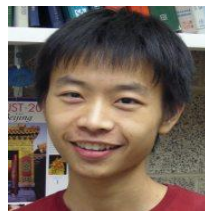


Exploring the thermodynamic landscape and kinetic pathways for metastable materials synthesis



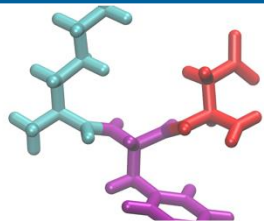
**S. Banik, R. Batra, T. Loeffler, H. Chan, S. Srinivasan, S. Manna,
A. Chandra, G. Chen, Chris Fry, Pierre Darancet,**

Subramanian Sankaranarayanan

Center for Nanoscale Materials
Argonne National Laboratory

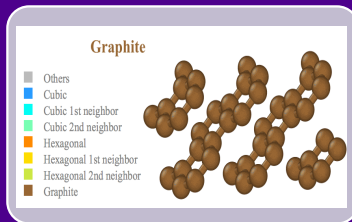
Department of Mechanical and Industrial Engineering,
University of Illinois Chicago

Overview - High-dimensional Search Problems in Materials Synthesis



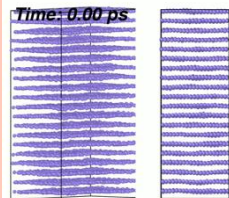
Sequence problem in peptide synthesis

- Large and discrete search space
- Overcoming human bias in peptide synthesis



Explore thermodynamic landscape of metastable materials

- Large and Continuous search space
- A phase diagram for metastable materials



Control kinetic pathways for synthesizing metastable materials

- Learn time-dependent synthesis protocols
- Controlled selection of metastable polymorphs

Sequence problem in peptide synthesis

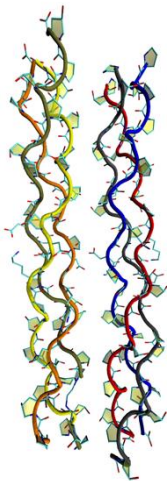
Amino acid sequence controls the self-assembly of peptides

Examples

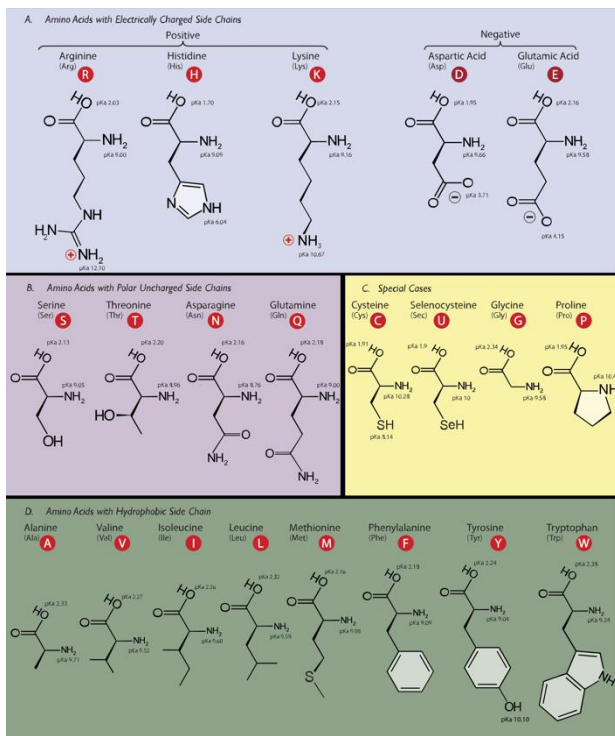
Spider Silk



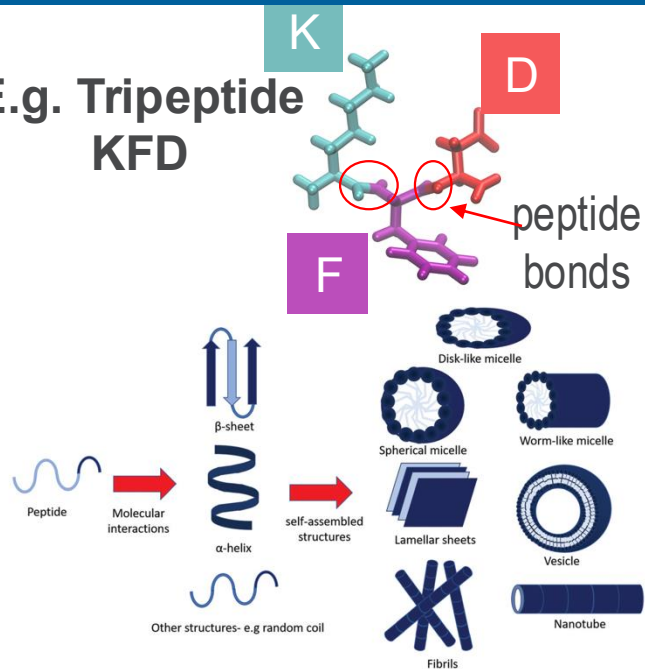
Collagen



20 Amino Acids



E.g. Tripeptide KFD



- Useful for catalysis, sensing, tissue engineering
- Easy tunable properties
- Huge diversity

Traditional Approaches to Peptide Design and Synthesis

Sequence Challenge in Peptide Discovery and Synthesis

of possibilities: 20^n

Empirical Rational Design

- Patterning (npnnpn)
- Hydrophobicity scales
- **Human bias; based on limited data**

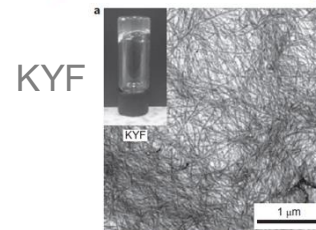
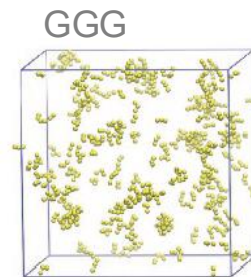
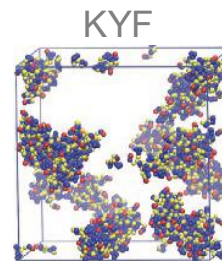
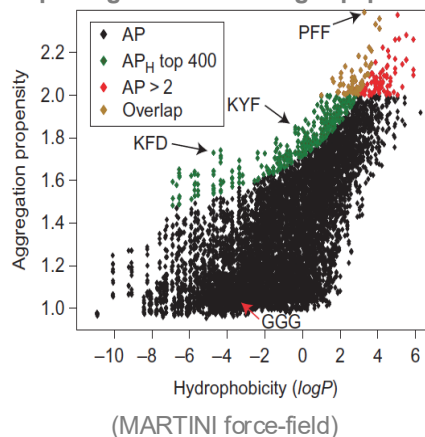
Computational Design

- MD simulation to estimate peptide properties
- Hydrophobicity scales
- **Brute-force search; non-scalable**

Can we do better?

Sequence length (n)	# of candidates	Simulation time
3	8000	3 weeks
5	3.2 M	20 years
8	25.6 B	Many years

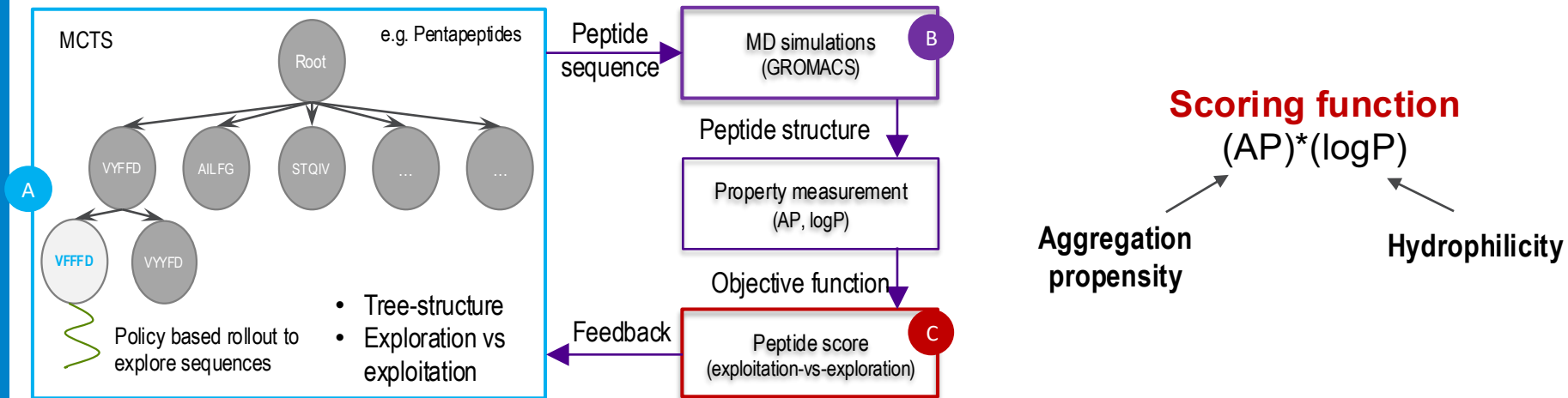
Exploring self-assembling tripeptides



P. W. J. M. Frederix et al., Nature Chemistry, 7, 30 (2015)

AI-Expert for Peptide Discovery

AI-guided autonomous computational search



Key components

A. Monte Carlo tree search (MCTS) → Promising sequence generation

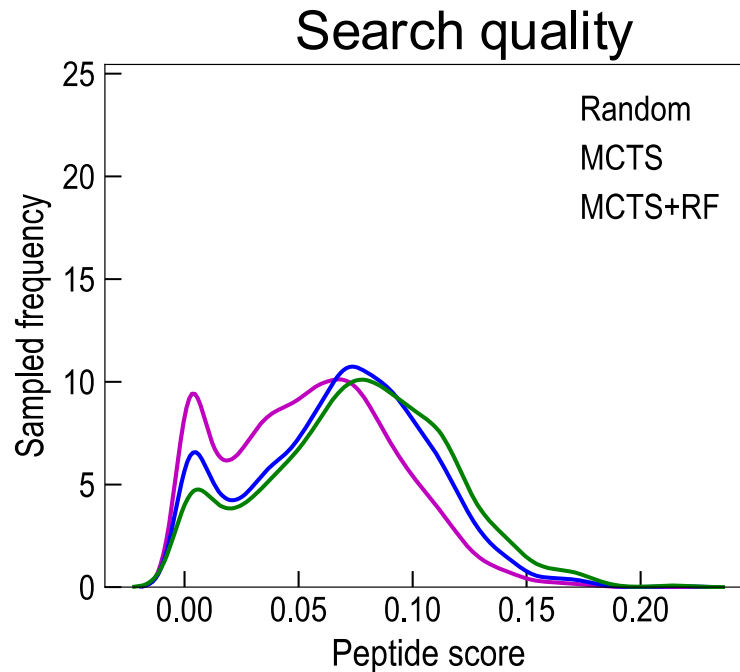
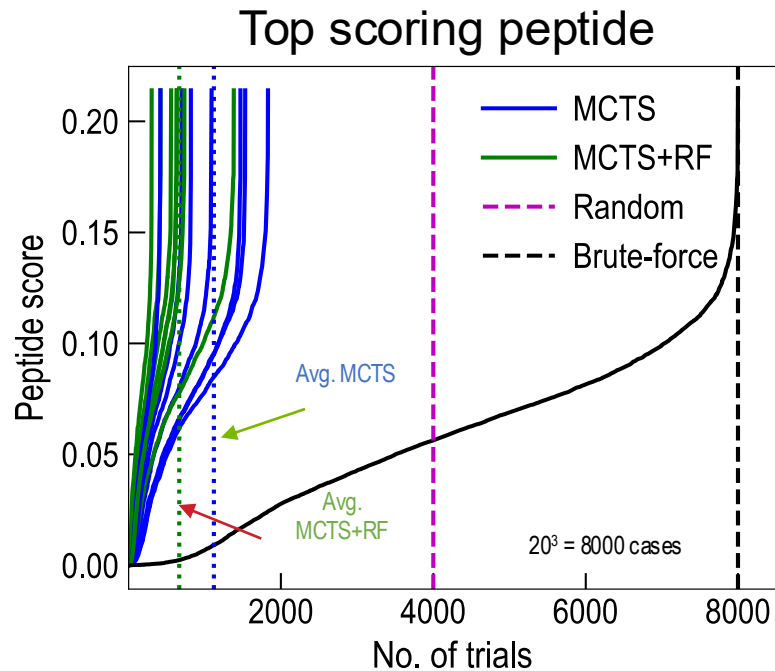
B. MD simulations → Model structure

C. Scoring function → Sequence evaluation

Search acceleration: Random forest ML model for efficient rollouts (MCTS+RF scheme)

Batra, R, et al. "Self-assembling Peptide Discovery: Overcoming Human Bias With Machine Learning." Nature Chem (2022) (12):1427-1435

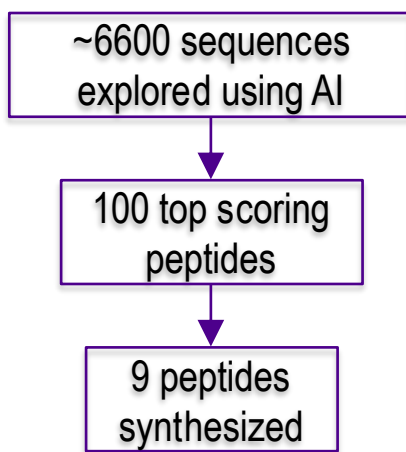
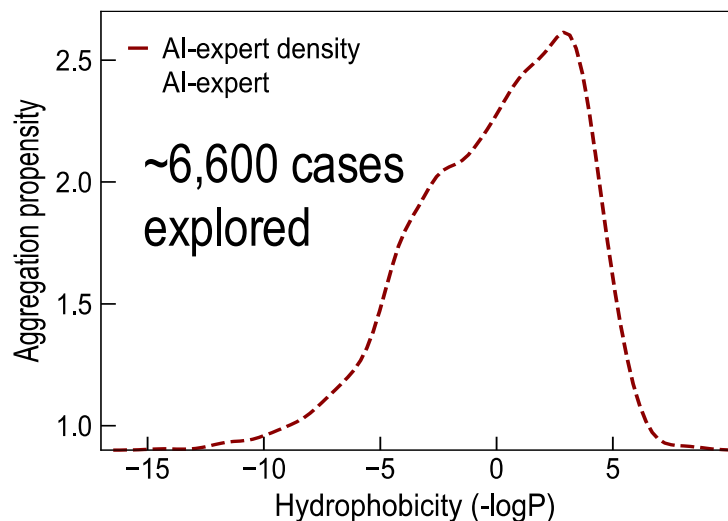
AI-Expert Results – Validation on Tripeptides



- Tripeptides offer a manageable space (8000 cases) for validation
- Superior performance of MCTS+RF, MCTS methodology

AI-expert with MCTS+RF scheme performs best!

AI-Expert Results – Screening of Pentapeptides



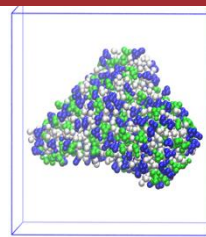
Example top candidates from AI-expert

pep	AP	logP	Score
PTPCY	2.54	-0.2	0.49
PPPHY	2.46	-0.18	0.44
SYCGY	2.42	0.17	0.42
FFEKF	2.24	1.30	0.34
KWEFY	2.20	1.92	0.33
FKIDF	2.17	1.90	0.31
WKPY Y	2.11	-0.57	0.26

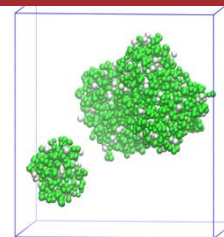
- Pentapeptides cannot be explored exhaustively (3.2 M cases)
- AI-expert candidates have **high scores** and **high diversity**

Are these candidates any good?

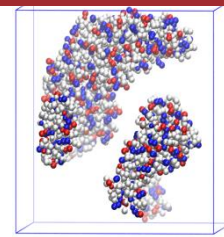
Computationally visible aggregation



PPPHY



SYCGY



FKIDF

Nature Chem (2022) (12):1427-1435

Comparison of AI-expert & Human Experts

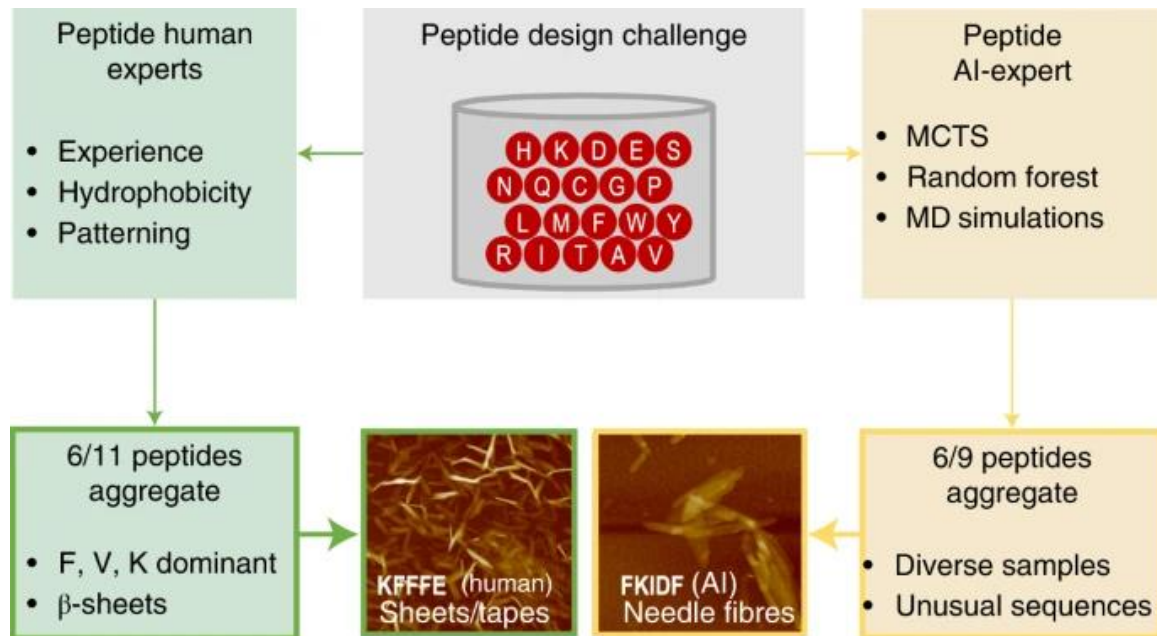
Requested sequences from six Human experts

- Leaders in peptide research
- Years of synthesis experience
- 11 (out of 23) sequences synthesized

Top scoring pentapeptides: Computational score

Rank	Peptide	Expert	Score
1	PPPHY	AI	0.44
2	SYCGY	AI	0.43
3	WKPY Y	AI	0.34
4	KWEFY	AI	0.33
5	FFEKF	AI	0.31
6	FKFEF	Human	0.28
7	FKIDF	AI	0.25
8	VVVV	Human	0.21

- **For aggregation, AI-expert >= Human-experts**
- Based on just computational scores, AI-expert performs better
- Low computational score = no aggregation (e.g. KVKVK)
- High computational score \neq aggregation (e.g. PPPHY)



Nature Chem (2022) (12):1427-1435

Key conclusions

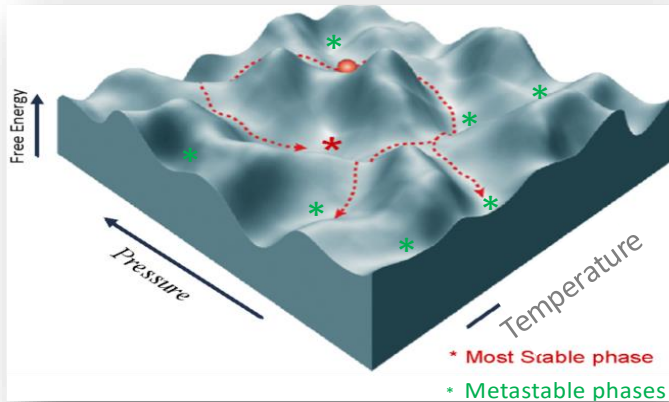
- Typically, peptide discovery research yields fewer than ten candidates per study and is subject to human bias.
- A fully autonomous computational search engine has been developed to reveal non-intuitive peptide sequences with high potential for self-assembly.
- The AI expert combines Monte Carlo tree search, molecular dynamics simulations, and a scoring function closely paired with experimental synthesis.
- AI-Expert screens millions of pentapeptide to efficiently determine 6,600 candidates for self-assembly
- While success percentages are similar between AI-Expert and Human Experts, the sequences that self-assemble are non-intuitive with high self-assembly propensity

Exploring Thermodynamic Landscape of Metastable Phases

Discrete → Continuous Search Space

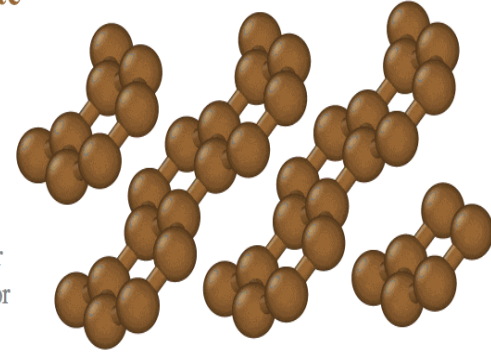
Synthesis of Metastable Materials

Opportunity: Metastable materials often have exotic properties but their synthesis pathways remain poorly understood



Graphite

- Others
- Cubic
- Cubic 1st neighbor
- Cubic 2nd neighbor
- Hexagonal
- Hexagonal 1st neighbor
- Hexagonal 2nd neighbor
- Graphite

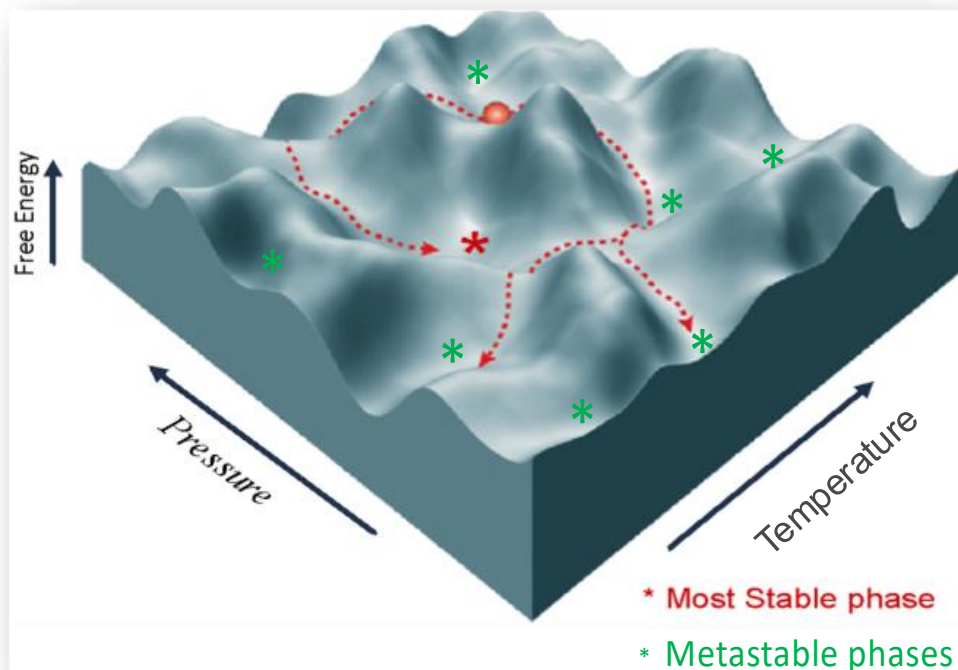


Materials Science Challenge: Thermodynamics & Kinetics

AI Challenge: Continuous space search, high-dimensional, multiple objectives/rewards, physical/chemical constraints to name a few

Grand Challenge and Beyond Exascale but Game Changer for Materials Science

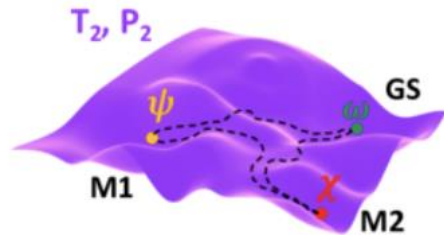
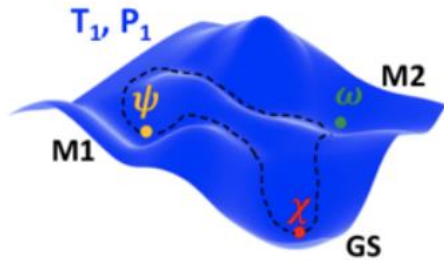
Equilibrium and metastable states during synthesis



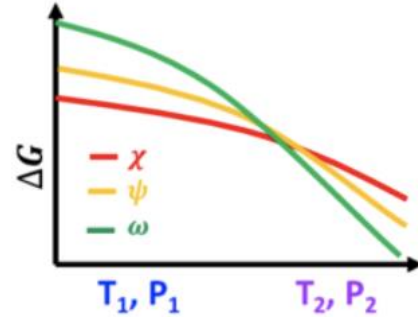
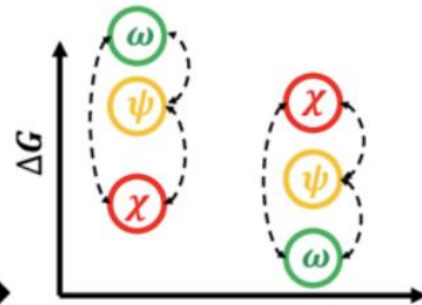
Metastable phase diagrams are data-intensive and offer infinite possibilities

Metastable Phase Diagram

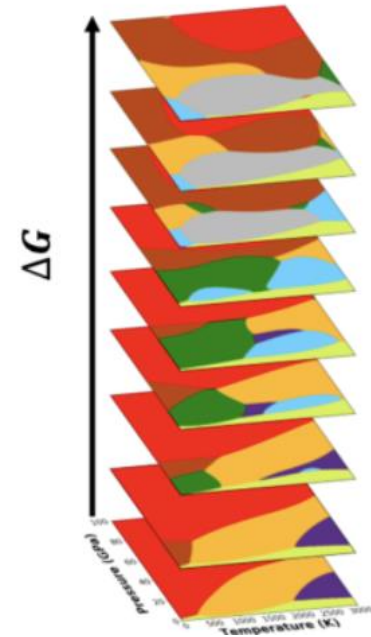
Configurational space
 $G(\{r_i\}, a, b, c, \alpha, \beta, \gamma)$



Thermodynamic space
 $G(T, P)$



Metastable phase diagram

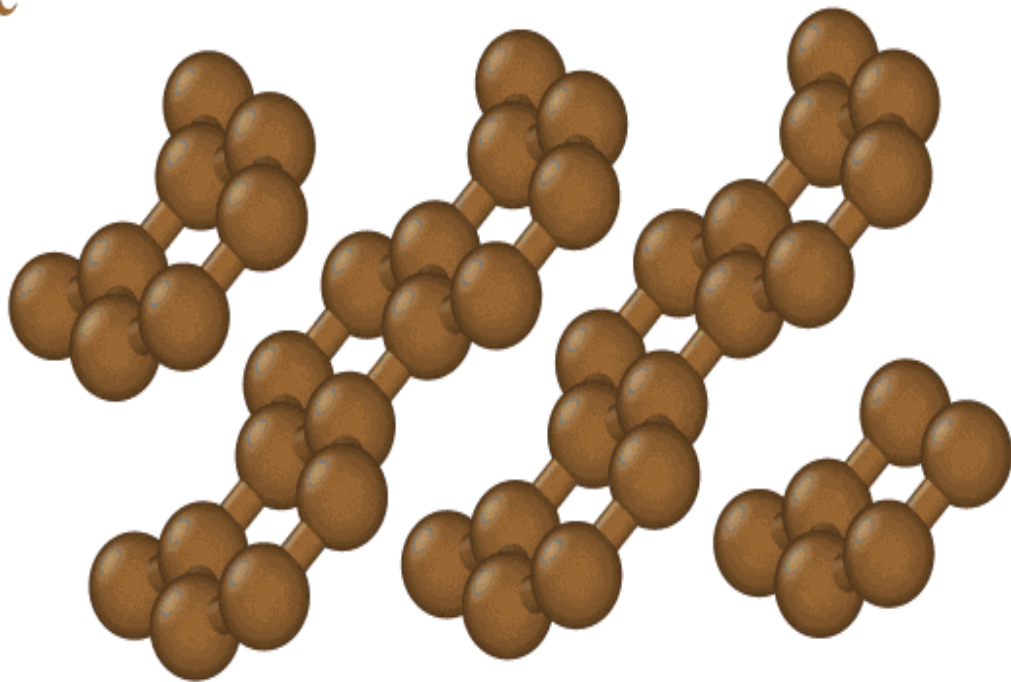


Sample and Discover Metastable Phases

Graphite

Carbon

- Others
- Cubic
- Cubic 1st neighbor
- Cubic 2nd neighbor
- Hexagonal
- Hexagonal 1st neighbor
- Hexagonal 2nd neighbor
- Graphite

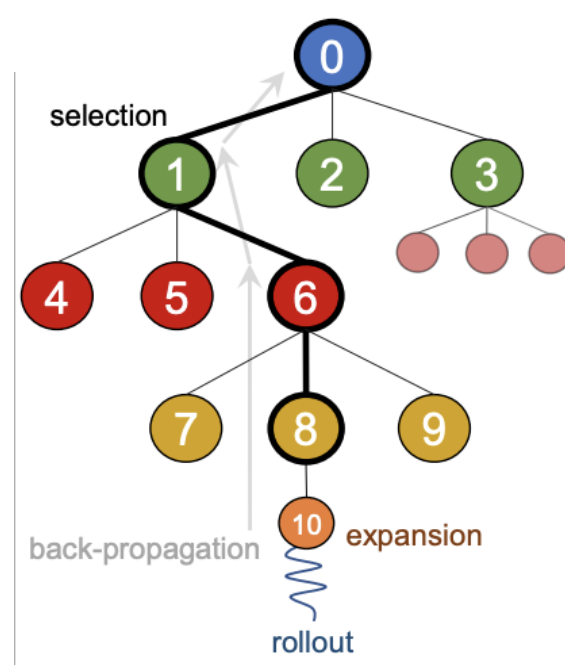


Continuous Search Space

High Dimensional Continuous Search Space

Important Modifications for Continuous MCTS

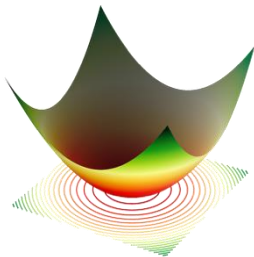
- (1) A uniqueness function to avoid degeneracy
- (2) Correlating tree depth to action space
- (3) Implementing an adaptive sampling of playouts



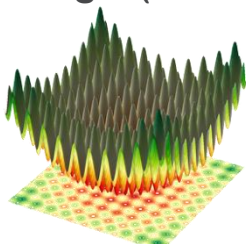
Nature Communications volume 13, Article number: 368 (2022)

Performance on Standard Test Functions

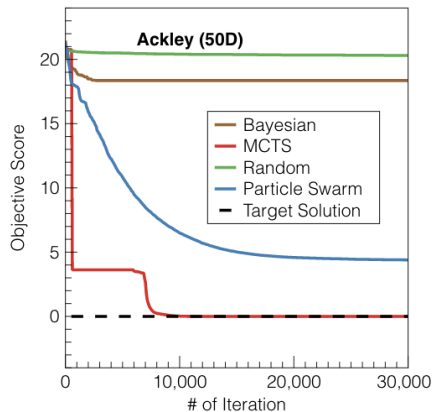
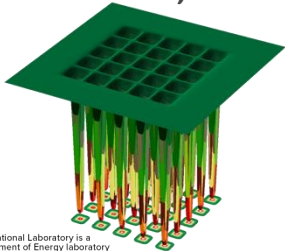
Sphere (Unimodal)



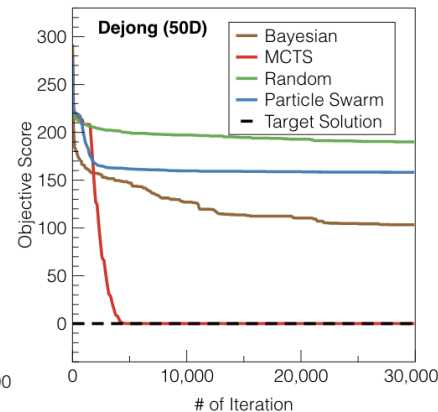
Rastrigin (Multimodal)



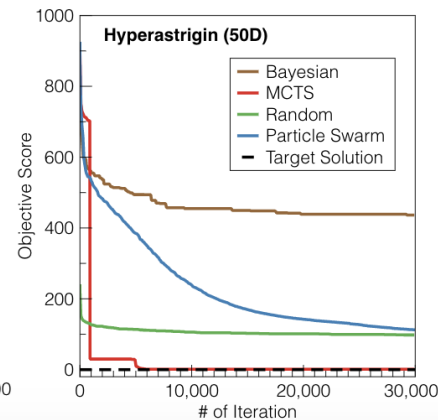
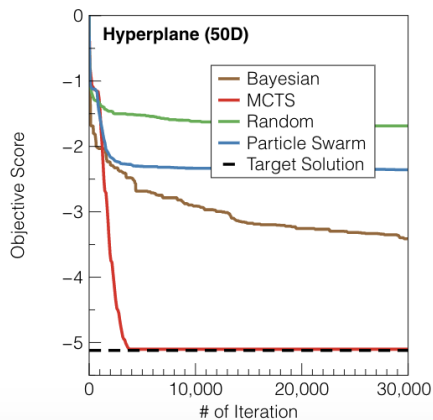
Dejong N5 (Fixed Dimension)



(a)

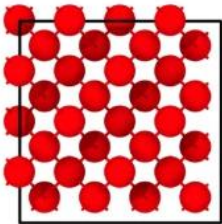


(b)

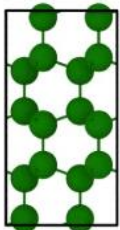


Discover metastable phases

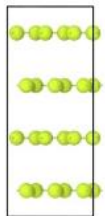
**Cubic
Diamond**



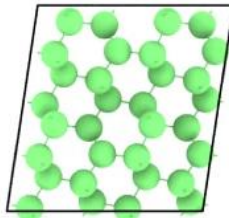
**Hexagonal
Diamond**



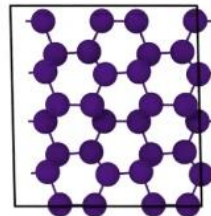
**Hexagonal
Graphite**



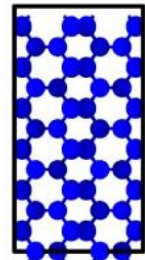
**n-diamond
(S291)**



**Diaphite
(S353)**



C2/m-16



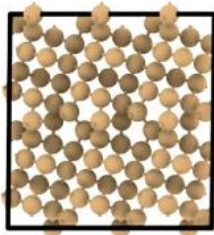
G92



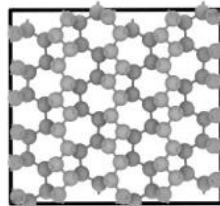
W-carbon



G173



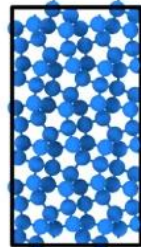
H-carbon



G178

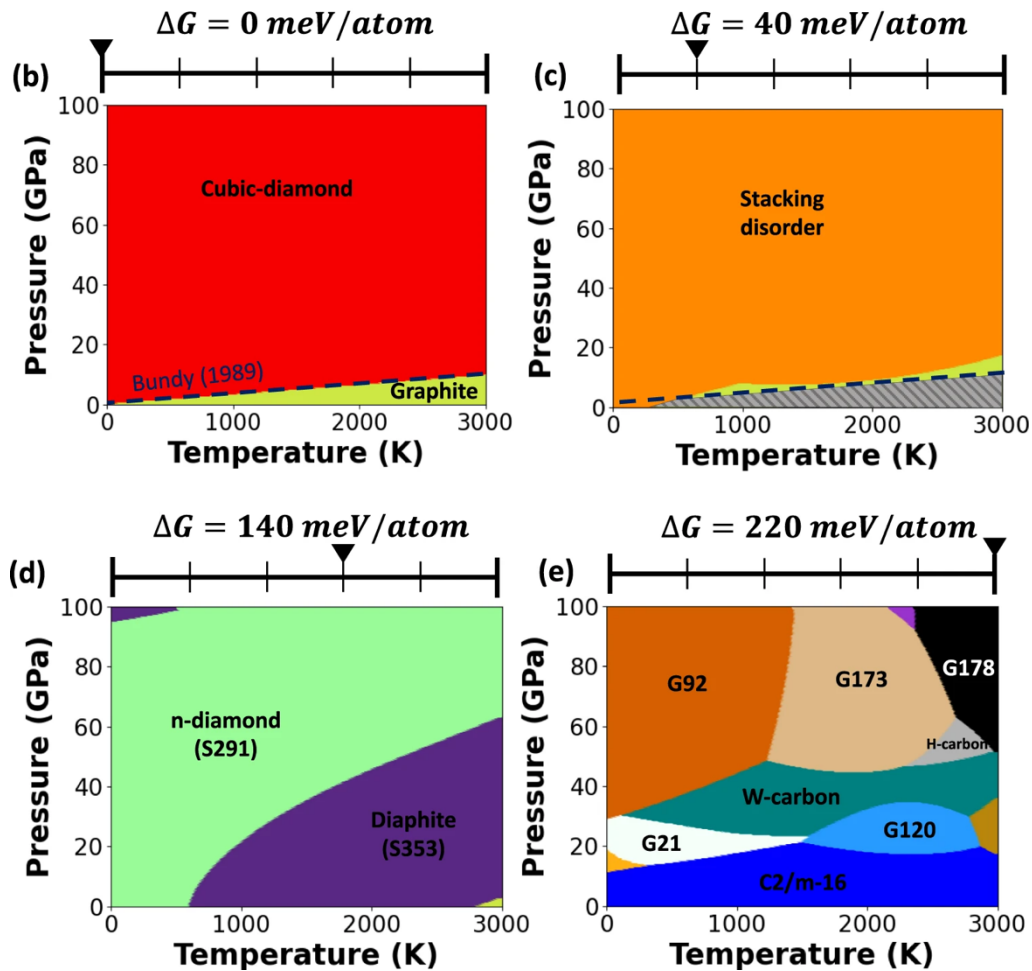


G120



~1000 metastable phases of carbon identified within 0.5 eV per atom

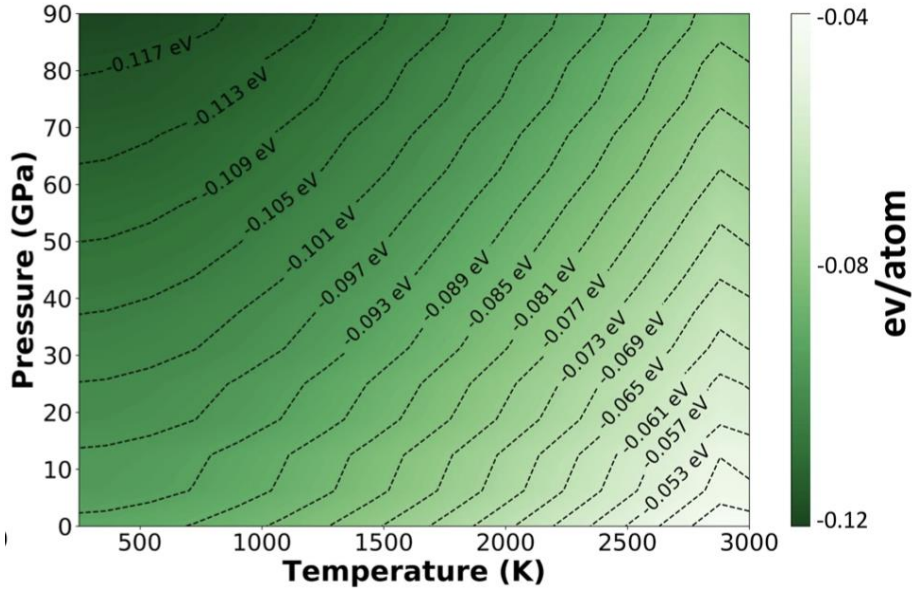
Construct Metastable Phase Diagram



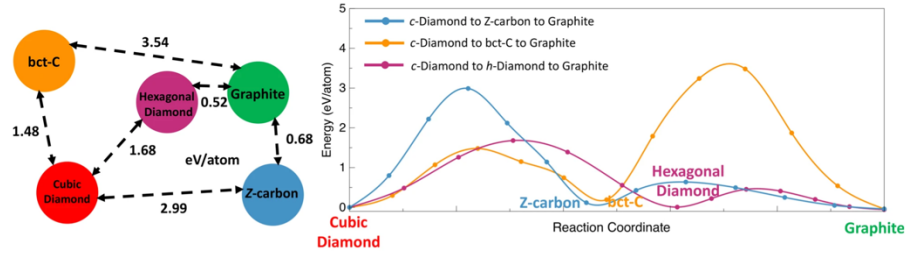
Post Processing – Synthesizability and Domains of relative phase stability

Relative stability of metastable phases

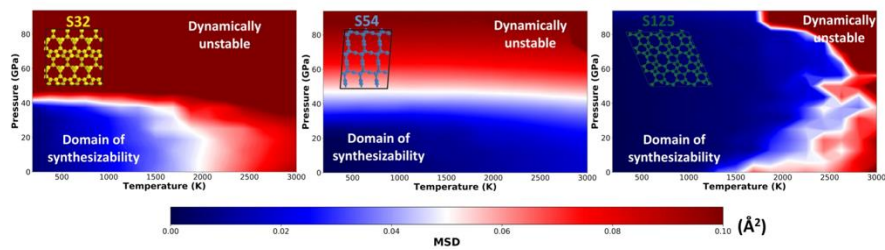
$$\Delta G = G_{\text{hexagonal-diamond}} - G_{\text{diaphite}}$$



Transformation barriers to metastable phases

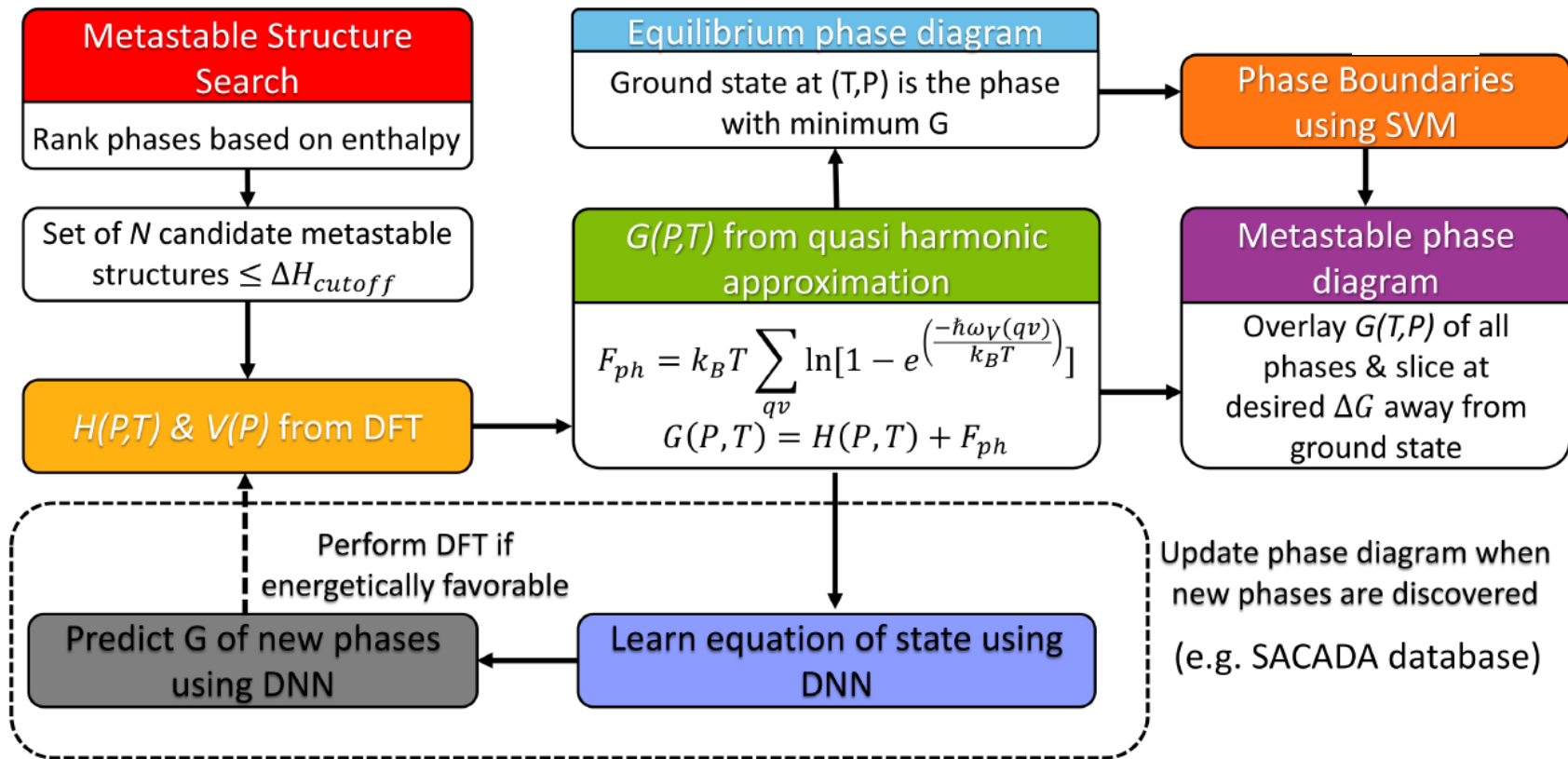


Domains of synthesizability



Nature Communications volume 13,
Article number: 3251 (2022)

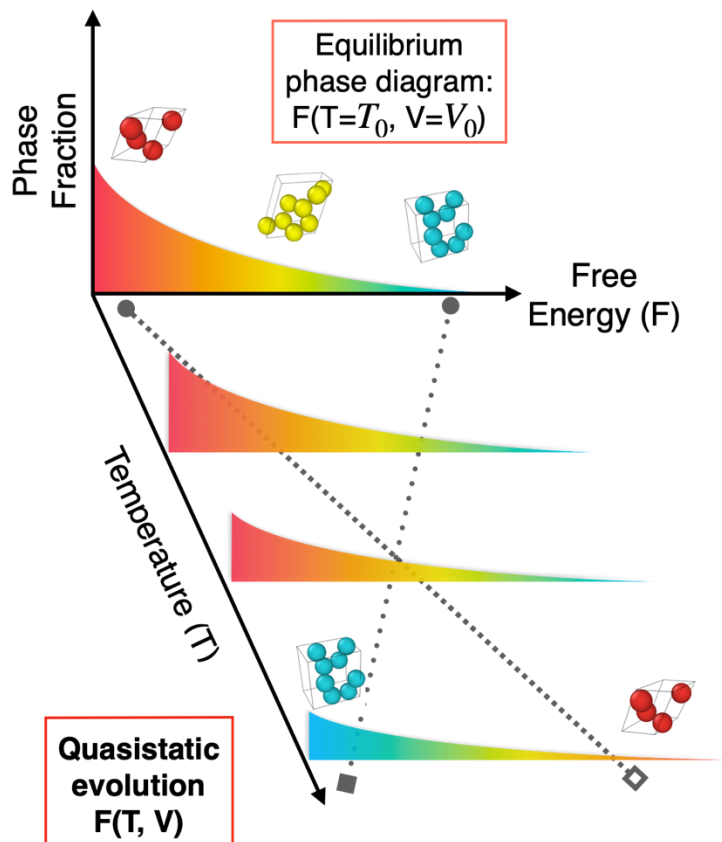
A workflow for metastable phase diagram construction



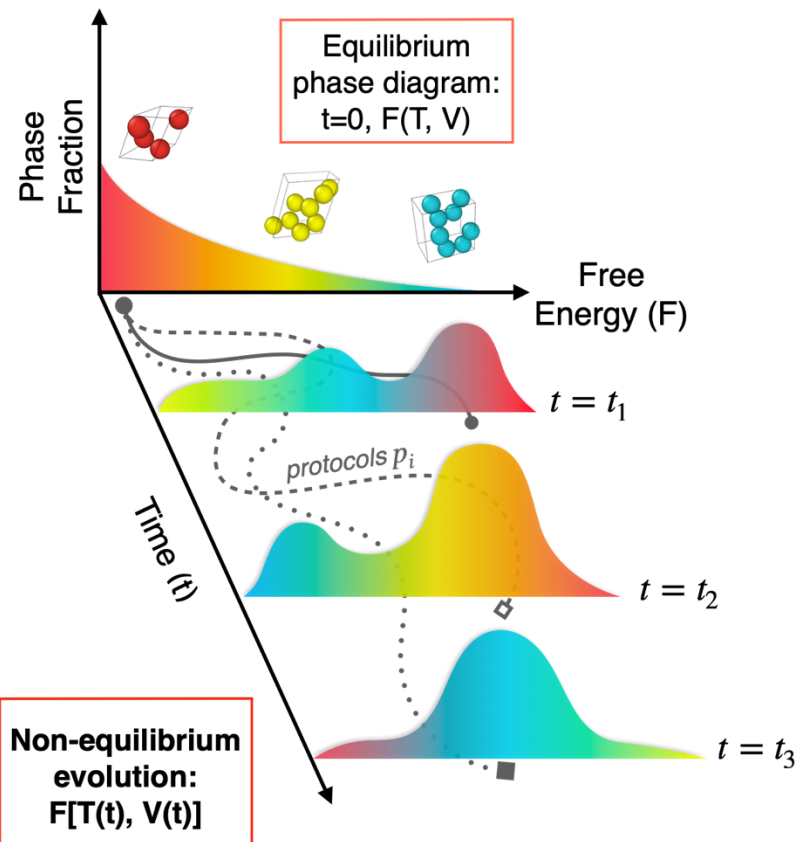
Learning kinetic pathways for synthesizing metastable materials

What about kinetics?

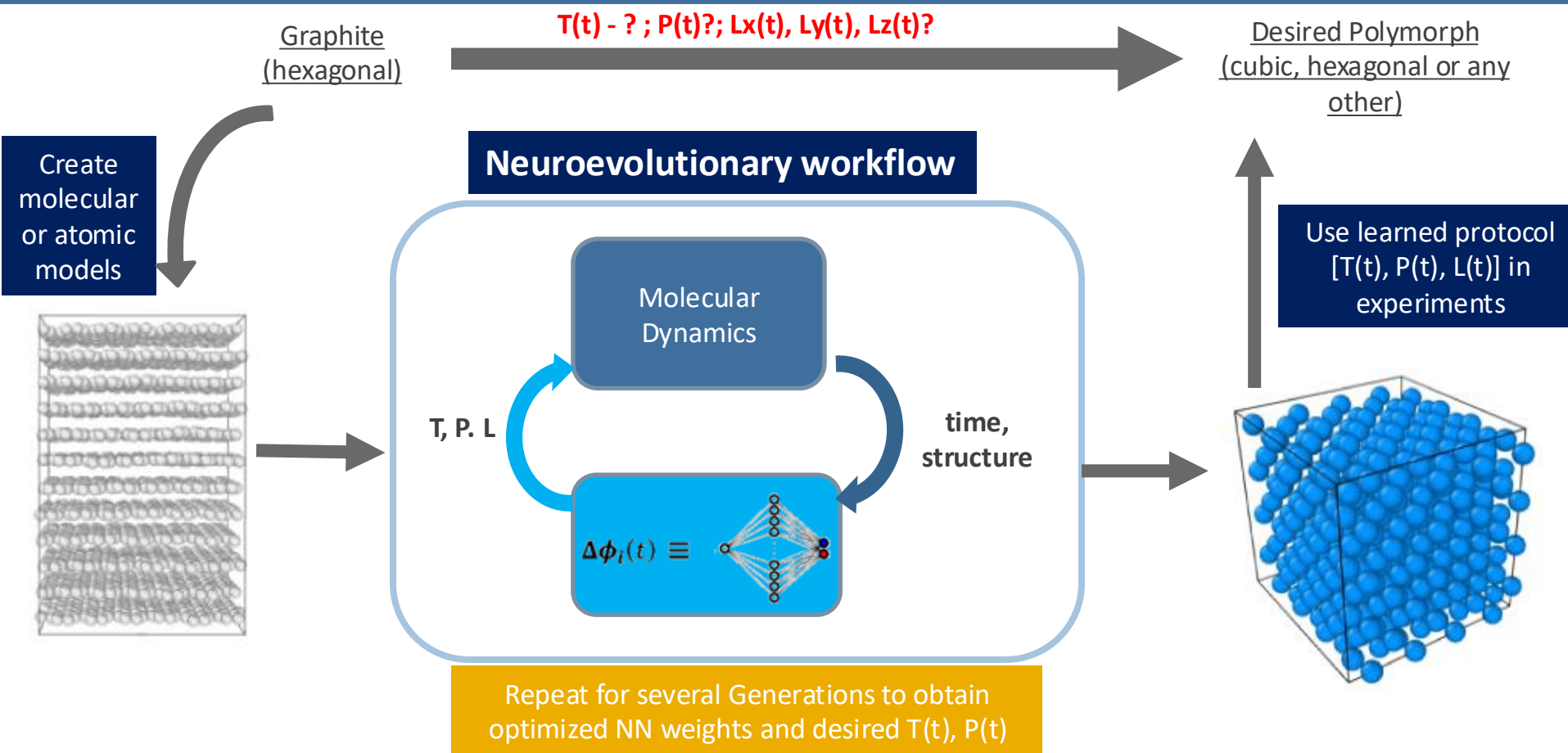
Thermodynamics



Kinetics



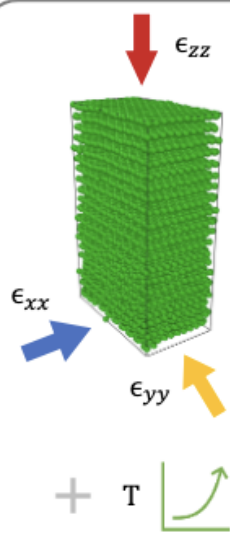
Can we learn time-dependent protocols for polymorph selection?



Learn time-dependent protocols to form Cubic Diamond

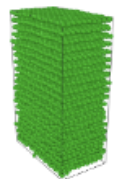
Synthesis with constant rate protocols

Graphite

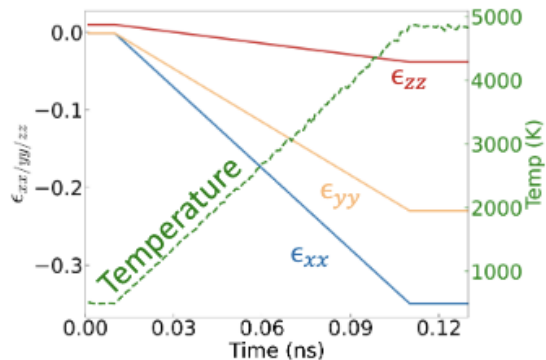


Successful search for constant rate protocols

t = 0ns



Graphite



t = 0.13ns

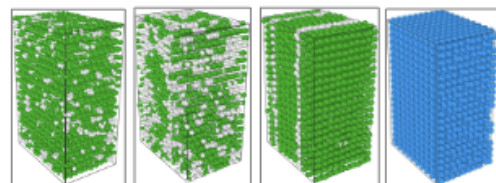
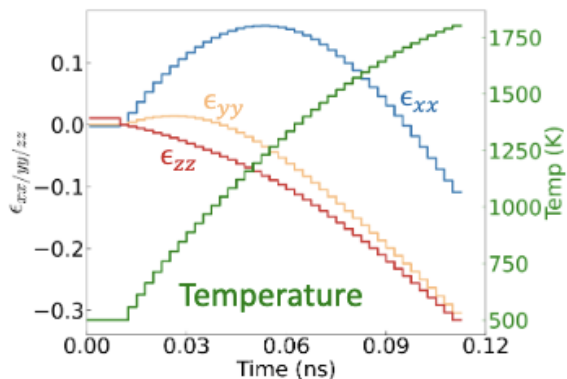


Cubic Diamond

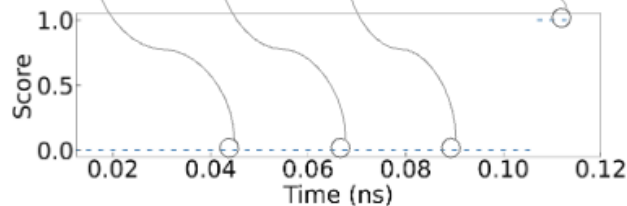
- Graphite
- Cubic Diamond
- Hexagonal Diamond
- Other States

Synthesis with time-dependent protocols

Successful search for time-dependent protocols



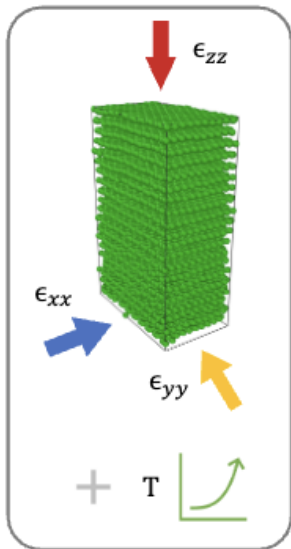
Cubic Diamond



Learn time-dependent protocols to form Hexagonal Diamond

Synthesis with constant rate protocols

Graphite



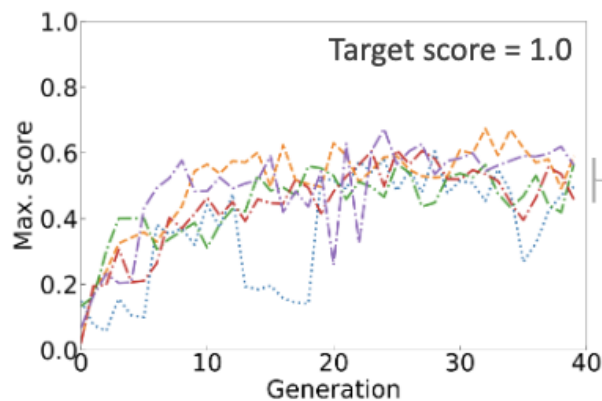
Unsuccessful
Search for
constant
rate
protocols



Successful
Search for
time-
dependent
protocols

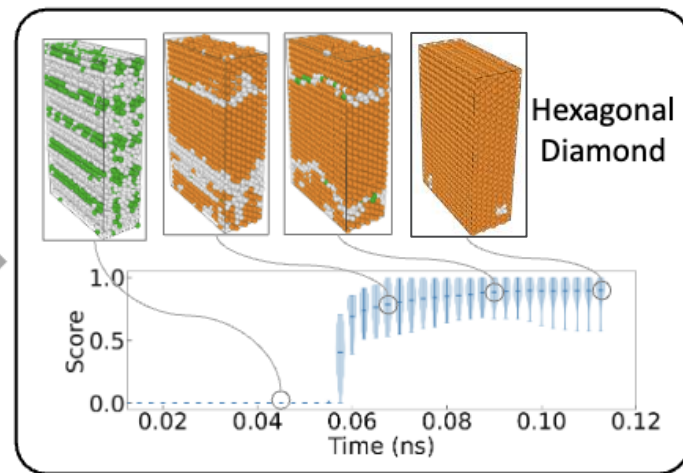
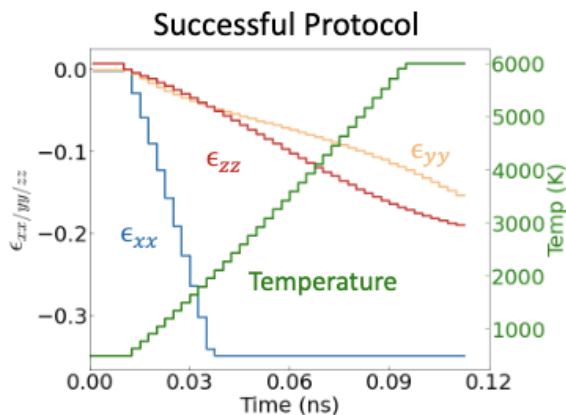


Synthesis with time-
dependent protocols

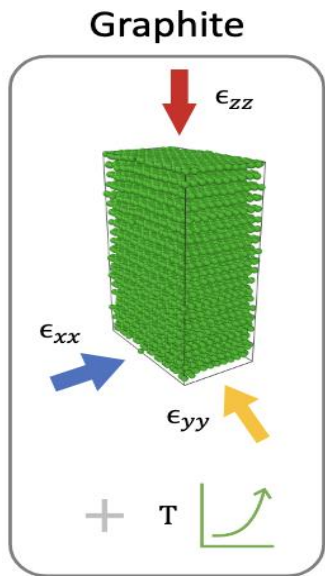


Maximum scores
achieved by constant
rate protocols across
different iterations of
the learning algorithm

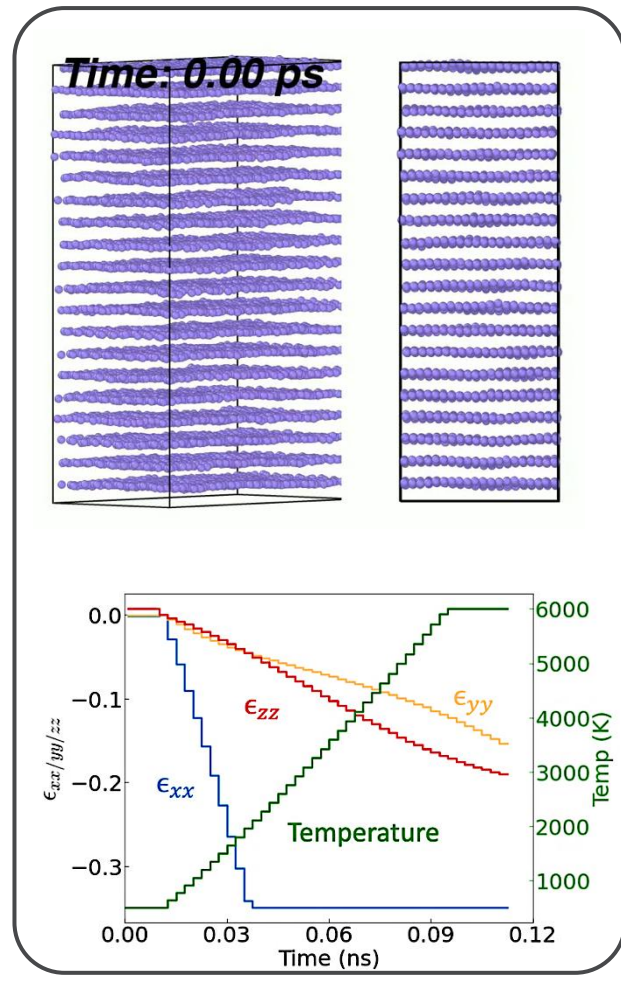
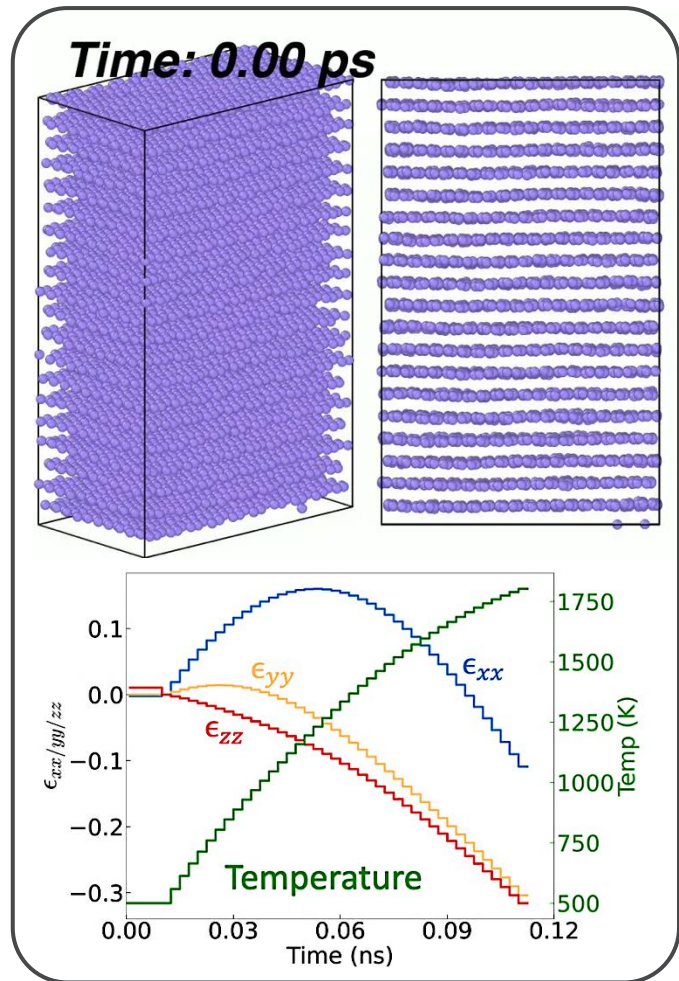
- Graphite
- Cubic Diamond
- Hexagonal Diamond
- Other States



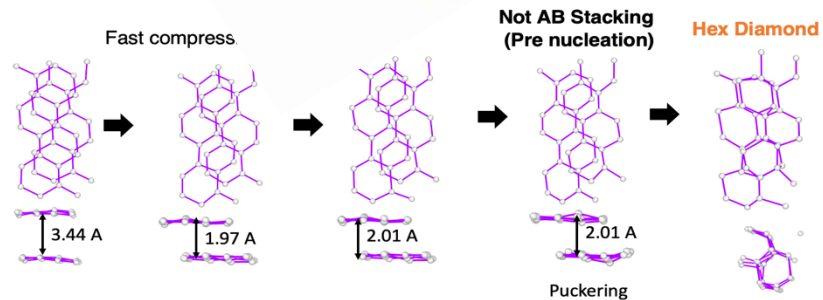
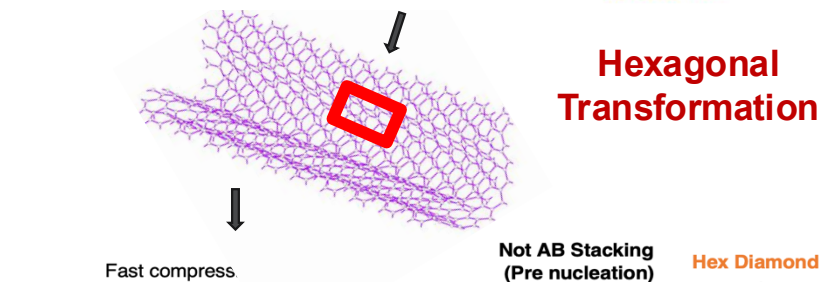
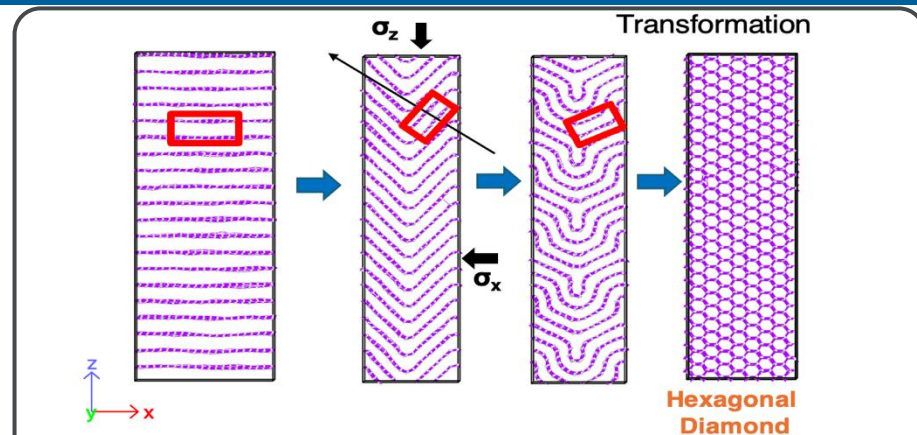
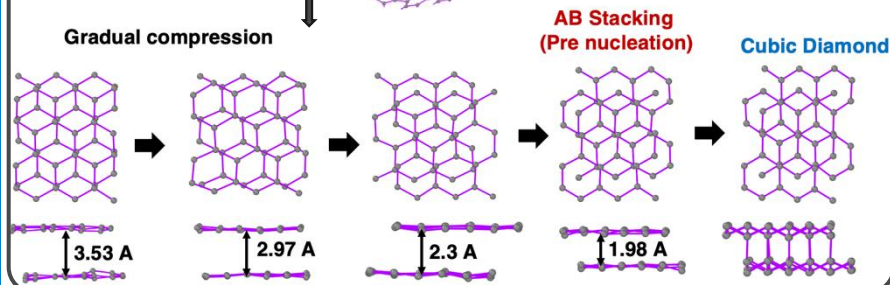
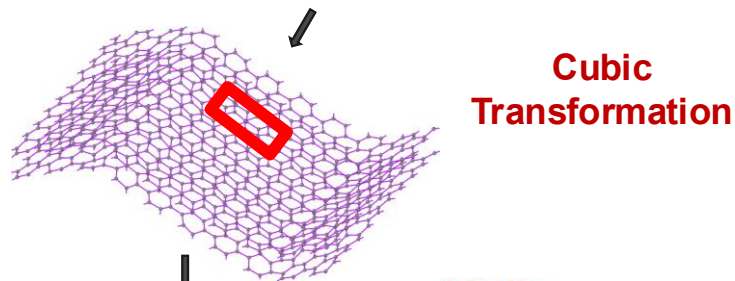
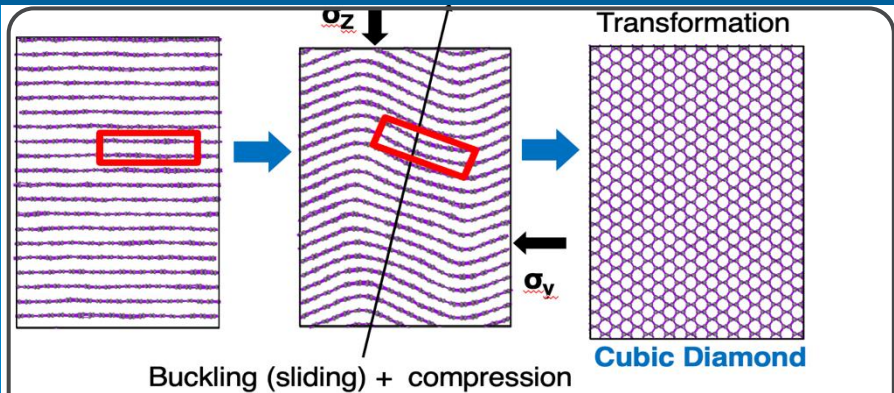
Mechanisms for transformation to Cubic vs. Hexagonal Diamond



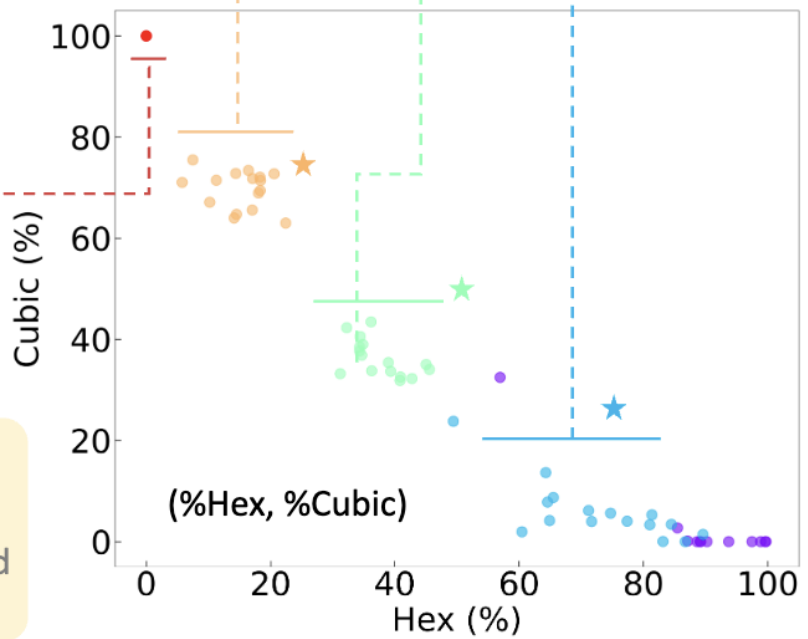
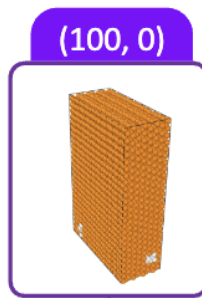
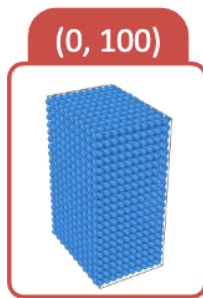
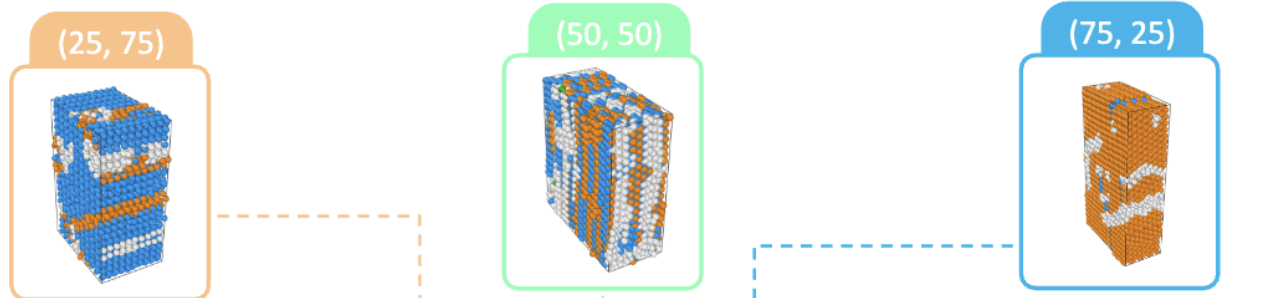
- Graphitic
- Cubic
- Hexagonal



Mechanisms for transformation to Cubic vs. Hexagonal Diamond



Synthesis of mixtures of cubic and hexagonal phases



- Graphite
- Cubic Diamond
- Hexagonal Diamond
- Other States

Future Outlook

- Extension to multicomponent systems
- Fast accurate models to represent atomistic and molecular interactions
- Fingerprinting or scoring function to identify emerging order
- Information extraction from multimodal experiments
- Digital Twins for synthesis of metastable materials
- Autonomous synthesis of materials beyond equilibrium

THANK YOU

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