

# Geometry-Driven Selectivity in Macrocyclization via Rigid-Body Simulation

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# ■ What are Macrocycles?

## □ Definition

- Ring-shaped molecules with  $\geq 12$  atoms
- Applications: drug discovery, catalysis, advanced materials

## □ Synthesis Challenge

- Various ring sizes form simultaneously
- Selective formation of specific size is difficult
- Especially challenging under irreversible conditions

# ■ Surprising Experimental Discovery

## □ This Experiment(Iwamoto et al. 2025)

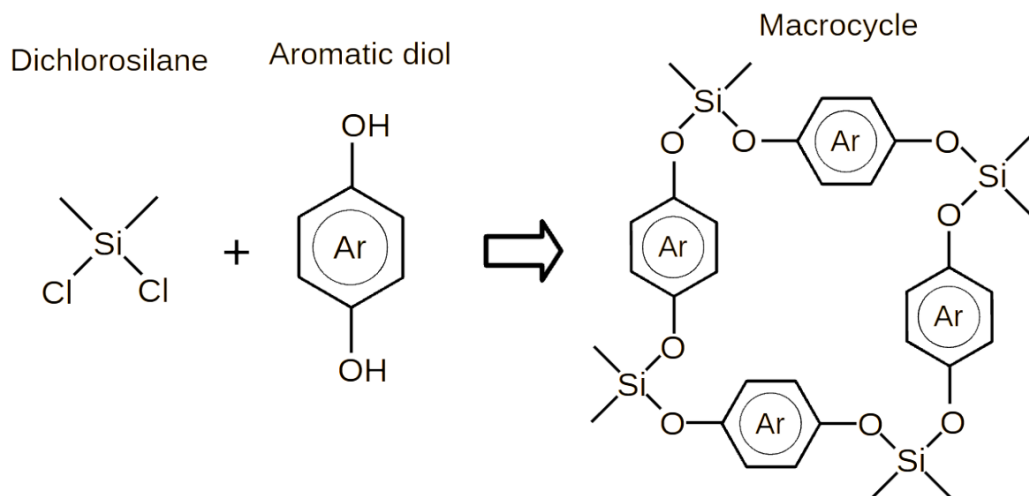
Only squares formed at ~100% yield

Completed within minutes

Works even at high concentration

## □ The Surprise

Selective formation under irreversible conditions — without any special mechanism



History-dependent  
assembly, NOT equilibrium

→ **Why does this happen?**

# ■ Research Objective

## □ RESEARCH QUESTION

**Can geometric constraints alone reproduce the observed selectivity?**

## □ KEY IDEA

Build a minimal model with only geometric rules — bond angles and coplanarity

Strip away chemical details to isolate geometry's role

## □ GOAL

Test if geometry alone can drive selective square formation

If yes → geometry is the key mechanism

→ First, let's define a coarse-grained model...

# ■ Coarse-Grained Rigid-Body Model

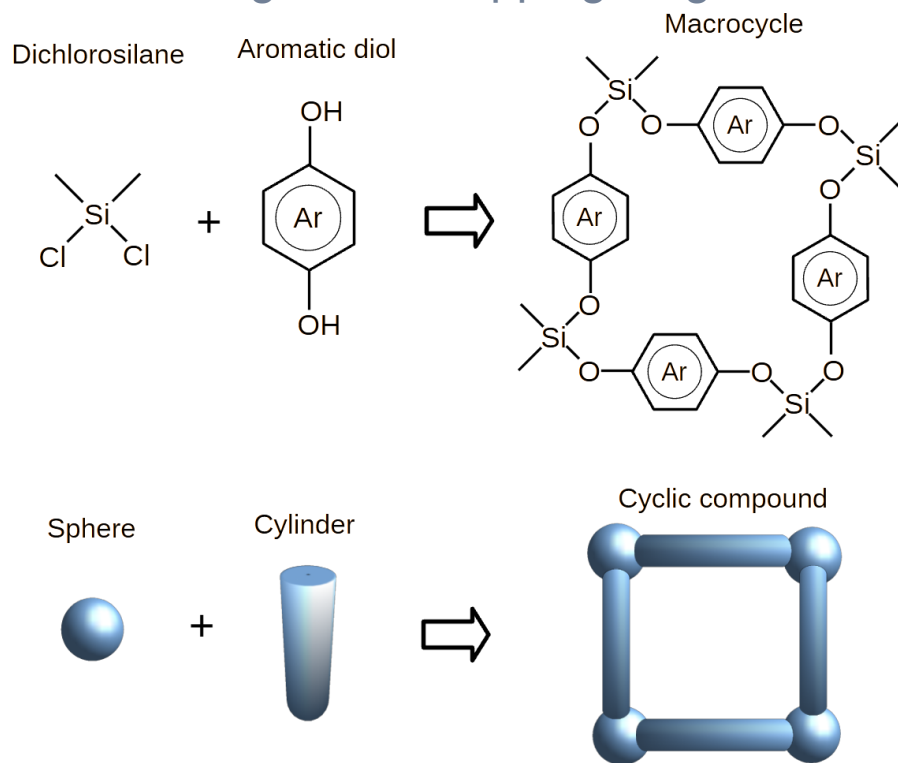
## Sphere (S): Organosilicon reagent

- Compact and isotropic
- Valence = 2

## Cylinder (L): Aromatic diol

- Short rigid rod
- Each end: Valence = 1

## Coarse-grained mapping diagram



## ■ Implementation

NVIDIA PhysX

- Rigid-body dynamics library
- Collision detection + constraint solving

→ Next, what geometric constraints do we impose?

# ■ Geometric Constraints on Bond Formation

## 1. Capture distance

Bond forms when distance  $< 0.1$

## 2. Valence

Sphere  $\leq 2$  bonds, Linker end  $\leq 1$  bond

## 3. Bond angle ( $\theta$ ): $90^\circ \pm 5^\circ$

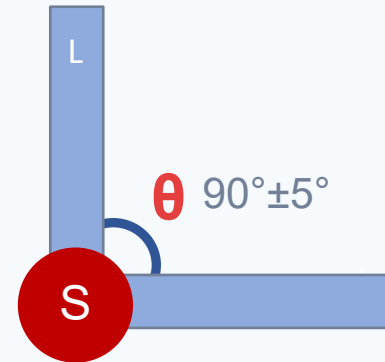
DFT calculation: Ar–O–Si–O–Ar  $\approx 86^\circ$

## 4. Coplanarity ( $\phi$ ): deviation $\leq 5^\circ$

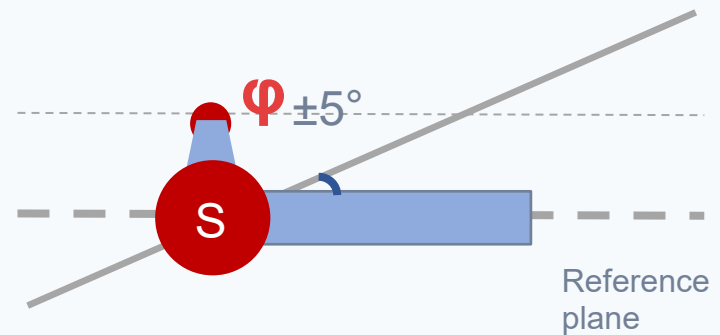
Enforces quasi-2D assembly

Once formed, bonds are irreversible

$\theta$  : Bond Angle



$\phi$  : Coplanarity



→ How do we set up the simulation?

# ■ Simulation Setup

## □ Configuration

- $N = S = 300$   
(Spheres : Linkers = 1:1)
- Box size:  $50^3$
- Reflective boundary conditions

## □ Statistics

- 1000 independent runs
- MSER for steady-state detection

Identifies when simulation reaches equilibrium

Only uses stable data for analysis

## □ Target

Square macrocycle (tetramer)

4 spheres + 4 linkers in alternating ring

## □ Evaluation Metrics

### Target-Object Fraction

→  $\Phi_{\text{target}} = N_{\text{target}} / N_{\text{total}}$

→ How many squares formed?

### Square Selectivity

→  $S_4 = N_4 / N_{\text{cycle}}$

→ How pure is the square yield?

### Viable-Object Fraction

→  $\Phi_{\text{viable}} = N_{\text{viable}} / N_{\text{total}}$

→ How many clusters can still grow?

### $\tau_{\text{target}}$

→ simulation time units

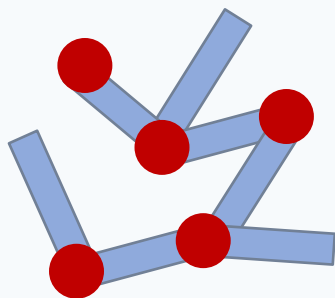
→ How fast did the first square form?

# ■ Passive Geometric Filters: Necessary but Insufficient

Scenario	Constraints	$\Phi_{\text{viable}}$	$S_4$	$\Phi_{\text{target}}$
1	Valence only	26%	24%	6%
2	+ Bond angle	48%	47%	6%
3	+ Coplanarity	94%	57%	<1%
4	Both	97%	80%	<1%

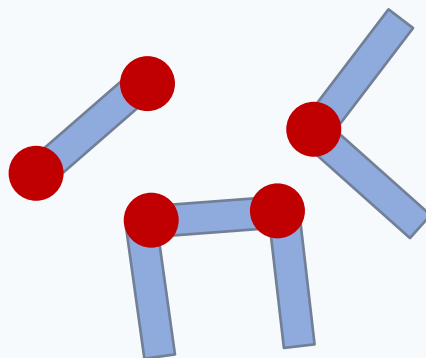
High  $S_4$  achieved, but  $\Phi_{\text{target}}$  remains very low

Without constraints



Chain / Branched

With constraints



Precursors (L-shape, U-shape, Square)

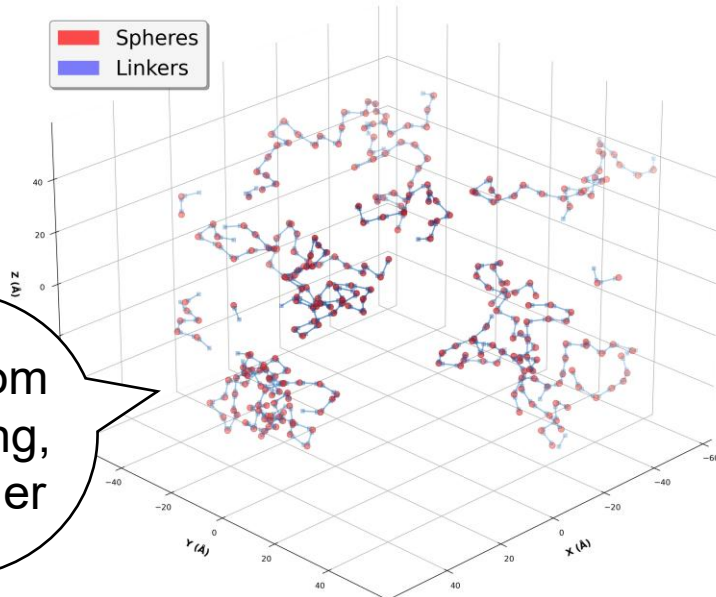
Can post-bond stabilization improve yield?



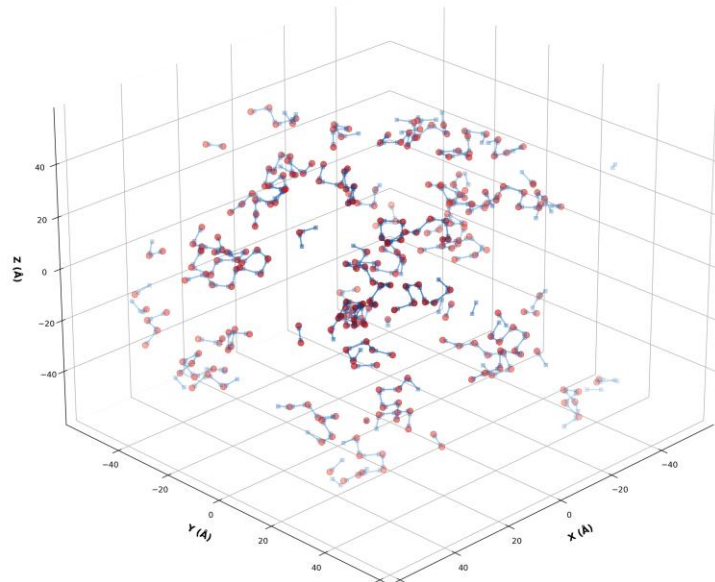
# ■ Scenario1-4: Steady State Snapshots

■ Spheres  
■ Linkers

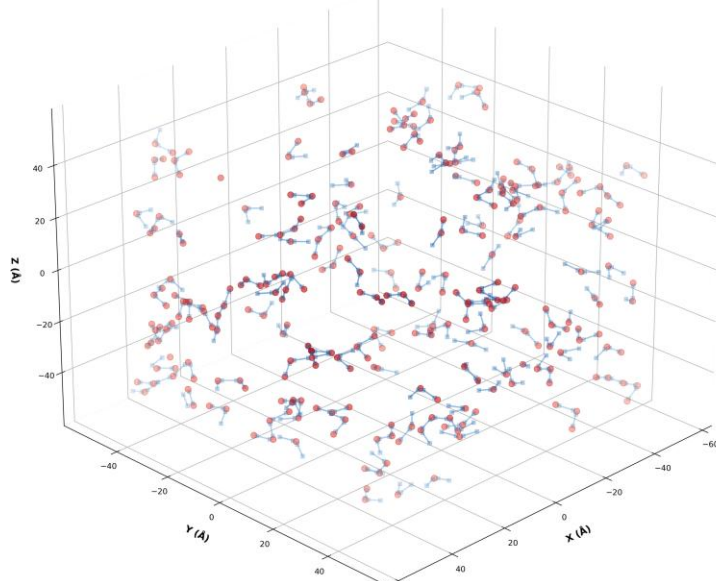
Random  
bonding,  
no order



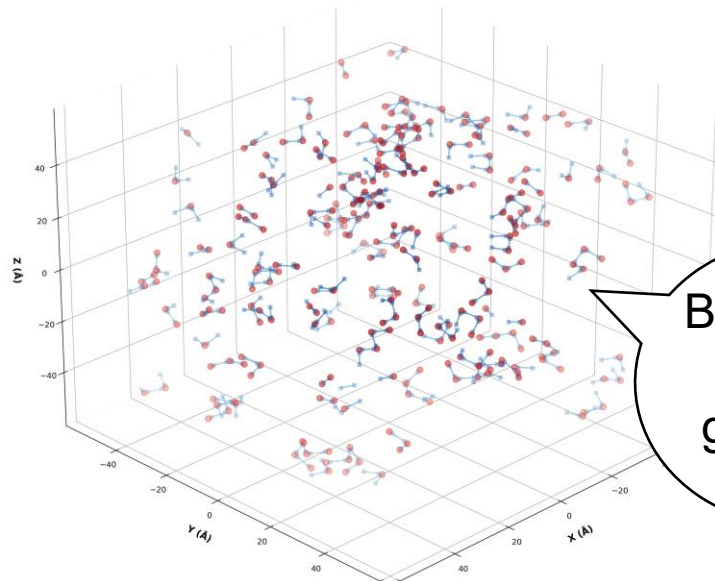
Scenario1: Valence only



Scenario2: +Bond Angle



Scenario3: +Coplanarity



Bonds form  
with  
geometric  
order

Scenario3: Both

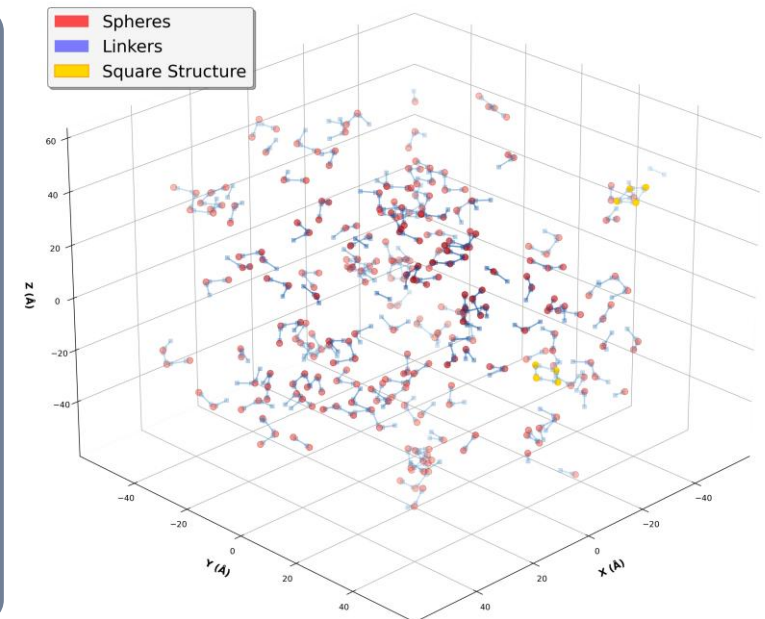
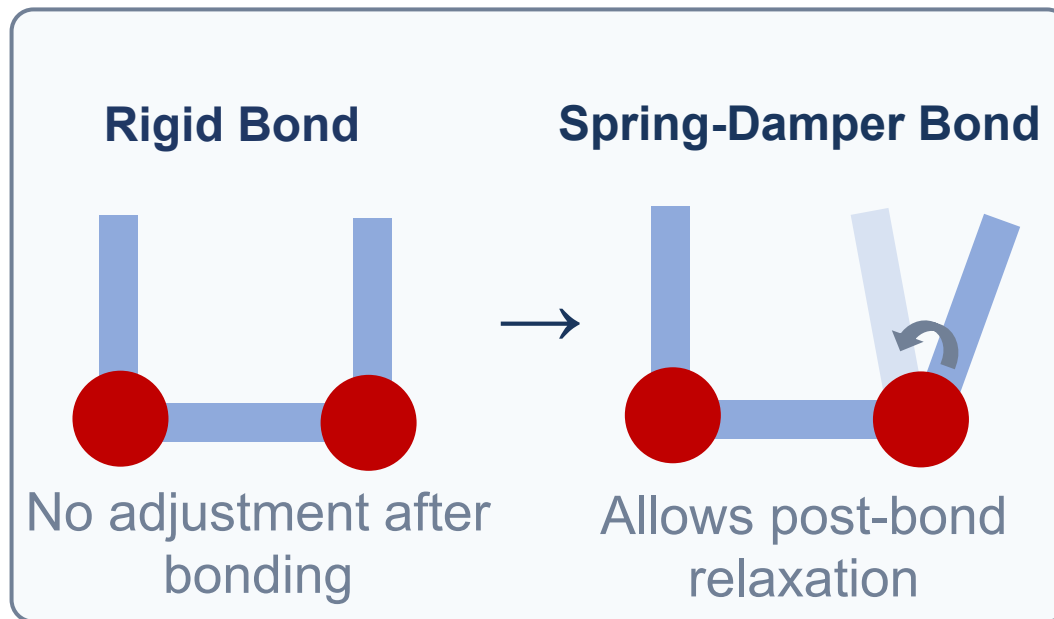
# ■ Compliant Bonds: 19× Target Yield Improvement

## □ Scenario 5: Spring-Damper Bonds

Bonds can flex and relax into stable configurations

Scenario	Description	$\Phi_{\text{viable}}$	$S_4$	$\Phi_{\text{target}}$
4	Rigid constraints	97%	80%	0.37%
5	+ Spring-damper	92%	85%	<b>6.97%</b>

**19× increase** in  $\Phi_{\text{target}}$  (0.37% → 6.97%)



→ What about collision frequency (density)?

# ■ High Density: Accelerated While Maintaining Selectivity

## □ Scenario 6: Density Sweep

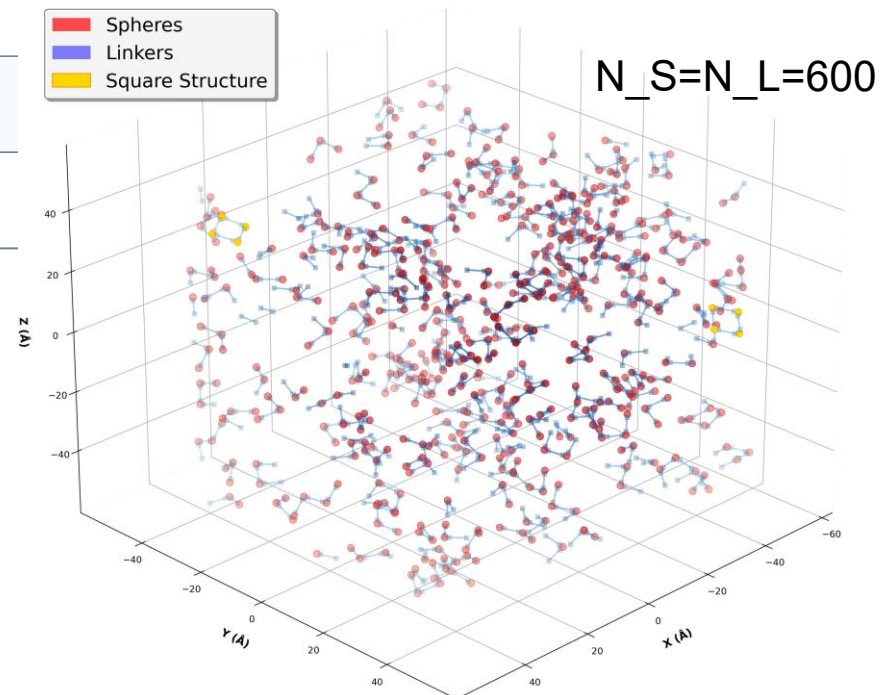
Same as Scenario 4 (rigid bonds +  $\theta$ ,  $\phi$  constraints)

Box size fixed at  $50^3$

Increase  $N_S$  and  $N_L \rightarrow$  higher density

$N_S = N_L$	$\tau_{\text{target}}$	$S_4$	$\Phi_{\text{viable}}$
150	3280	84%	100%
300	2340	80%	97%
600	2040	83%	95%

Higher density  $\rightarrow$  faster  $\tau$   
(target formation)  
 $S_4$  stable at 80–84%  
Viable fraction remains high



$\rightarrow$  These results suggest synergy of three elements...

# ■ Three Elements for Selective Macrocyclization

**1**

## Strict Local Geometry

Bond angle + Coplanarity

$\Phi_{\text{viable}}$ : 26%  $\rightarrow$  97%

$S_4$ : 80%

**2**

## Post-Bond Stabilization

Spring-Damper mechanism

$\Phi_{\text{target}}$ : 0.4%  $\rightarrow$  7%

$\tau$  reduced 4×

**3**

## Appropriate Collision Frequency

Density control

Accelerated formation

$S_4$  maintained: 80–84%

**Selectivity emerges from the synergy of all three elements**

# ■ Future Directions

## □ Parameter Mapping to Experiments

Correspondence with angular rigidity, hemilabile lifetime, linker ratios

## □ Extended Architectures

Multi-component systems, templates, alternative ring sizes

## □ Pathway Analysis

Trajectory descriptors: path entropy, kinetic funneling

This minimal model framework can be extended to diverse assembly problems

# ■ References

- [1] Martí-Centelles, V., Pandey, M.D., Burguete, M.I., Luis, S.V.: Macrocyclization Reactions: The Importance of Conformational, Configurational, and Template-Induced Preorganization. *Chemical Reviews* 115, 8736–8834 (2015)
- [2] Iwamoto, T., Amano, S., Maeda, K., et al.: Exclusive macrocyclization through multiple Si–O bond formations from diol and dichlorosilane. *Chemical Communications* 61, 8180–8183 (2025)
- [3] NVIDIA PhysX. <https://github.com/NVIDIA-Omniverse/PhysX/>
- [4] Chodera, J.D.: A Simple Method for Automated Equilibration Detection in Molecular Simulations. *J. Chem. Theory Comput.* 12, 1799–1805 (2016)