

Apply quantum algorithms to protein structure prediction

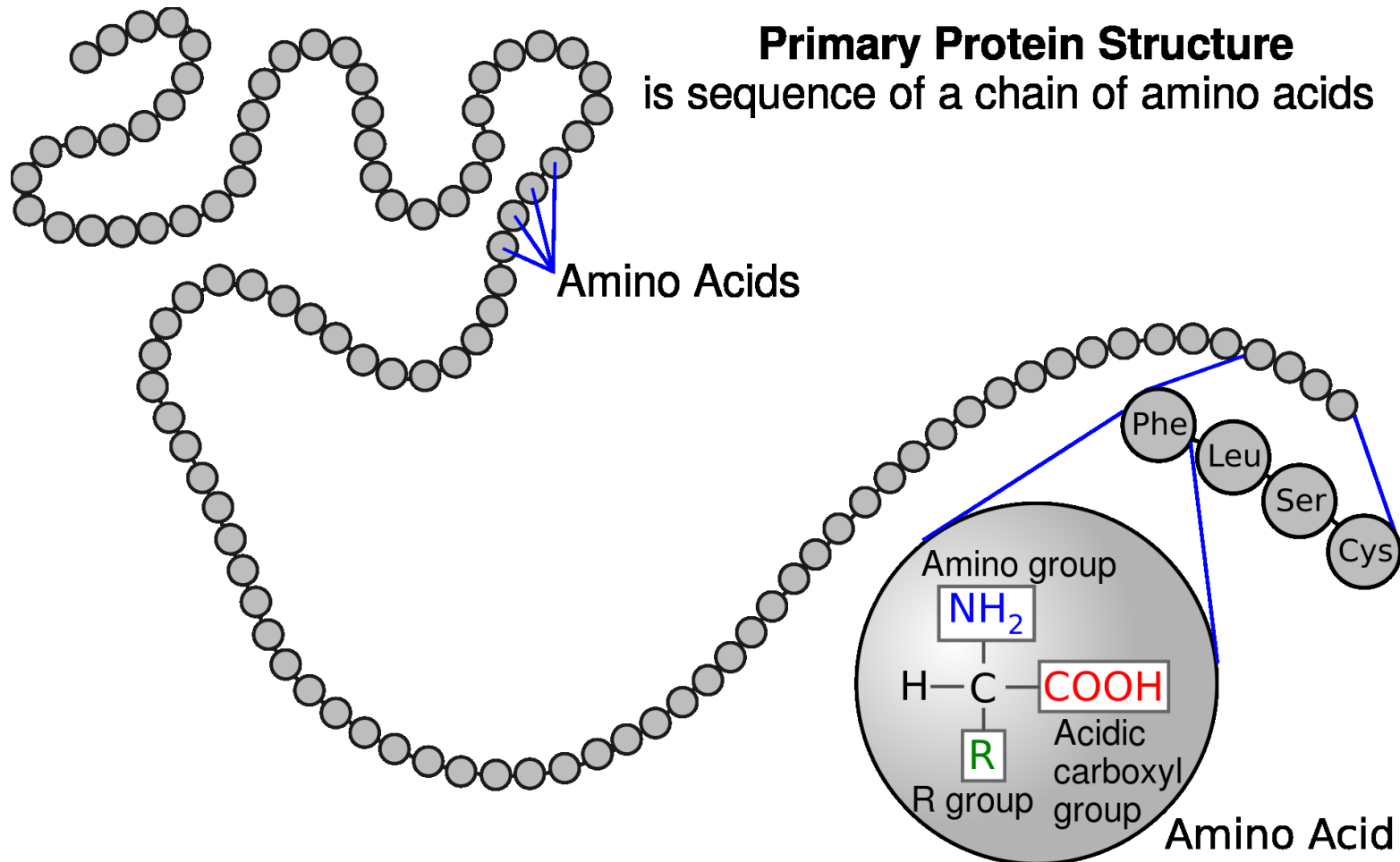
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What is a Protein?

- ▶ **Protein:** macromolecules consisting of **amino acids**
- ▶ **Peptide:** a linear chain of amino acid residues



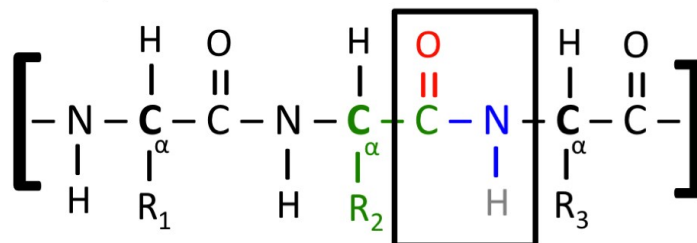
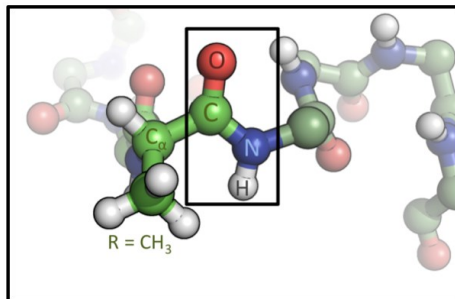
Amino Acids

► Amino acids consist of:

- ❑ amino (NH_3^+)
- ❑ carboxylate (COOH)
- ❑ R group (C, H)

► 20 amino acids

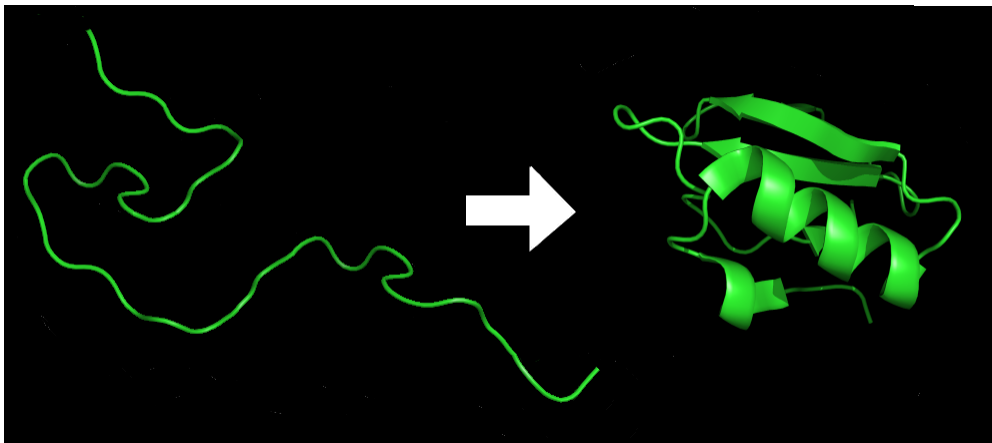
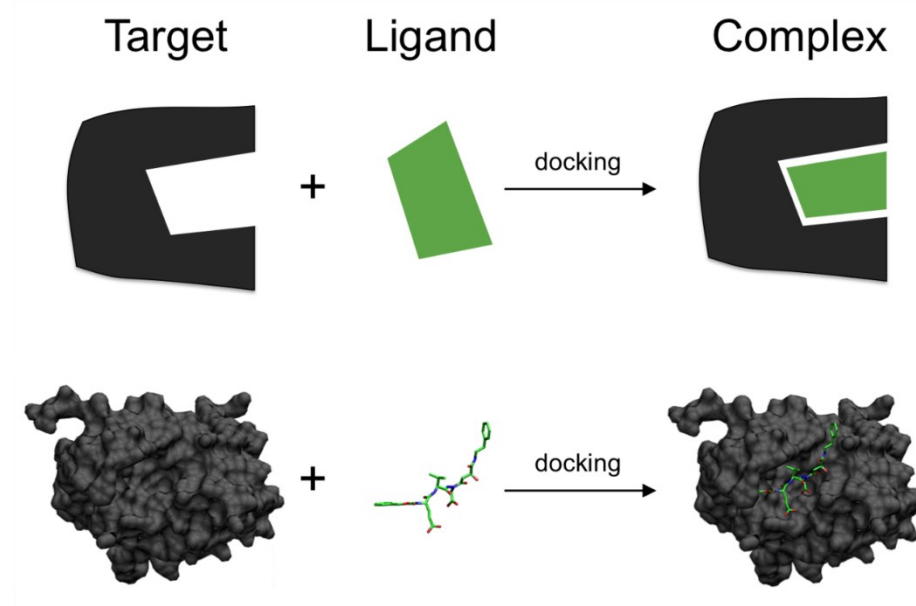
► <https://www.rcsb.org/3d-view/1GWD>



#	Amini acid	3-letter symbols	1-letter symbols
1	Alanine	Ala	A
2	Arginine	Arg	R
3	Asparagine	Asn	N
4	Aspartate	Asp	D
5	Cysteine	Cys	C
6	Glutamine	Gln	Q
7	Glutamate	Glu	E
8	Glycine	Gly	G
9	Histidine	His	H
10	Isoleucine	Ile	I
11	Leucine	Leu	L
12	Lysine	Lys	K
13	Methionine	Met	M
14	Phenylalanine	Phe	F
15	Proline	Pro	P
16	Serine	Ser	S
17	Threonine	Thr	T
18	Tryptophan	Trp	W
19	Tyrosine	Tyr	Y
20	Valine	Val	V

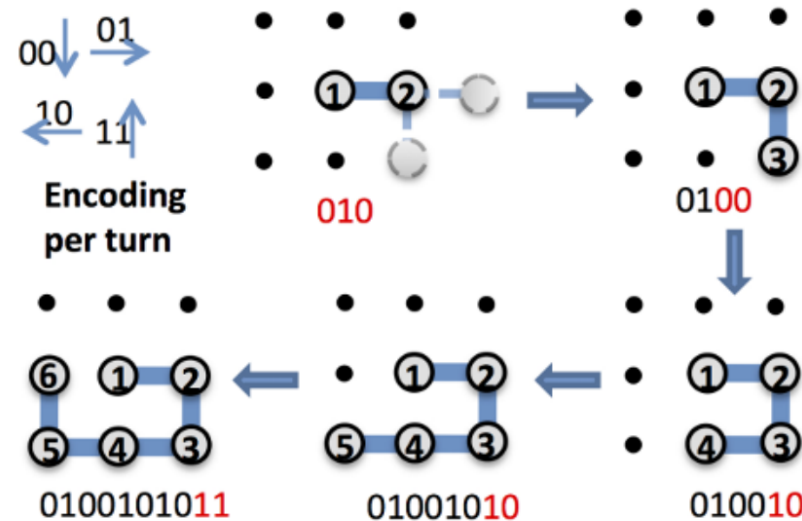
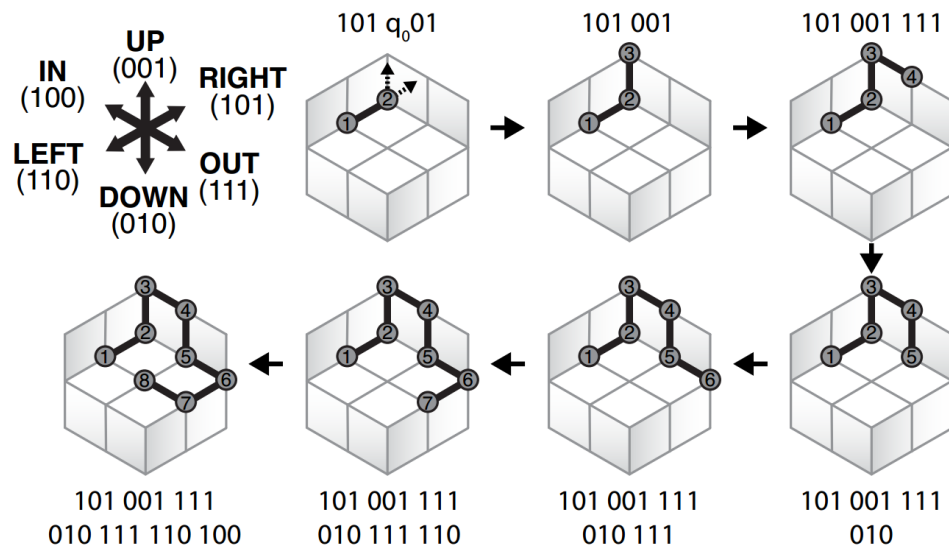
Quantum algorithms for protein problems

- ▶ Protein Folding (Quantum Annealing / QAOA / VQE)
- ▶ Protein Ligand Docking (VQE)
- ▶ Protein Design (Grover)



Protein Folding Problem

- ▶ Definition: The protein folding problem is finding the lowest energy conformations of a protein given its primary sequence of amino acids.
- ▶ large-scale problem
- ▶ **Lattice Models**
 - self-avoiding walk (SAW)
- ▶ Coarse-grained representation
 - Points: amino acids



Previous works

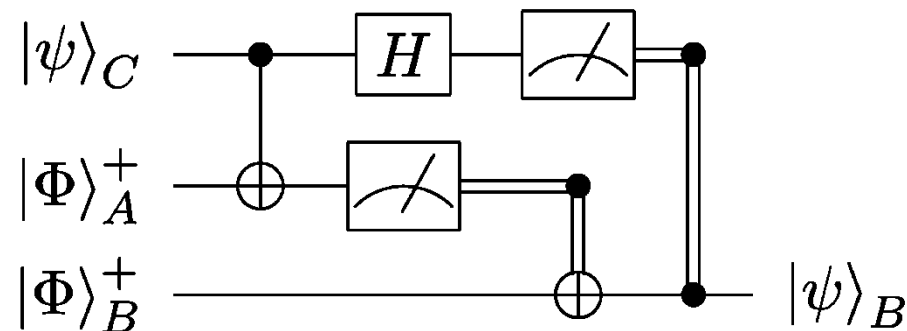
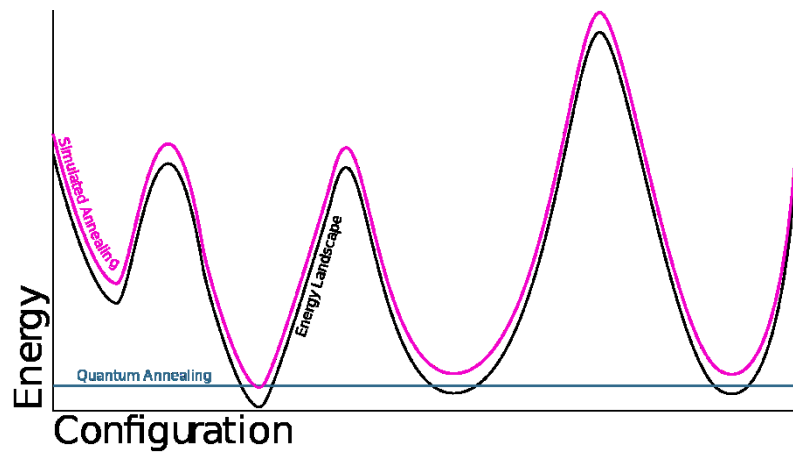
- There are several lattice models, but only two of them are used in quantum computing [2].
 - ❑ Hydrophobic-polar model (HP) [1]
 - ❑ Miyazawa-Jernigan model [3]

Table 3. Contact energies in RT units; e_{ij} for upper half and diagonal and e'_{ij} for lower half

	Cys	Met	Phe	Ile	Leu	Val	Trp	Tyr	Ala	Gly	Thr	Ser	Asn	Gln	Asp	Glu	His	Arg	Lys	Pro	
Cys	-5.44	-4.99	-5.80	-5.50	-5.83	-4.96	-4.95	-4.16	-3.57	-3.16	-3.11	-2.86	-2.59	-2.85	-2.41	-2.27	-3.60	-2.57	-1.95	-3.07	Cys
Met	0.46	-5.46	-6.56	-6.02	-6.41	-5.32	-5.55	-4.91	-3.94	-3.39	-3.51	-3.03	-2.95	-3.30	-2.57	-2.89	-3.98	-3.12	-2.48	-3.45	Met
Phe	0.54	-0.20	-7.26	-6.84	-7.28	-6.29	-6.16	-5.66	-4.81	-4.13	-4.28	-4.02	-3.75	-4.10	-3.48	-3.56	-4.77	-3.98	-3.36	-4.25	Phe
Ile	0.49	-0.01	0.06	-6.54	-7.04	-6.05	-5.78	-5.25	-4.58	-3.78	-4.03	-3.52	-3.24	-3.67	-3.17	-3.27	-4.14	-3.63	-3.01	-3.76	Ile
Leu	0.57	0.01	0.03	-0.08	-7.37	-6.48	-6.14	-5.67	-4.91	-4.16	-4.34	-3.92	-3.74	-4.04	-3.40	-3.59	-4.54	-4.03	-3.37	-4.20	Leu
Val	0.52	0.18	0.10	-0.01	-0.04	-5.52	-5.18	-4.62	-4.04	-3.38	-3.46	-3.05	-2.83	-3.07	-2.48	-2.67	-3.58	-3.07	-2.49	-3.32	Val
Trp	0.30	-0.29	0.00	0.02	0.08	0.11	-5.06	-4.66	-3.82	-3.42	-3.22	-2.99	-3.07	-3.11	-2.84	-2.99	-3.98	-3.41	-2.69	-3.73	Trp
Tyr	0.64	-0.10	0.05	0.11	0.10	0.23	-0.04	-4.17	-3.36	-3.01	-3.01	-2.78	-2.76	-2.97	-2.76	-2.79	-3.52	-3.16	-2.60	-3.19	Tyr
Ala	0.51	0.15	0.17	0.05	0.13	0.08	0.07	0.09	-2.72	-2.31	-2.32	-2.01	-1.84	-1.89	-1.70	-1.51	-2.41	-1.83	-1.31	-2.03	Ala
Gly	0.68	0.46	0.62	0.62	0.65	0.51	0.24	0.20	0.18	-2.24	-2.08	-1.82	-1.74	-1.66	-1.59	-1.22	-2.15	-1.72	-1.15	-1.87	Gly
Thr	0.67	0.28	0.41	0.30	0.40	0.36	0.37	0.13	0.10	0.10	-2.12	-1.96	-1.88	-1.90	-1.80	-1.74	-2.42	-1.90	-1.31	-1.90	Thr
Ser	0.69	0.53	0.44	0.59	0.60	0.55	0.38	0.14	0.18	0.14	-0.06	-1.67	-1.58	-1.49	-1.63	-1.48	-2.11	-1.62	-1.05	-1.57	Ser
Asn	0.97	0.62	0.72	0.87	0.79	0.77	0.30	0.17	0.36	0.22	0.02	0.10	-1.68	-1.71	-1.68	-1.51	-2.08	-1.64	-1.21	-1.53	Asn
Gln	0.64	0.20	0.30	0.37	0.42	0.46	0.19	-0.12	0.24	0.24	-0.08	0.11	-0.10	-1.54	-1.46	-1.42	-1.98	-1.80	-1.29	-1.73	Gln
Asp	0.91	0.77	0.75	0.71	0.89	0.89	0.30	-0.07	0.26	0.13	-0.14	-0.19	-0.24	-0.09	-1.21	-1.02	-2.32	-2.29	-1.68	-1.33	Asp
Glu	0.91	0.30	0.52	0.46	0.55	0.55	0.00	-0.25	0.30	0.36	-0.22	-0.19	-0.21	-0.19	0.05	-0.91	-2.15	-2.27	-1.80	-1.26	Glu
His	0.65	0.28	0.39	0.66	0.67	0.70	0.08	0.09	0.47	0.50	0.16	0.26	0.29	0.31	-0.19	-0.16	-3.05	-2.16	-1.35	-2.25	His
Arg	0.93	0.38	0.42	0.41	0.43	0.47	-0.11	-0.30	0.30	0.18	-0.07	-0.01	-0.02	-0.26	-0.91	-1.04	0.14	-1.55	-0.59	-1.70	Arg
Lys	0.83	0.31	0.33	0.32	0.37	0.33	-0.10	-0.46	0.11	0.03	-0.19	-0.15	-0.30	-0.46	-1.01	-1.28	0.23	0.24	-0.12	-0.97	Lys
Pro	0.53	0.16	0.25	0.39	0.35	0.31	-0.33	-0.23	0.20	0.13	0.04	0.14	0.18	-0.08	0.14	0.07	0.15	-0.05	-0.04	-1.75	Pro
$e_{rr} - 2.55$	e_{rr}	-3.57	-3.92	-4.76	-4.42	-4.81	-3.89	-3.81	-3.41	-2.57	-2.19	-2.29	-1.98	-1.92	-2.00	-1.84	-1.79	-2.56	-2.11	-1.52	-2.09
$e_r - 3.60$	e_r	-4.29	-4.73	-5.57	-5.29	-5.71	-4.72	-4.41	-3.87	-3.17	-2.53	-2.63	-2.27	-2.14	-2.35	-2.02	-2.07	-2.94	-2.43	-1.82	-2.53
$f_r - 3.60$	f_r	-5.58	-6.14	-7.39	-7.09	-7.88	-6.15	-5.34	-4.60	-3.24	-2.22	-2.48	-1.92	-1.74	-1.93	-1.54	-1.49	-2.91	-2.07	-1.17	-1.97
N_H/N_L	2.096	2.723	2.722	2.780	2.811	2.893	2.728	2.537	2.493	2.143	1.840	1.973	1.771	1.699	1.720	1.598	1.508	2.075	1.787	1.343	1.629
q_i	7.162	6.281	6.646	6.137	5.870	6.042	6.087	6.155	5.793	6.037	6.334	6.284	6.486	6.582	6.574	6.469	6.487	6.235	6.241	6.318	6.569

Quantum Computing

- ▶ Richard Feynman: "Nature isn't classical, and if you want to make a simulation of nature, you'd better make it quantum mechanical"
- ▶ Quantum computing models:
 - ▣ Adiabatic Quantum Computing (Quantum Annealing) [6-9]
 - ▣ Digital Quantum Computing (Gate-based model) [10-11]



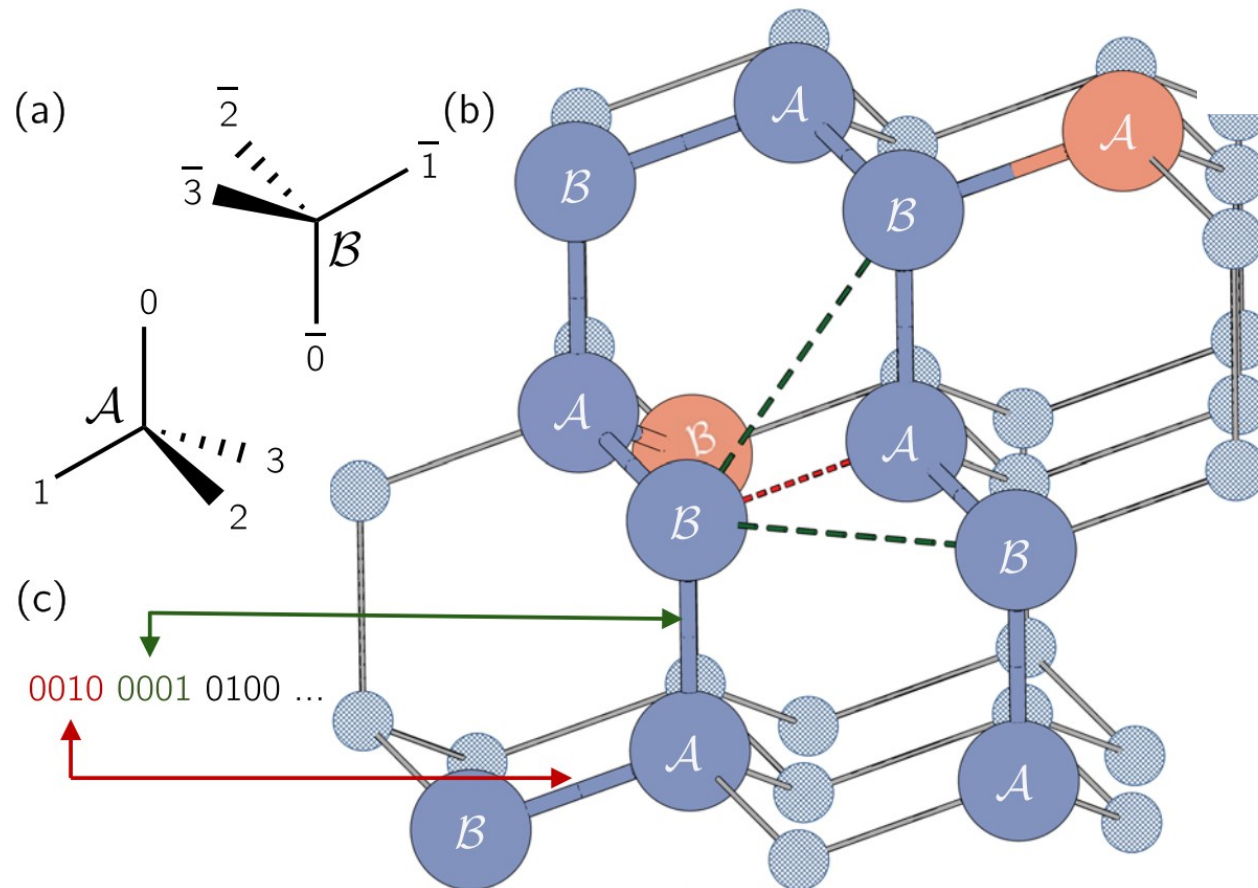
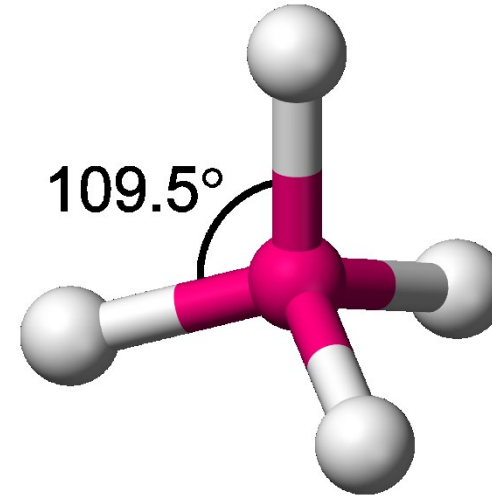
Gate-based model

- ▶ Variational Quantum Algorithms (VQA):
 - ❑ Quantum Approximate Optimization Algorithm (QAOA) [10]
 - ❑ Variational Quantum Eigensolver (VQE) [11]
- ▶ Ground State Problem (finding eigenvalue):
 - ❑ Quantum Phase Estimation (QPE)
 - ❑ Variational Quantum Eigensolver (VQE)
- ▶ Noisy Intermediate-Scale Quantum (NISQ) era

Qiskit Nature

► Tetrahedral lattice model

□ angle(s): $\approx 109.5^\circ$



VQE algorithm

► VQE algorithm can solve the Schrödinger equation:

► $H|\psi(\vec{\theta})\rangle = E|\psi(\vec{\theta})\rangle$

where:

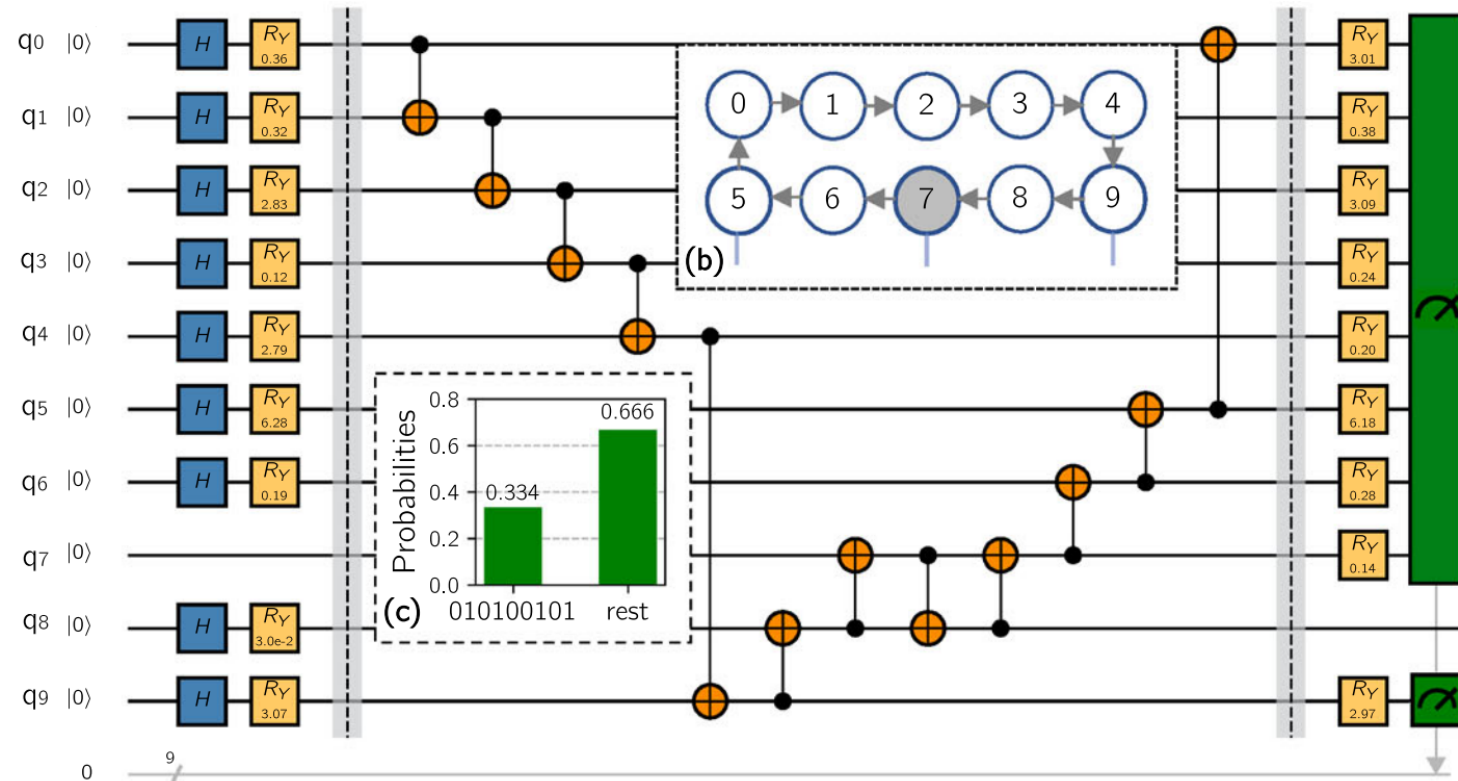
- E is energy / a number
- E_0 is the lowest energy / eigenvalue of H / a number
- $\psi(\vec{\theta})$ is the eigenstate of H / a vector
- H is the Hamiltonian / an operator / a matrix
- $\vec{\theta}$ is a vector of independent parameters and $\vec{\theta} = \{(\theta_1, \dots, \theta_n)\}^T$.

The lowest answer for this equation is the ground state (E_0).

$$E_0 \leq \langle \psi(\vec{\theta}) | H | \psi(\vec{\theta}) \rangle$$

VQE algorithm

- ▶ VQE algorithm consists of two important parts:
 - ❑ A quantum circuit (ansatz)
 - ❑ An optimization method (classical computers)



Hamiltonian

- ▶ They defined two kinds of qubits:

1. q_{cf} : Configuration qubits that are used for configurations and the relative position of the amino acids
2. q_{in} : Interaction qubits that encode interactions between the amino acids

- ▶ The Hamiltonian consists of some parts:

$H = H_{gc}(q_{cf}) + H_{ch}(q_{cf}) + H_{in}(q_{cf}, q_{in})$ where:

- H_{gc} is the geometrical constraint term
- H_{ch} is the chirality constraint
- H_{in} is the interaction energy terms of the system.

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Thanks For Attention