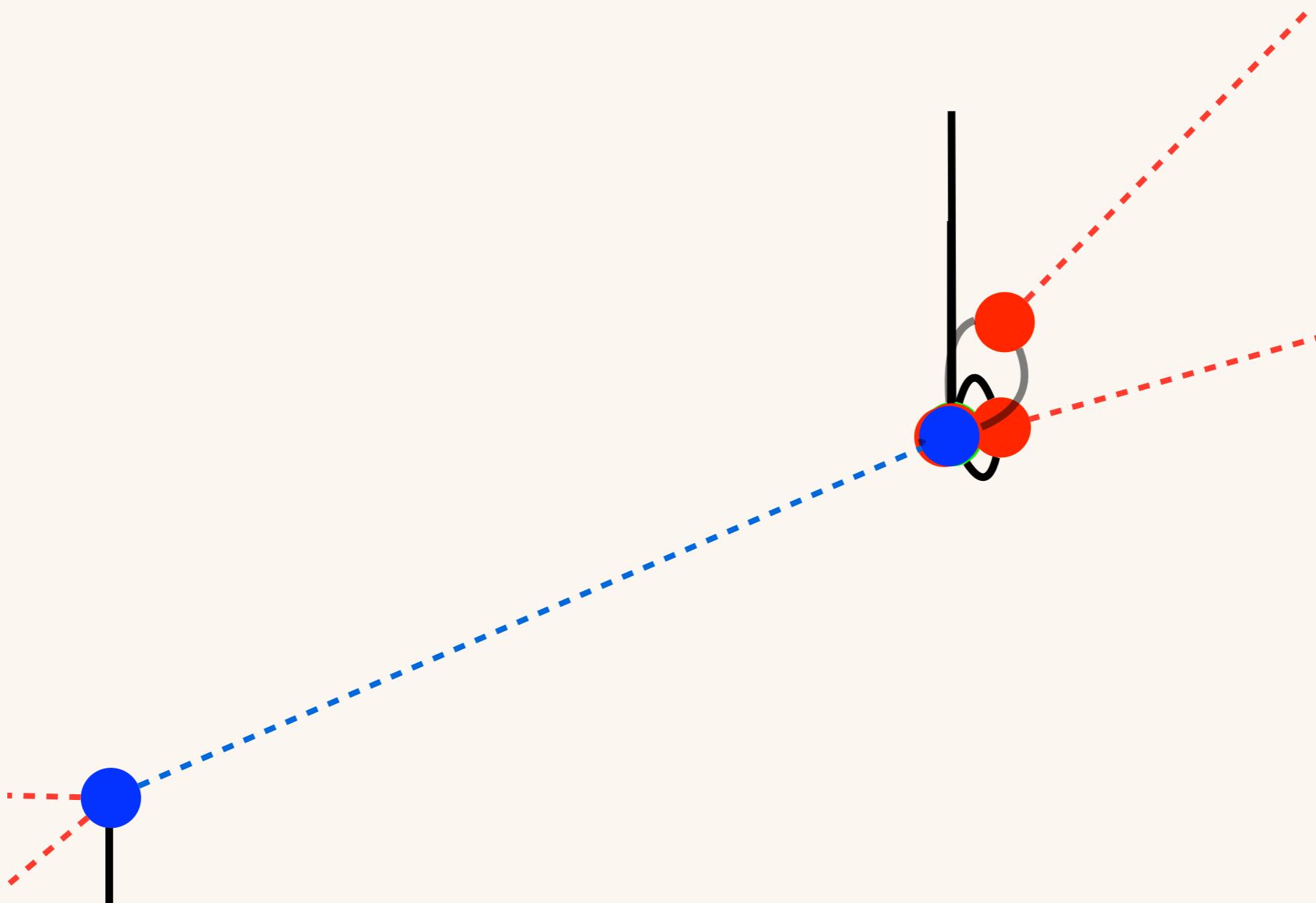




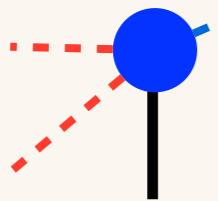




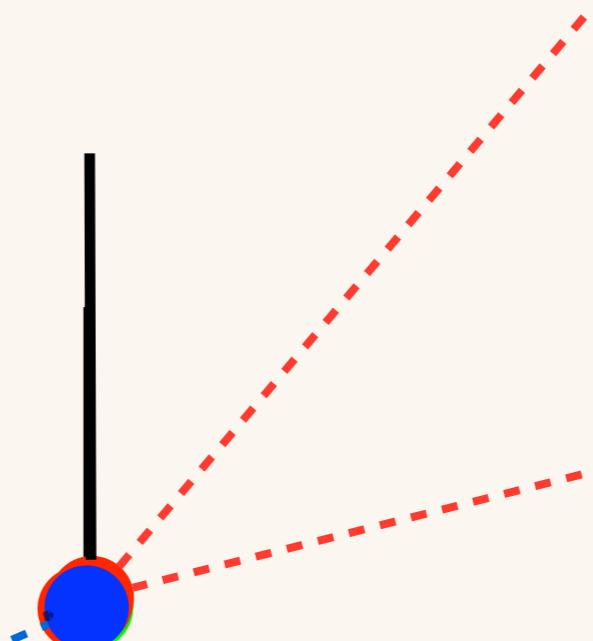




Q



Λ

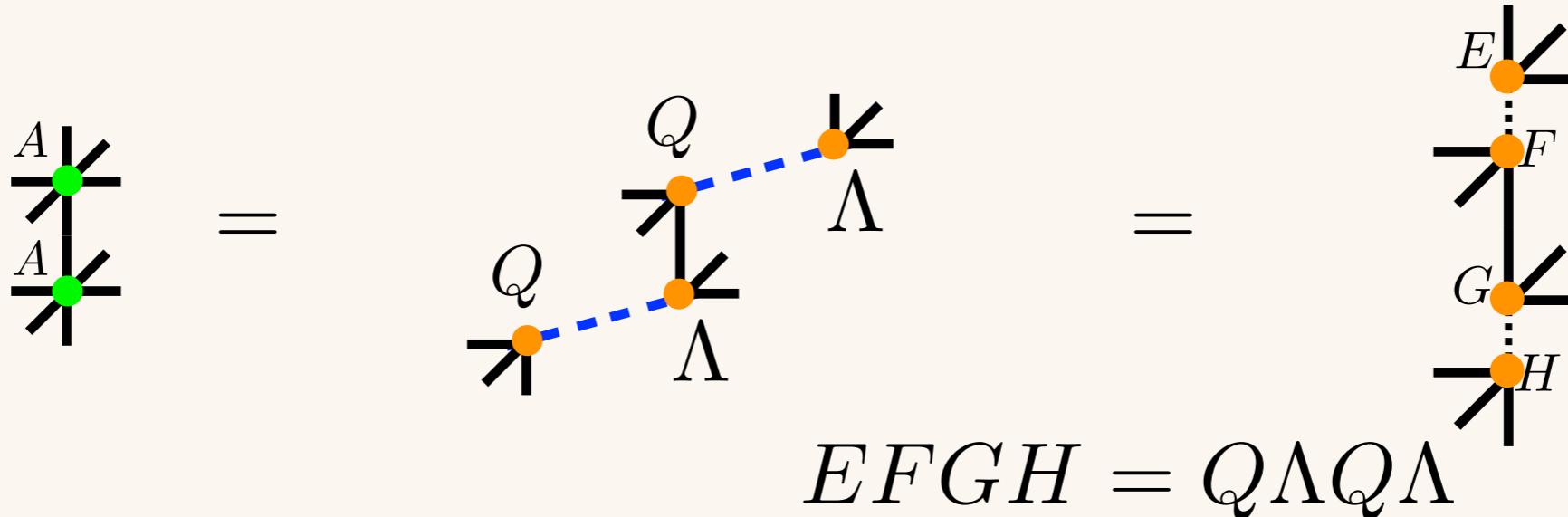


Cost and systematic error reduction

● 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 $d+1$

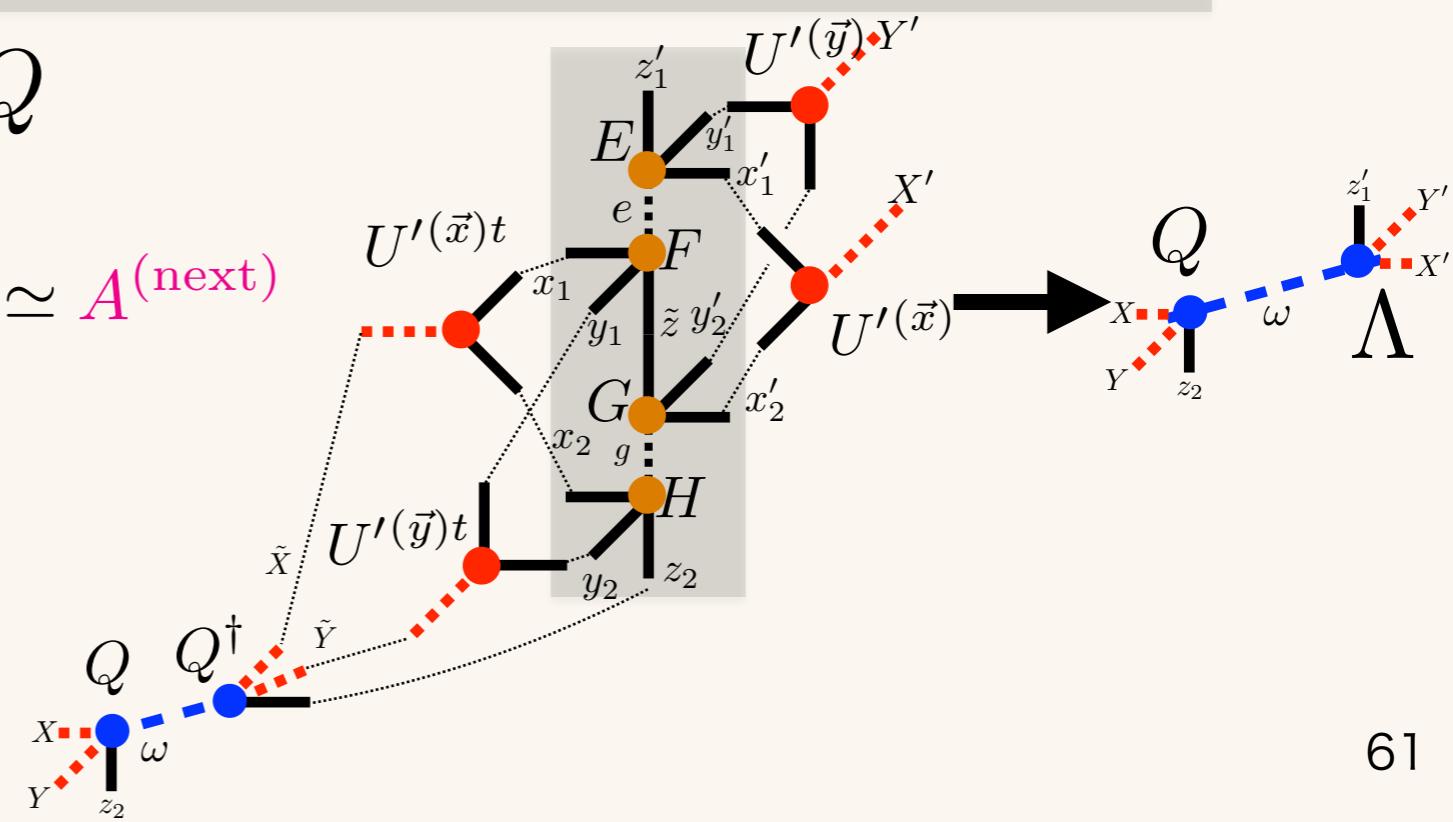
Q, Λ :Order $d+1$

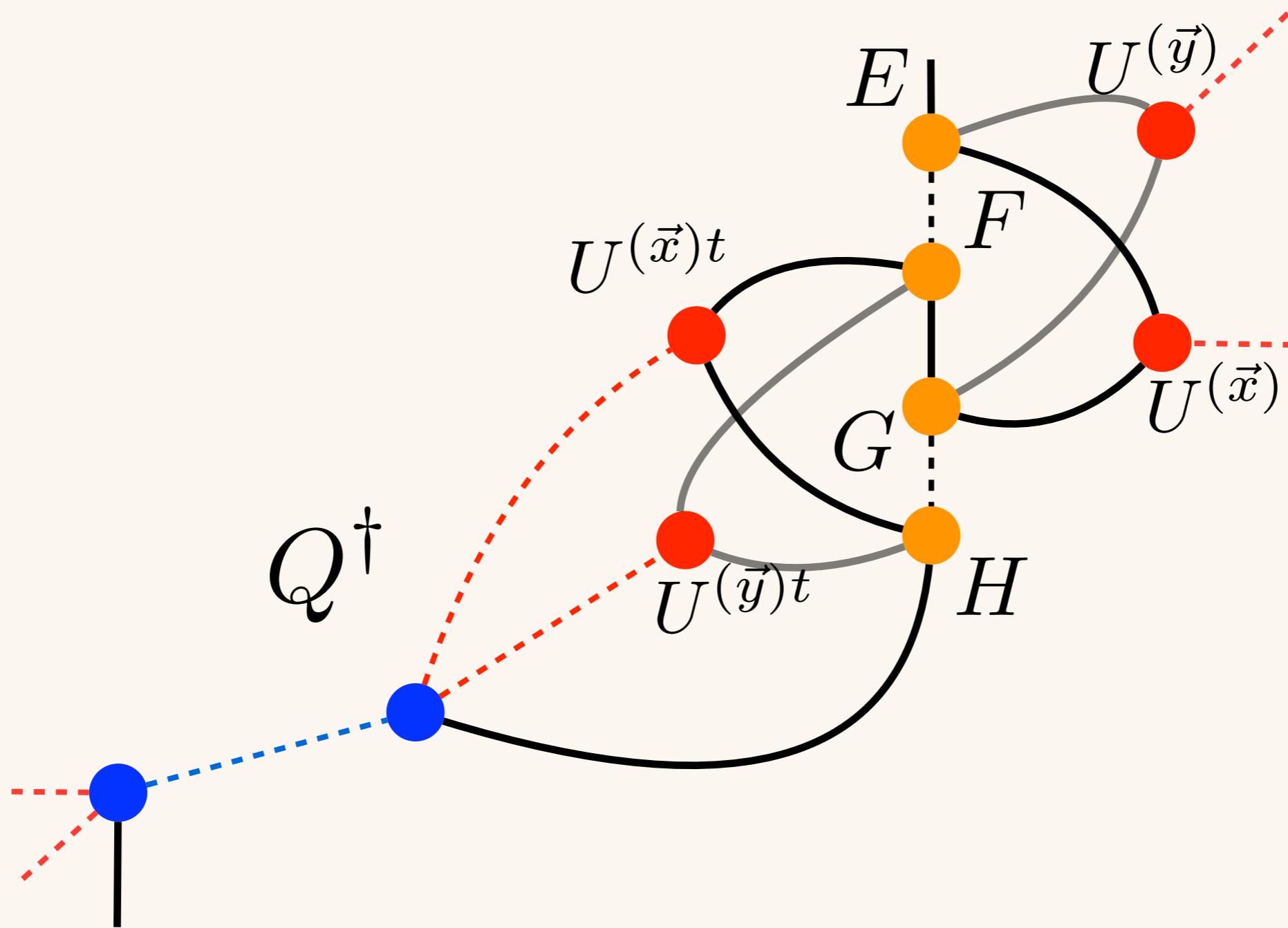
◇ Contraction with $EFGH, Q$

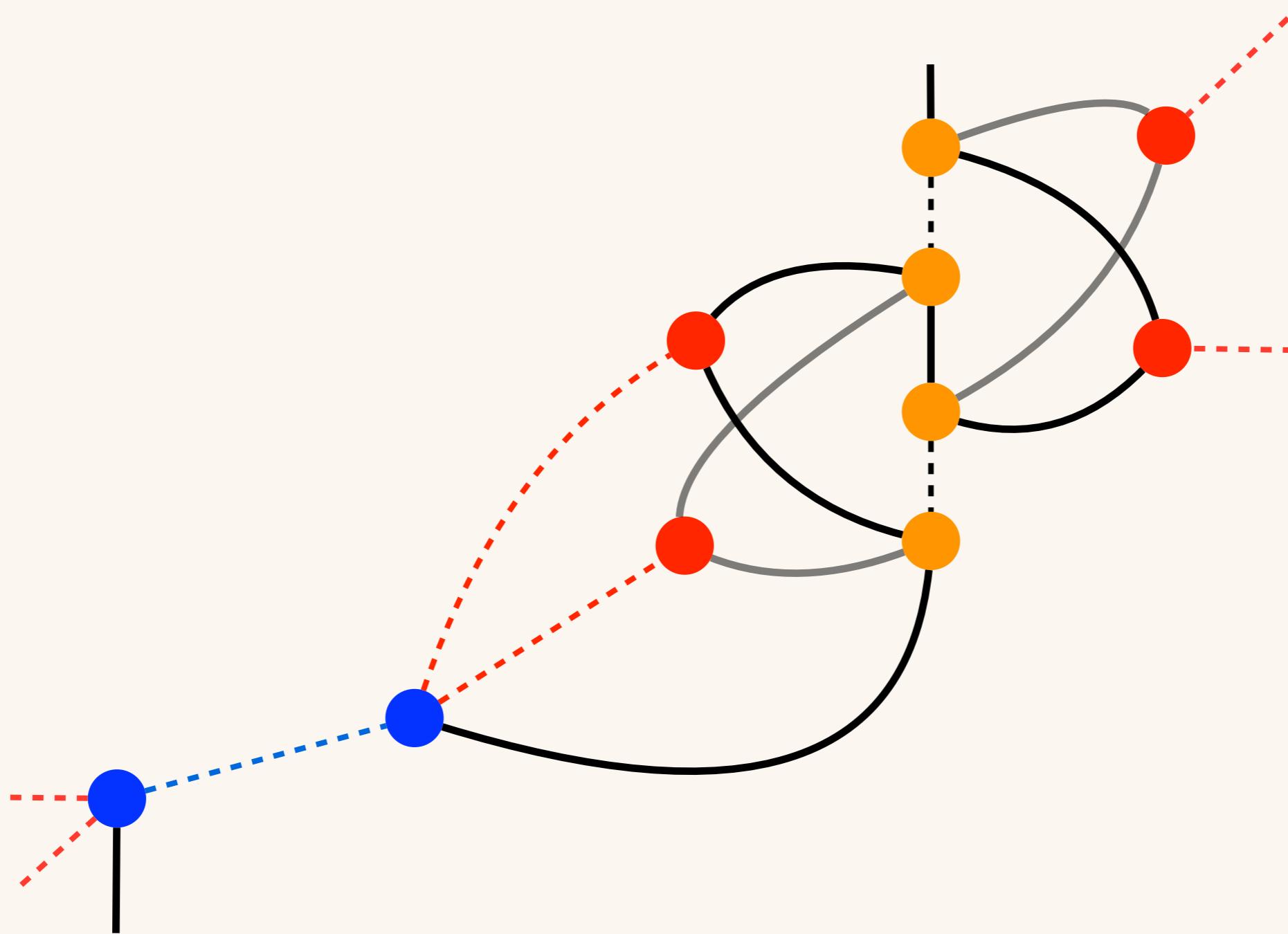
$$QQ^\dagger U'(\vec{y})^t U'(\vec{x})^t EFGH U'(\vec{x}) U'(\vec{y}) = Q\Lambda \simeq A^{(\text{next})}$$

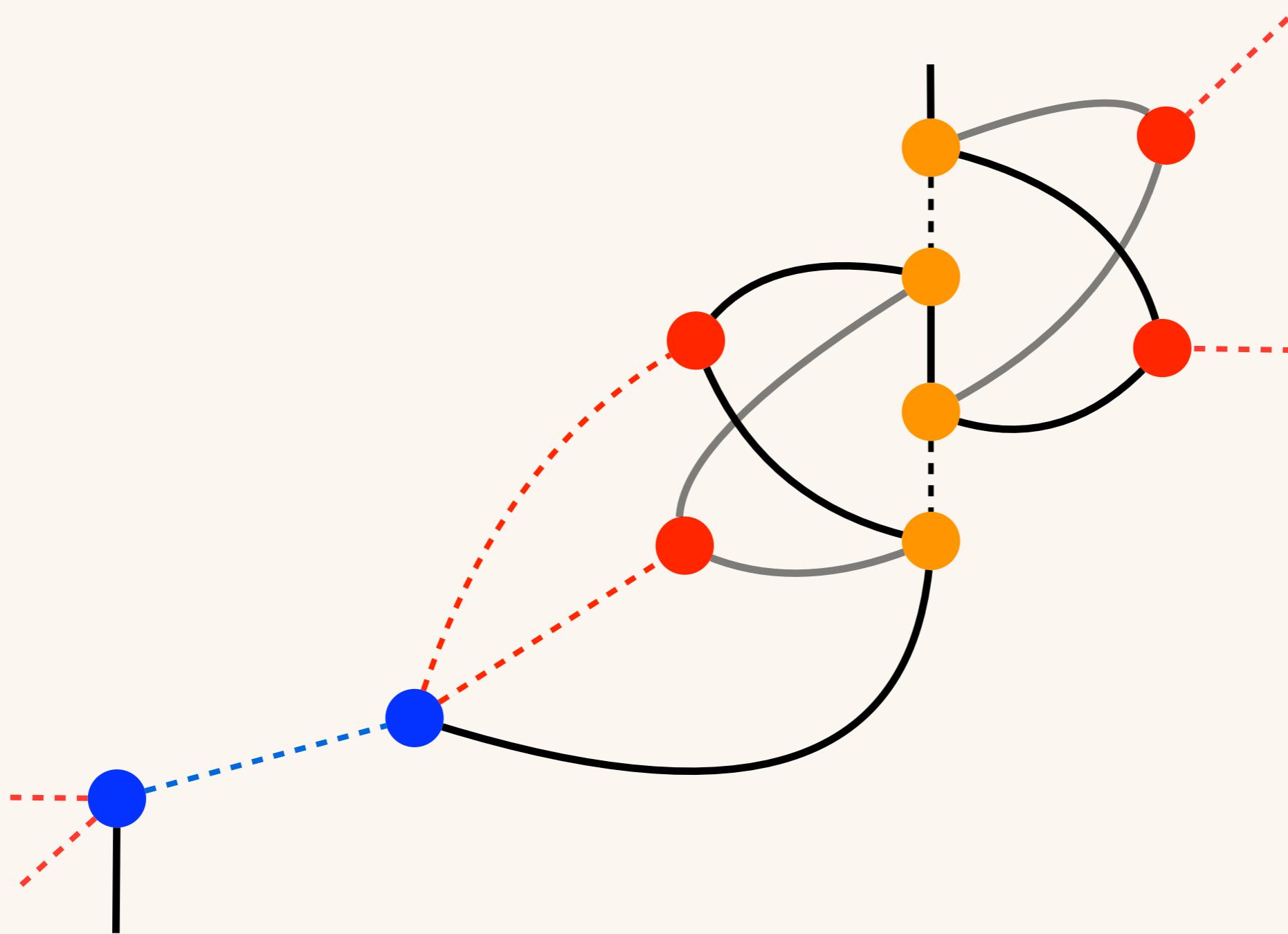
◇ Cost reduction

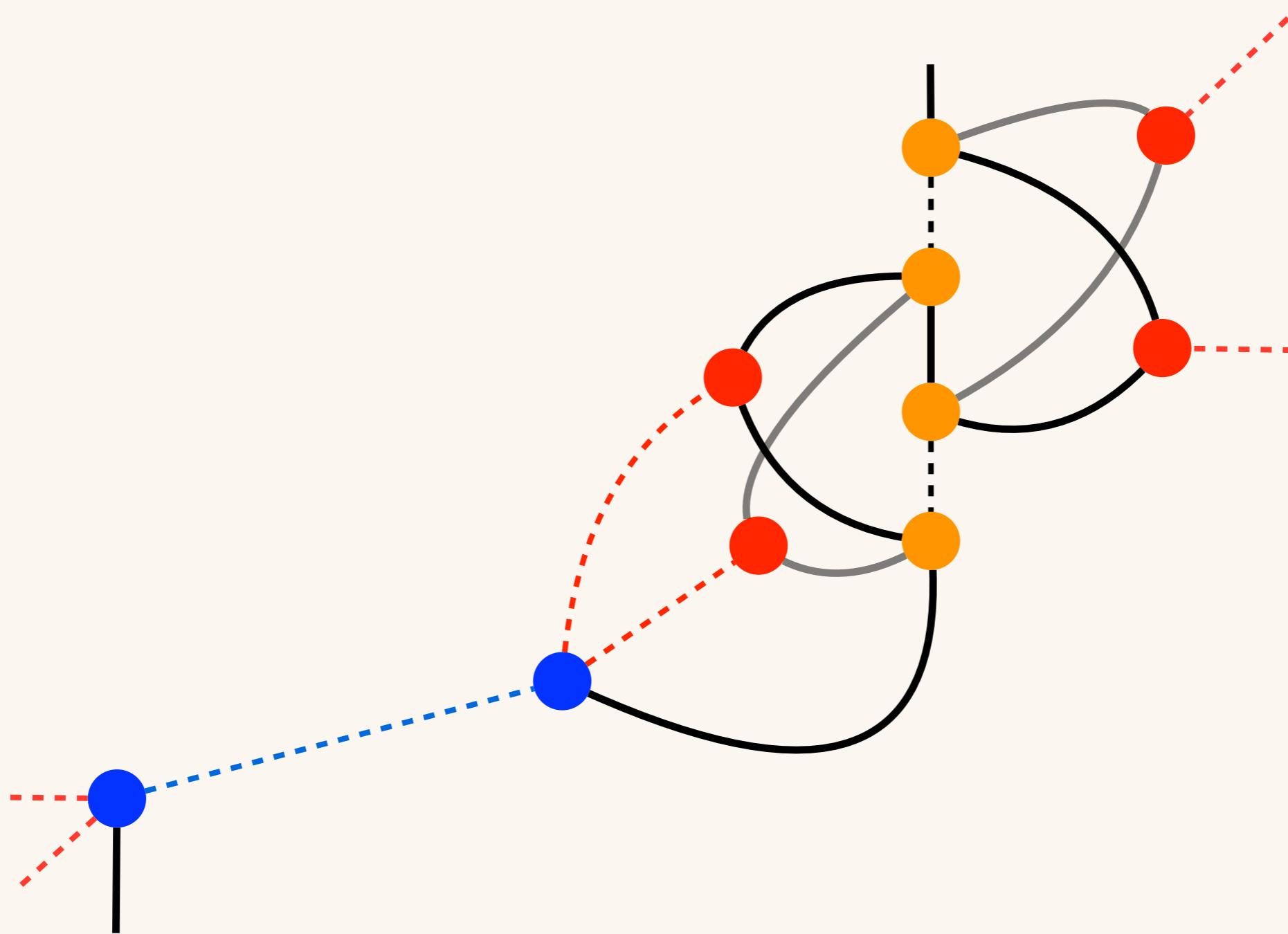
$$O(D^{3d}) \rightarrow O(D^{2d+1})$$

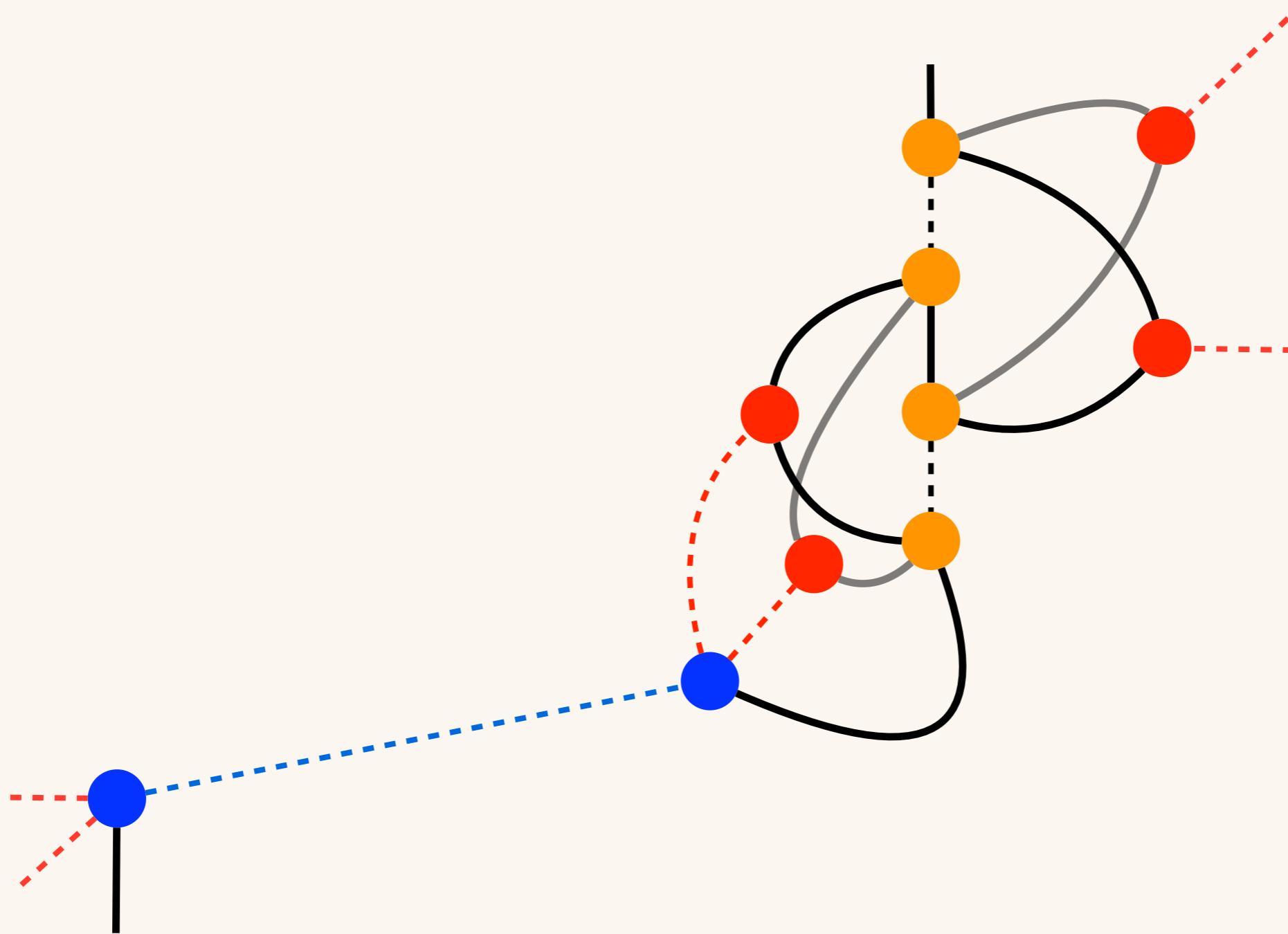


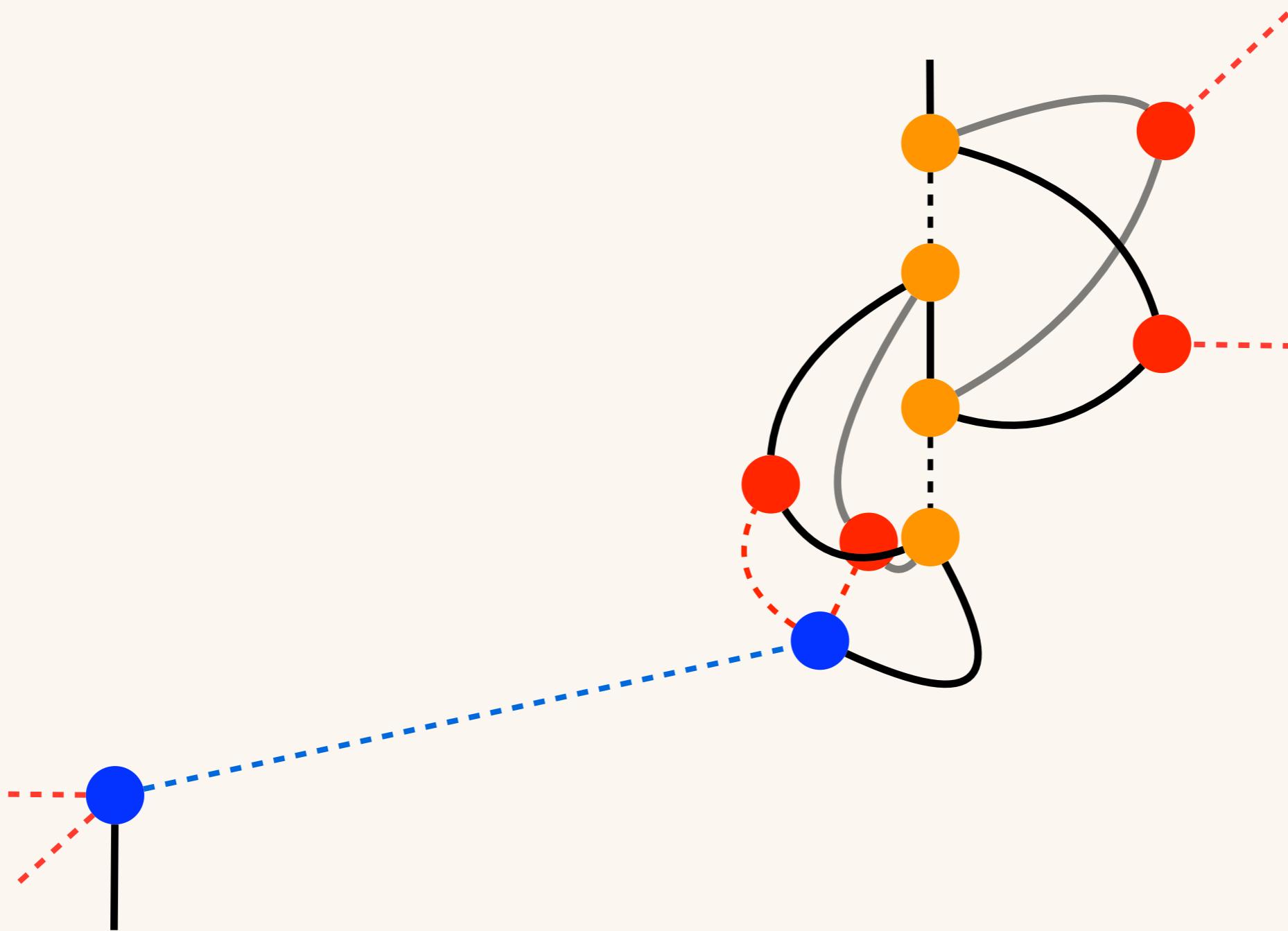


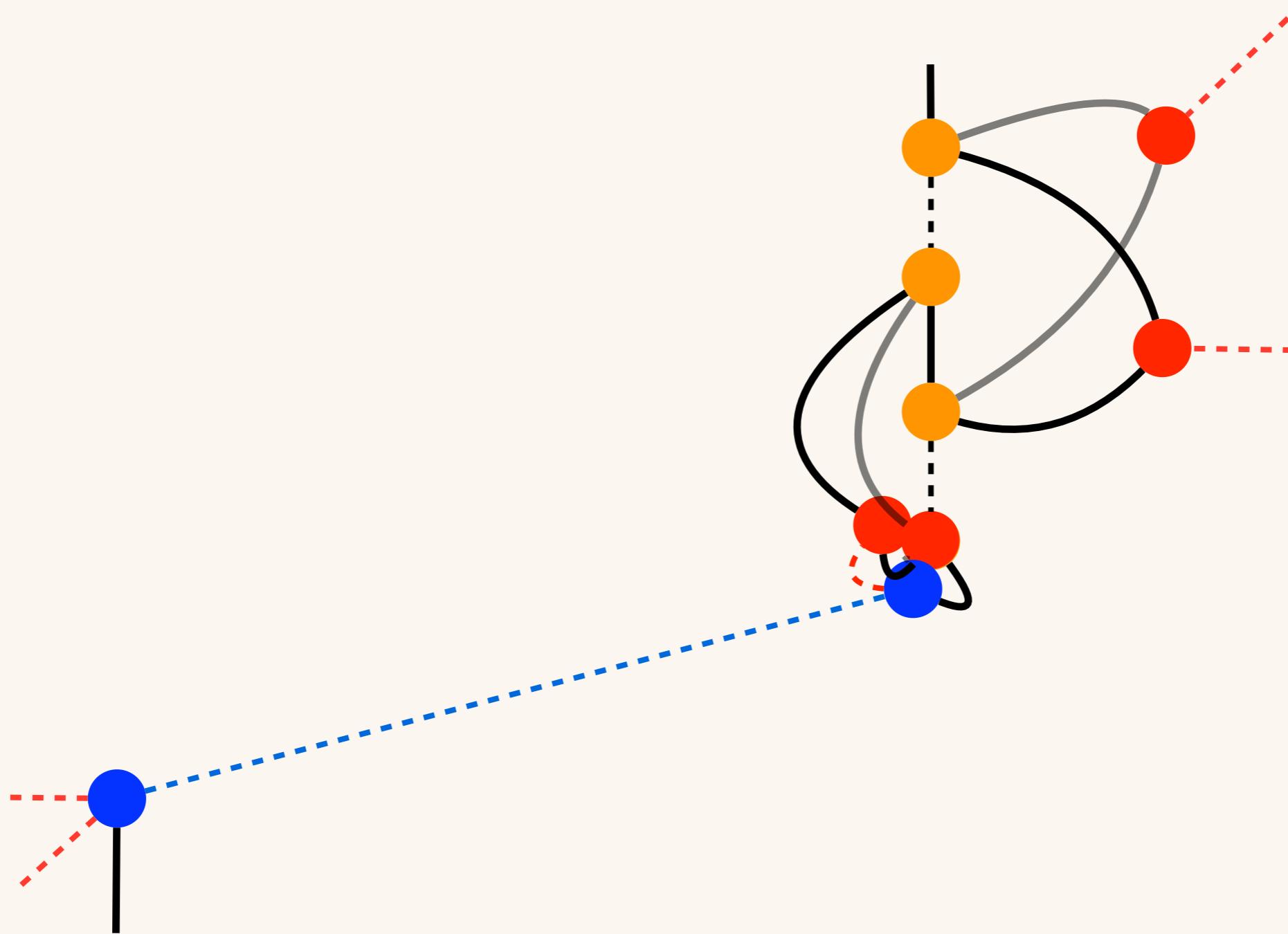


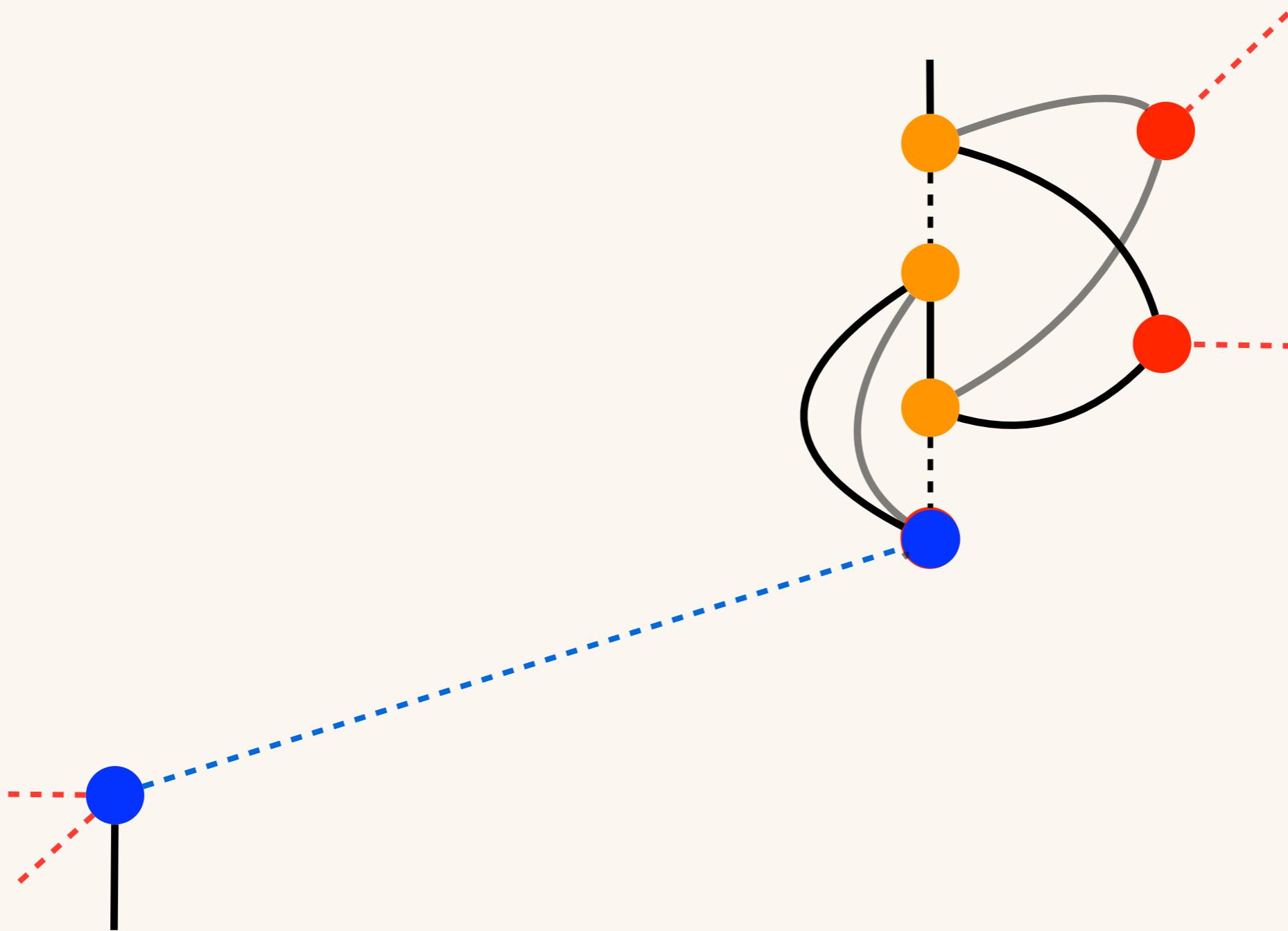


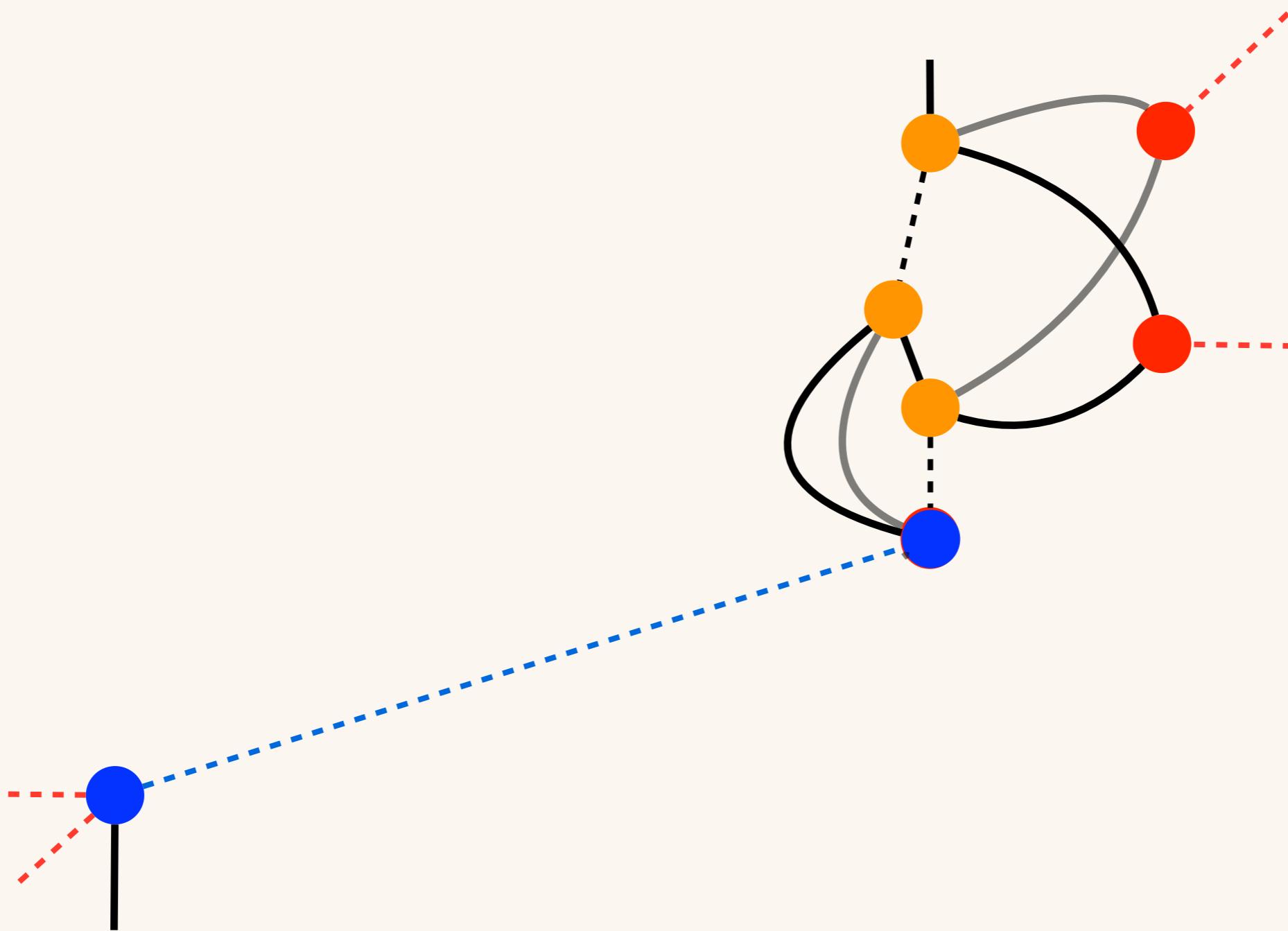


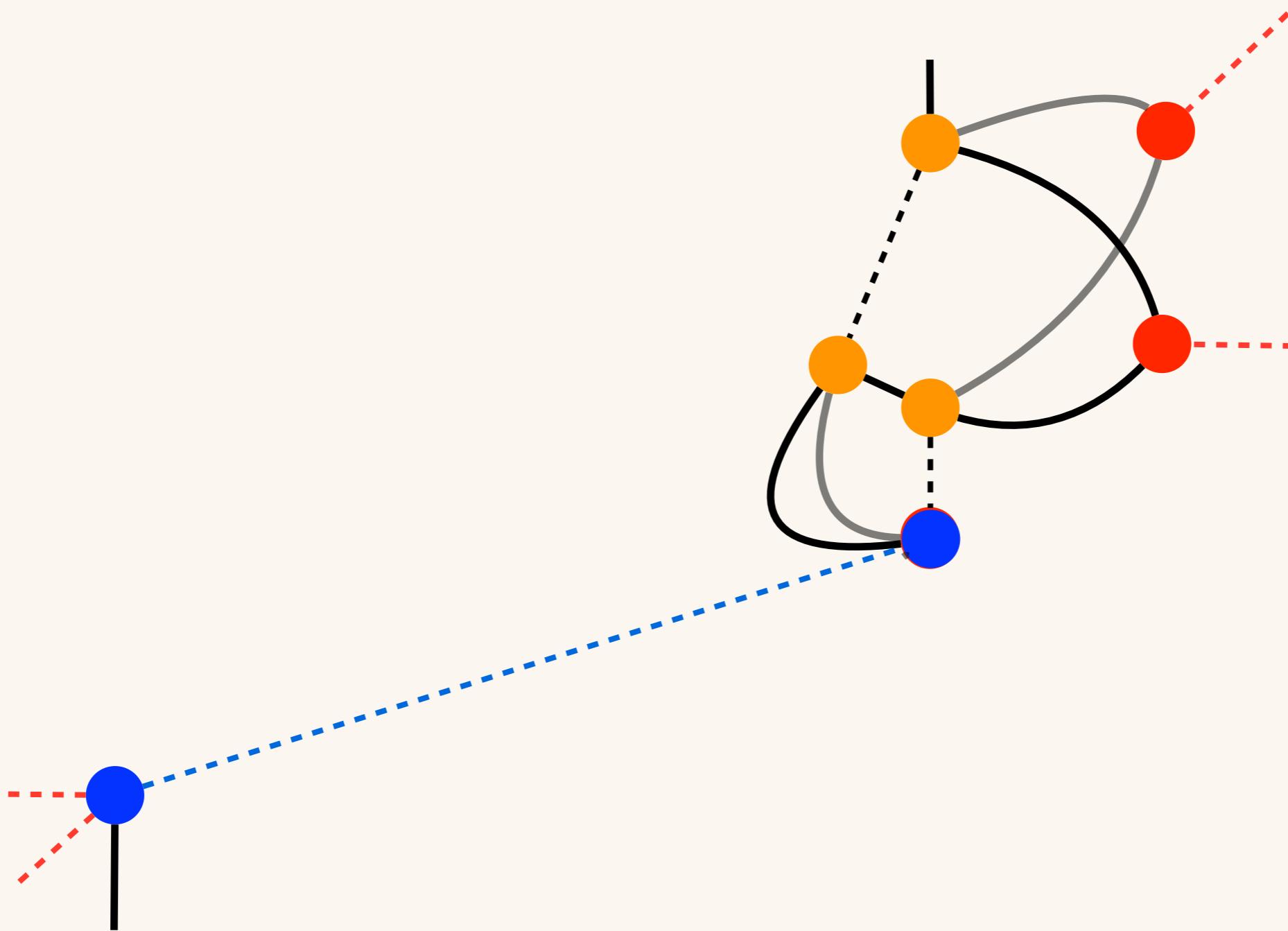


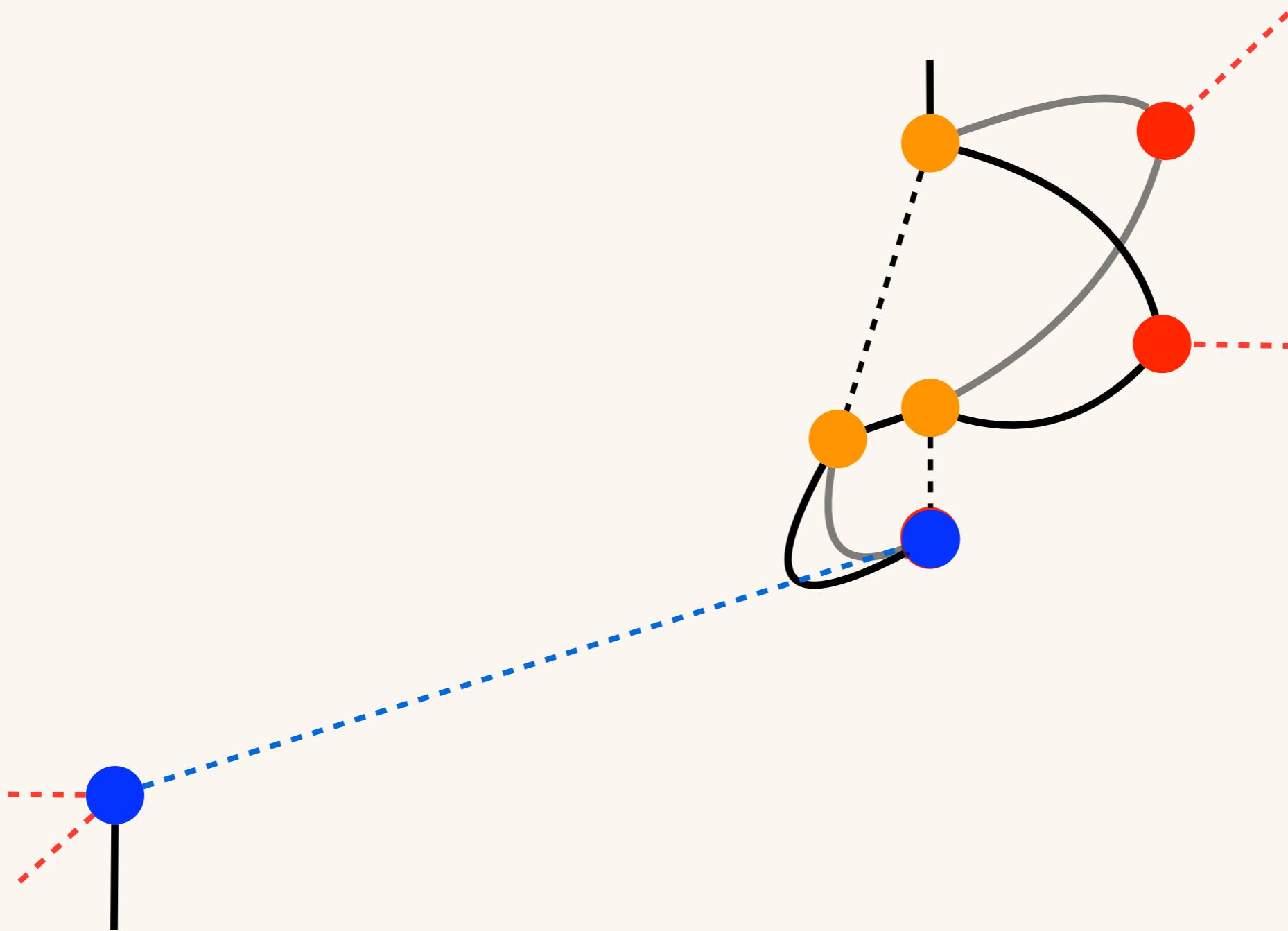


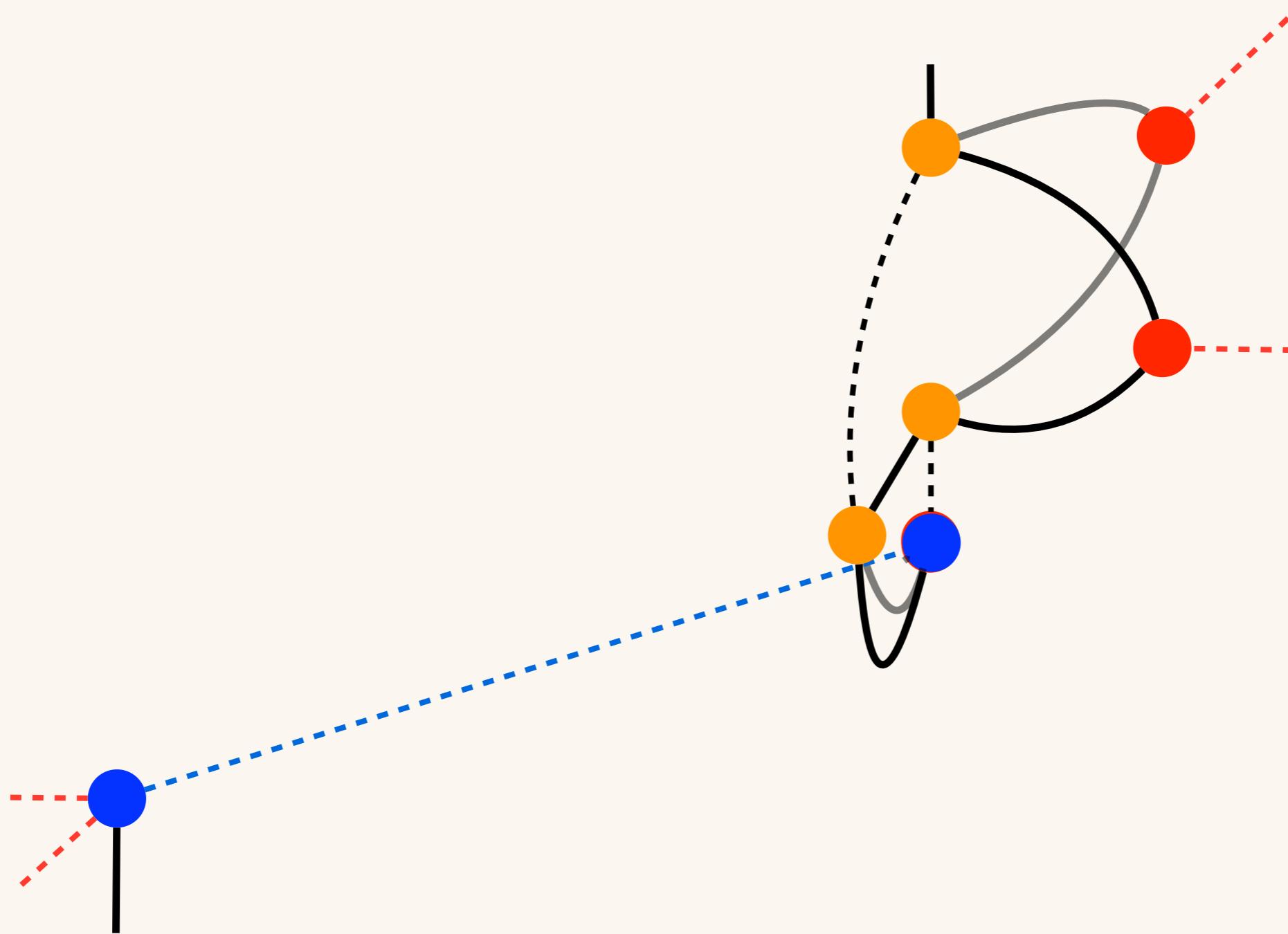


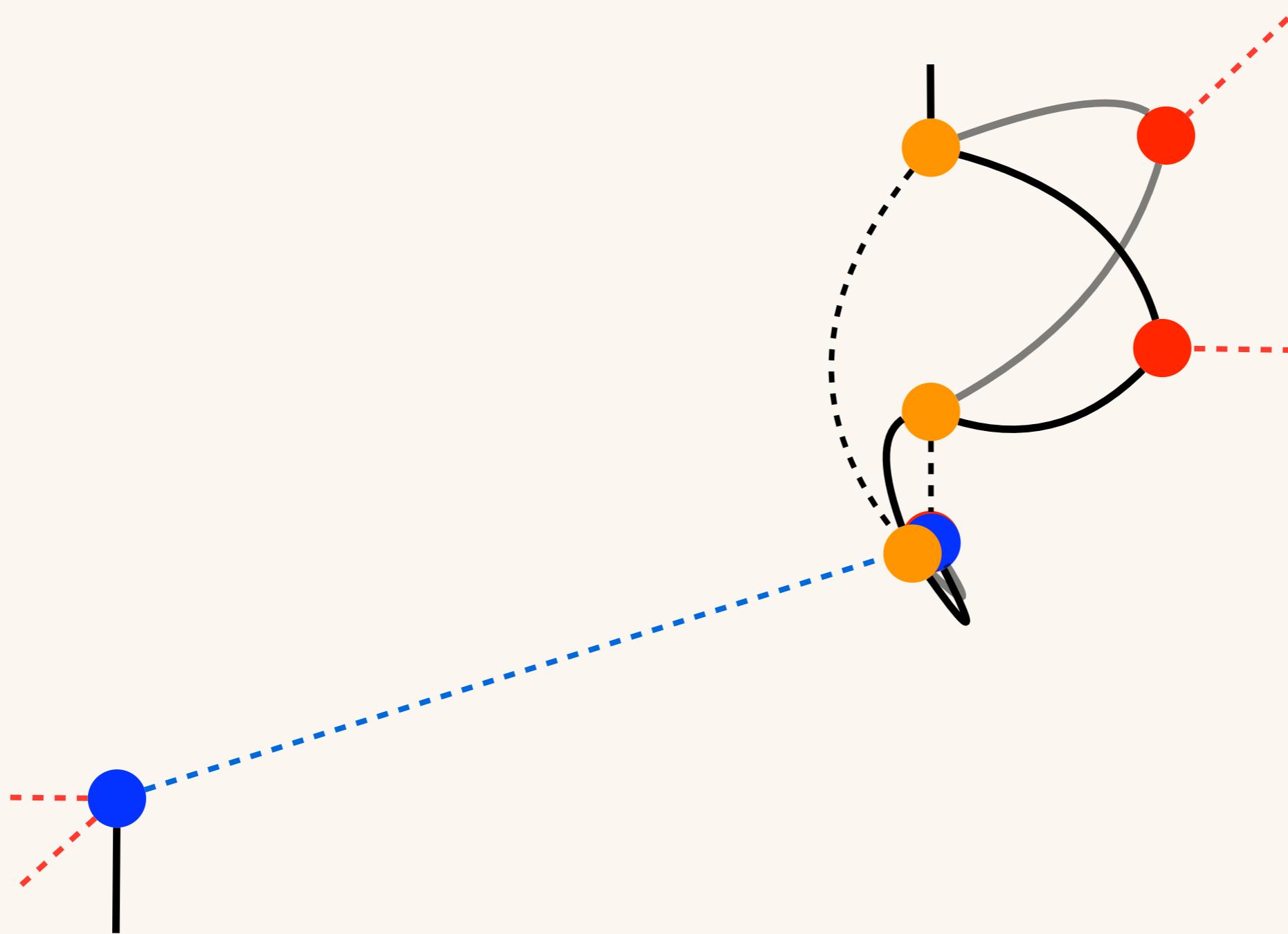


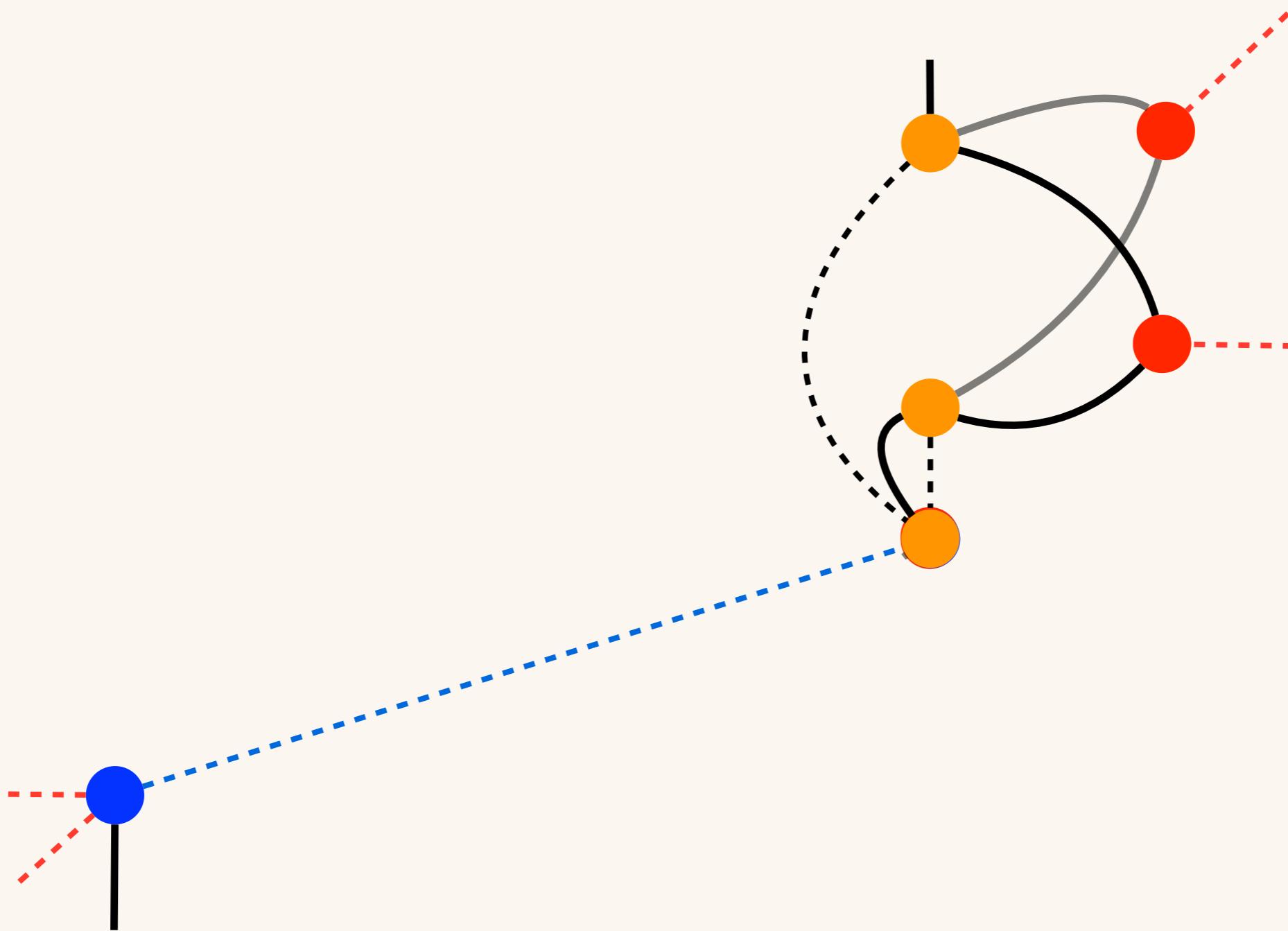


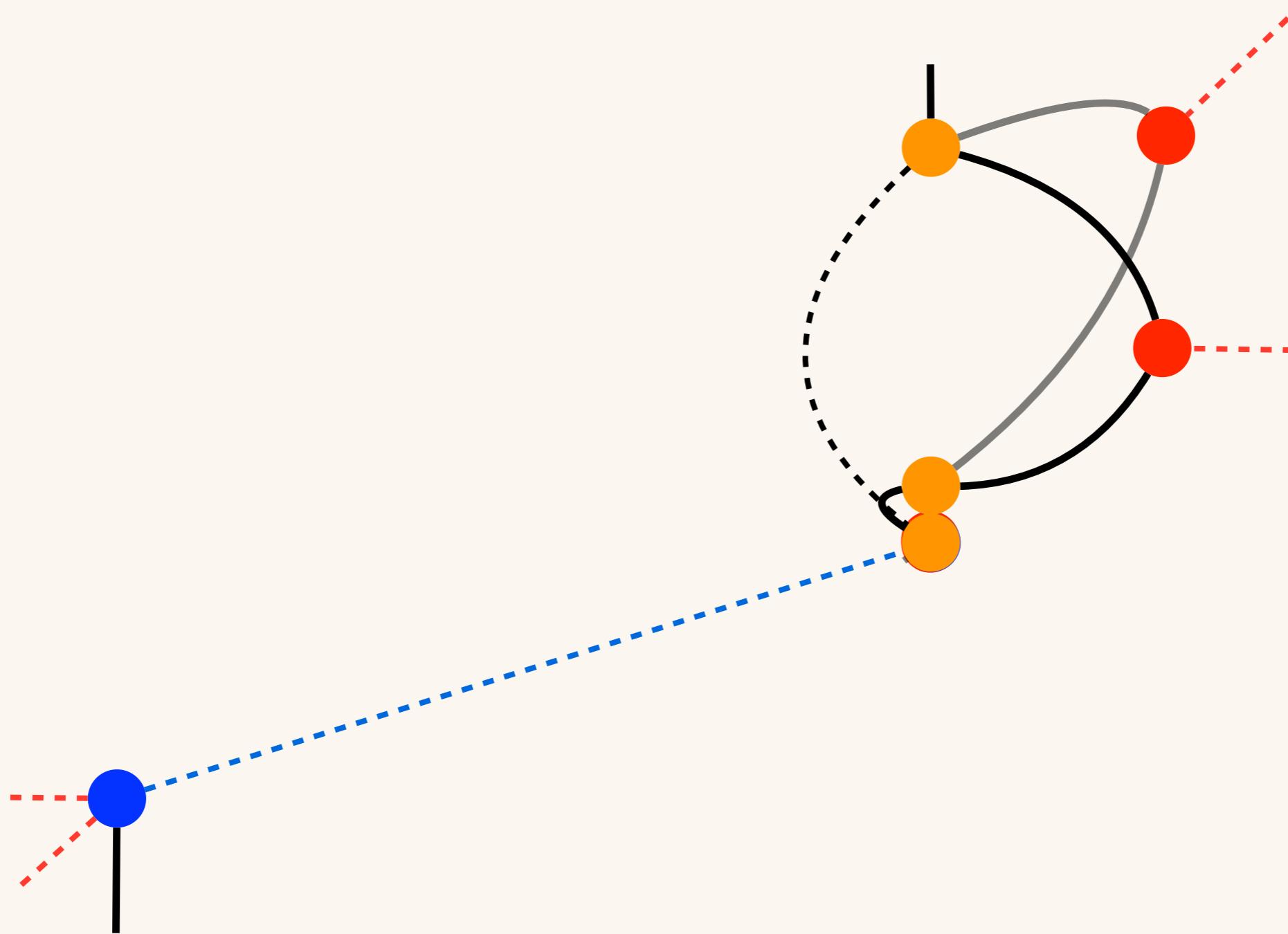


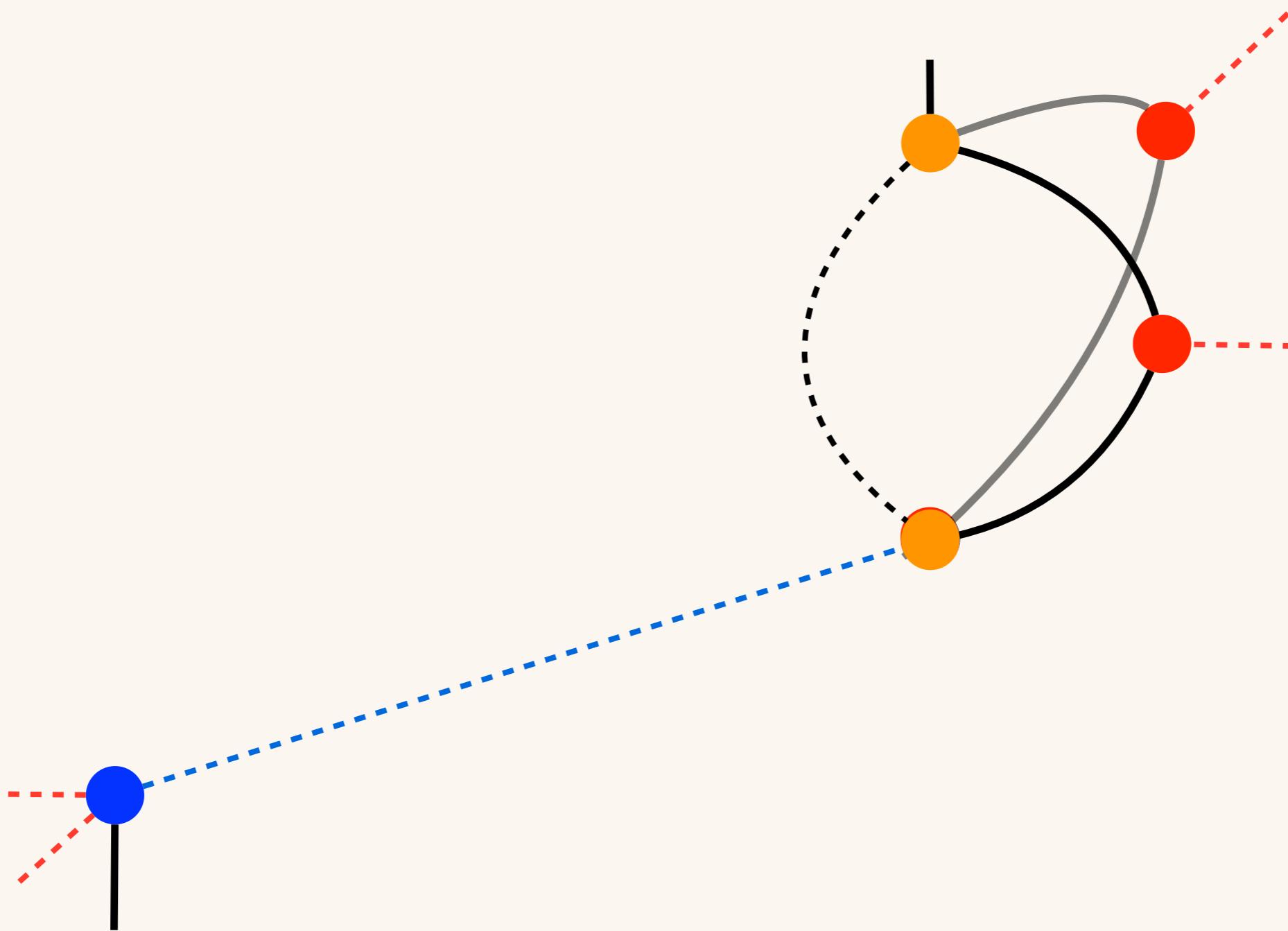


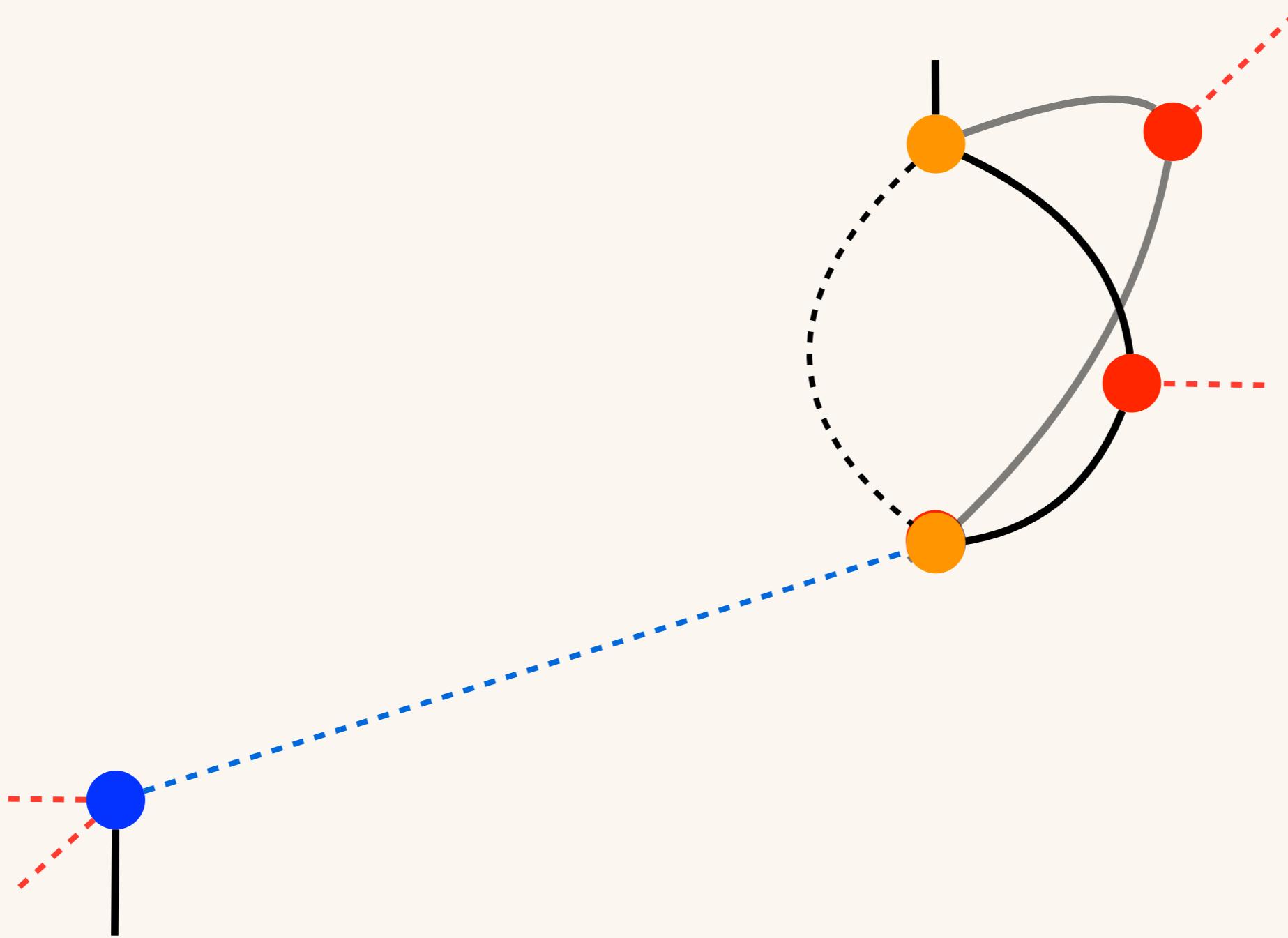


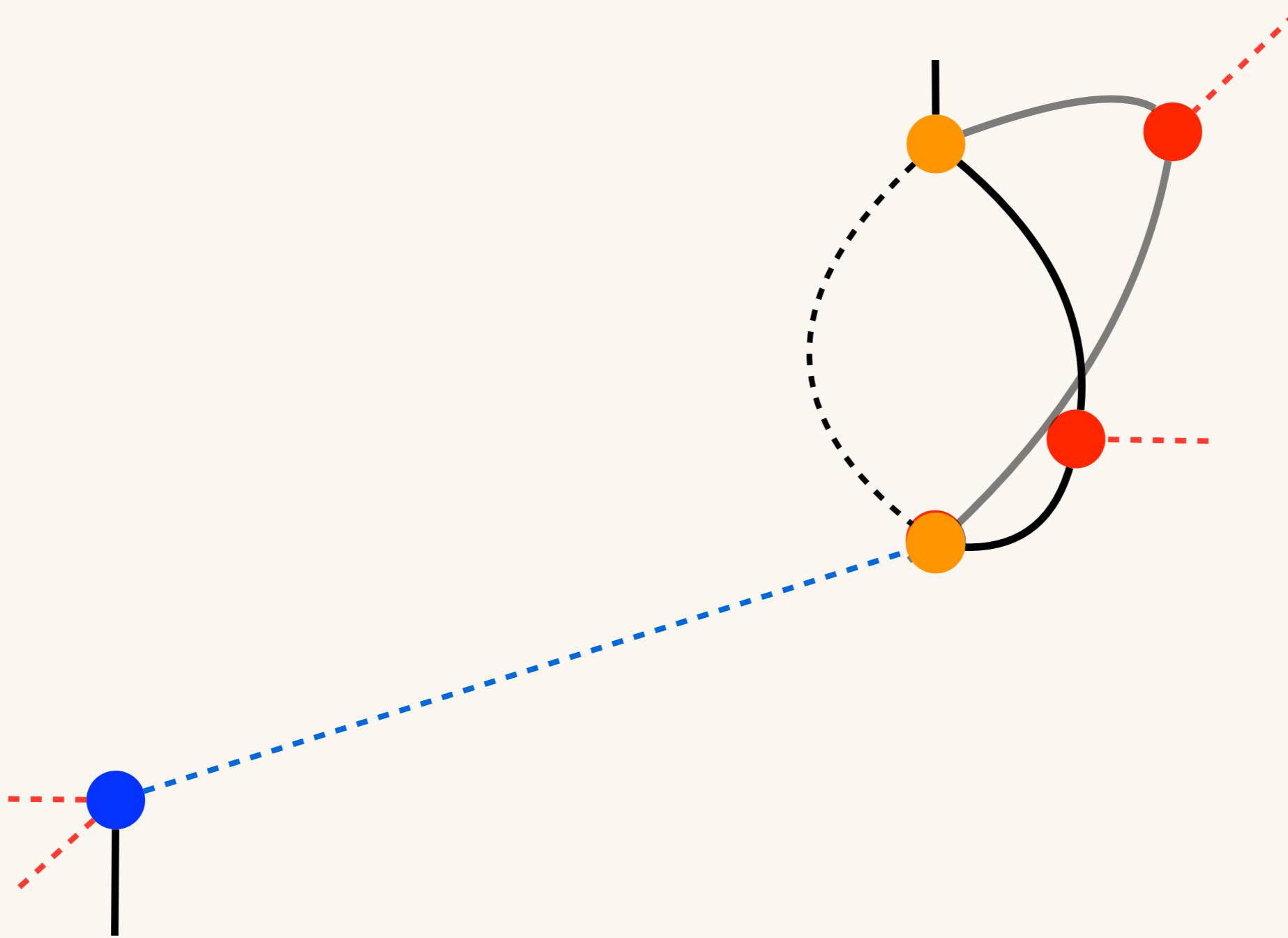


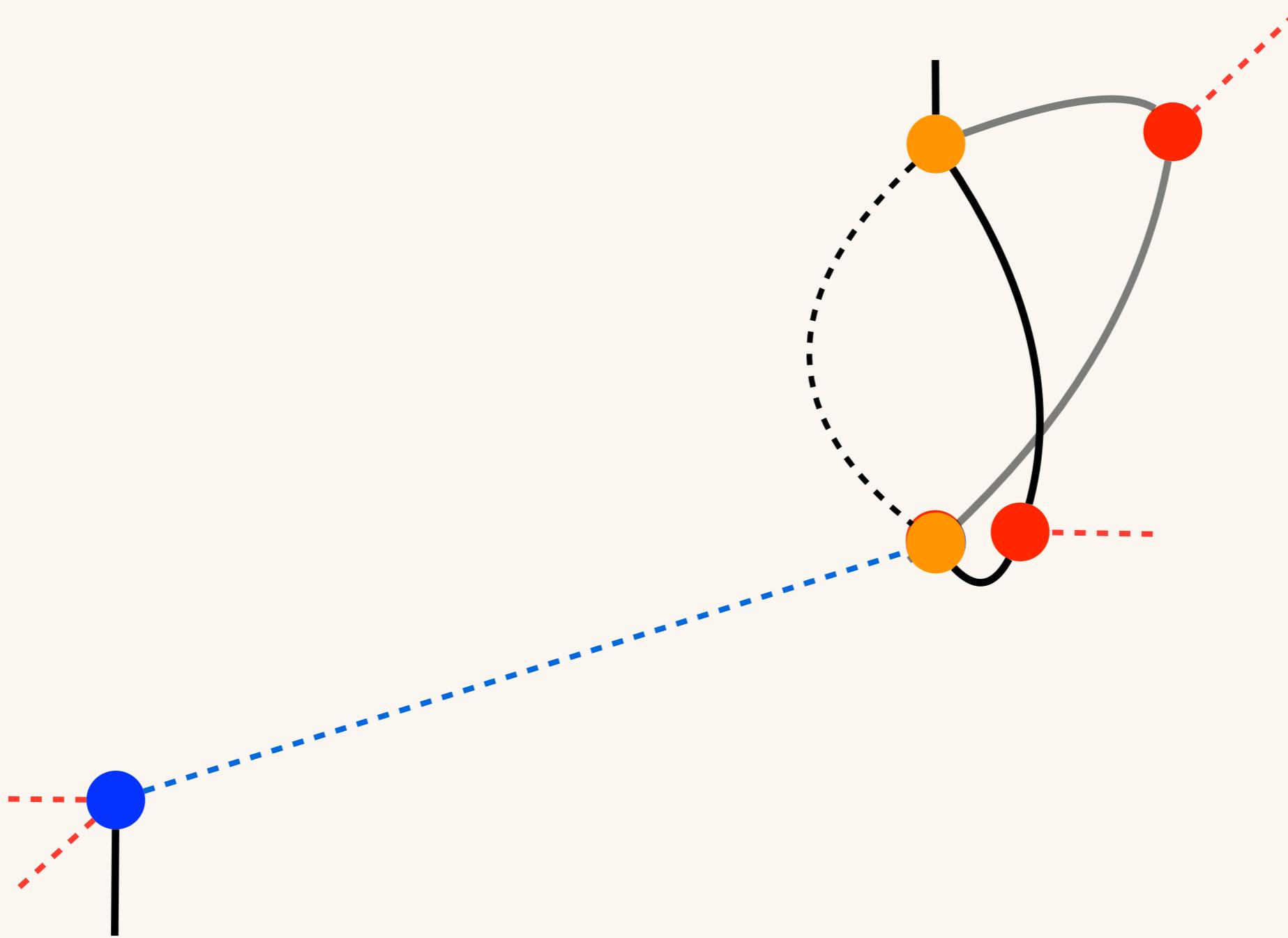


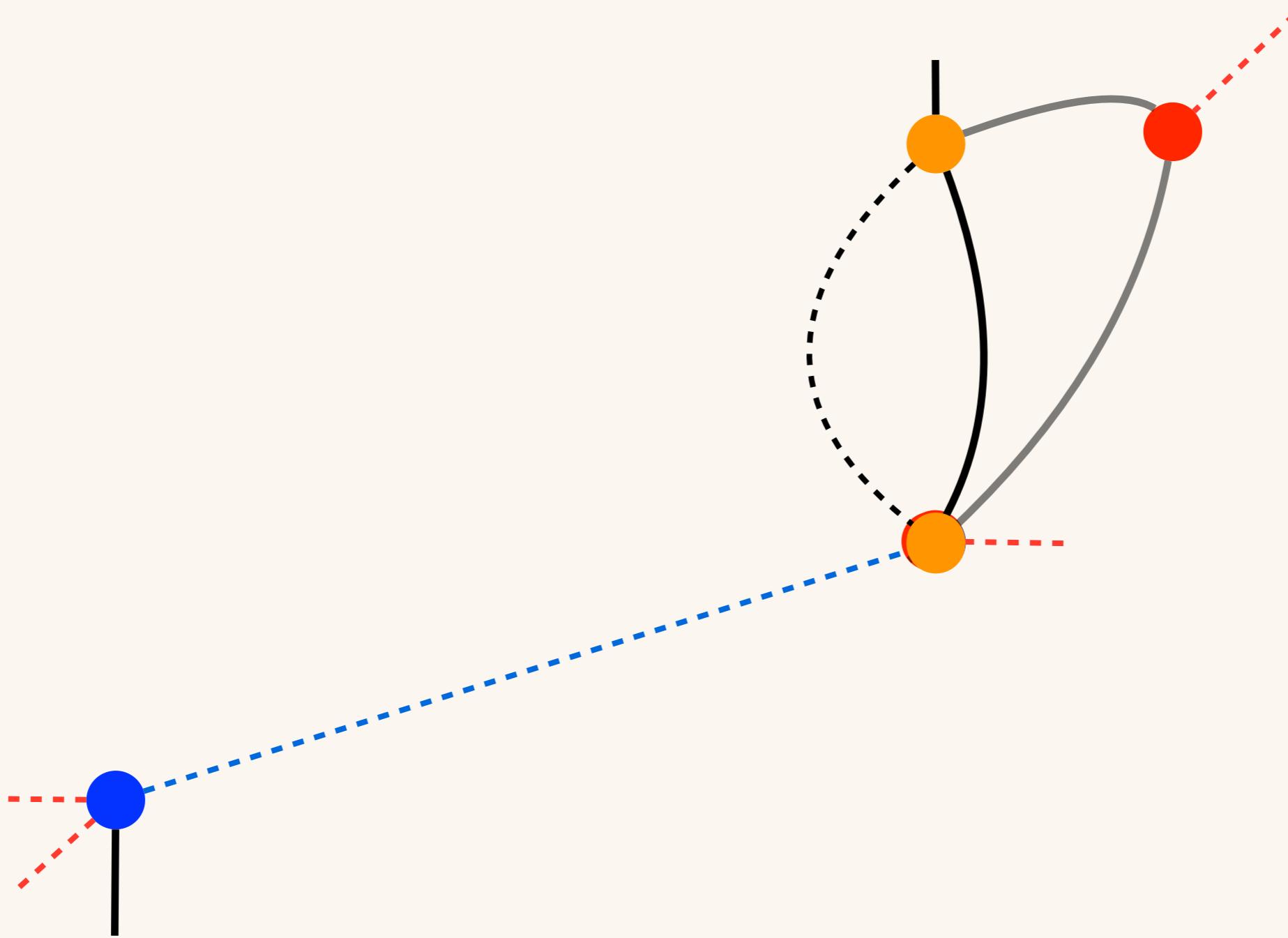


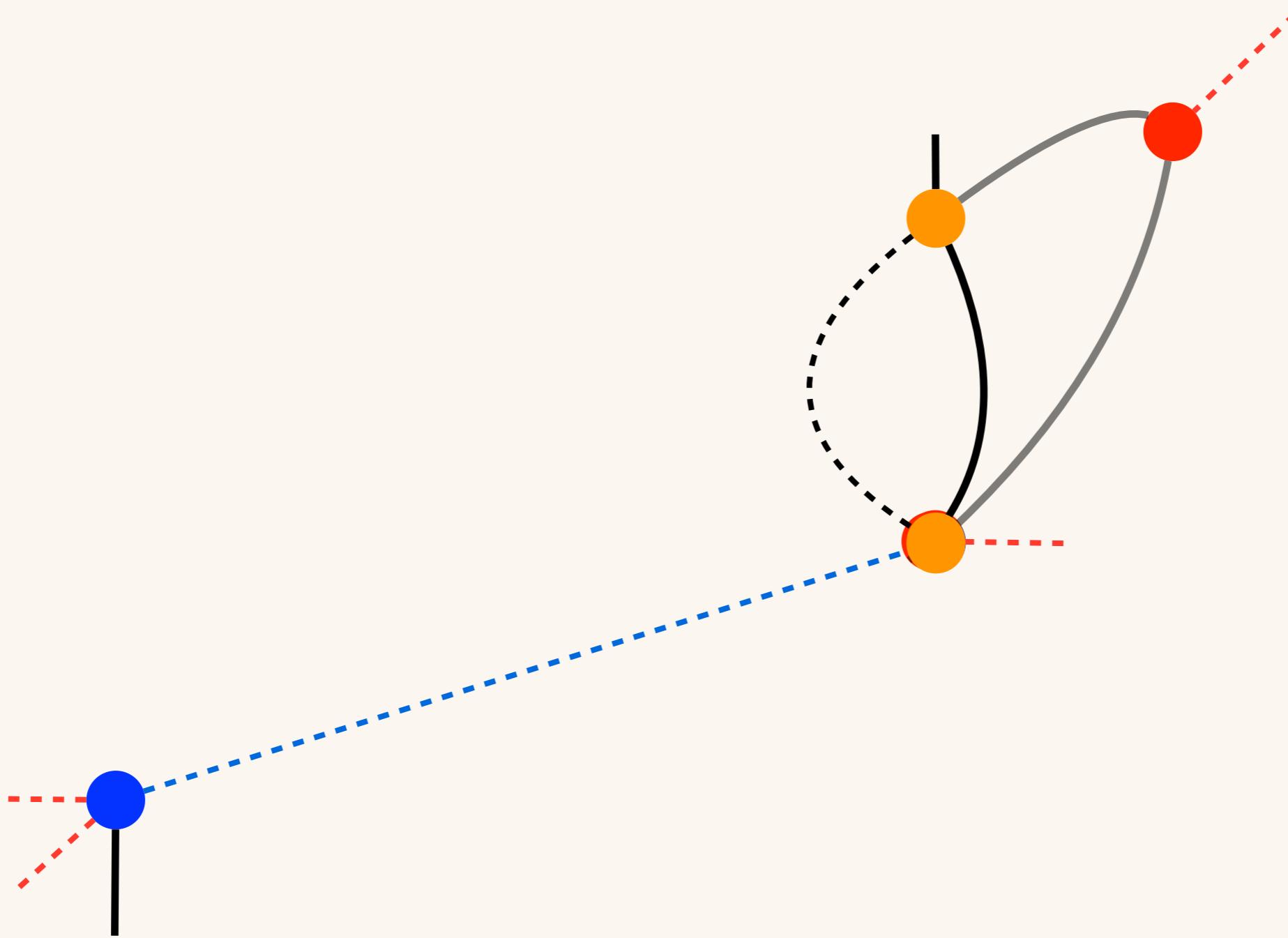


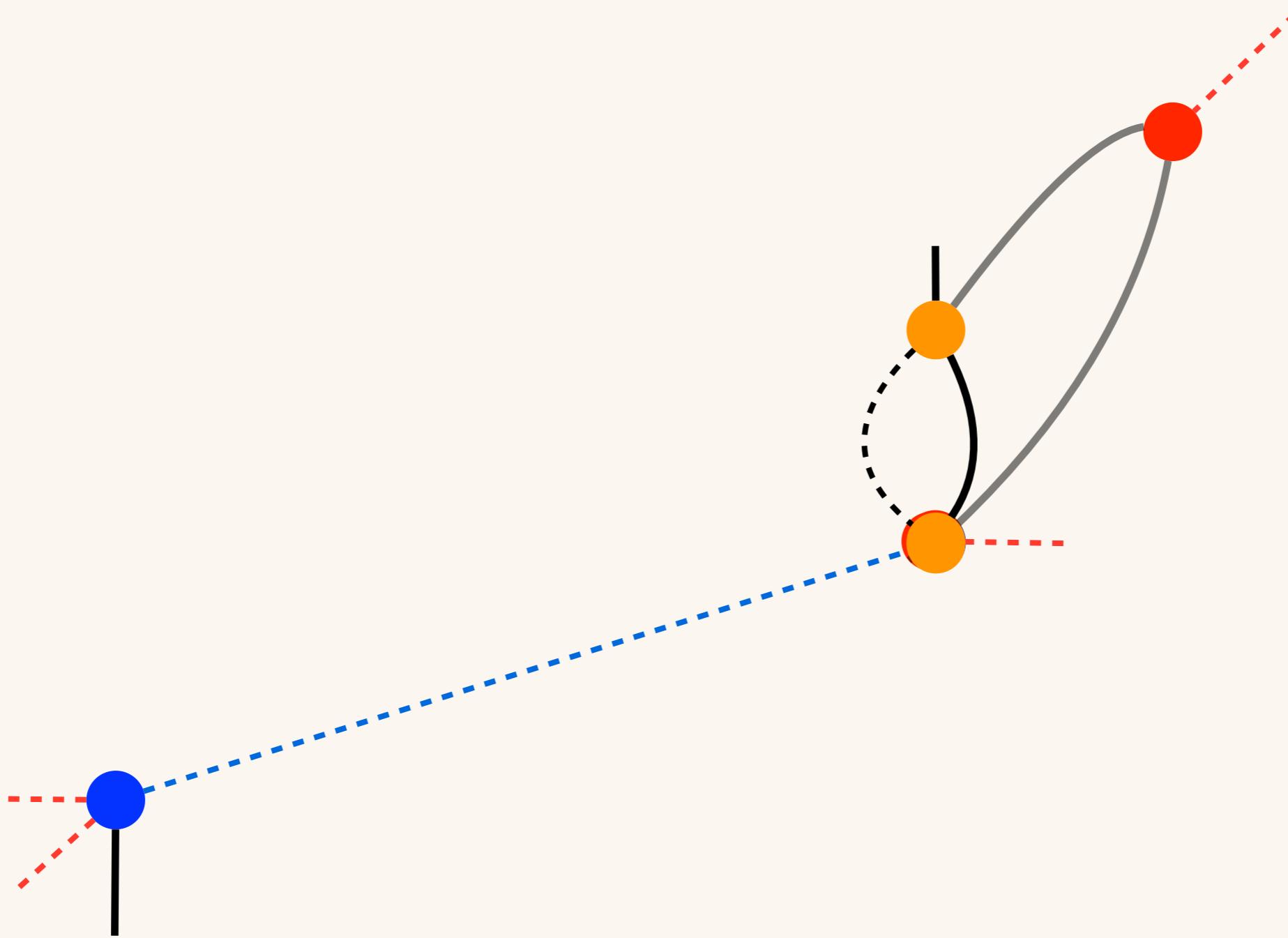


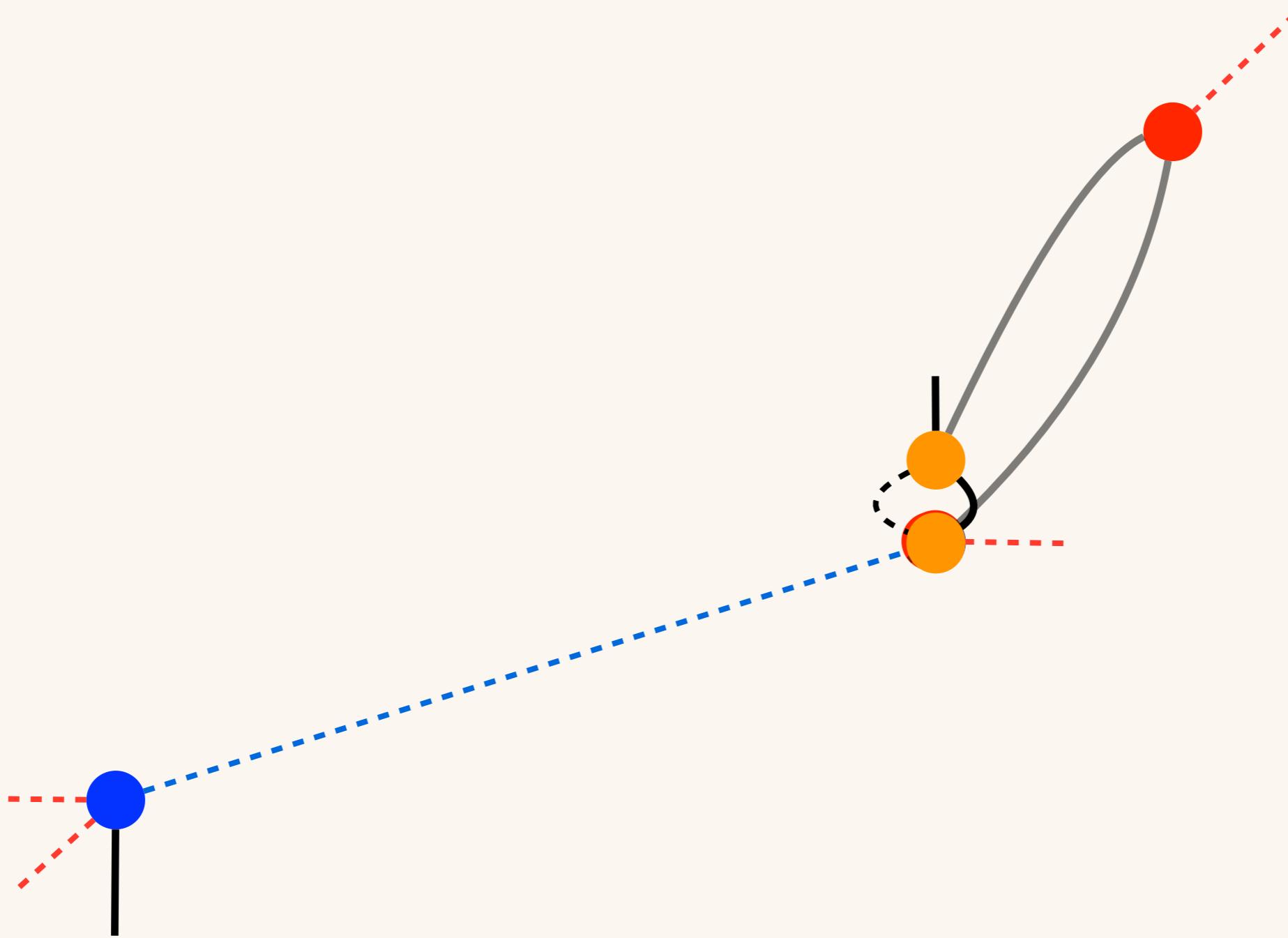


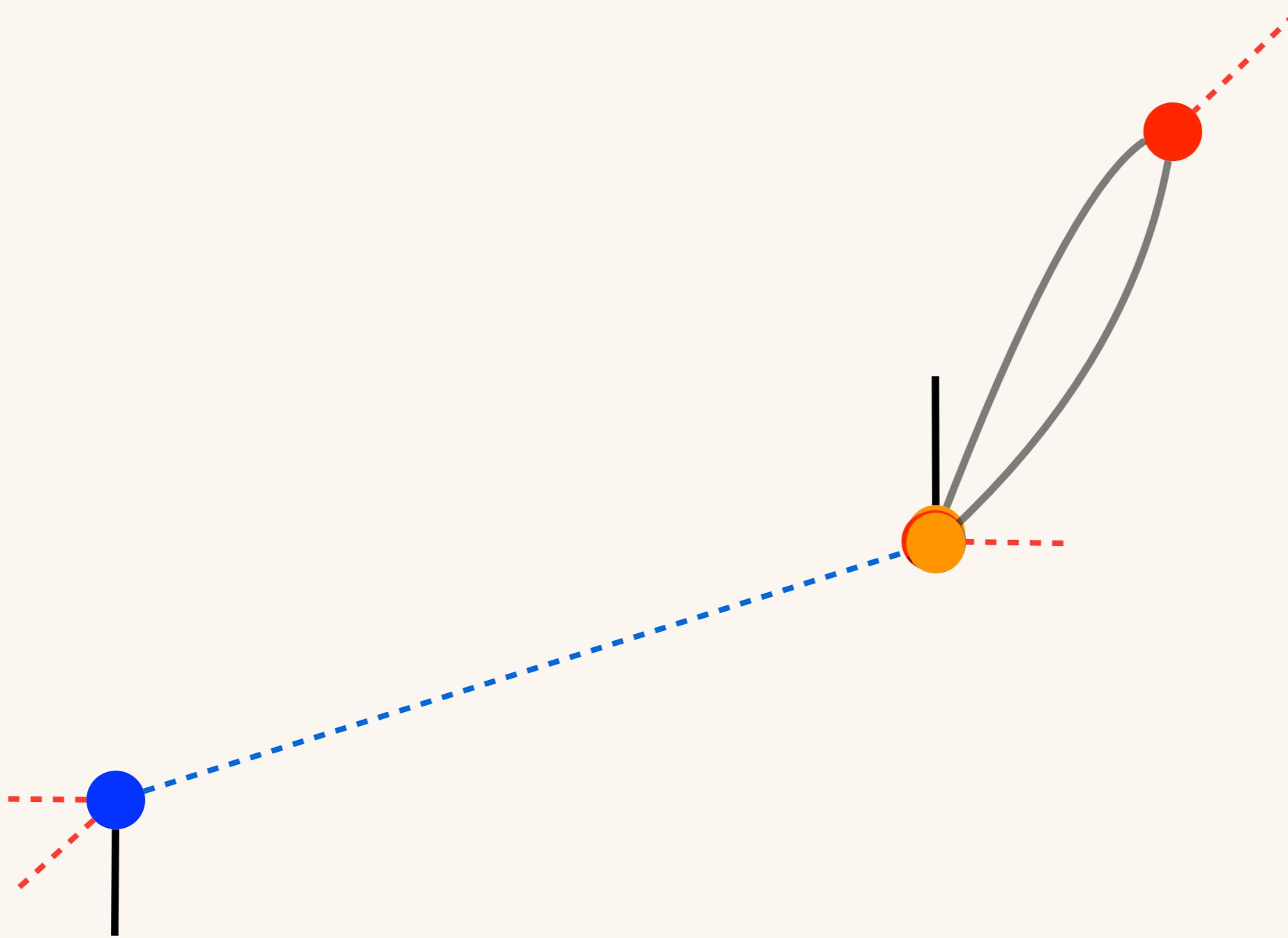


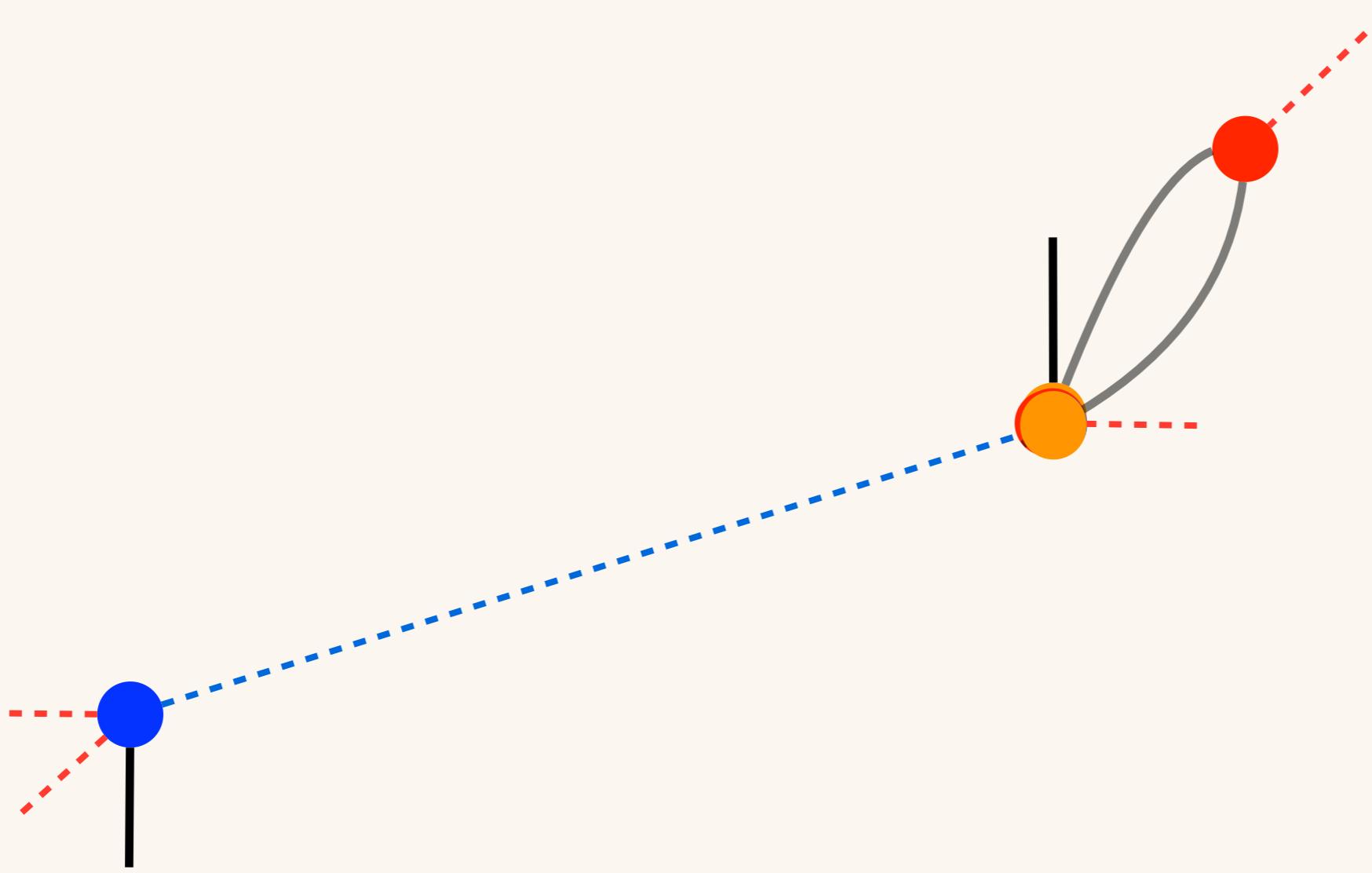


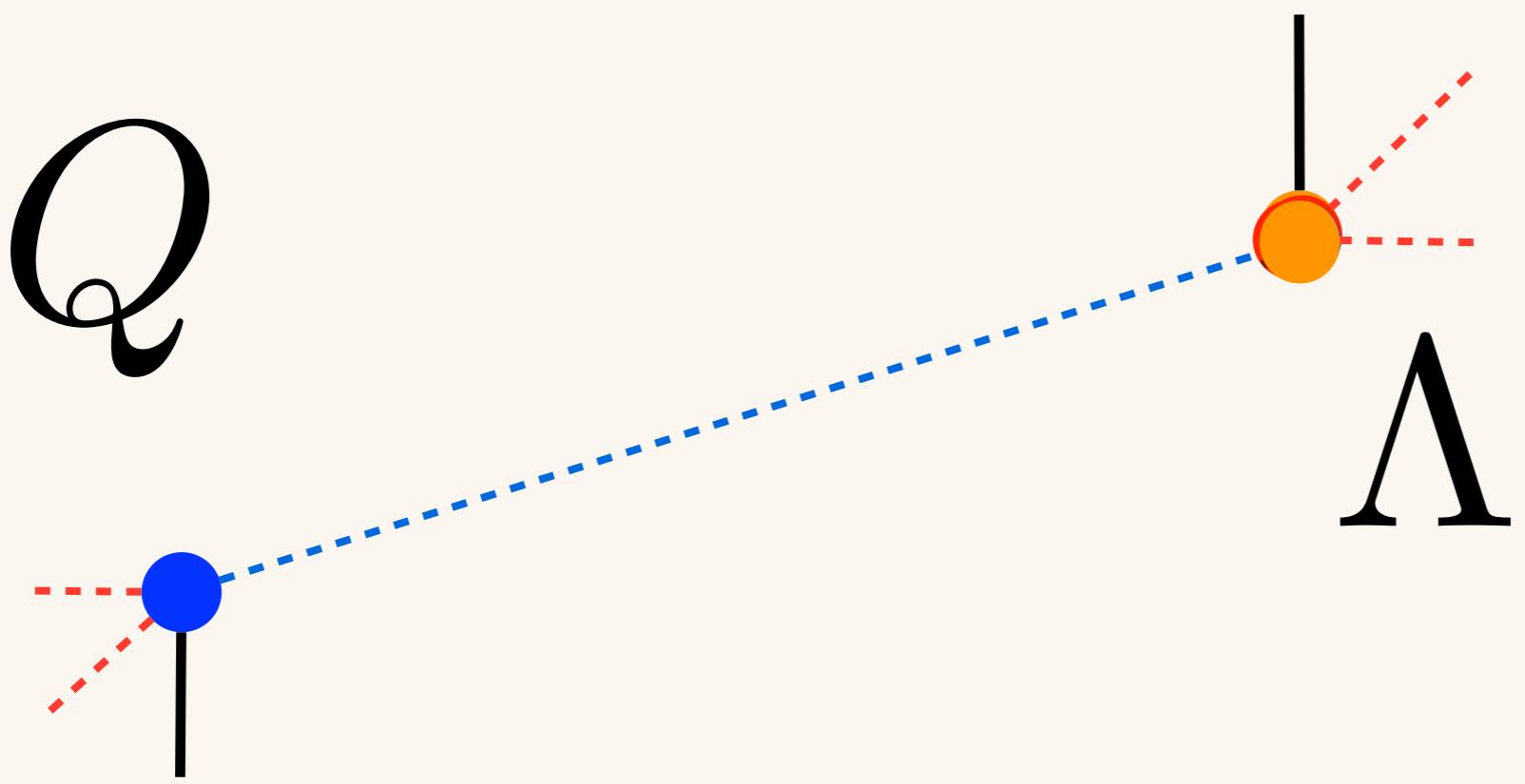






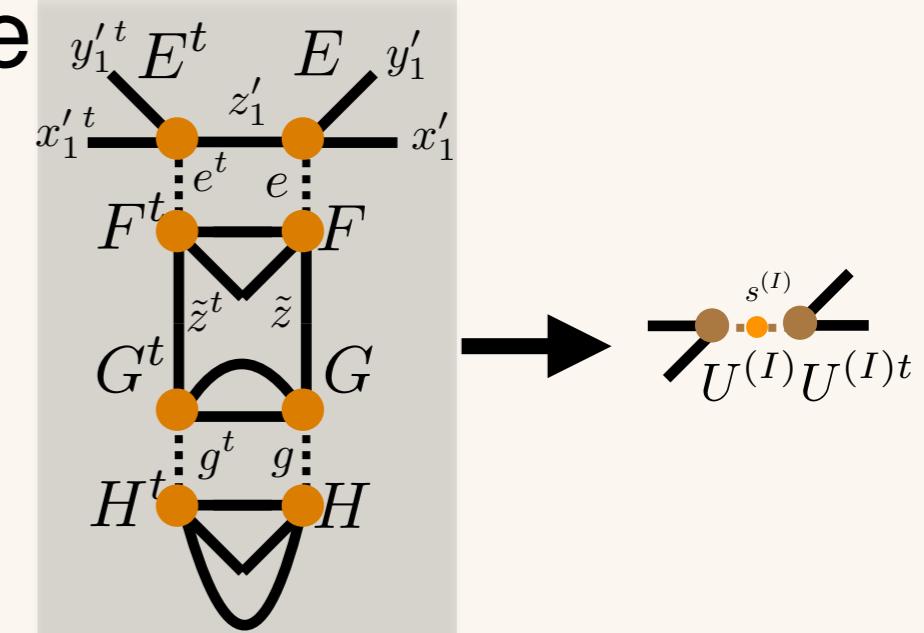
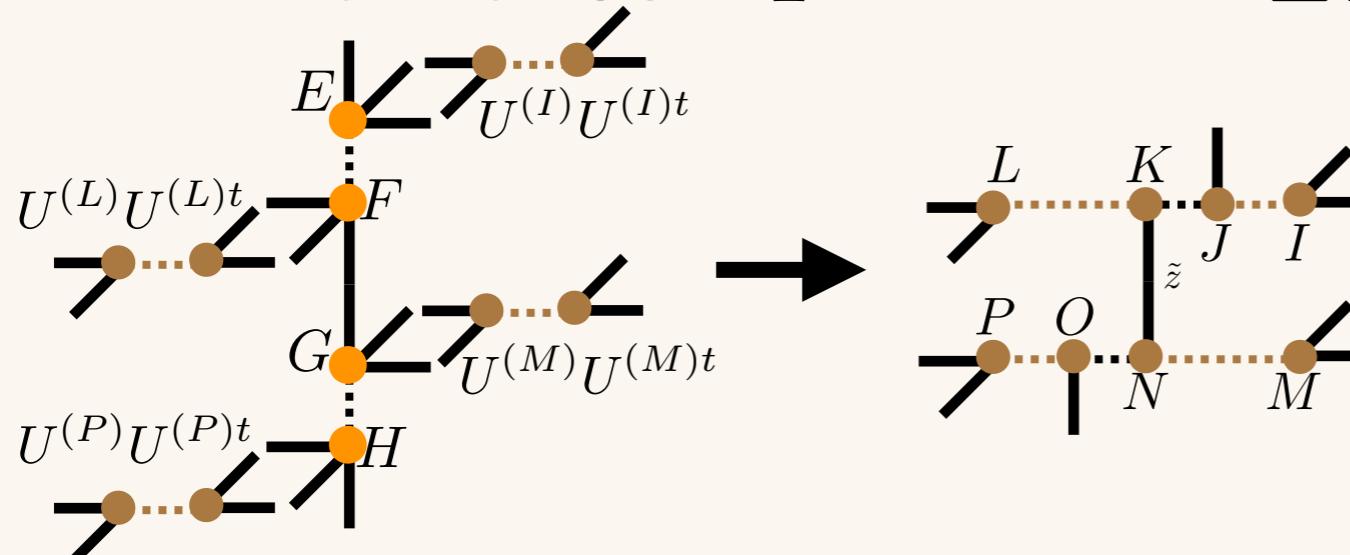






● Minimally-decomposed TRG on triad rep.

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



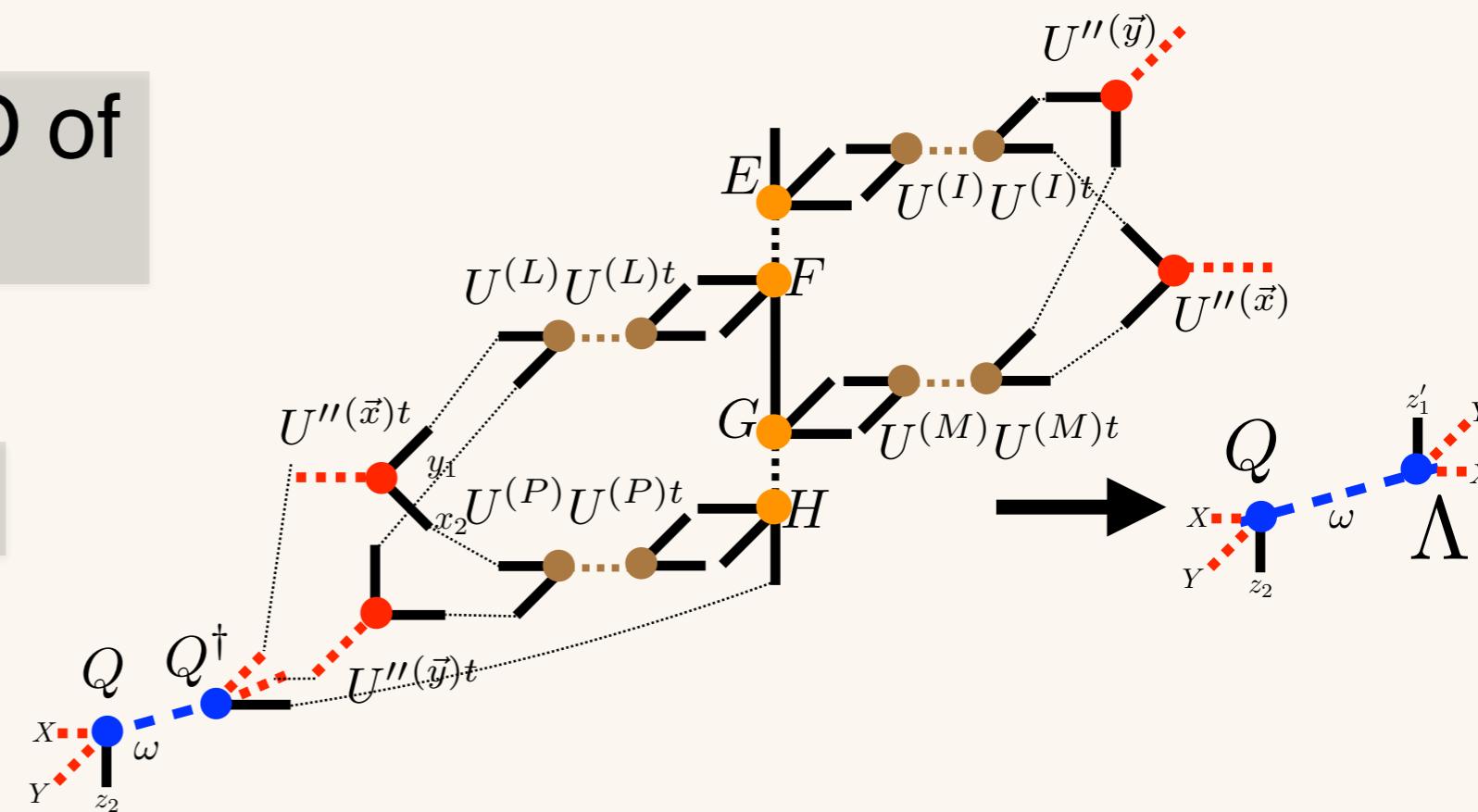
- ◇ We do NOT use SVD of E, F, G and H.

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

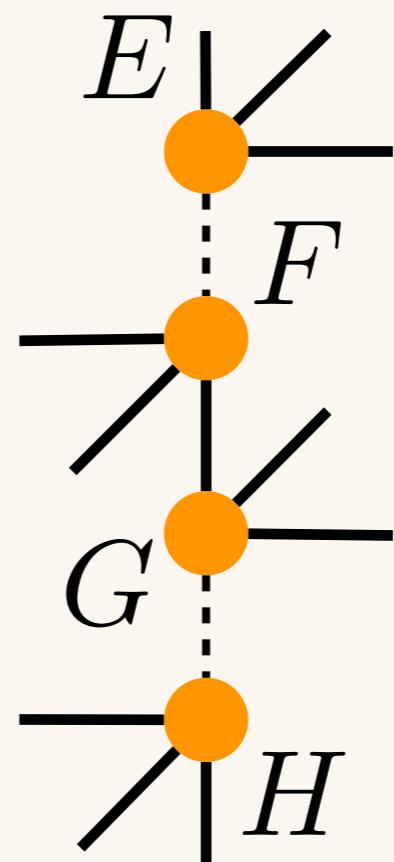
- ◇ We use SVD of $\Gamma\Gamma^t$.

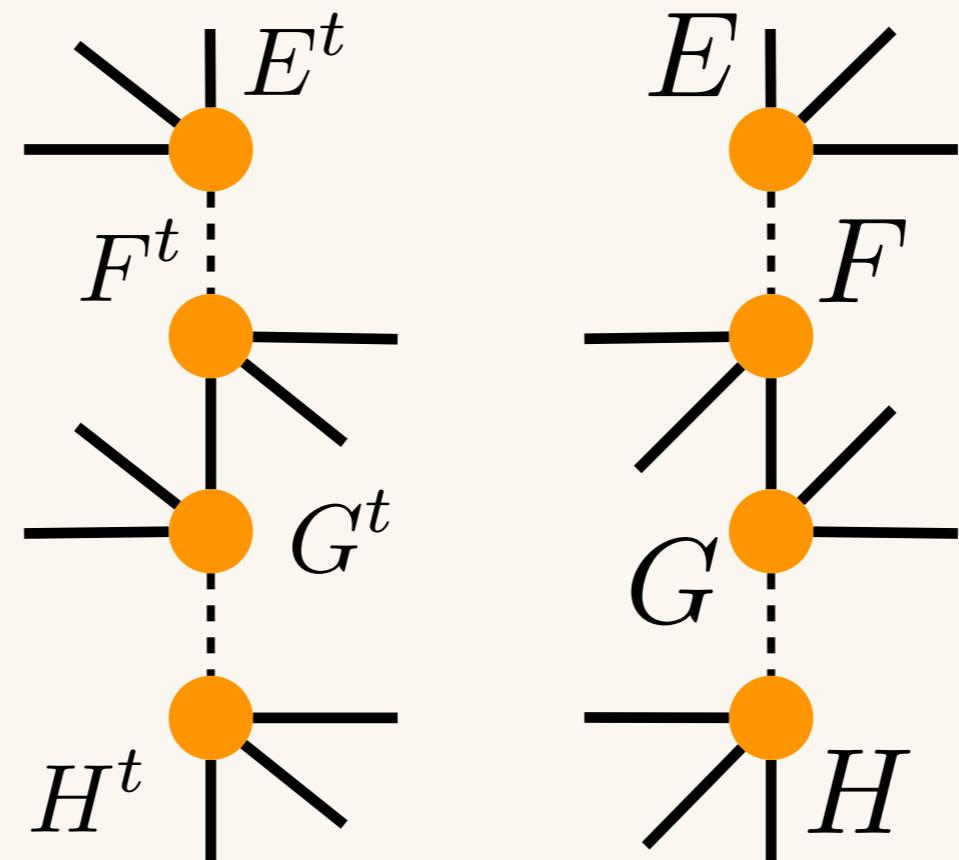
- ◇ Cost reduction

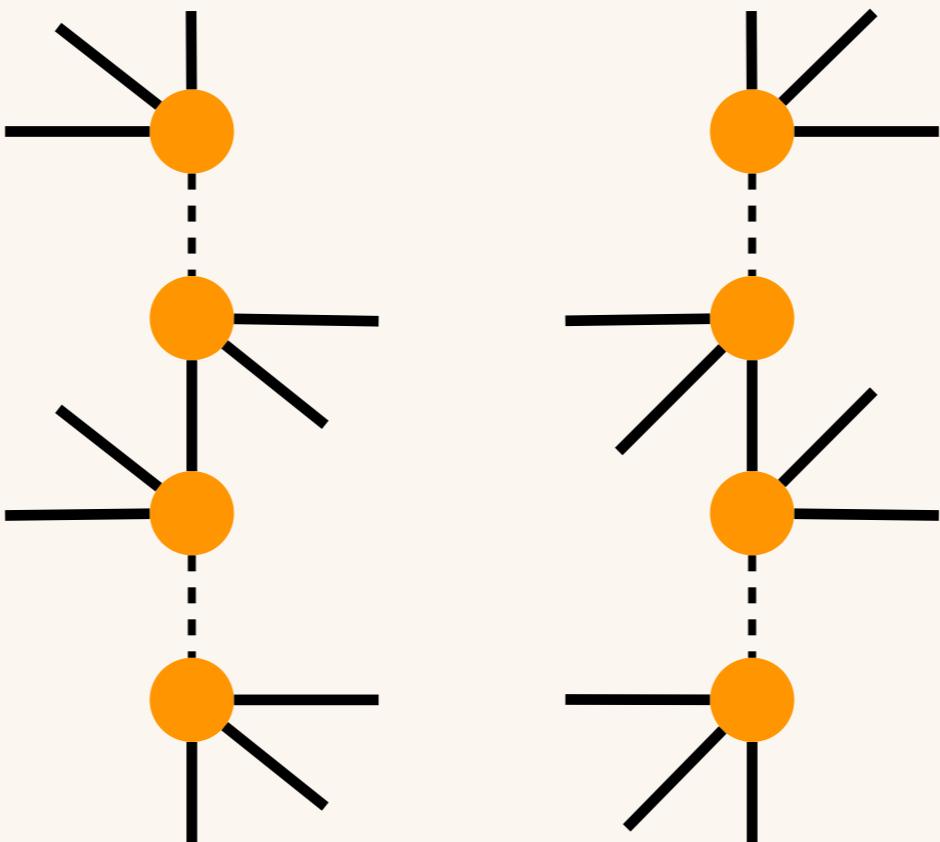
$$O(D^{2d+1}) \rightarrow O(D^{d+3})$$

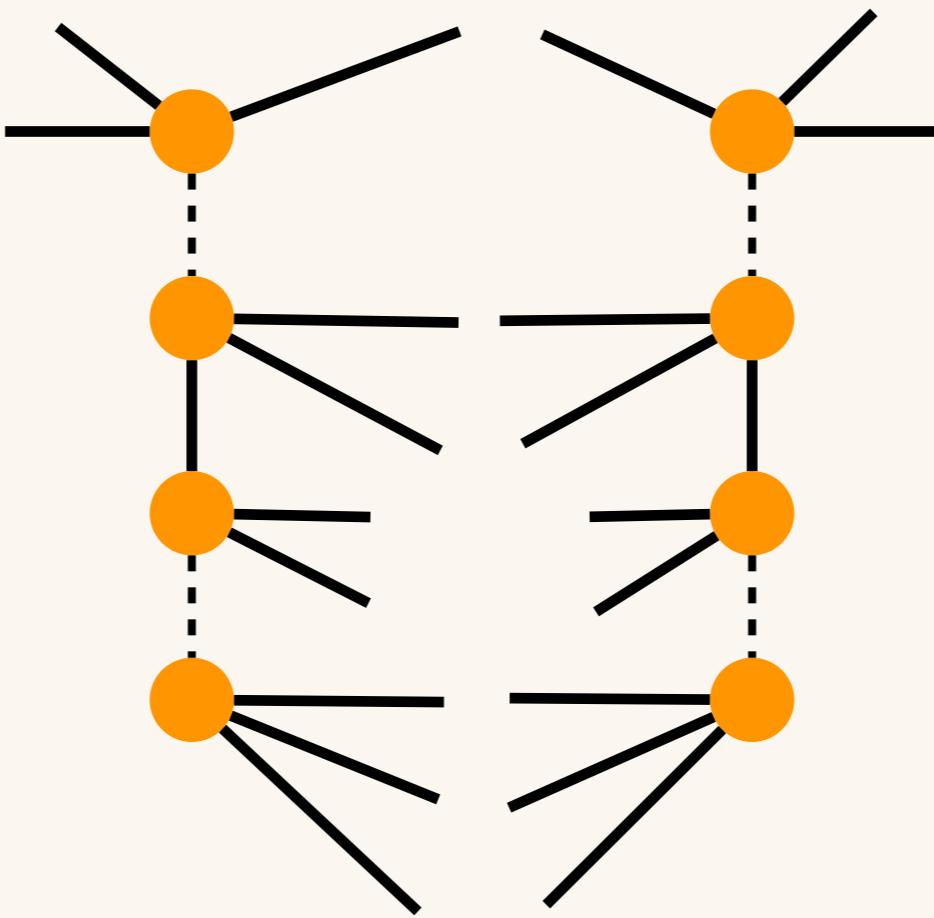


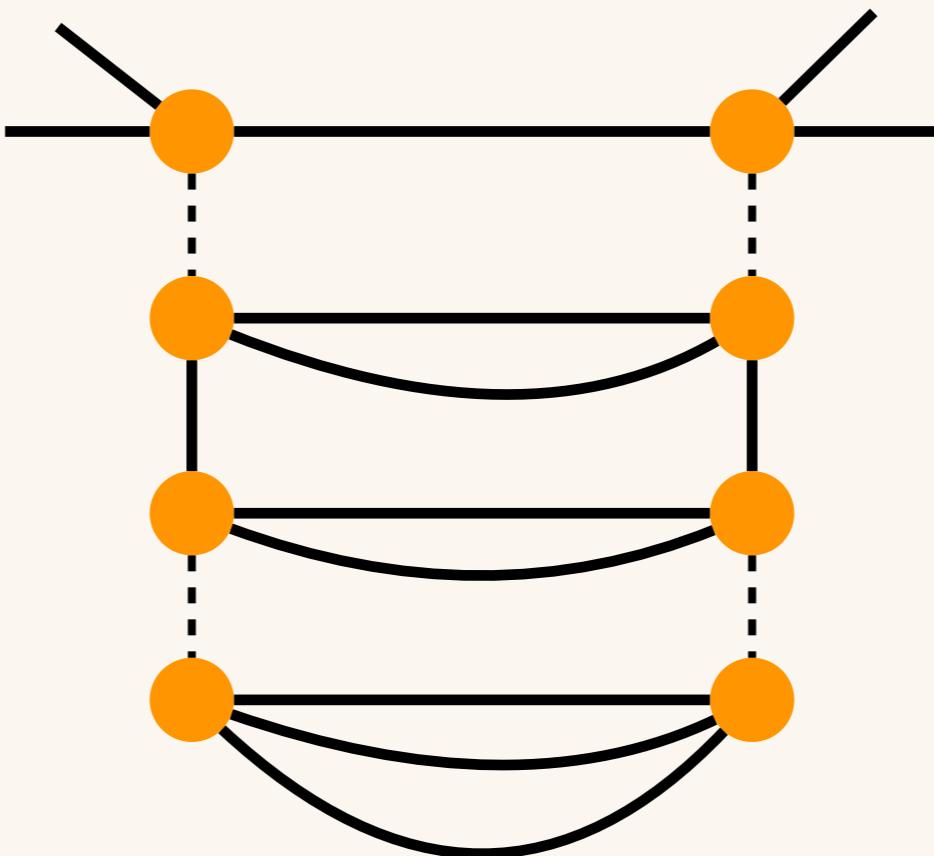
$$QQ^\dagger U''(\vec{y})t U''(\vec{x})t U^{(LP)}U^{(LP)t} EFGHU^{(IM)}U^{(IM)t} U''(\vec{x})U''(\vec{y}) = Q\Lambda \simeq A^{(\text{next})}$$

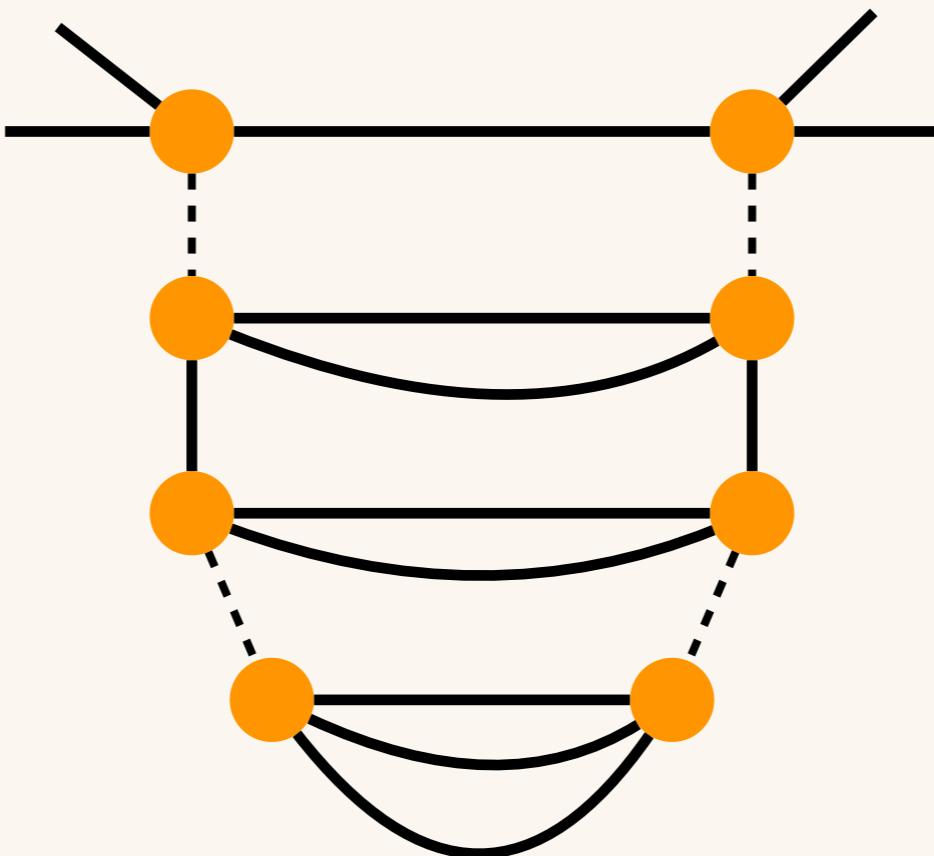


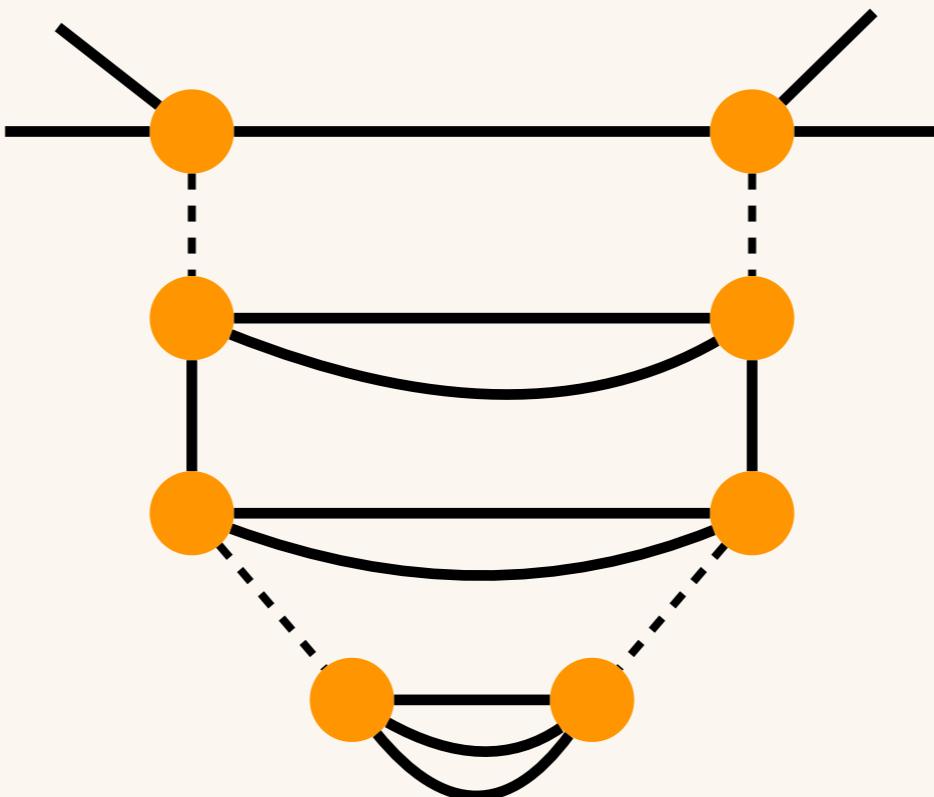


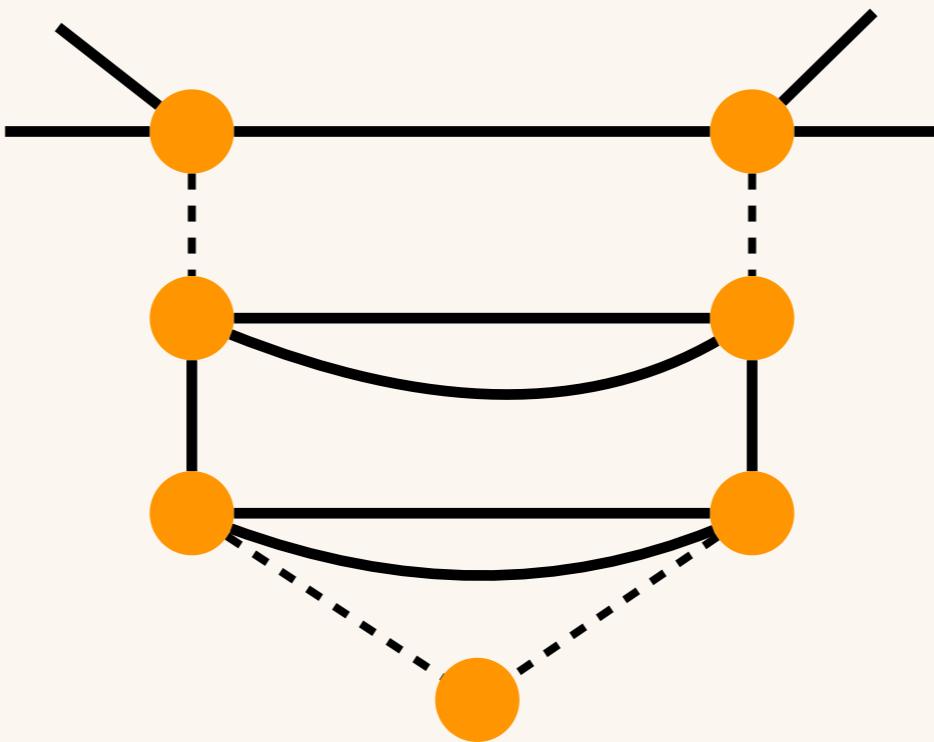


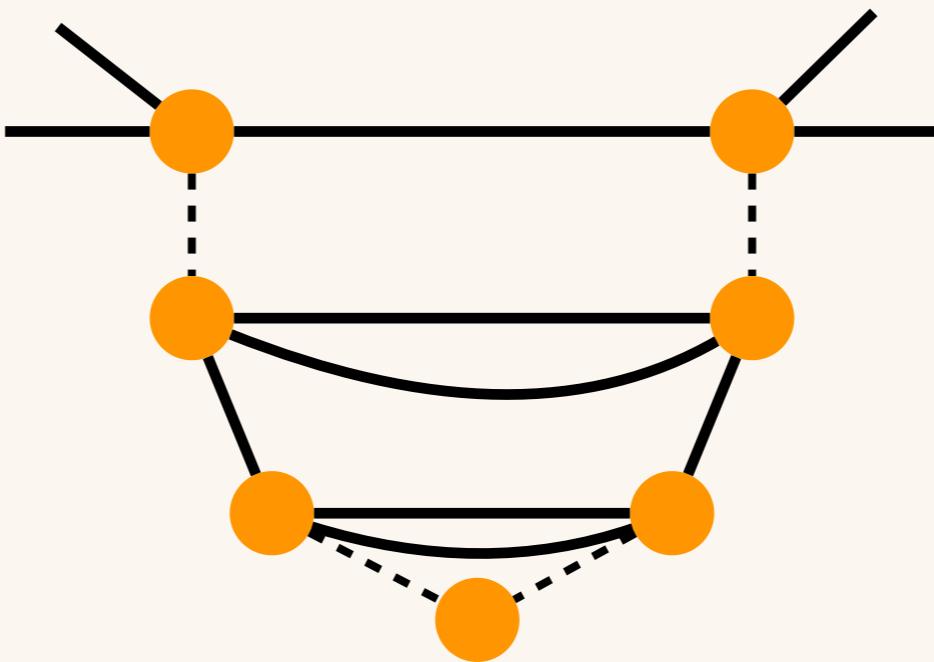


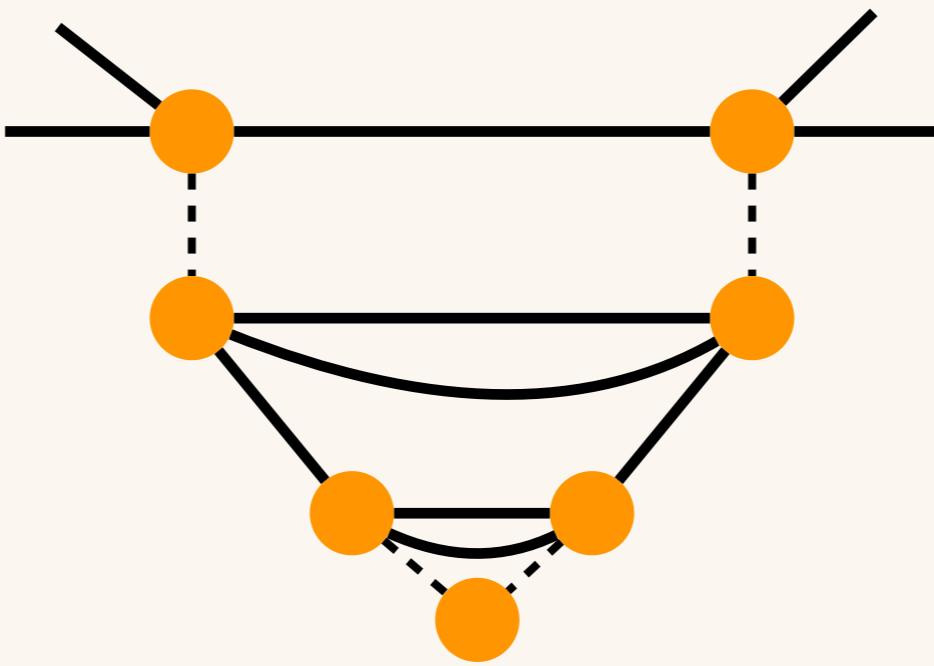


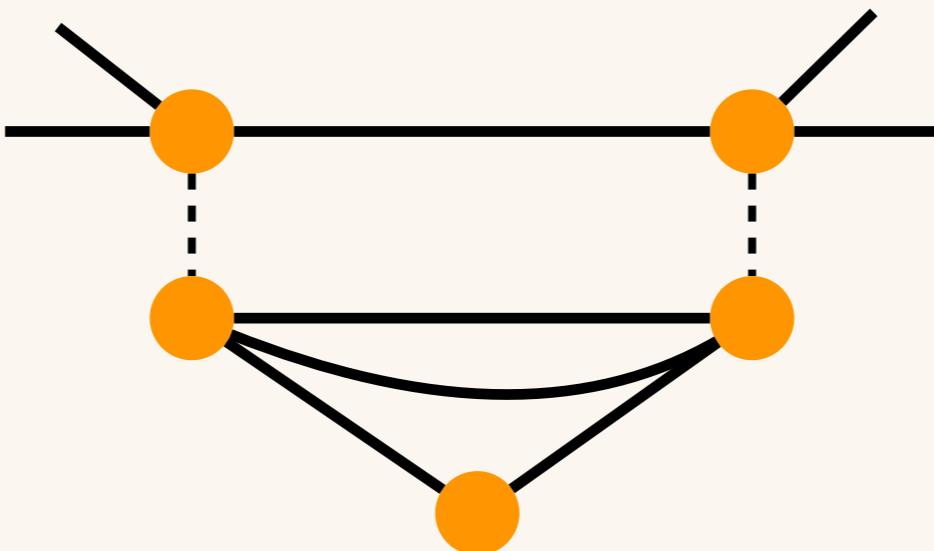


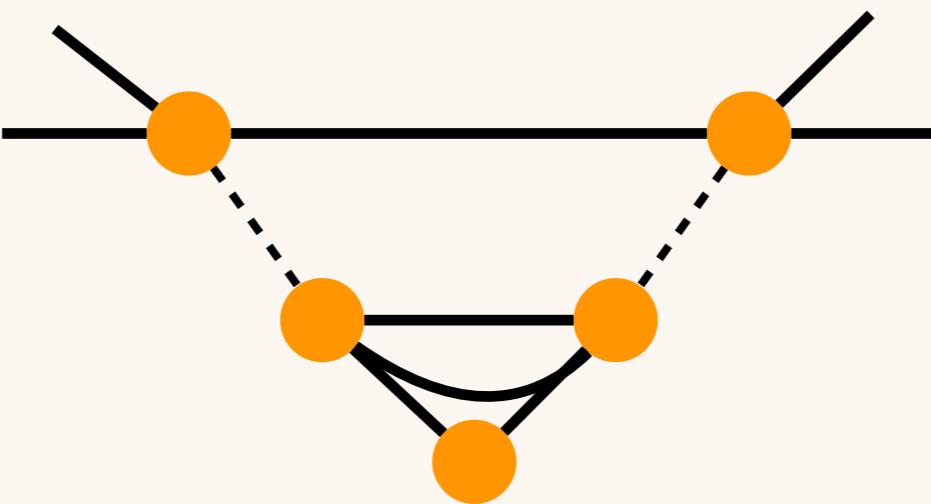


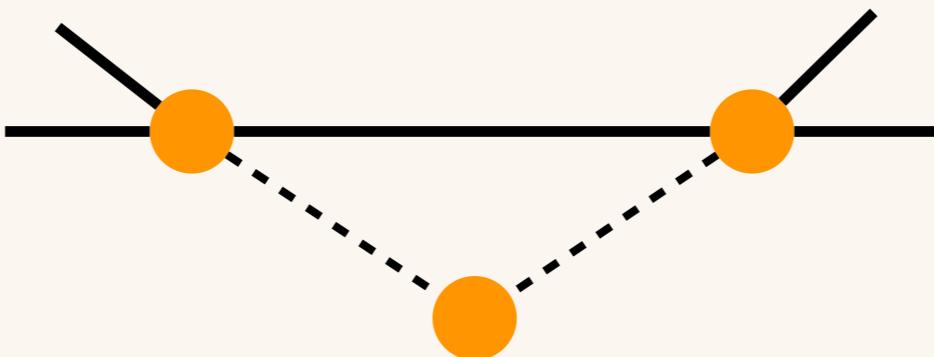


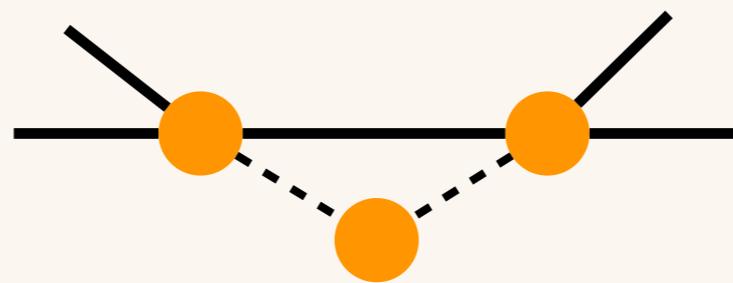


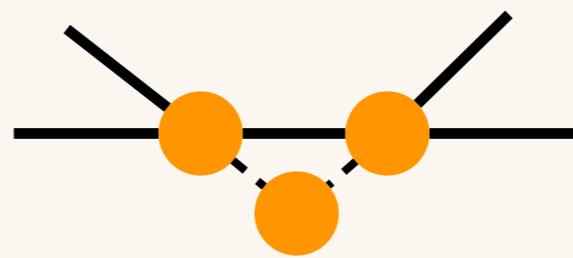


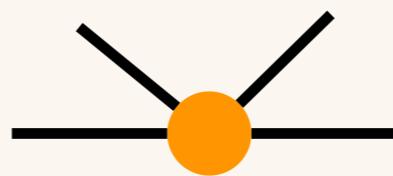




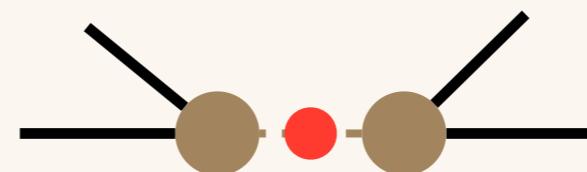




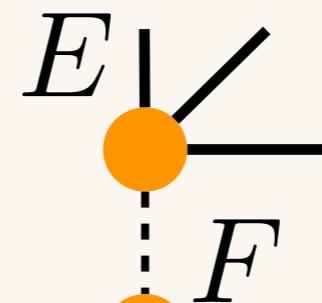
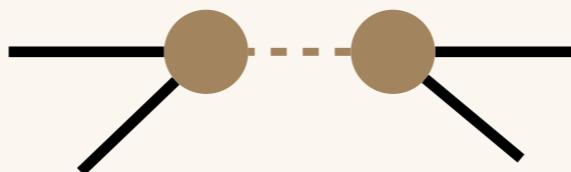
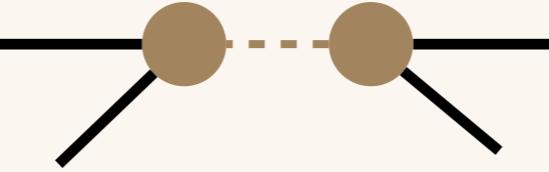
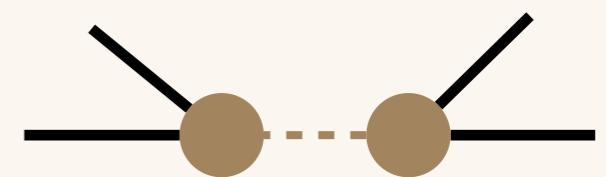


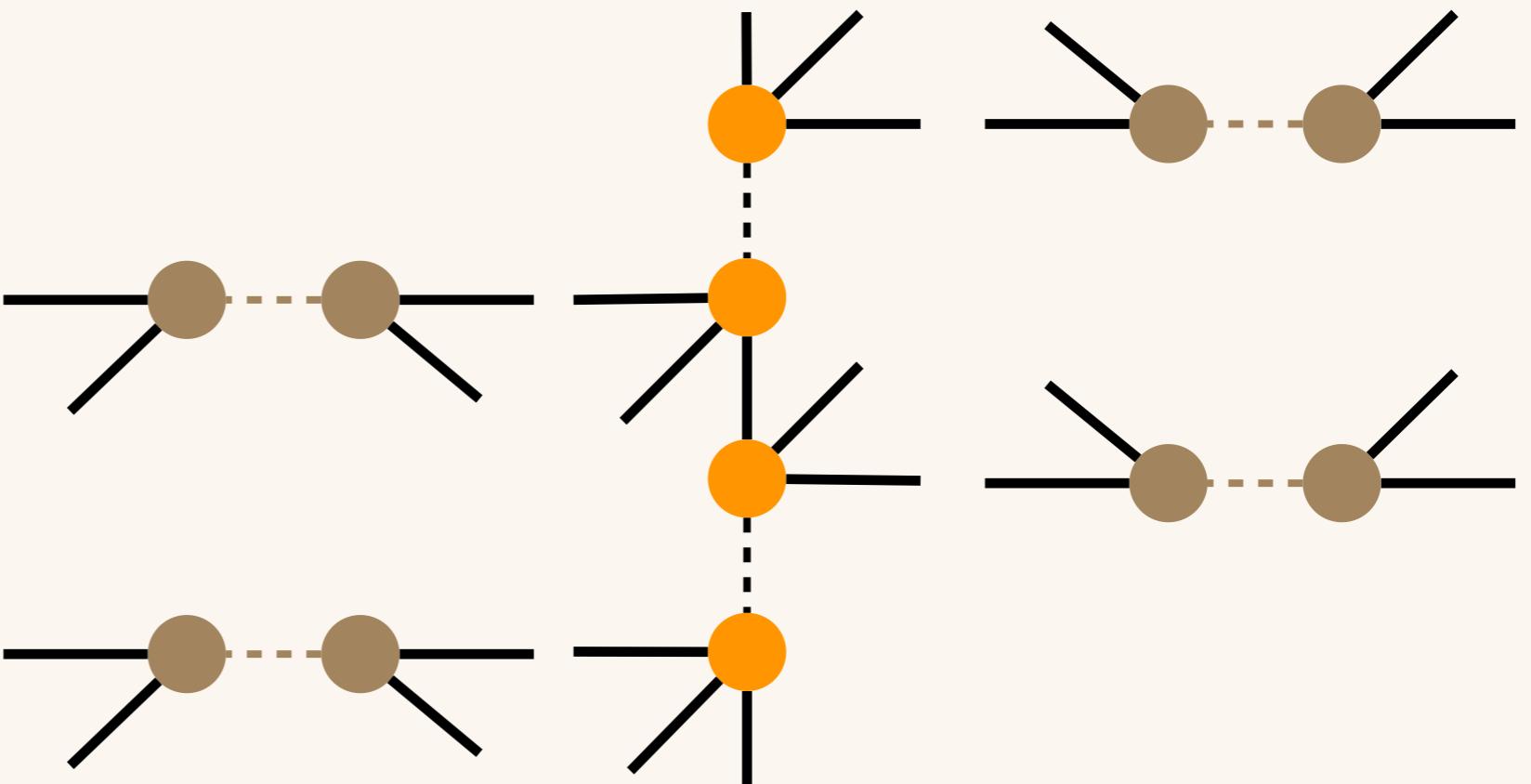


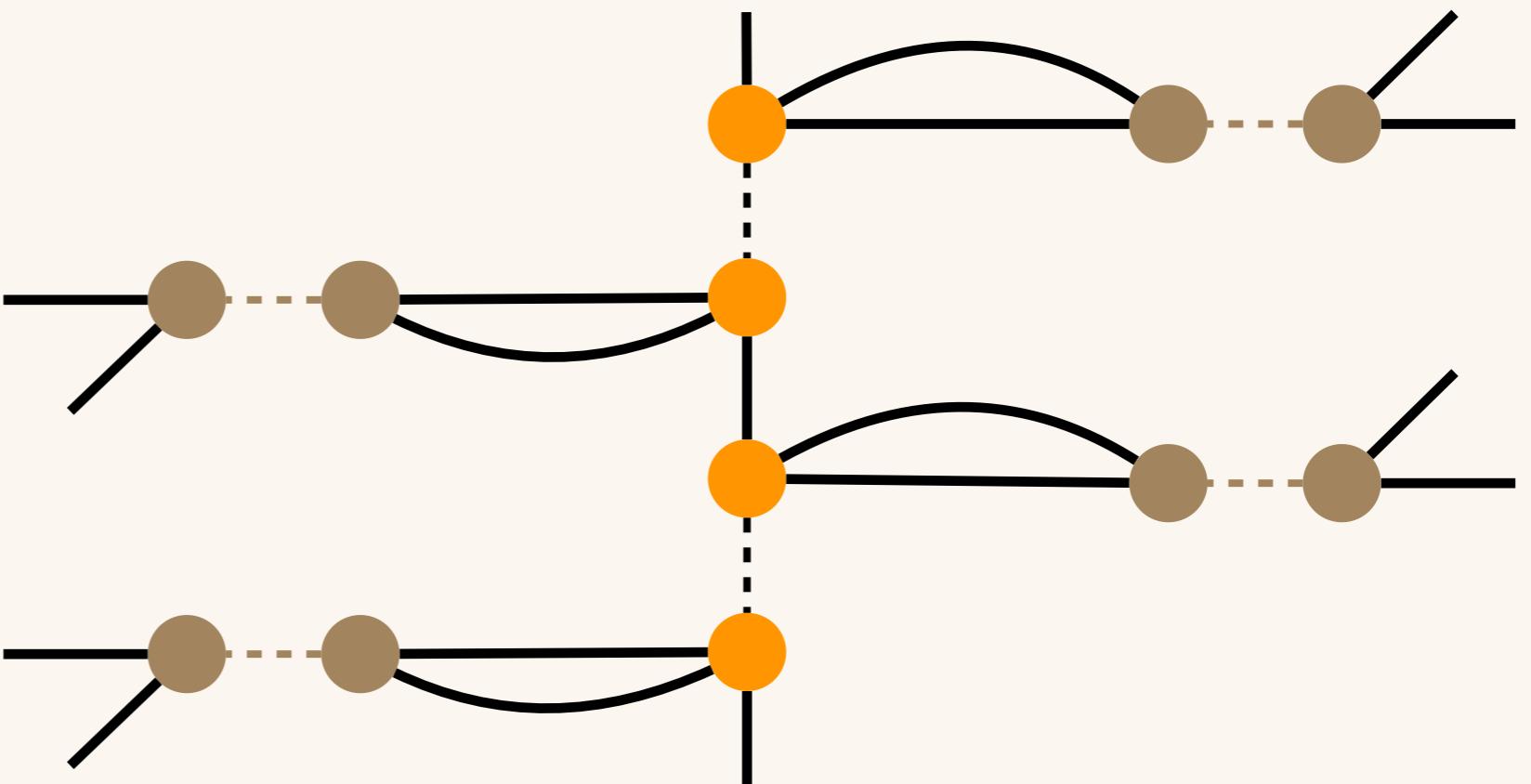
$$U^{(I)} U^{(I)t}$$

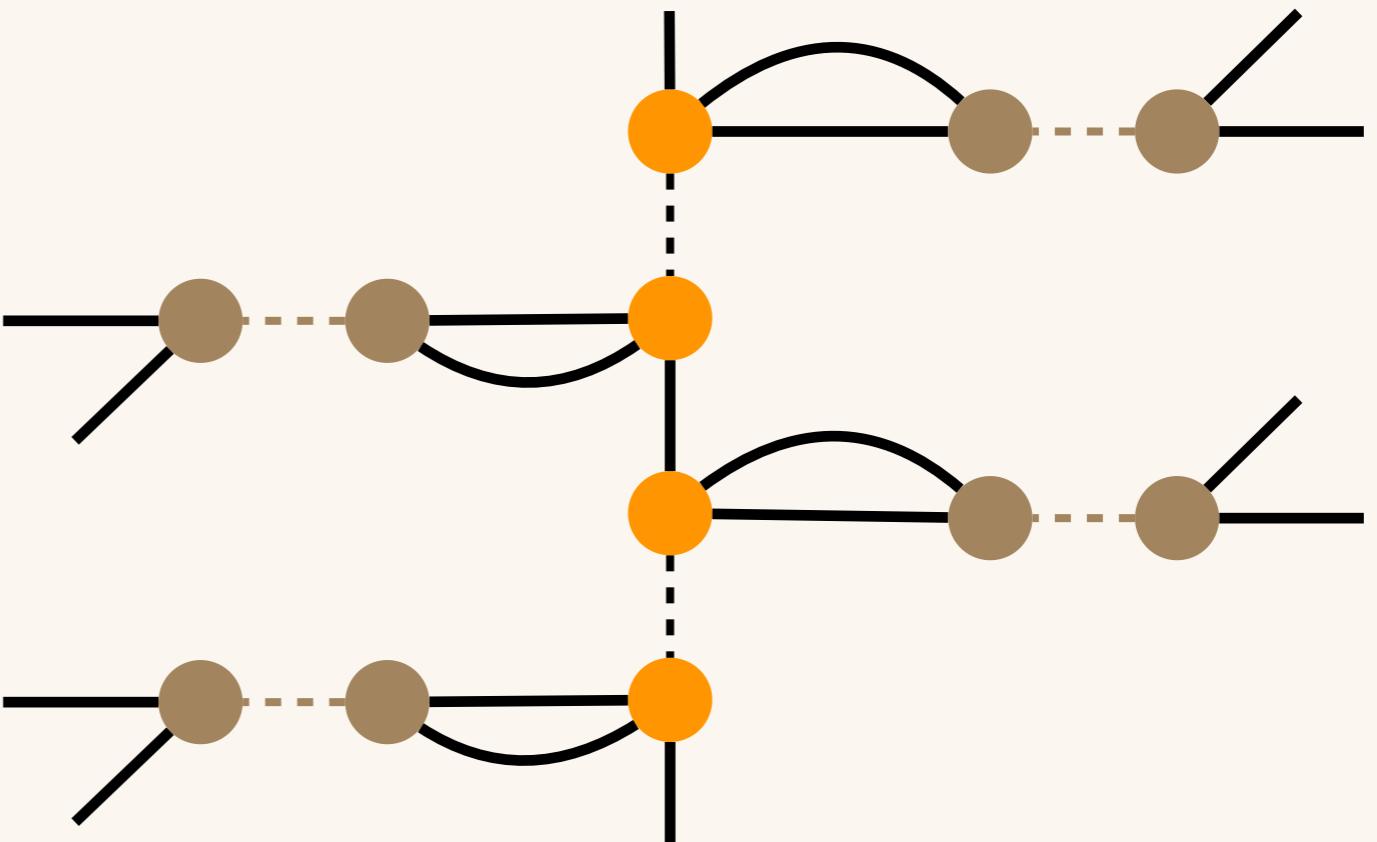


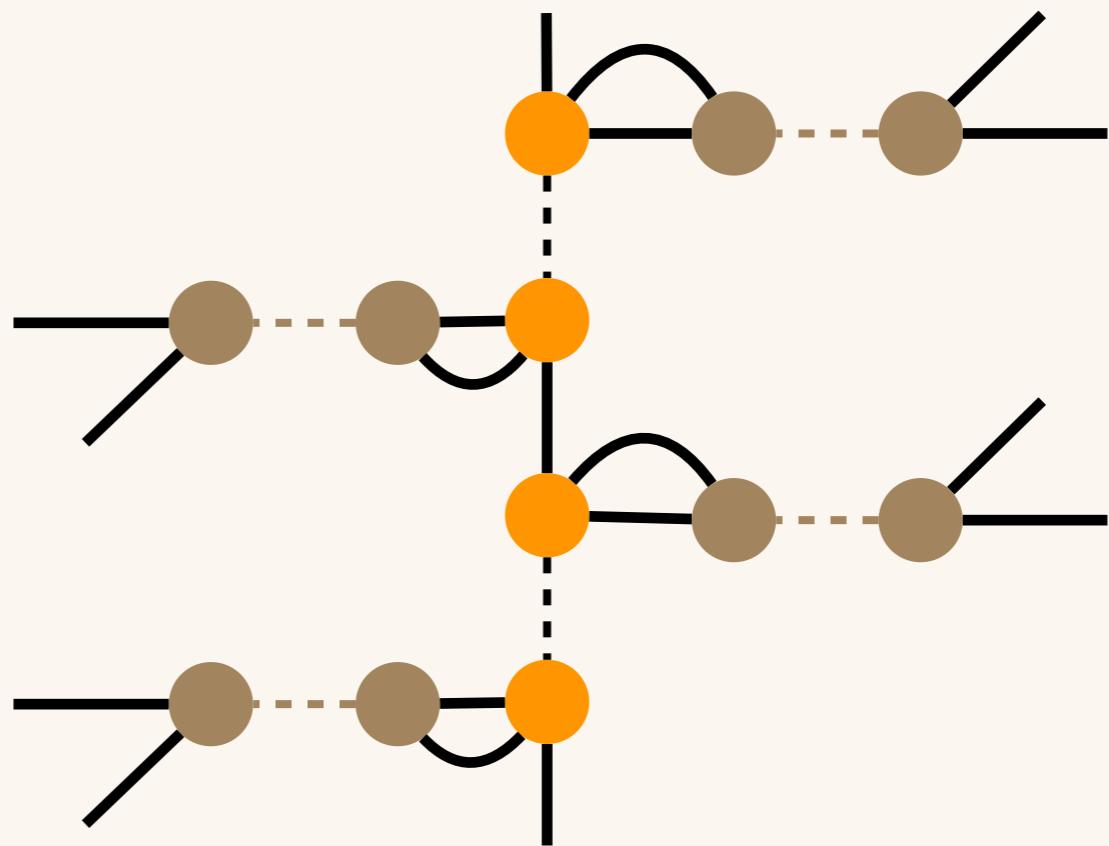
$$\lambda^{(A)}$$

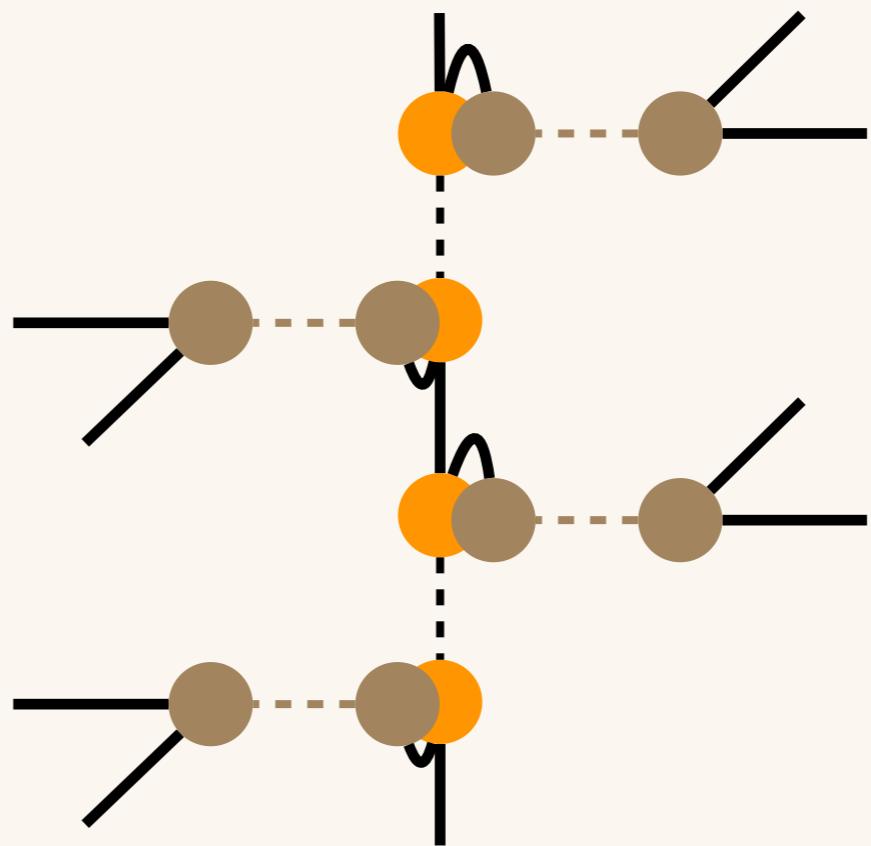
$U^{(I)} U^{(I)t}$  $U^{(L)} U^{(L)t}$  $U^{(M)} U^{(M)t}$  $U^{(P)} U^{(P)t}$

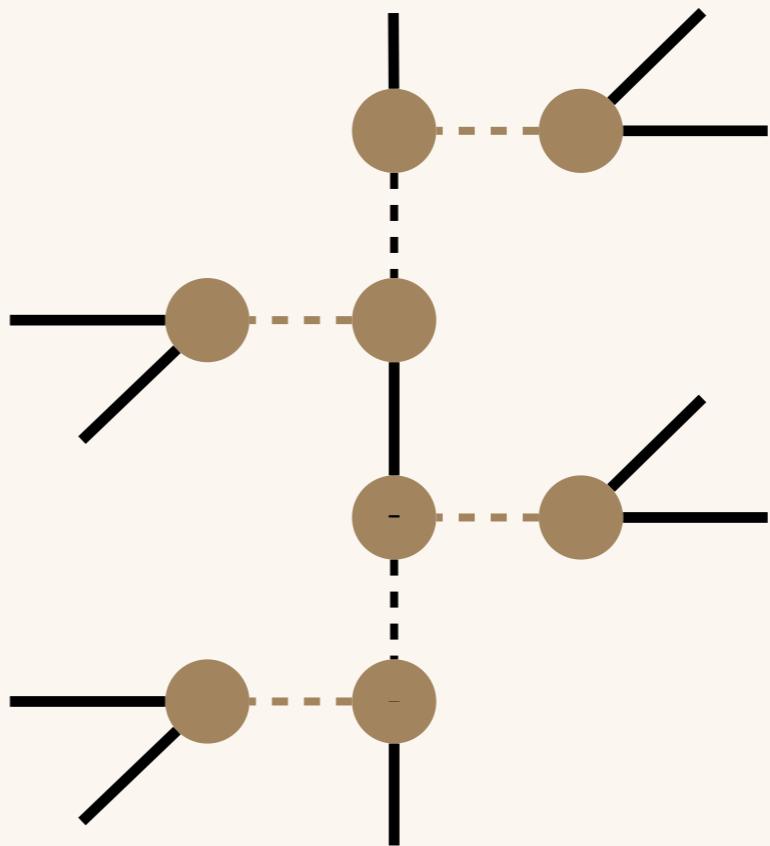


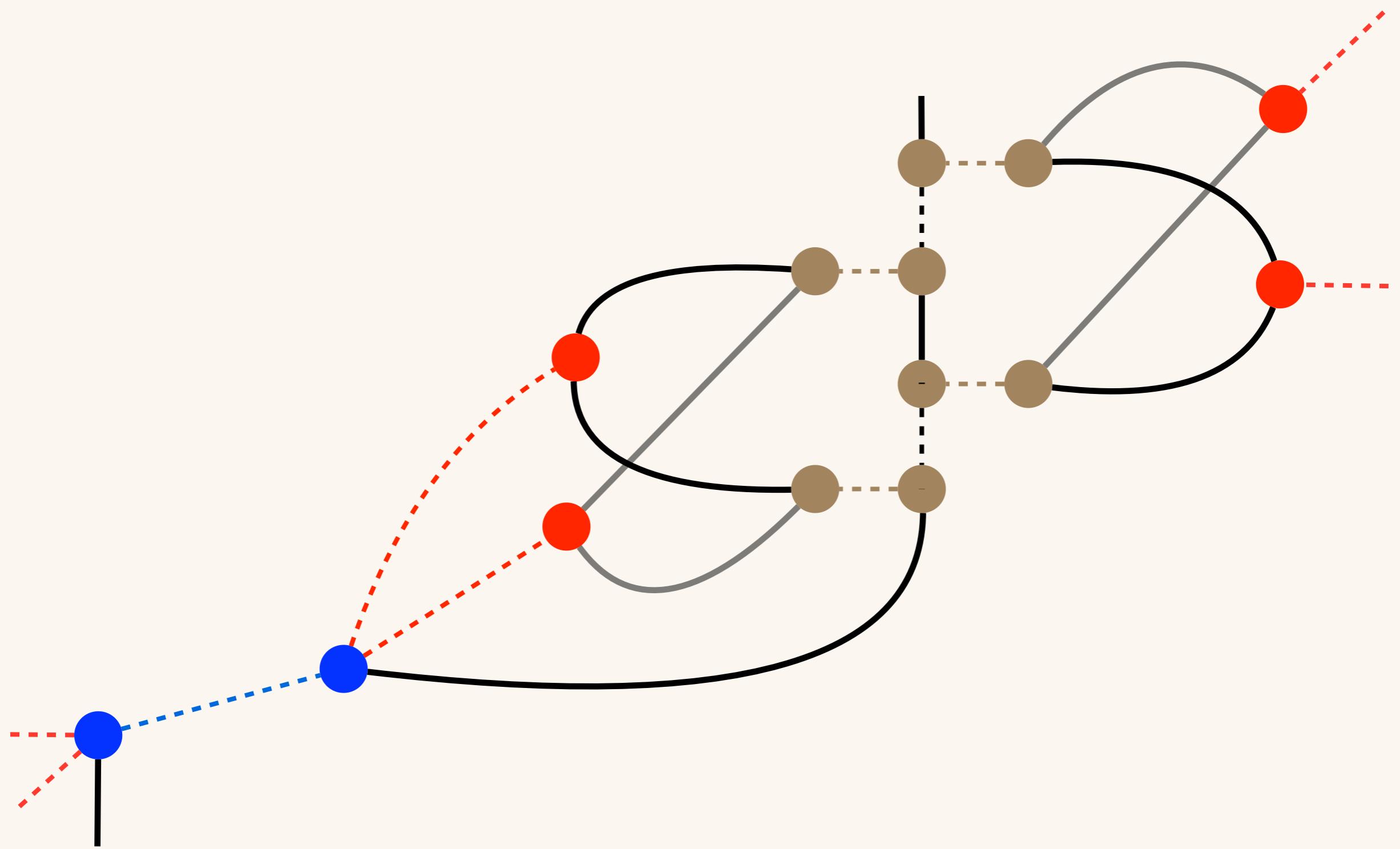


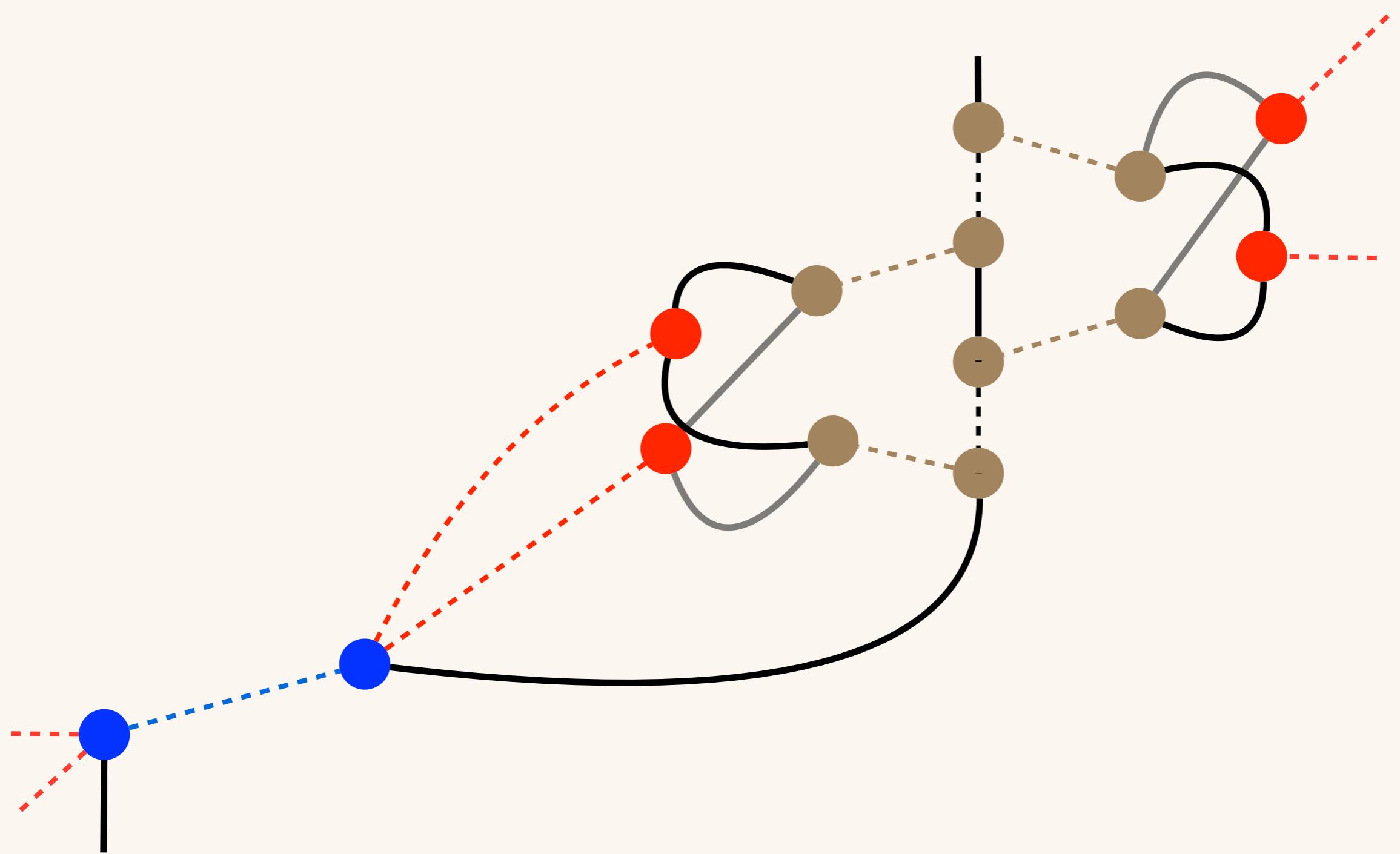


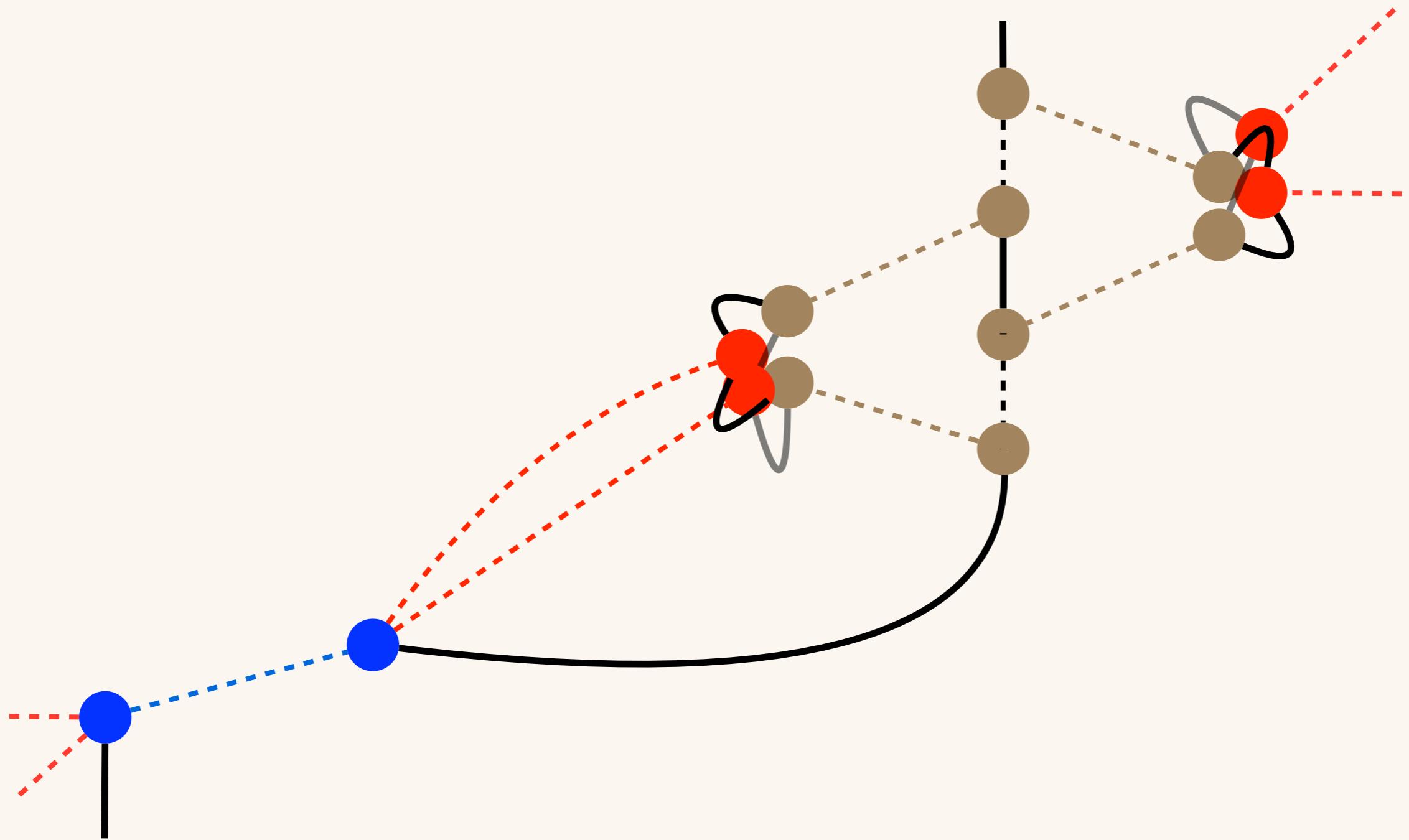


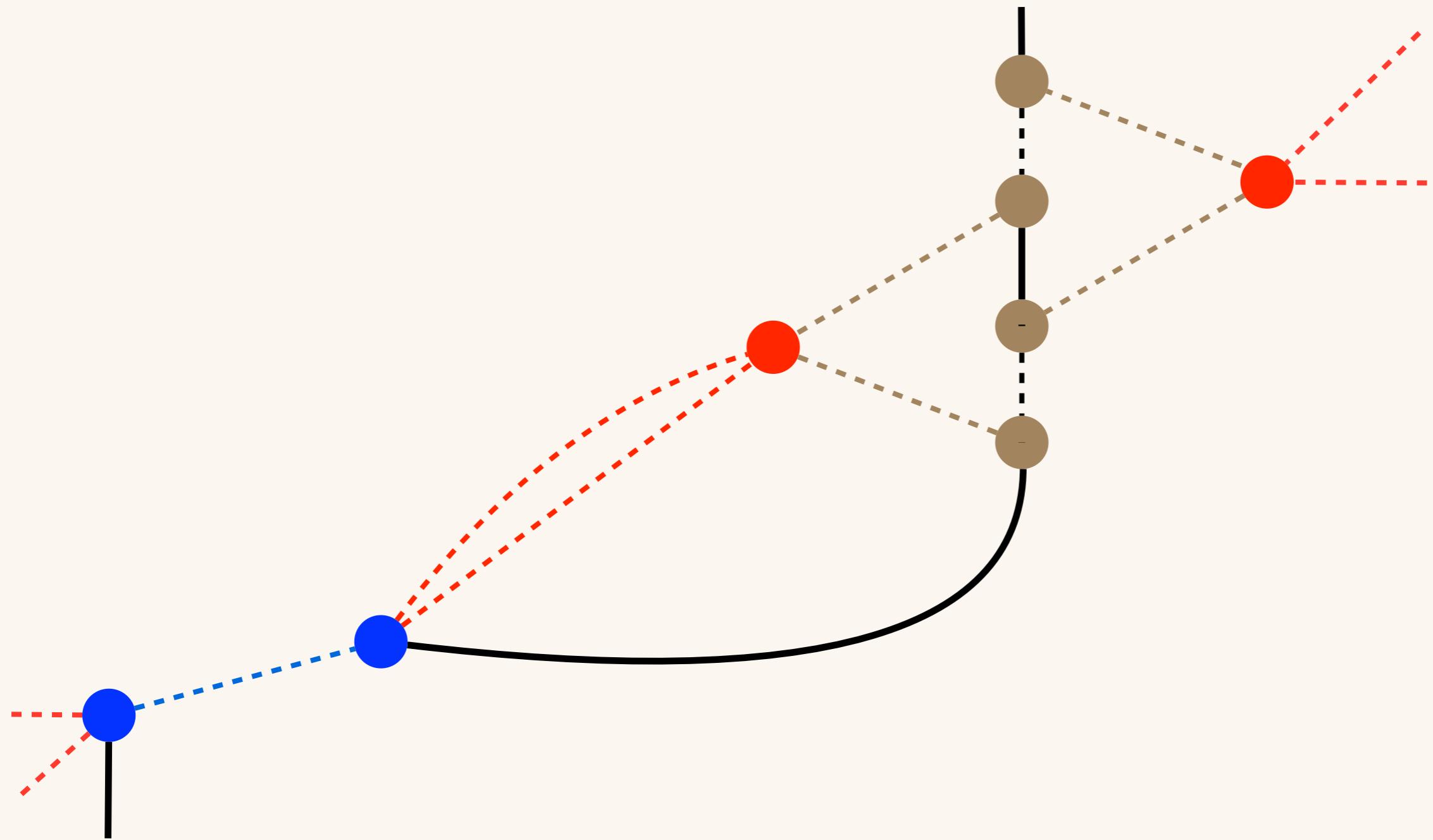


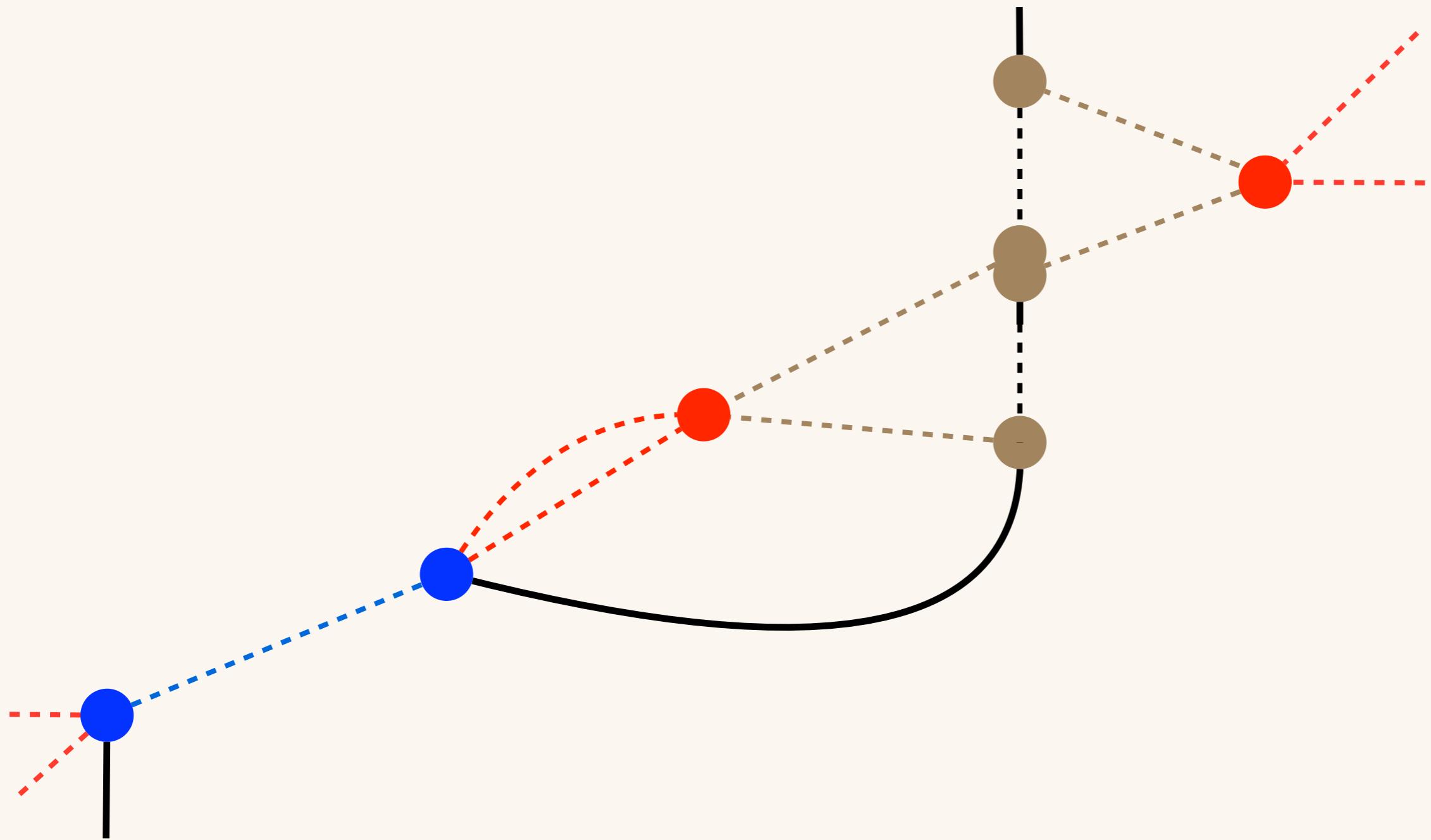


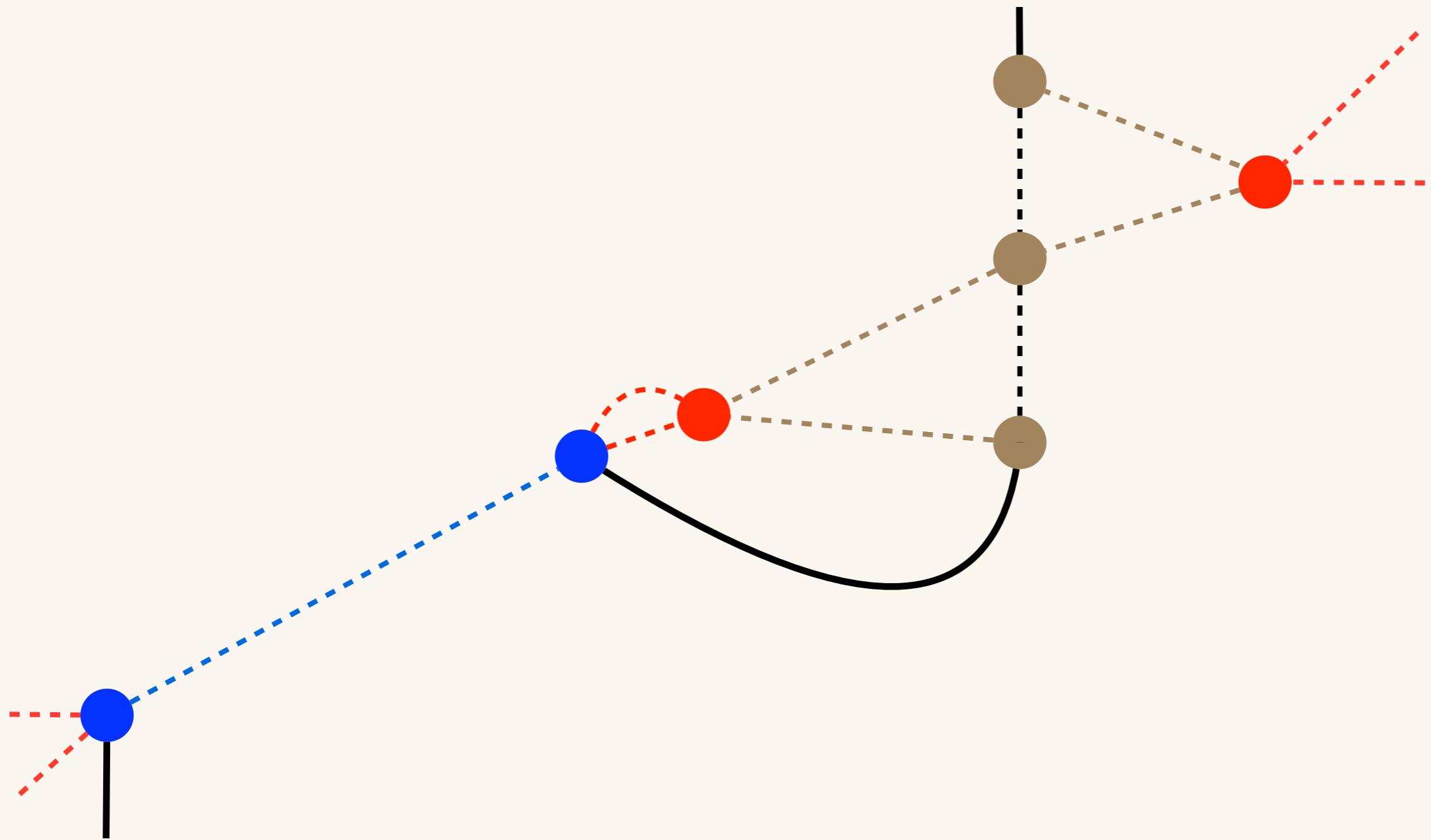


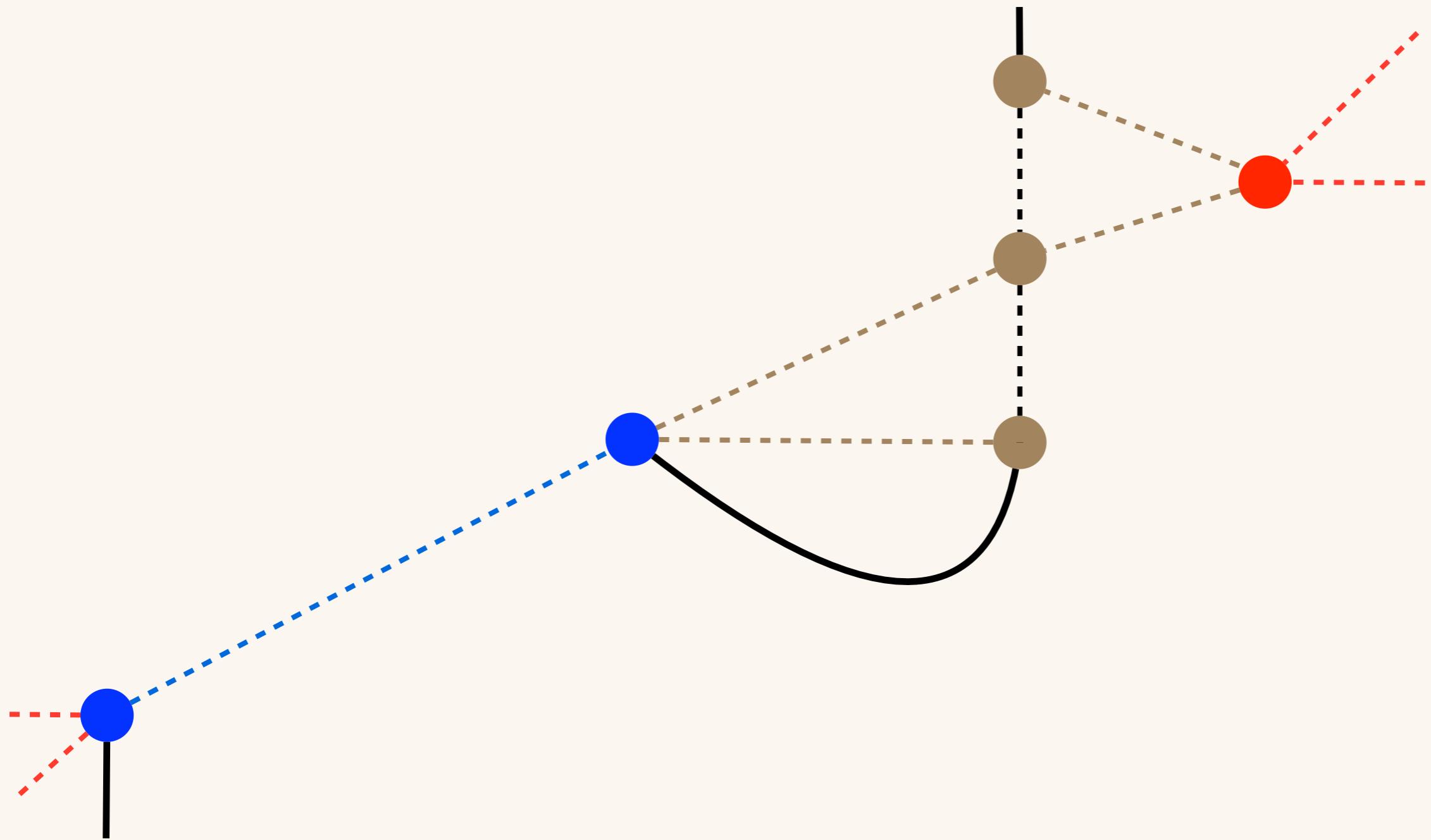


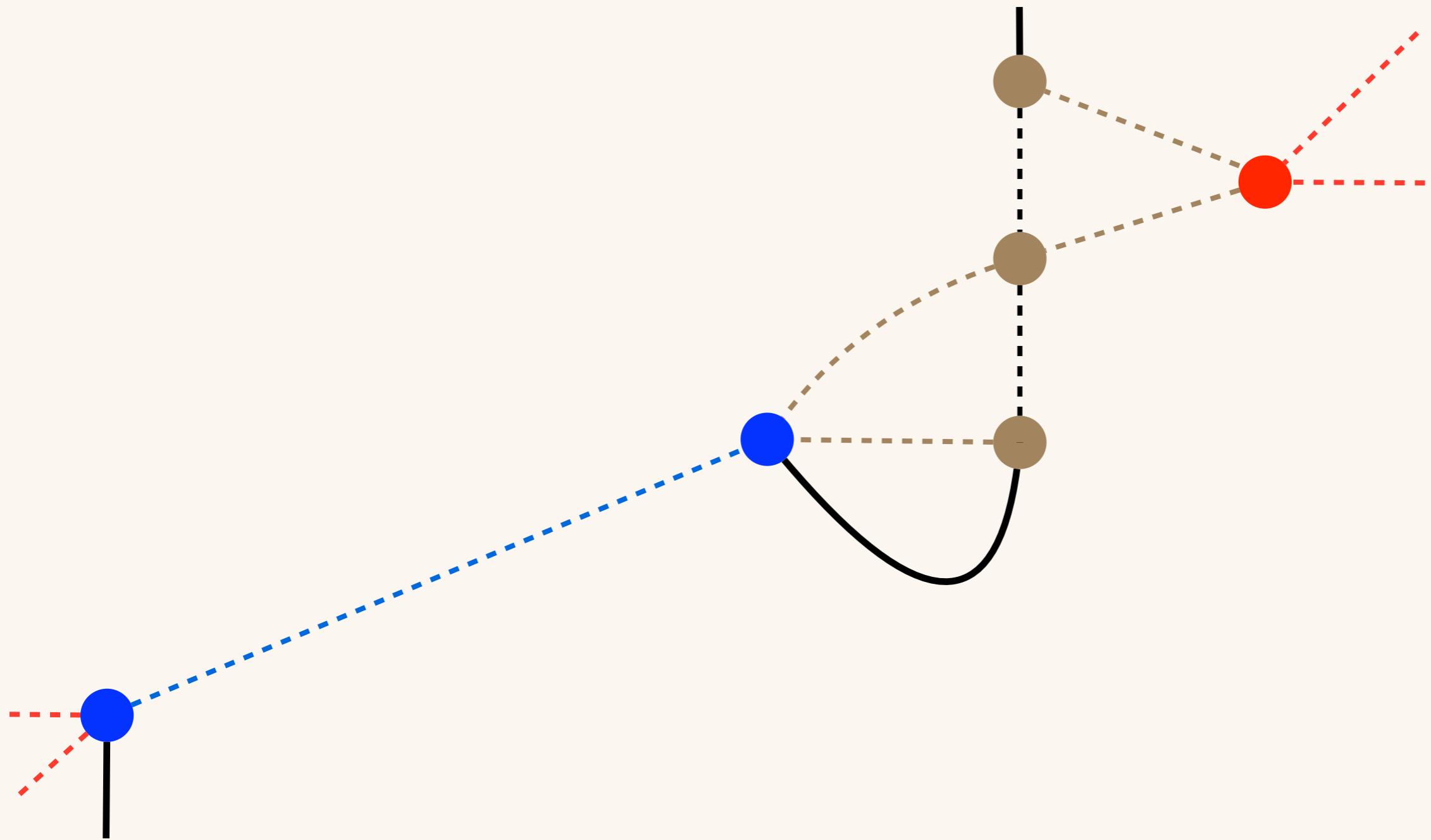


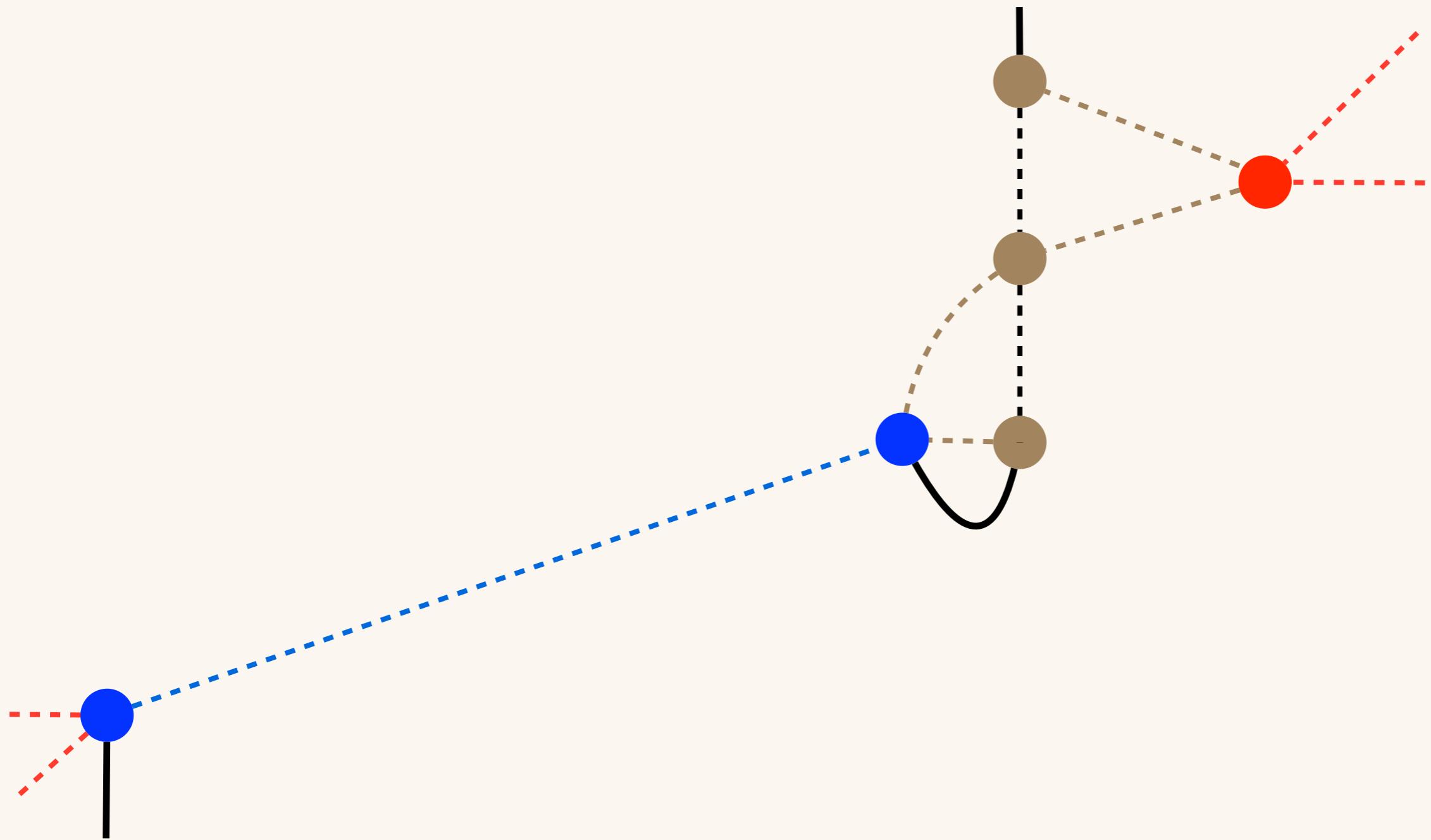


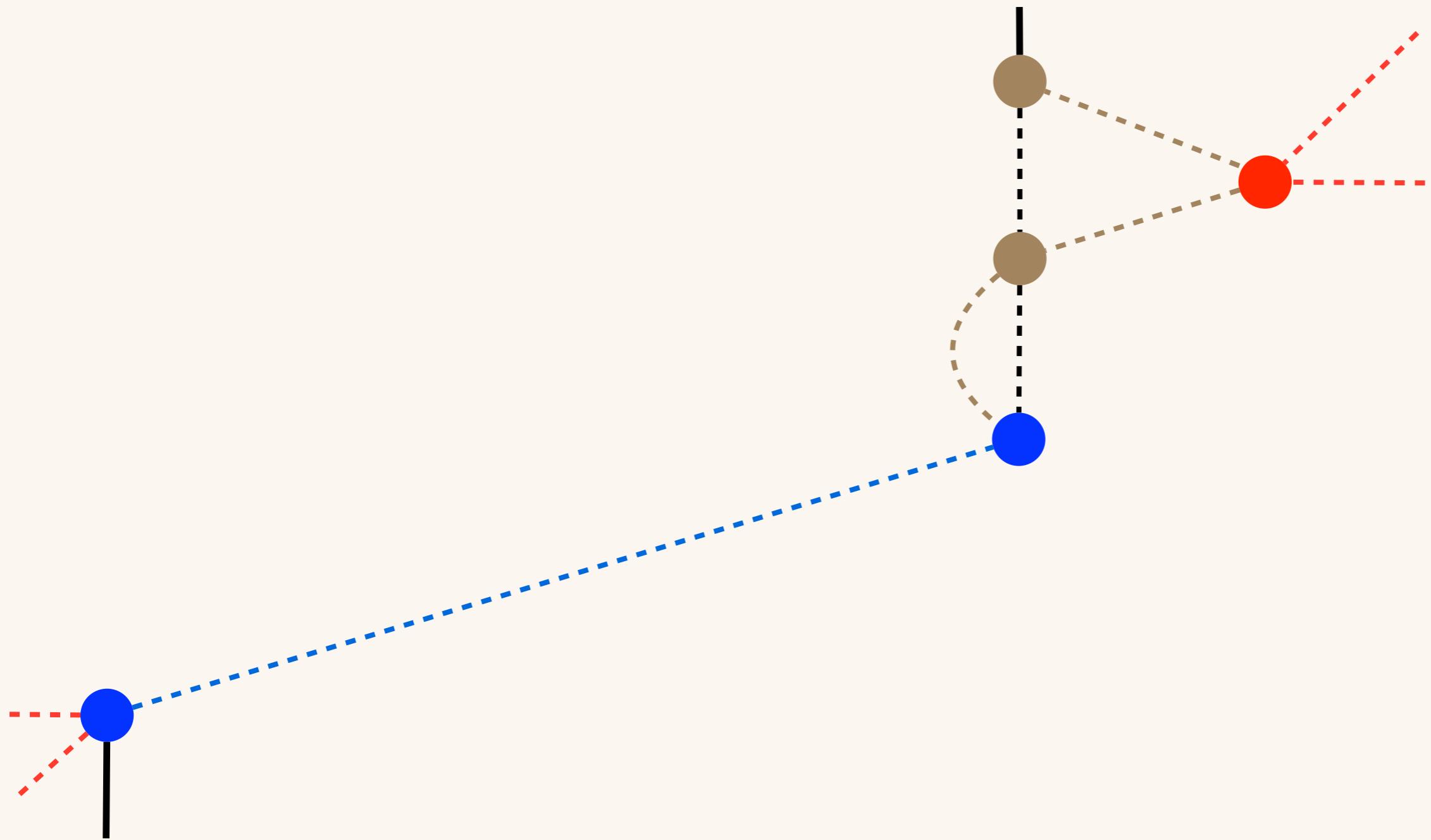


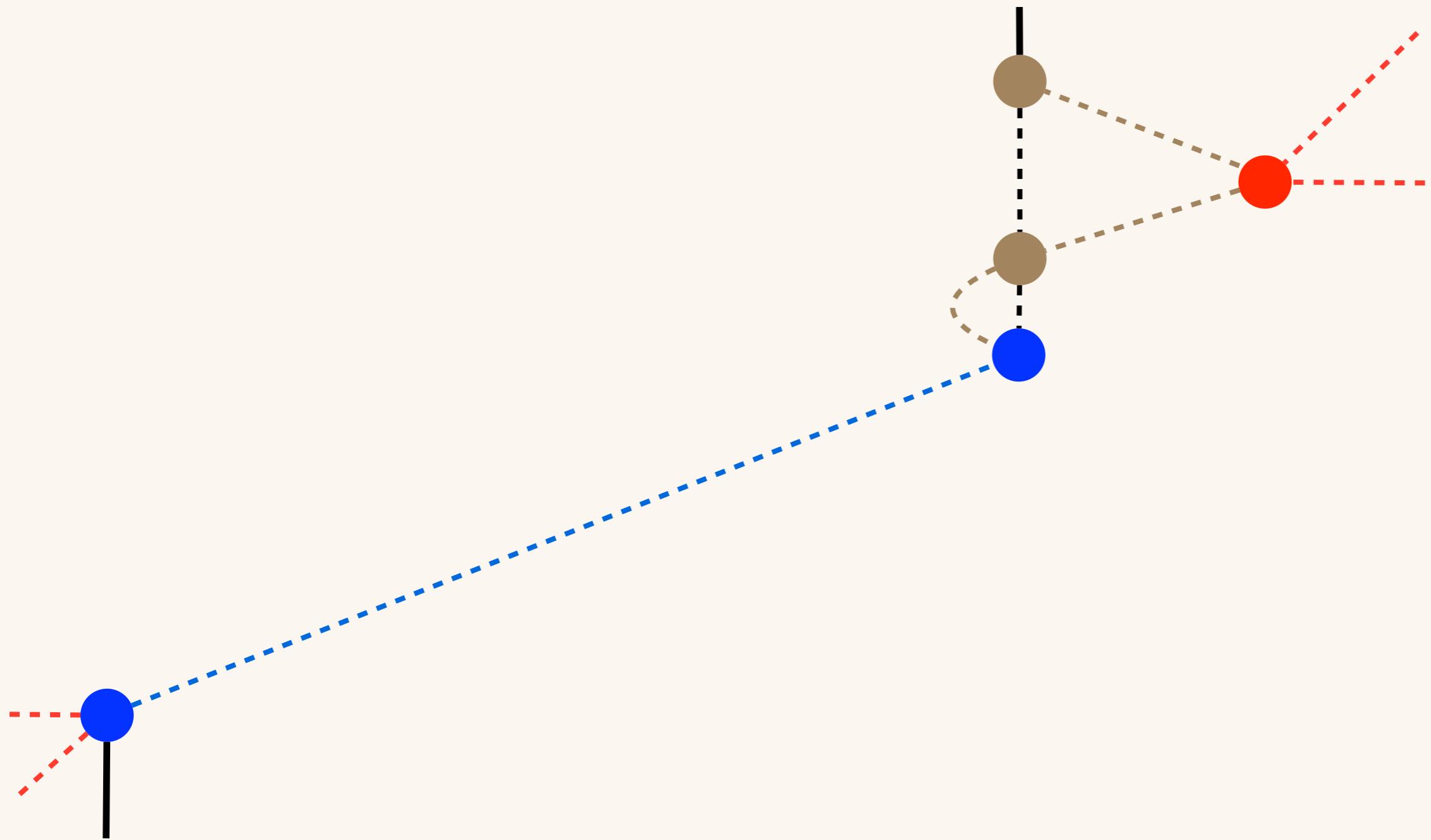


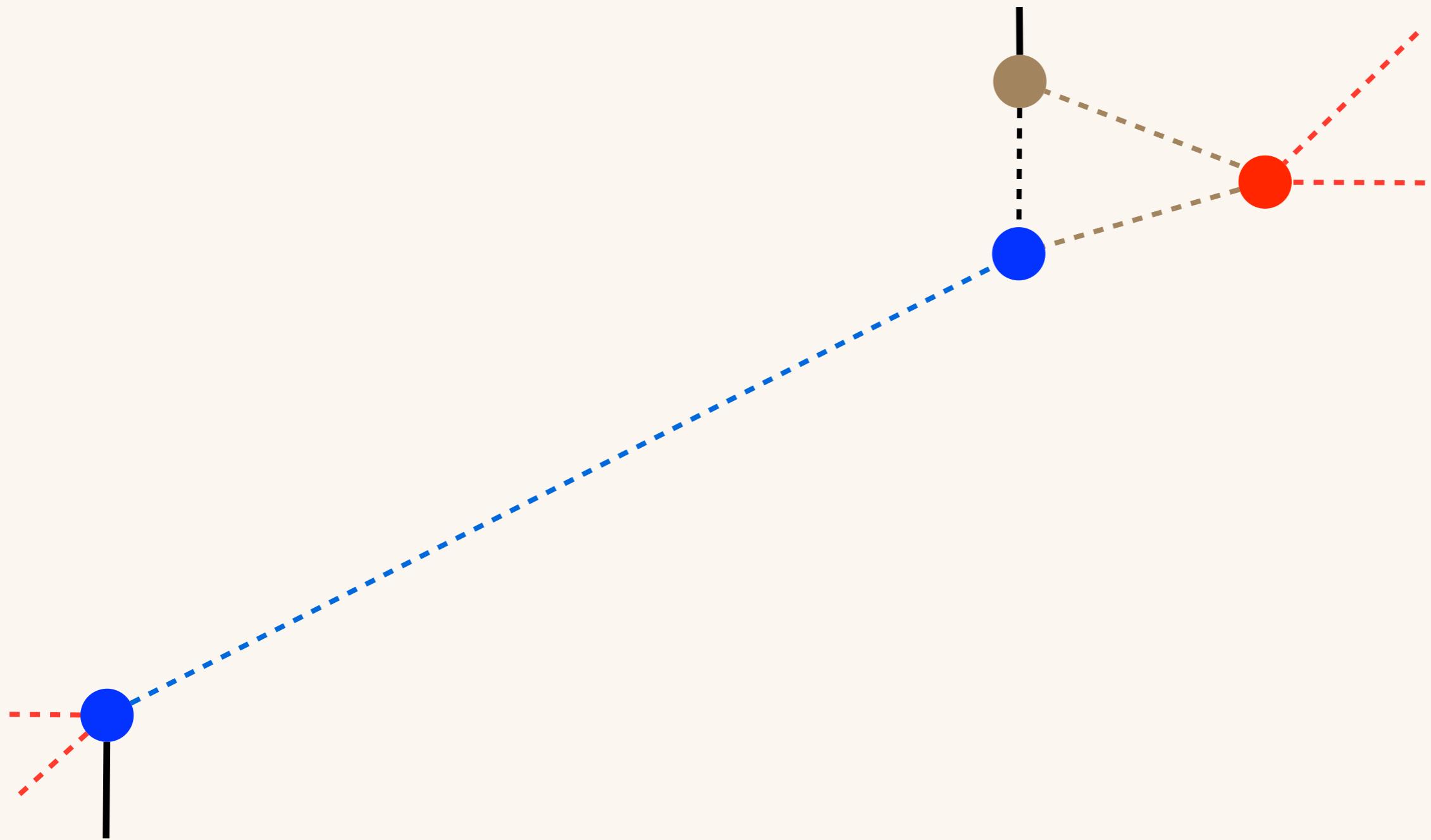


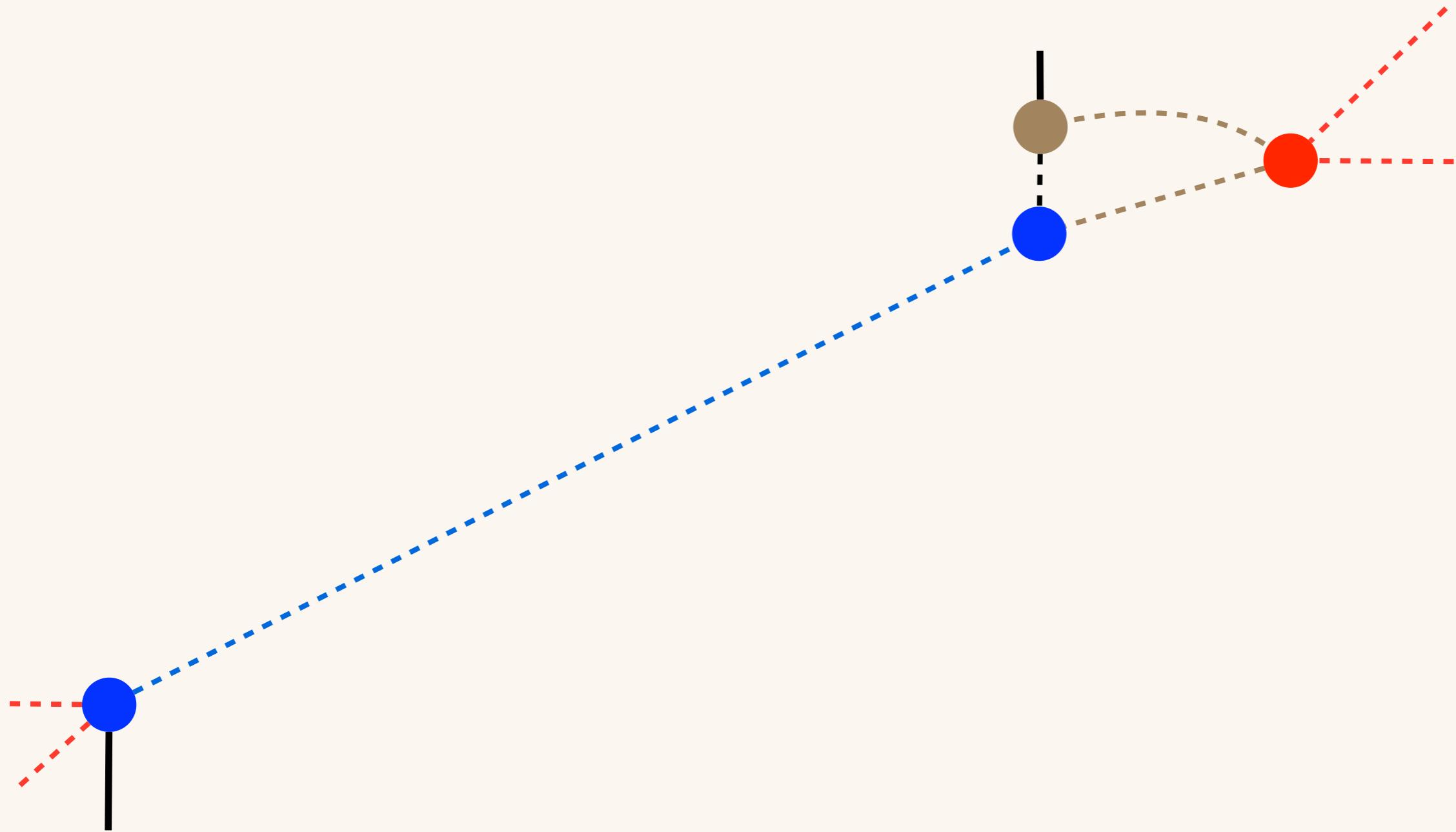


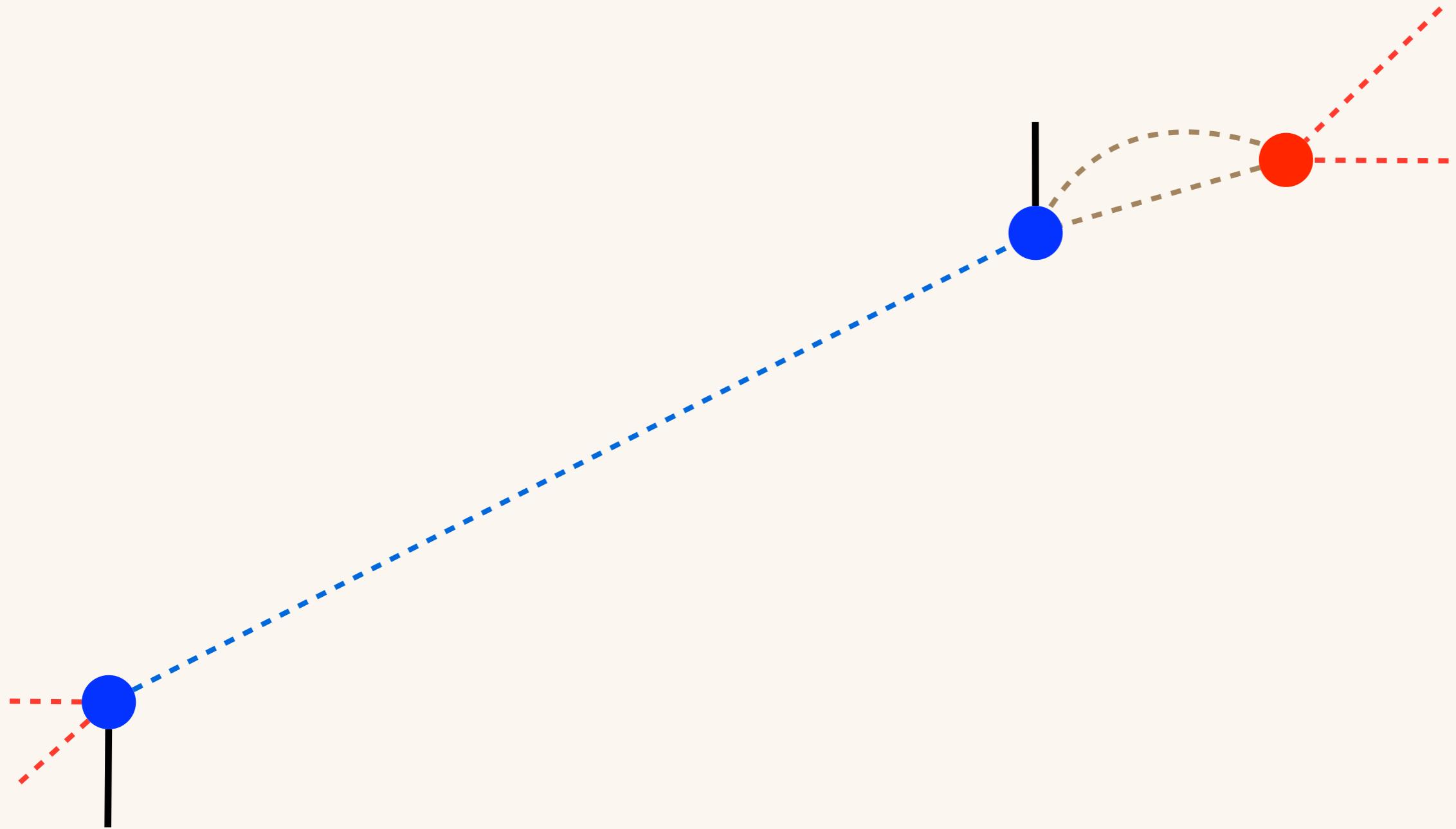


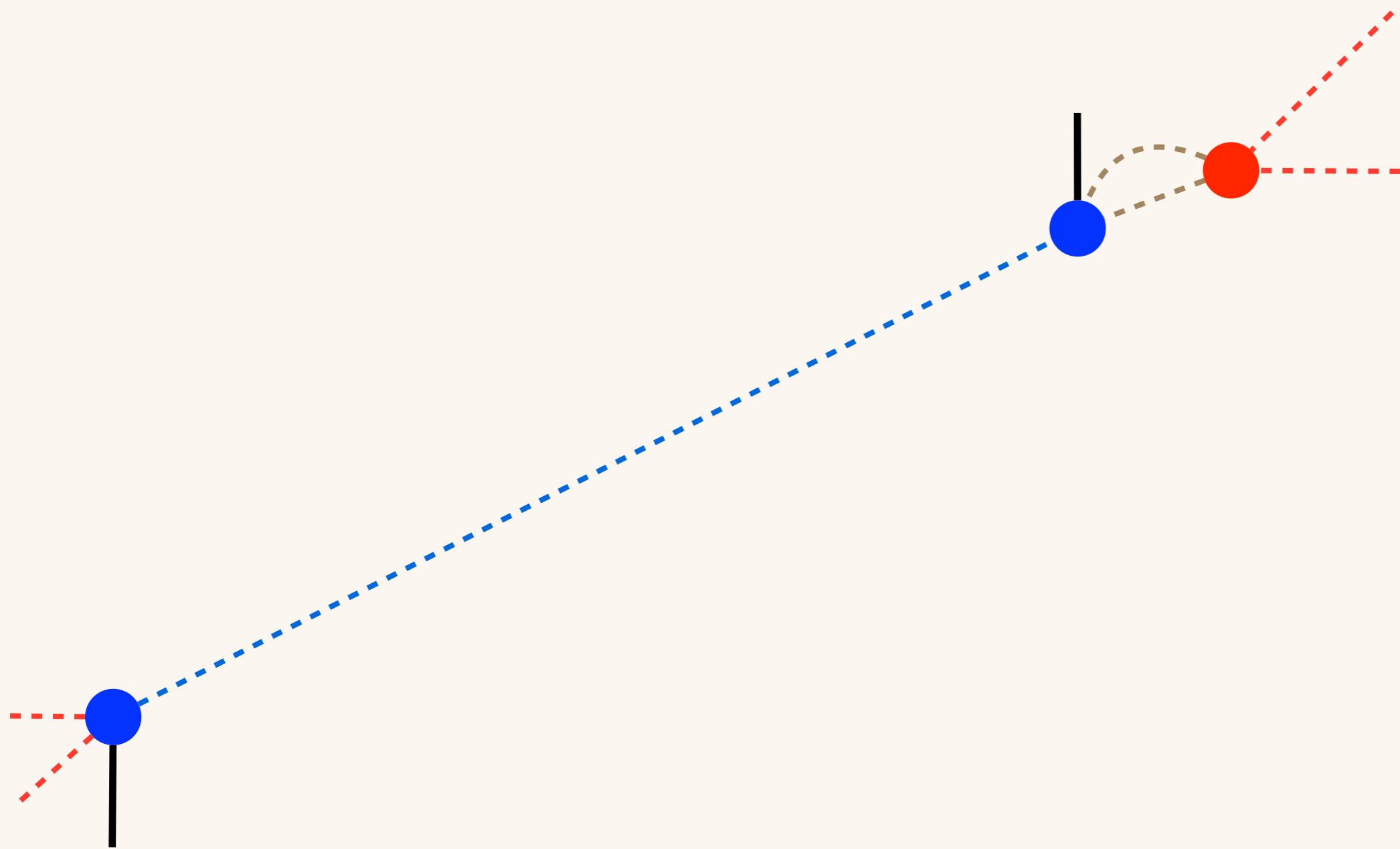


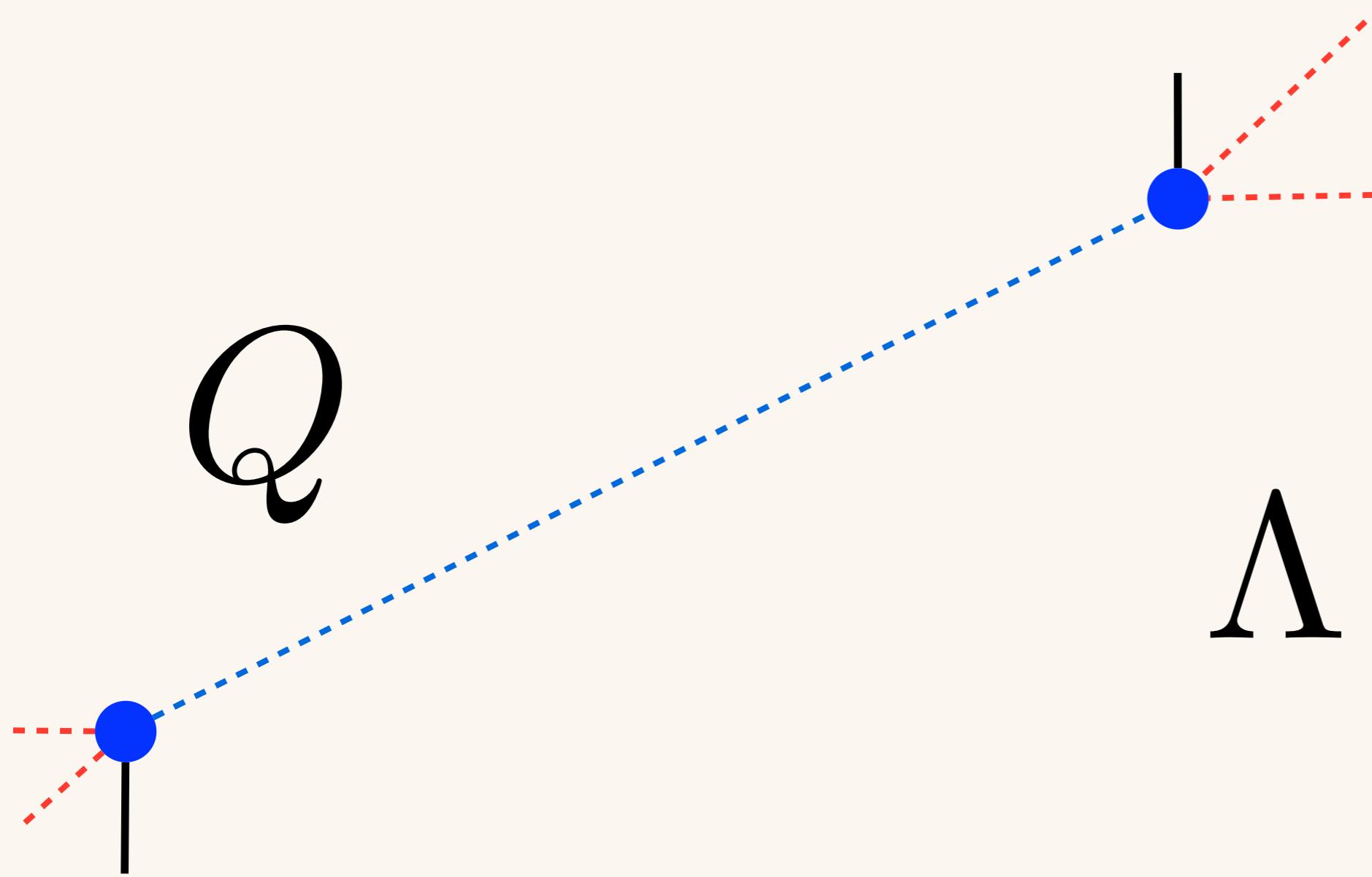




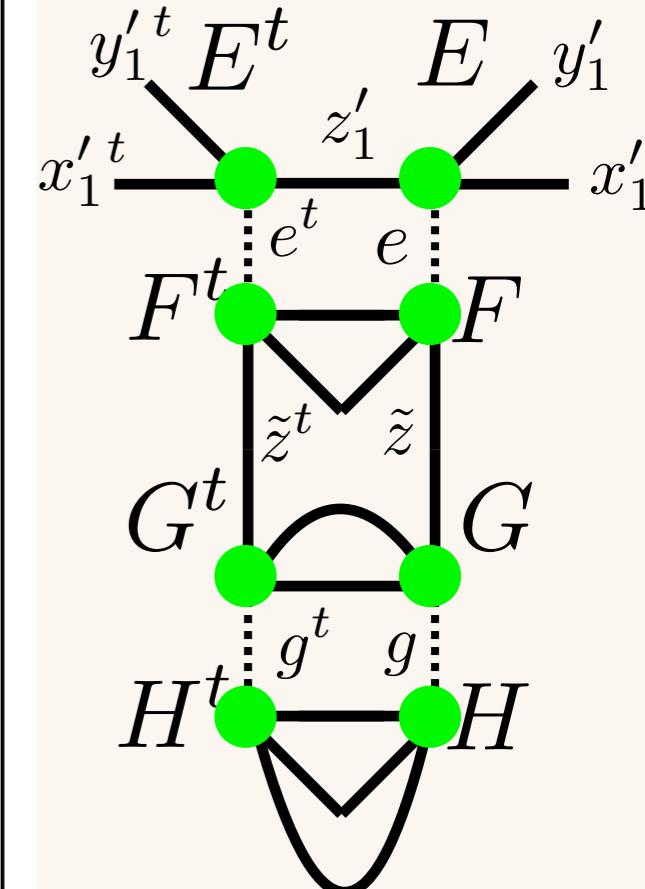
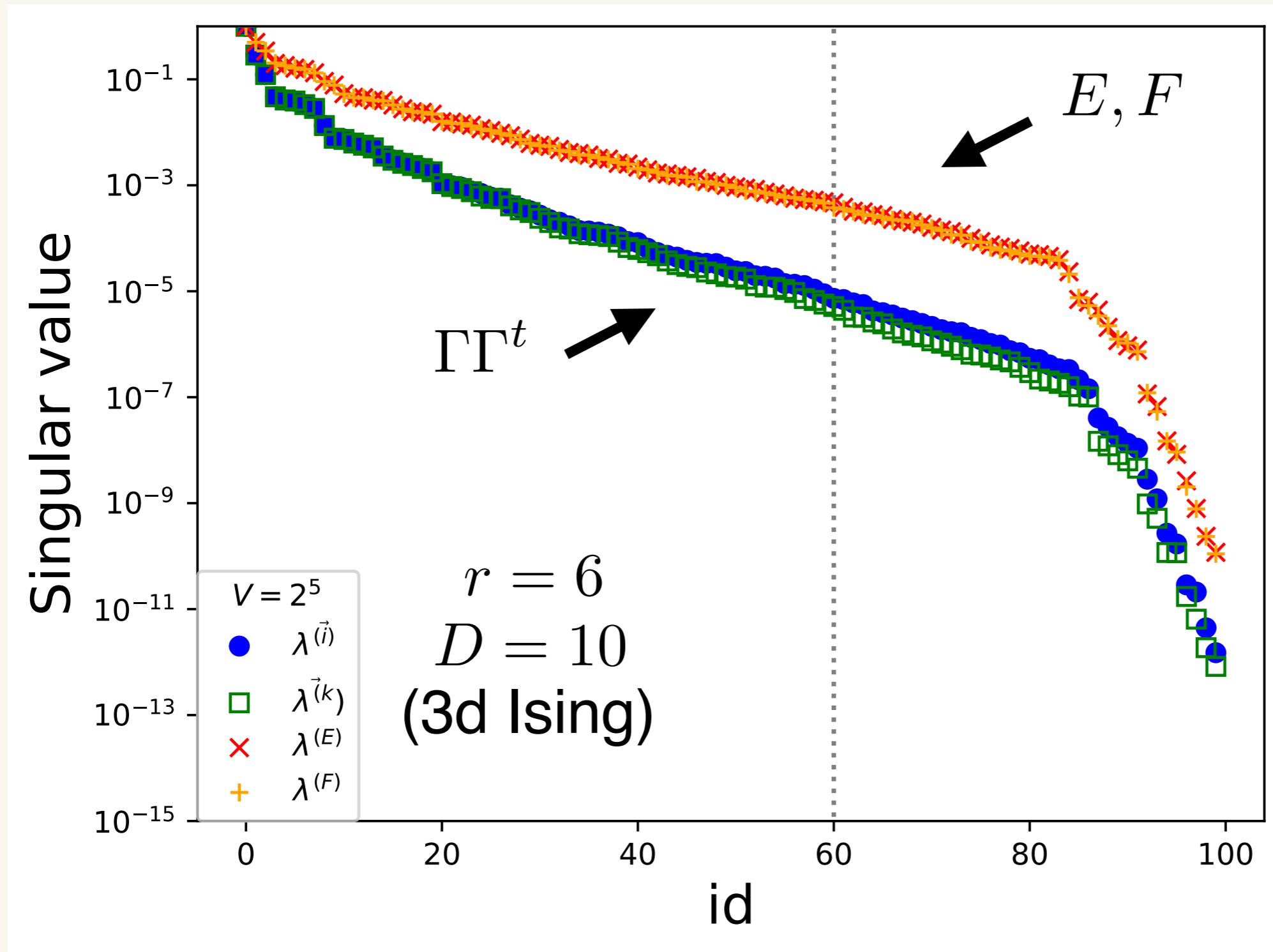








● Isometry for unit-cell tensor



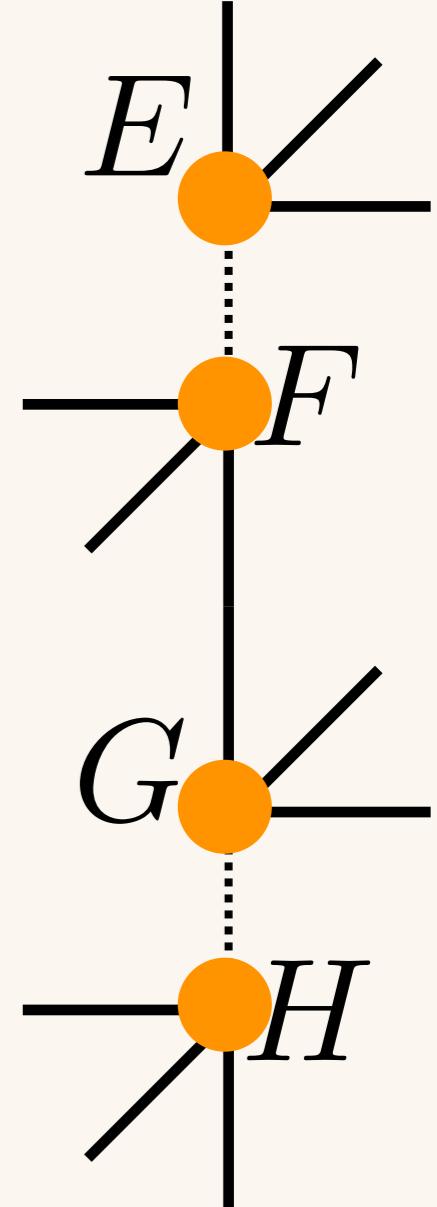
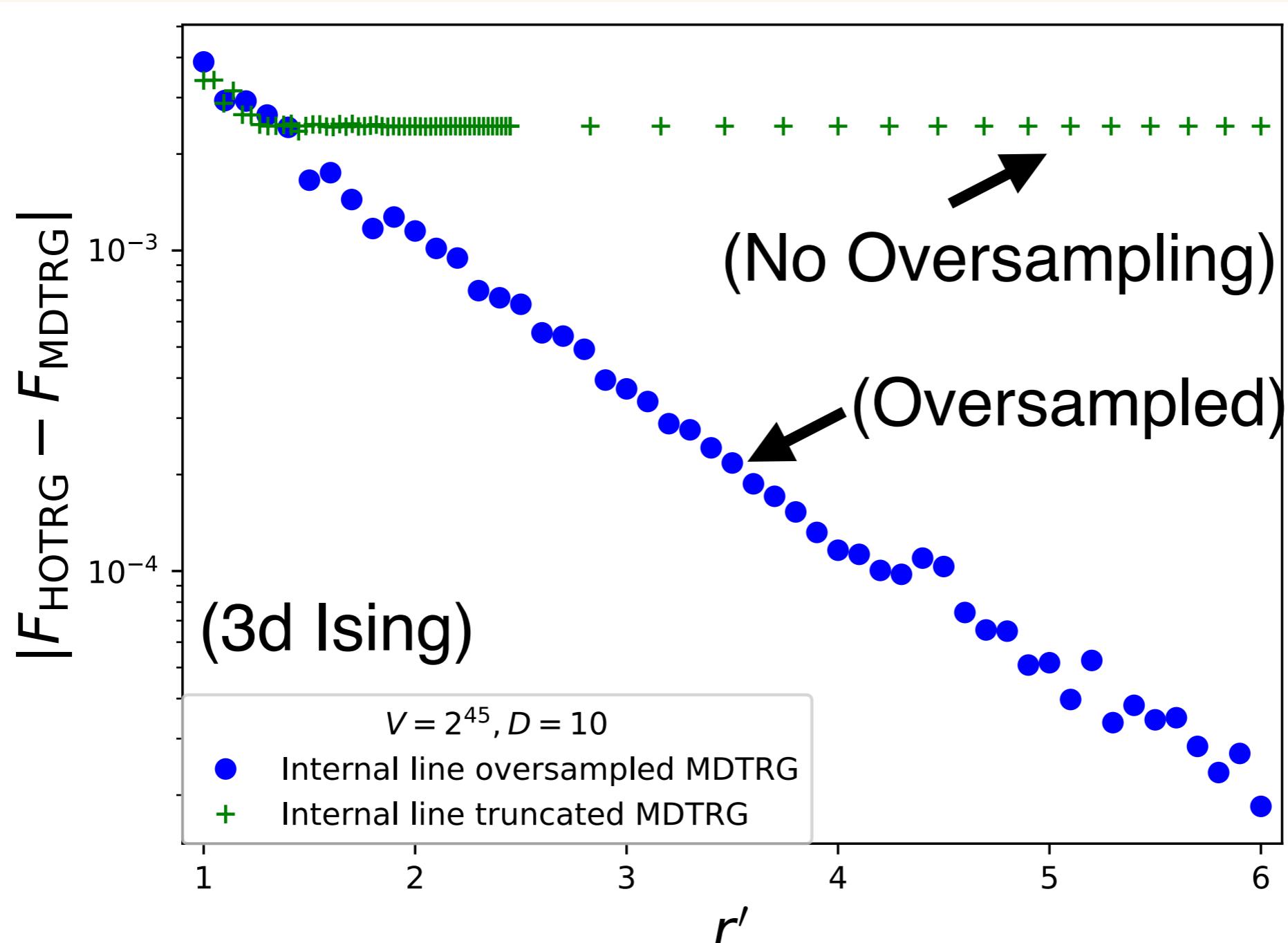
→ More global isometry reduces systematic error.

● 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.

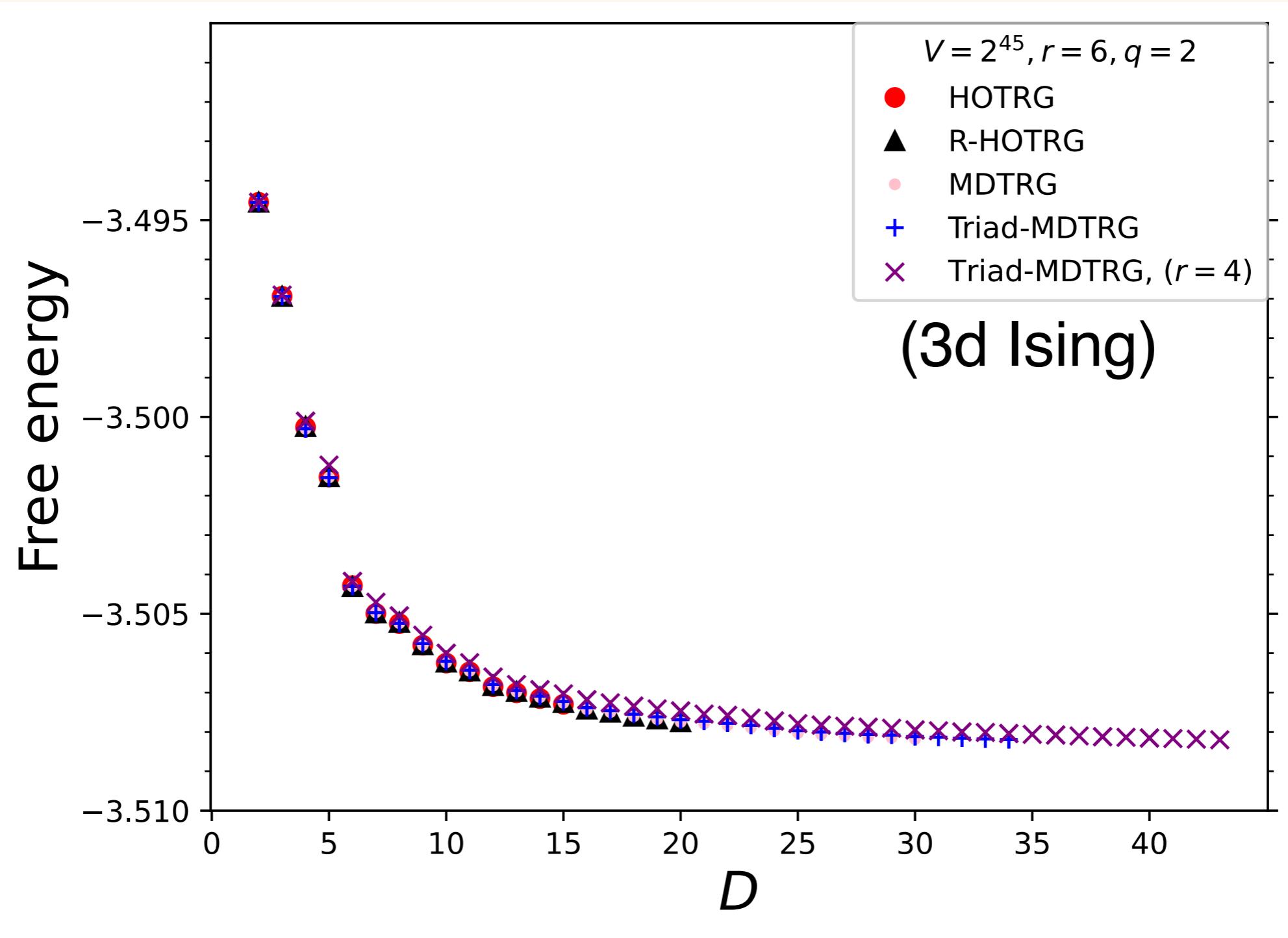
● Internal line oversampling



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

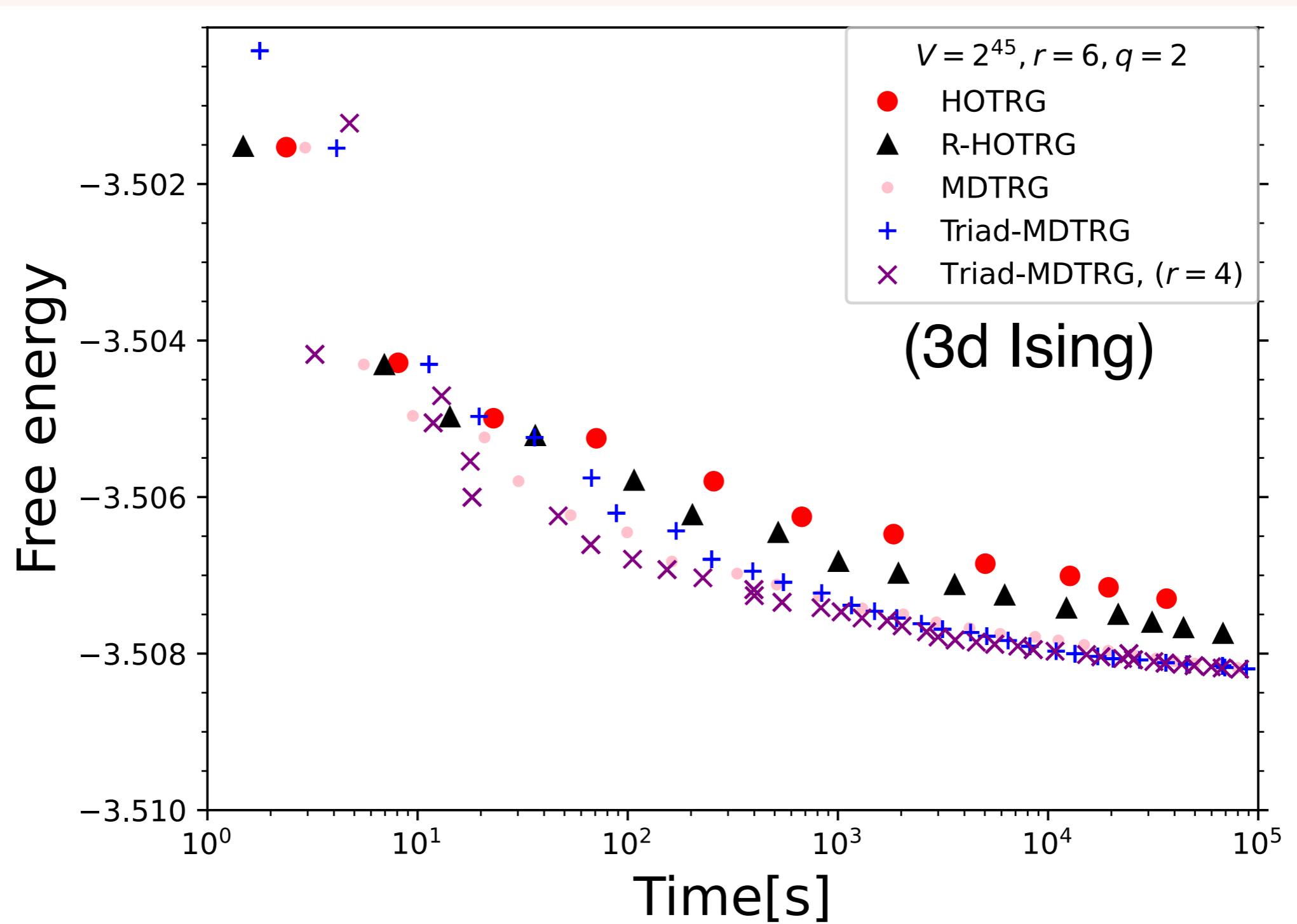
→ Internal lines (dotted lines) have to be oversampled ($D \rightarrow rD$) to reduce the systematic error from R-SVD.

Free energy density of 3d-Ising model



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

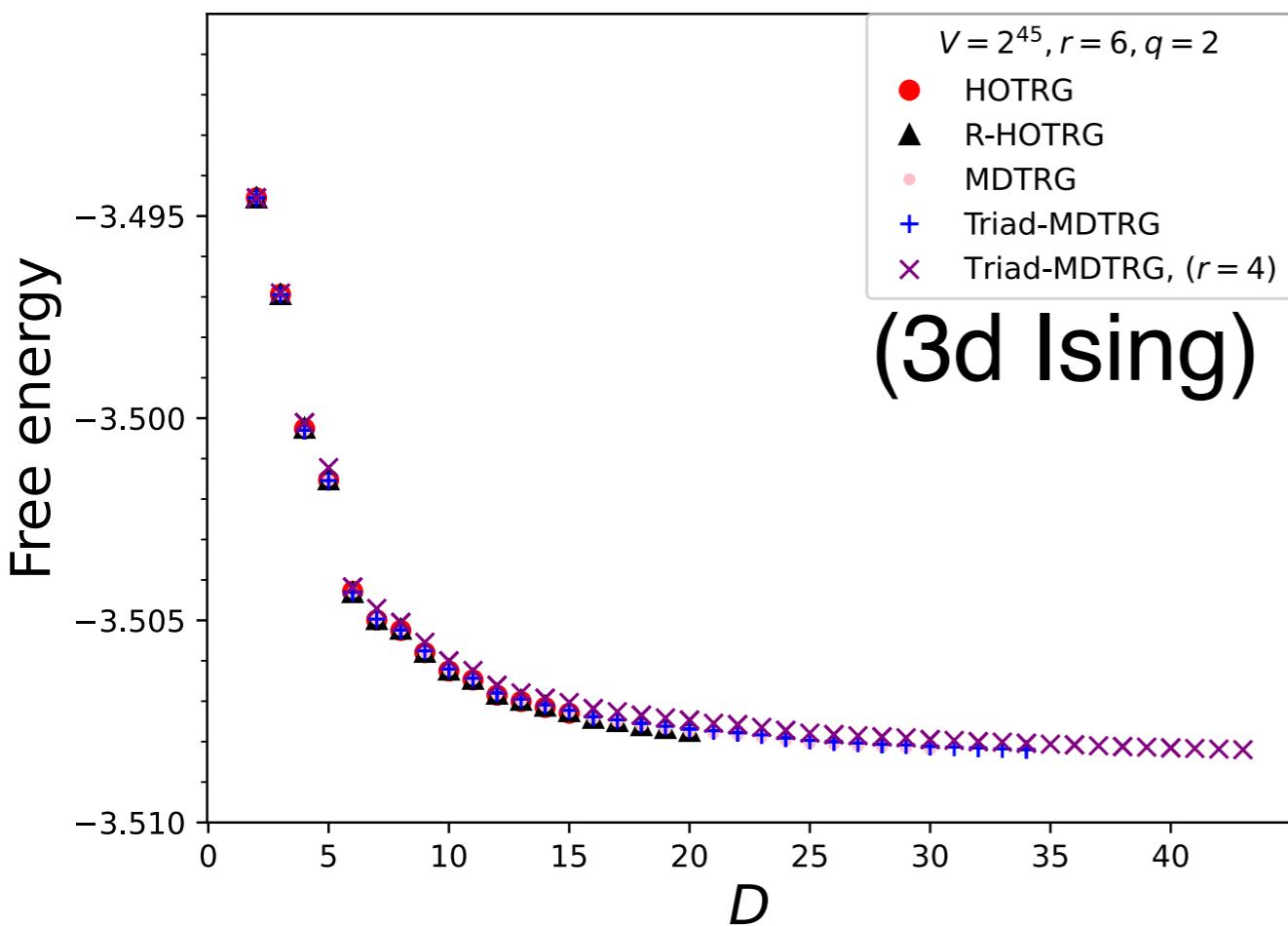
Free energy density of 3d-Ising model



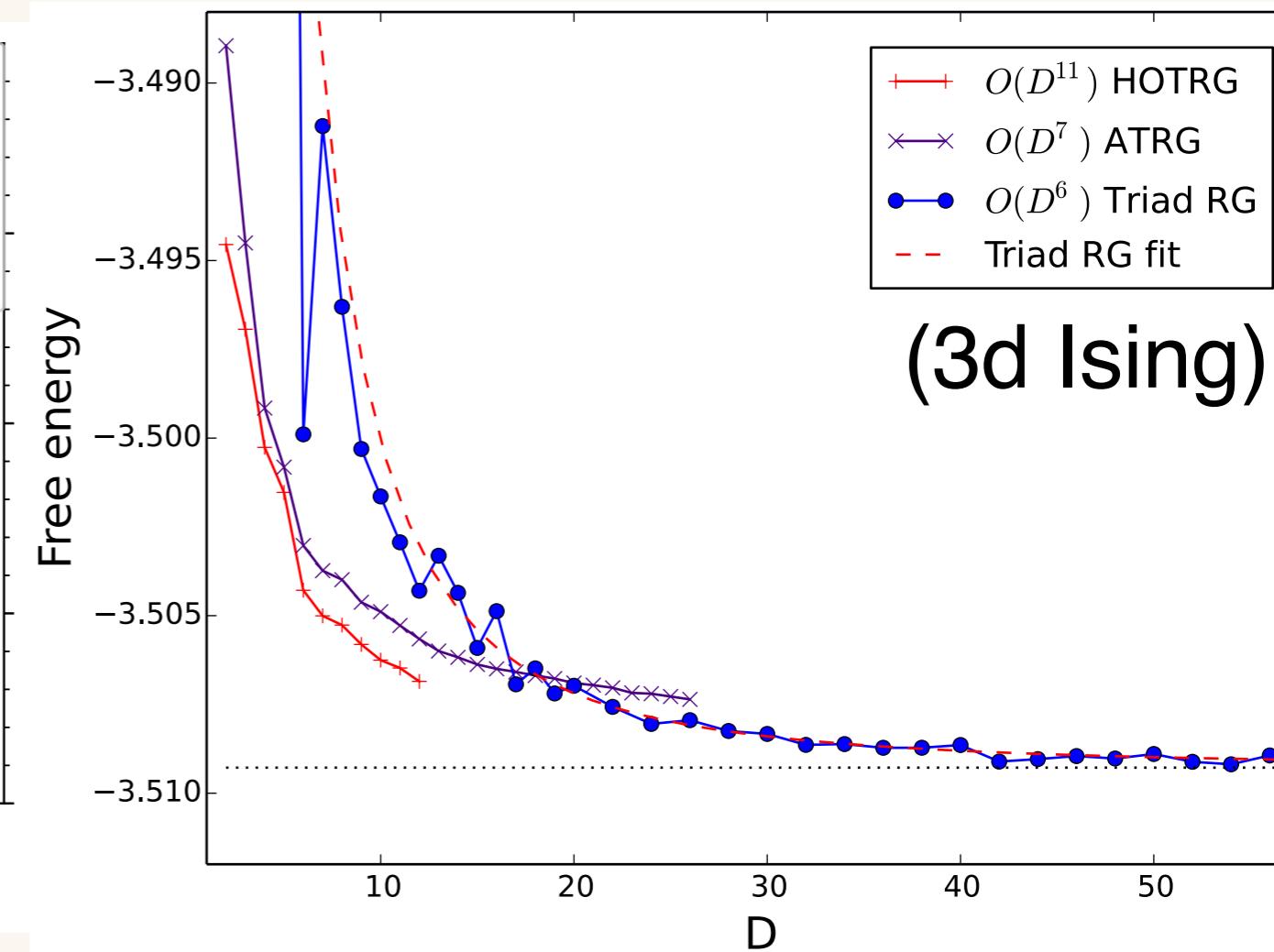
→ R-HOTRG, MDTRG, Triad-MDTRG results are consistent with HOTRG (additional systematic error is not dominant).

Free energy density of 3d-Ising model

[K.N. arXiv:2307.14191]

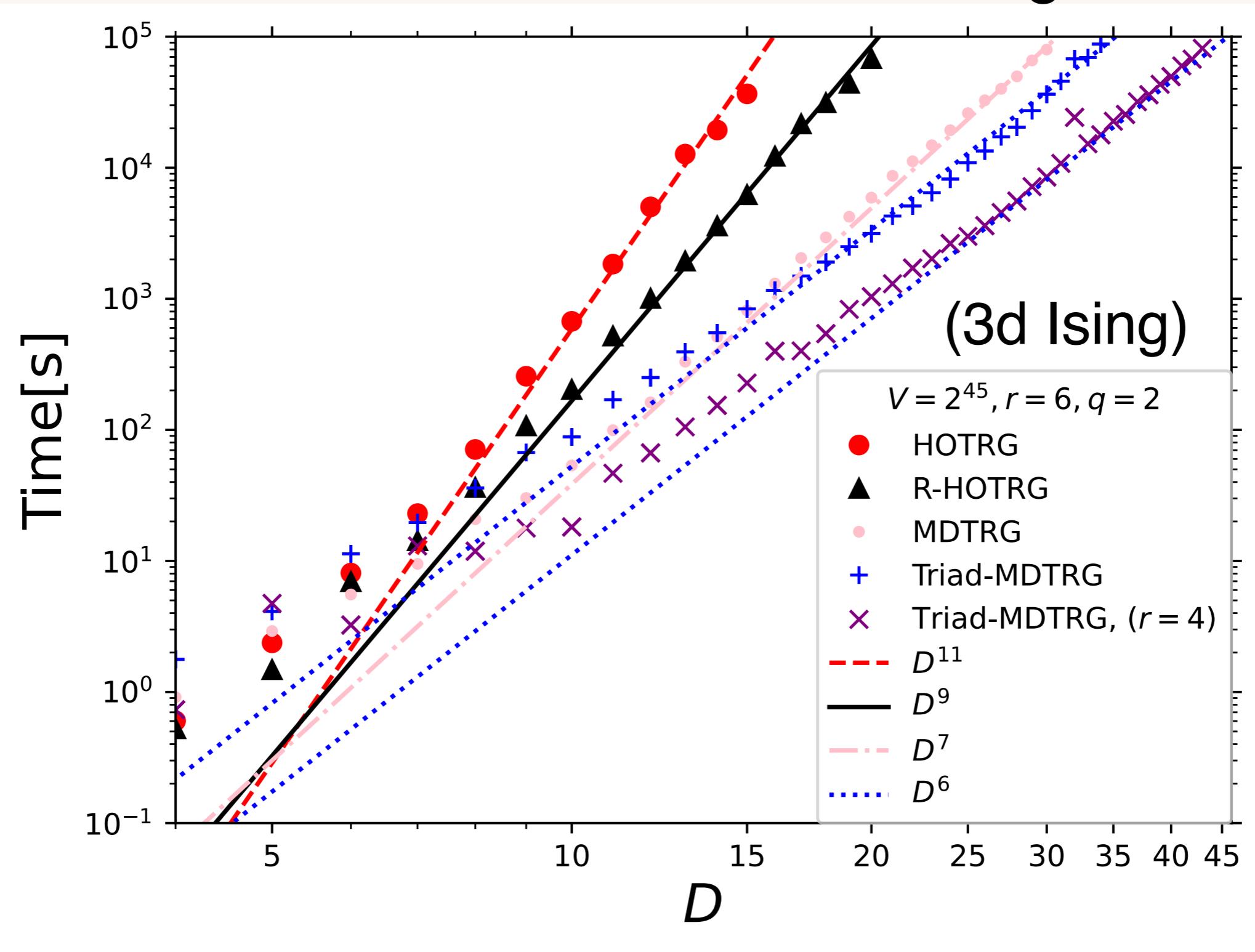


[D. Kadoh, K.N. arXiv:1912.02414]



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

● 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	$O(D^{3d})$	$O(D^{4d-1})$	$\nleq 2d$
◊ MDTRG	$O(D^{2d+1})$	$O(D^{3d})$	$\not\leq d+1$
◊ Triad-MDTRG	$O(D^{d+3})$	$O(D^{3d})$	$\not\leq d+1$

- ◊ 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.