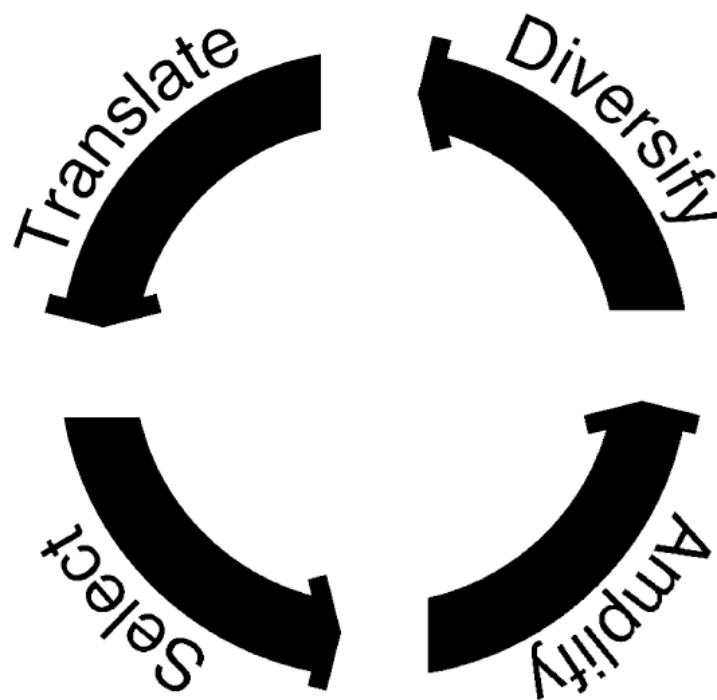


Chemistry from Nature

~ the Application of
Evolutional Principals~

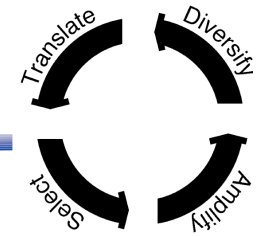


Literature Seminar 3

11/4/26

Yutaka Saga (D1)

Today's Contents



0. Introduction

1. Genetic Algorithms

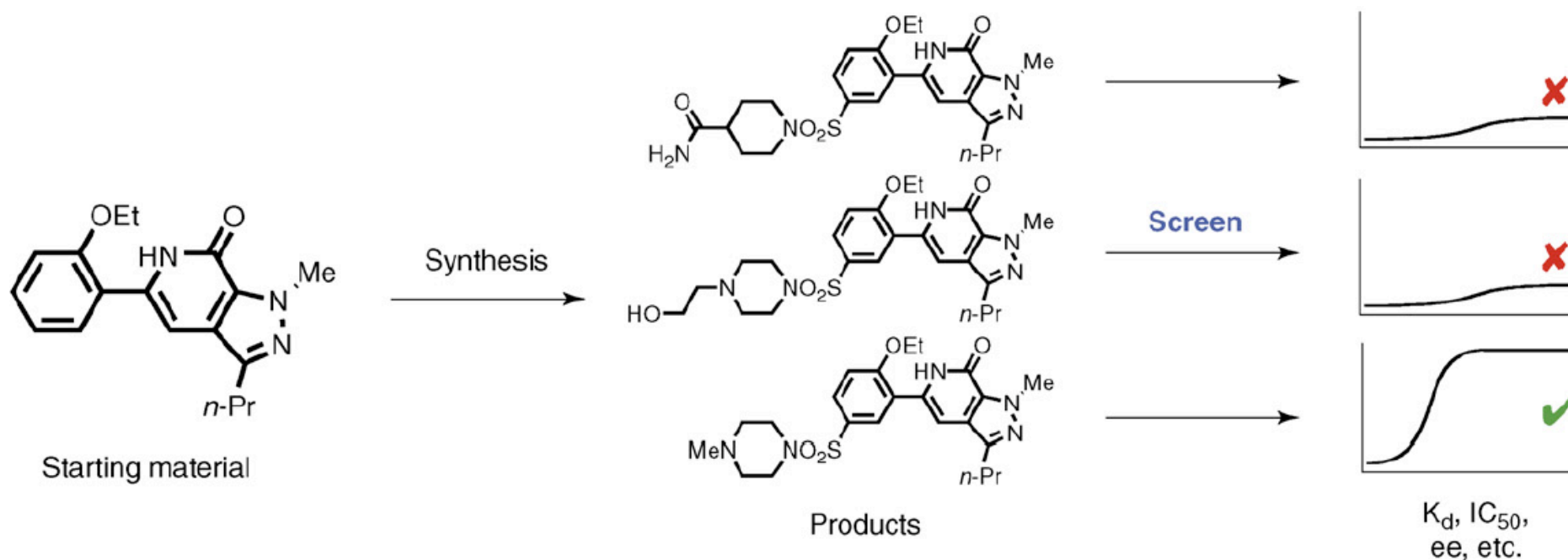
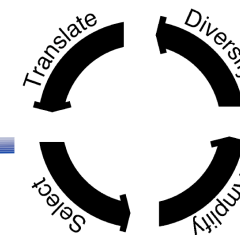
2. Dynamic Combinatorial Chemistry

3. DNA-Templated Synthesis

4. Future Direction

0. *Introduction*

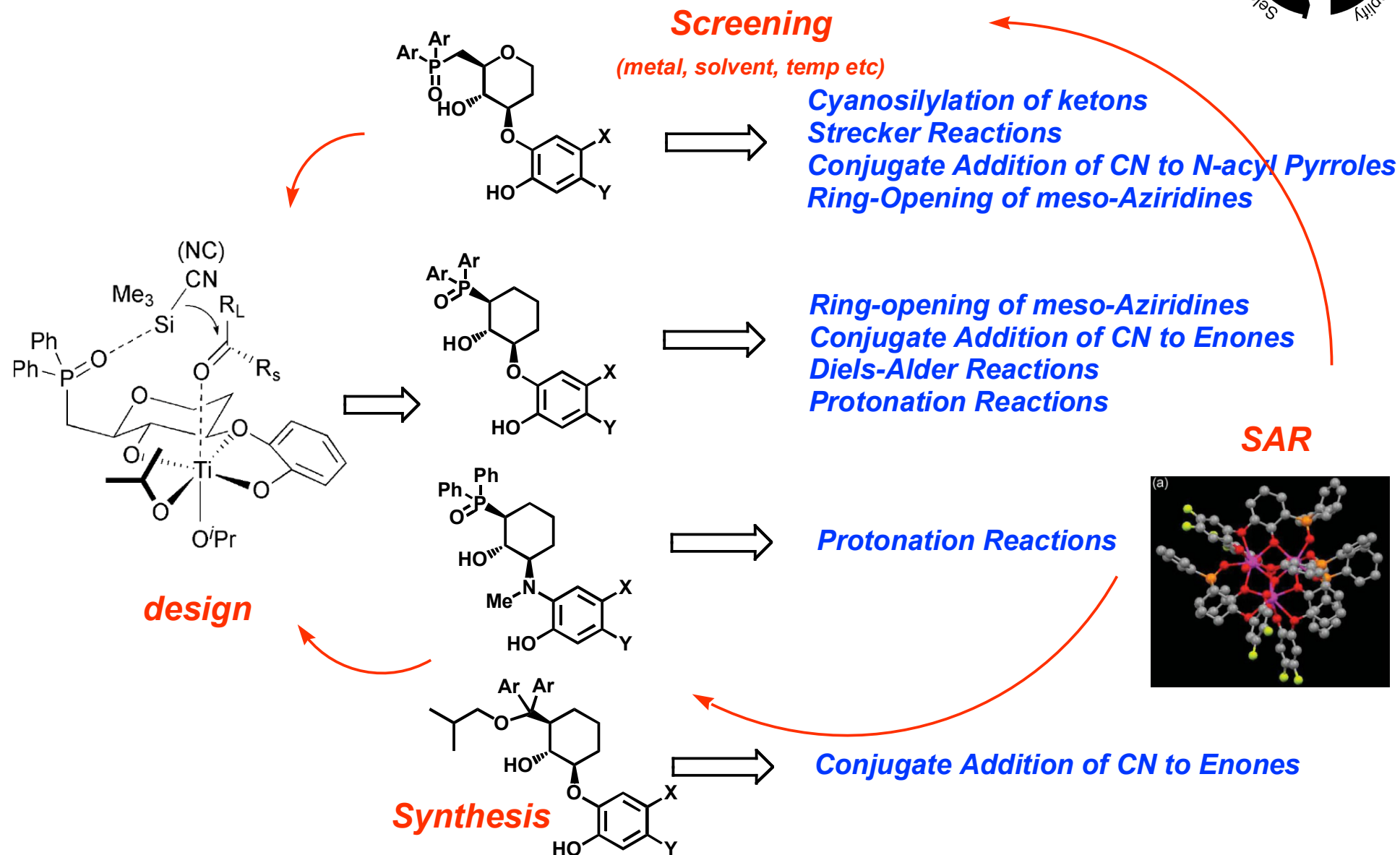
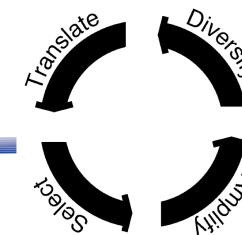
Traditional Chemical Approaches



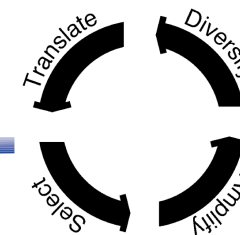
➤ Chemists specifically have been interested in the discovery of *functional molecules (catalysts, drugs, materials etc)*.

➤ Iterated cycle
design, synthesis, activity assay, SAR(structure activity relationship)

Chemistry of GluCAPO & FujiCAPO

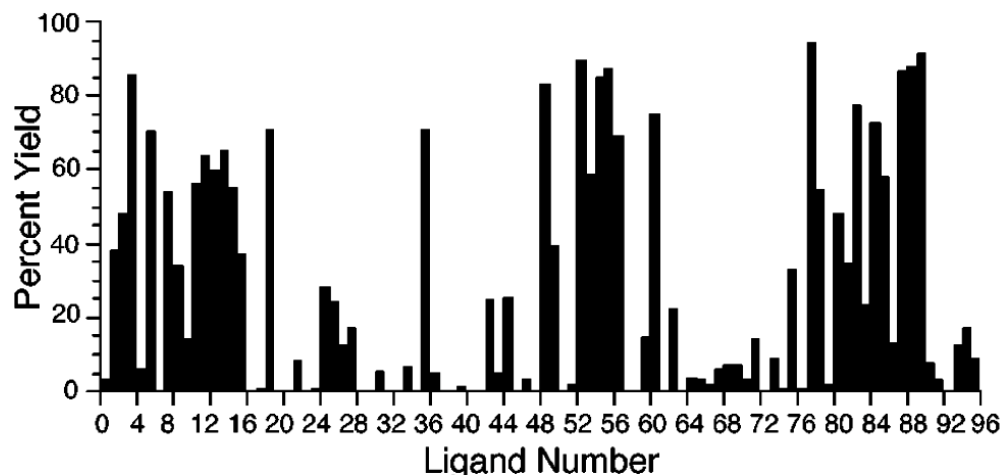
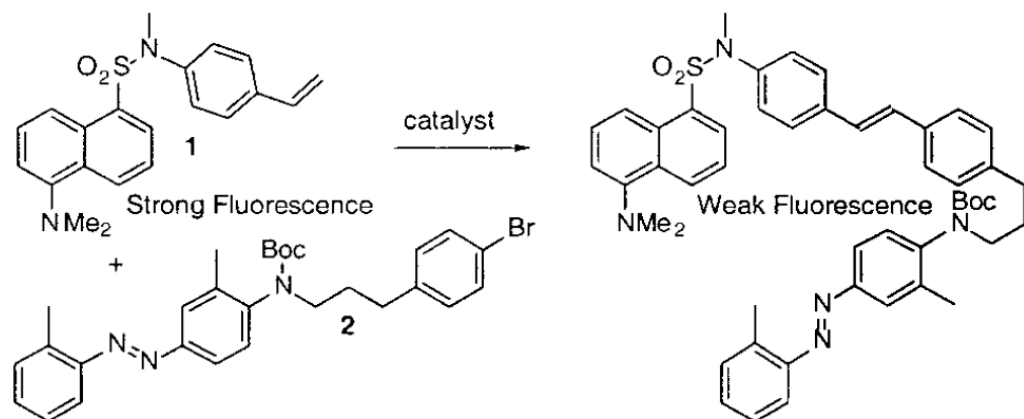


Overcoming the Limitations (1)



HTS (High-Throughput Screening) assays

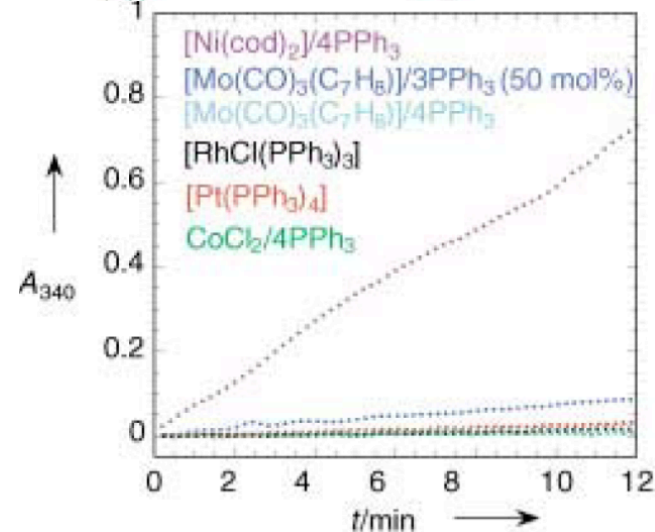
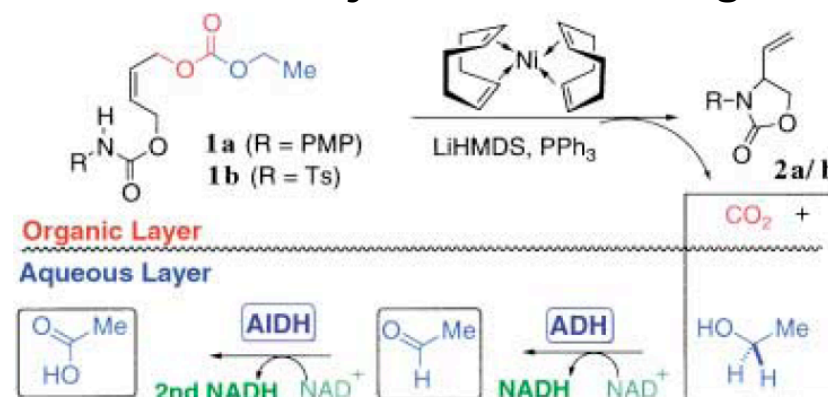
HTS of Heck reactions by using **FRET**



J. F. Hartwig *et al.* *JACS.*, 2001, 123, 2677

With **HPLC**, **MS**, **colorimetric**, **IR**, etc

In Situ Enzymatic Screening

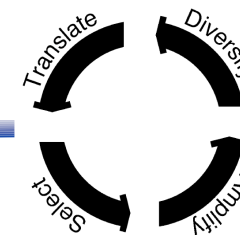


D.B. Berkowitz *et al.* *ACIE.*, 2002, 41, 1603

J. F. Hartwig *et al.* *Curr. Opin. Chem. Biol.*, 2003, 7, 420

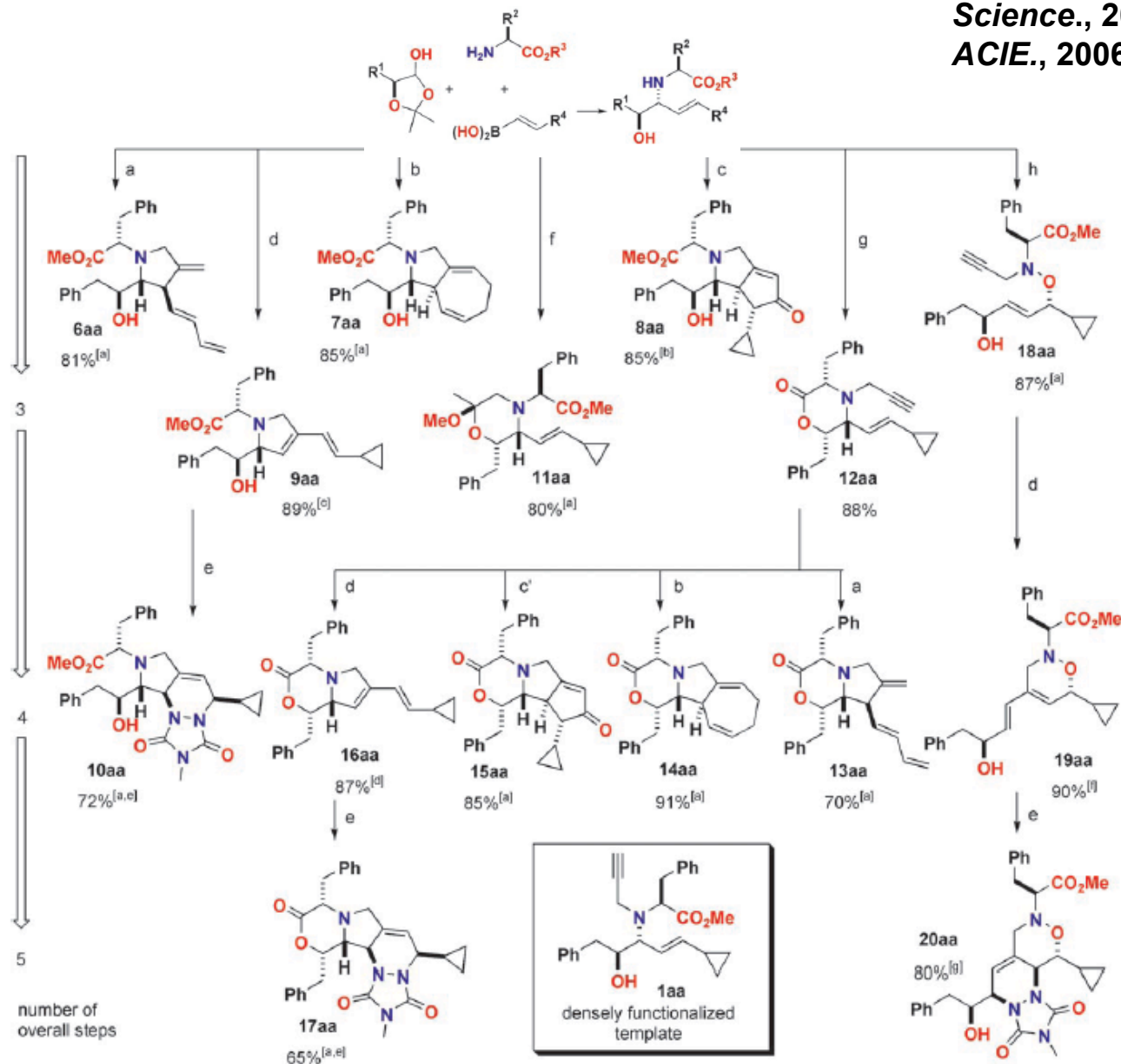
Overcoming the Limitations (2)

Diversity-Oriented Synthesis



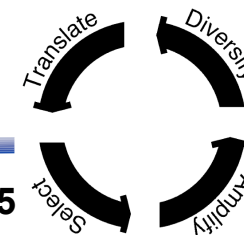
S.L. Schreiber *et al.*
Science., 2000, 287, 1964
ACIE., 2006, 45, 3635

Simple SM



Complexity,
Diversity

Overcoming the Limitations (3)

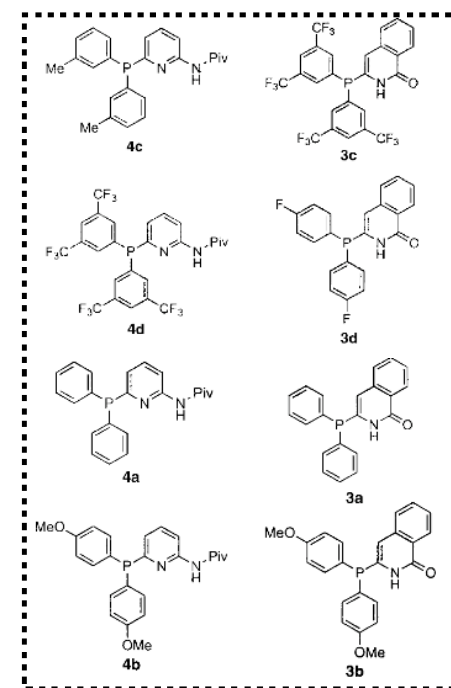
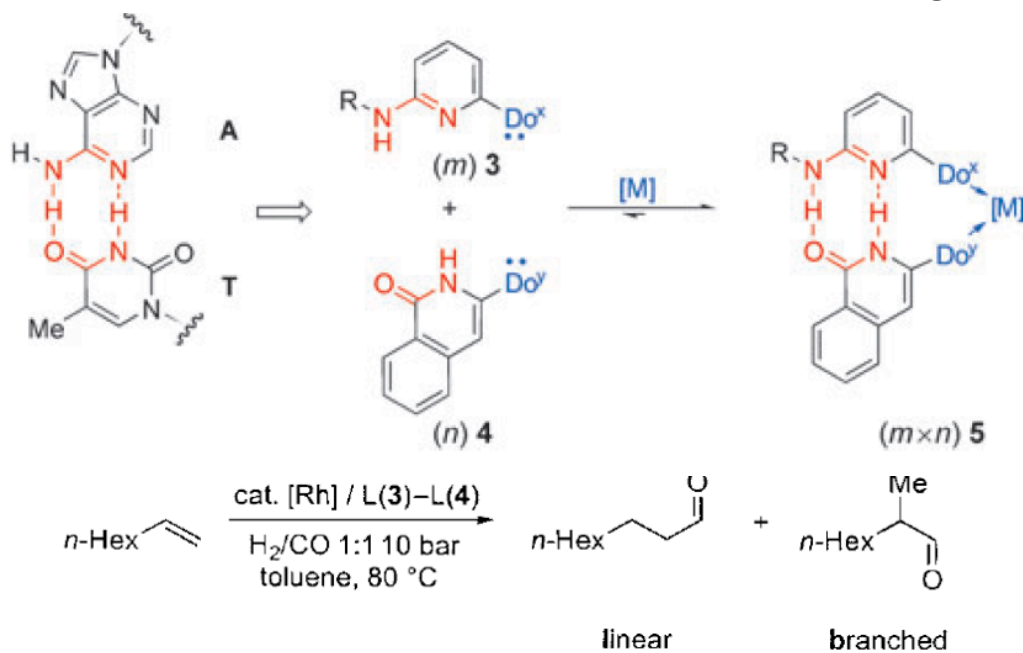


Self-Assembled Libraries

N.H. Reek *et al. Nat. Chem.*, 2010, 2, 615

B.Breit *et al. ACIE.*, 2005, 44, 1640

Dr. Shimizu's Lit. Seminar (D3)

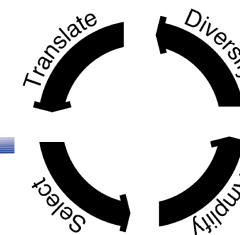


L(4)	L(3)			
	3 a	3 b	3 c	3 d
4 a	2425 h ⁻¹ [b] 94:6[c]	1040 h ⁻¹ 94:6	2732 h ⁻¹ 96:4	2559 h ⁻¹ 95:5
4 b	2033 h ⁻¹ 93:7	1058 h ⁻¹ 92:8	1281 h ⁻¹ 96:4	1772 h ⁻¹ 94:6
4 c	3537 h ⁻¹ 94:6	1842 h ⁻¹ 93:7	1808 h ⁻¹ 96:4	2287 h ⁻¹ 94:6
4 d	7439 h ⁻¹ 96:4	2695 h ⁻¹ 95:5	7465 h ⁻¹ 94:6	8643 h ⁻¹ 96:4

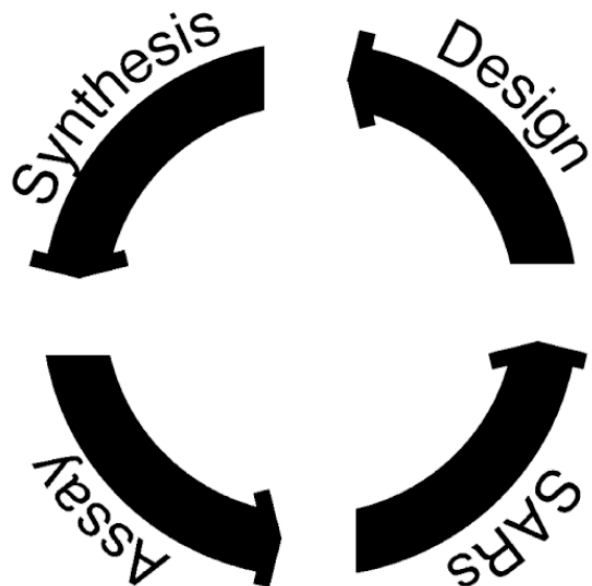
- > large catalyst libraries
- > just mixing components
- > Not covalently bonded, with multiple weak interactions

Also see: T. Ooi *et al. Science.*, 2009, 326, 120

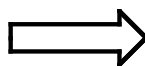
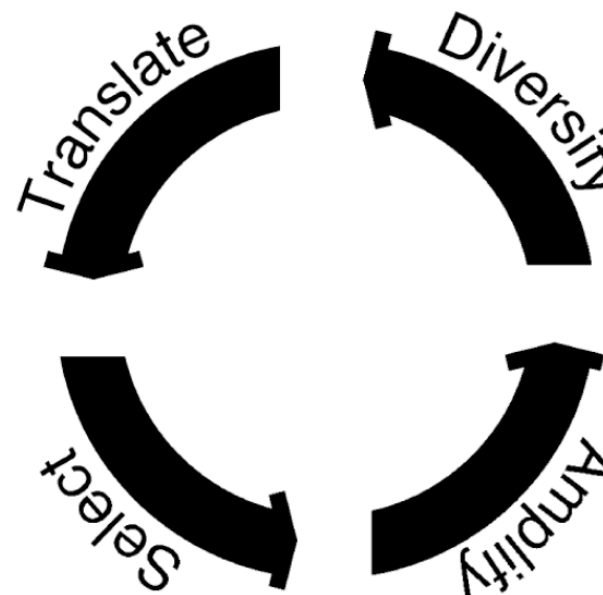
Toward Evolutional Approaches (1)



Chemists' Approaches



Nature's Approaches



> **limited** diversities and complexities

> **individually** evaluated (homogeneous)

> **spatial separation**

> **high** conc

> **vast magnitude** of diversities and complexities

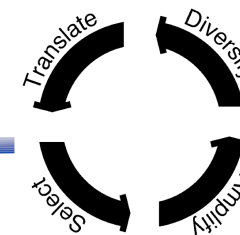
> **in one-pot** evaluated (heterogeneous)

> **without** spatial separation

> **very low** conc

> **new reaction discovery ???**

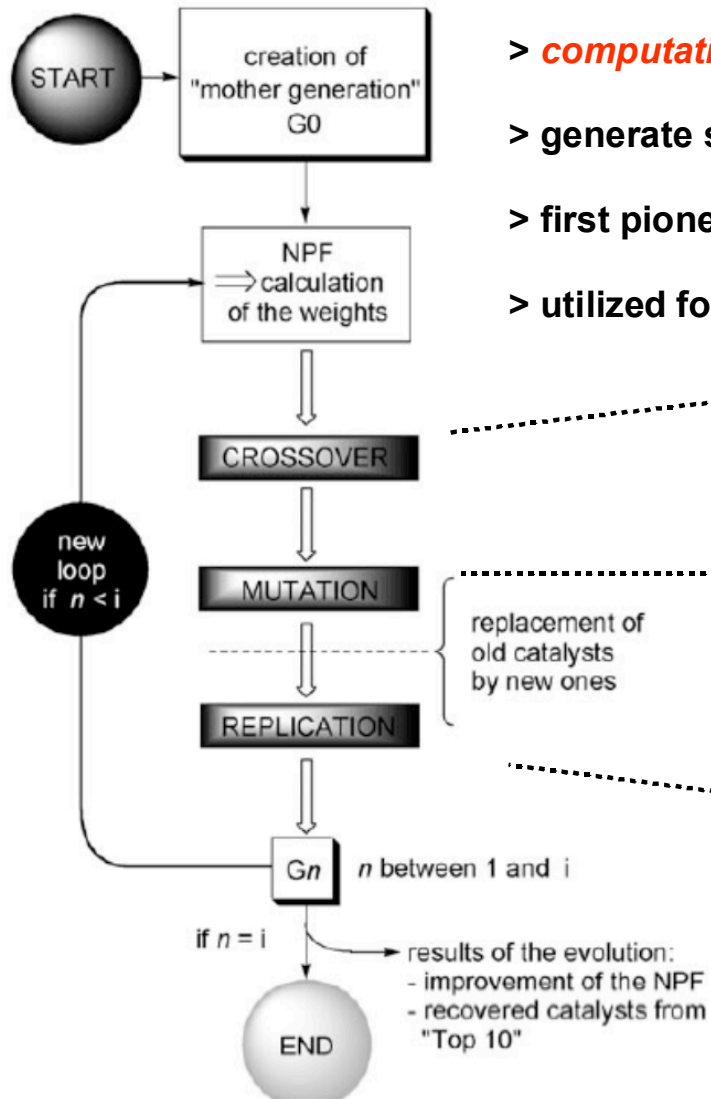
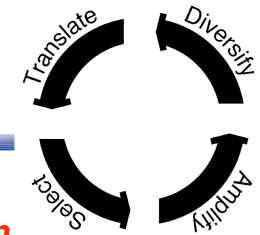
Toward Evolutional Approaches (2)



	<u>Chemist's Approach</u>	<u>Nature's Approach</u>
Identifying Hits	individually in spatially separated <i>assays</i>	one pot <i>selections</i>
Material Quantity	$> \sim 10^{12}$ copies of each molecule	$< \sim 1,000$ copies of each molecule
Sample Size	$\leq 10^6$ members per screen (CAS $< 10^8$)	$\leq 10^{15}$ members per selection
Generality	whatever can be synthesized	??? nucleic acids & proteins

1. Genetic Algorithms

GA (Genetic Algorithm)



- > a search heuristic that *mimics the process of natural evolution*.
- > *computationally mimic* Darwinian evolution
- > generate solutions to optimization problems inspired by natural evolution
- > first pioneered by Holland in the 1970s
- > utilized for *numerous fields* (computer, mathematics, biology, economics etc)

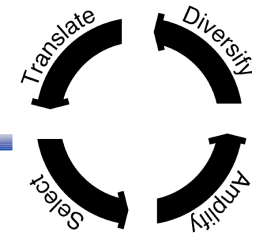


- > adjusting several parameters for the evolution process (G_0, R, RE, RM, RR, NI etc)

J.H.Holland *Adaptation in Natural and Artificial Systems*
Univ.of Michigan Press, 1975.

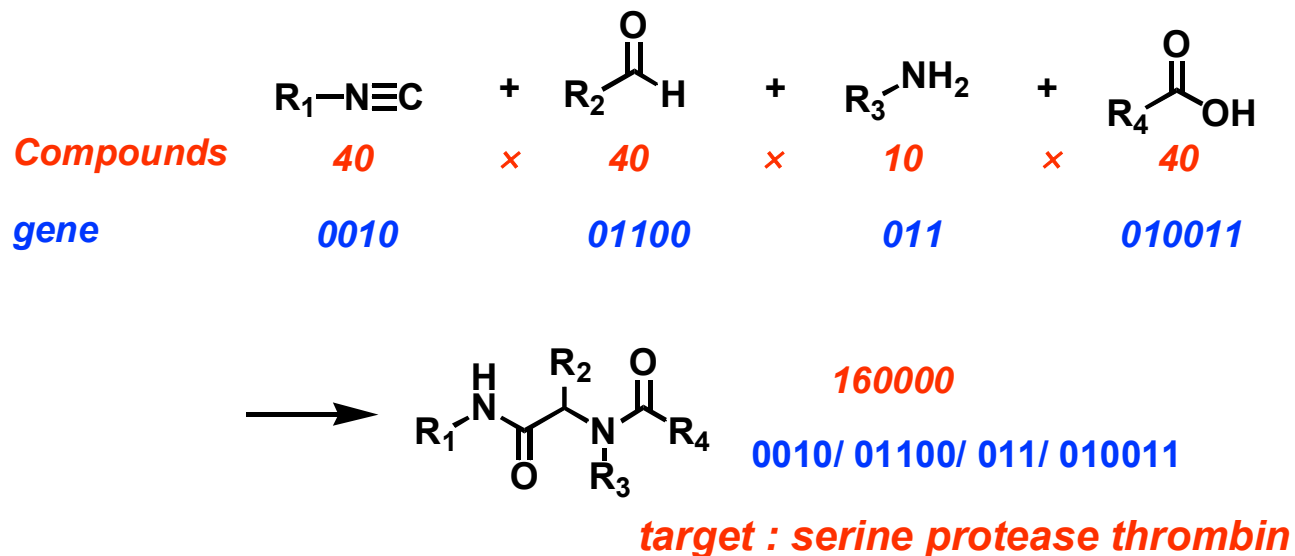
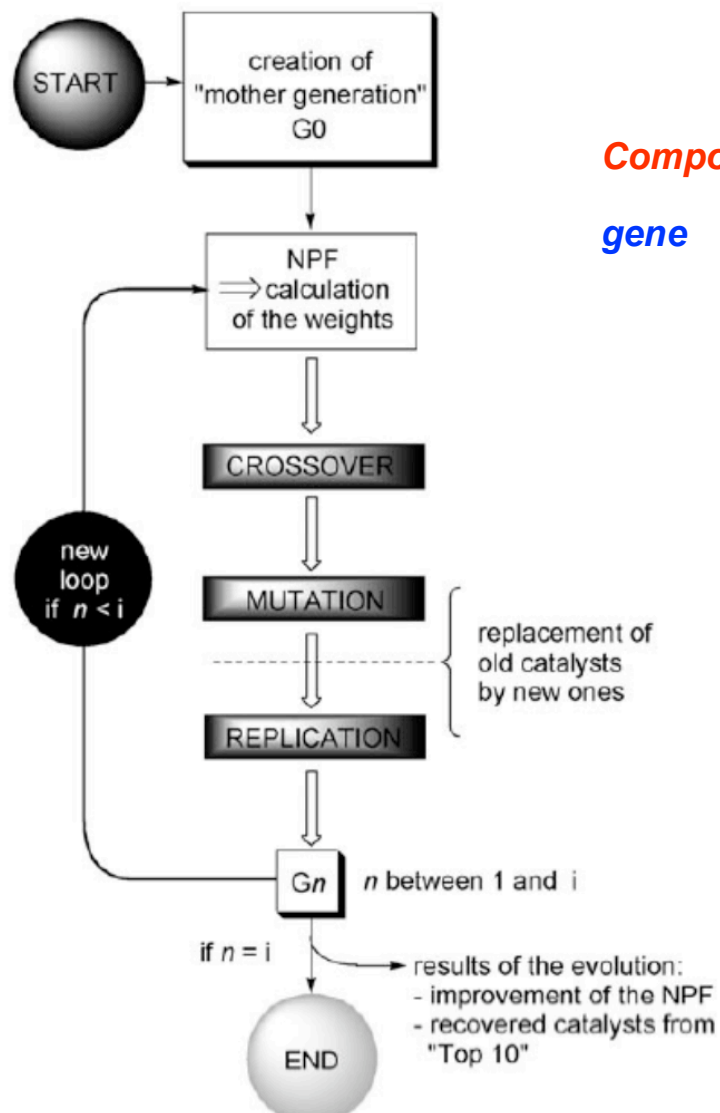
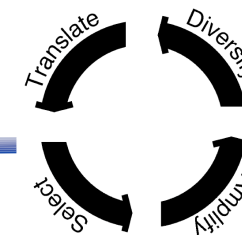
L. Weber et al. *Curr. Opin. Chem. Biol.*, 1998, 2, 381

GA (Genetic Algorithm) for Japanese Game



- the player cultivating vegetables on a fictional, futuristic star system.
- the objective is to win vegetable contests.
- pest called Baboo undergo the evolution based on **'genetic algorithm'**

Ambitious Application of GA (1)



Crossover

0010/ 01100/ 011/ 010011 \rightarrow 0010/ 01100/ 011/ 100011
 0010/ 01100/ 011/ 100011 \rightarrow 0010/ 01100/ 011/ 010011

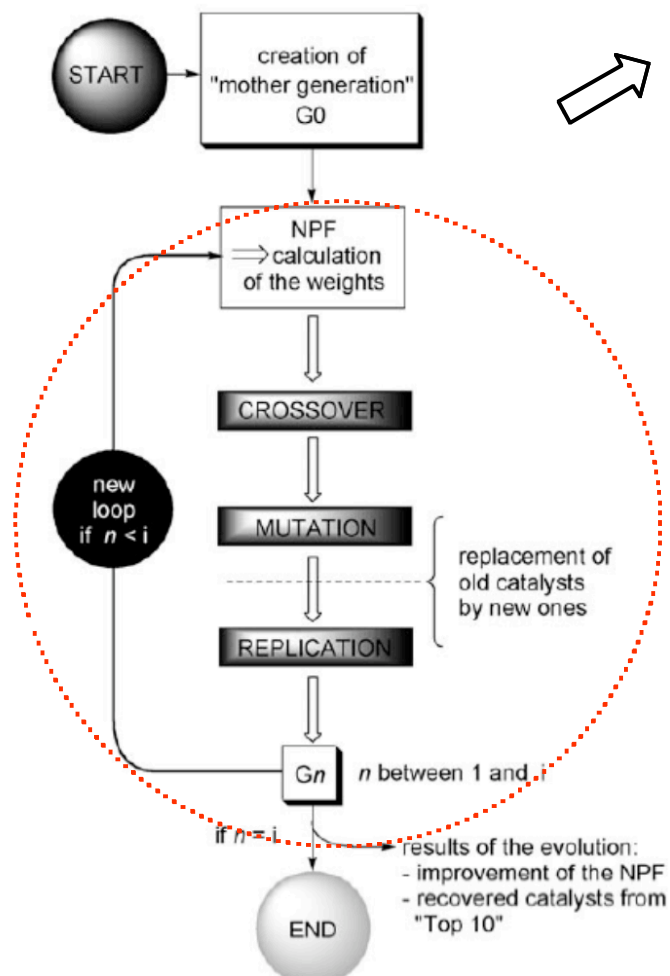
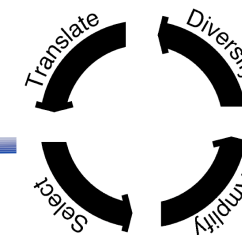
Mutation

0010/ 01100/ 011/ 010011 \rightarrow 0010/ 01100/ 011/ 000111

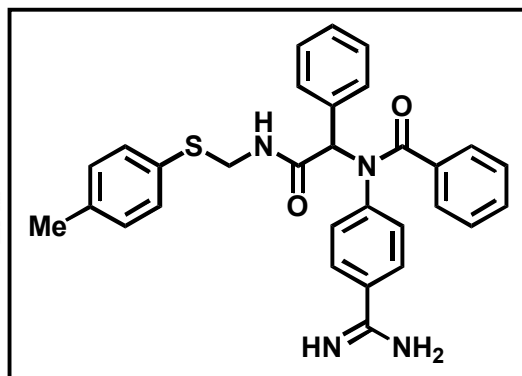
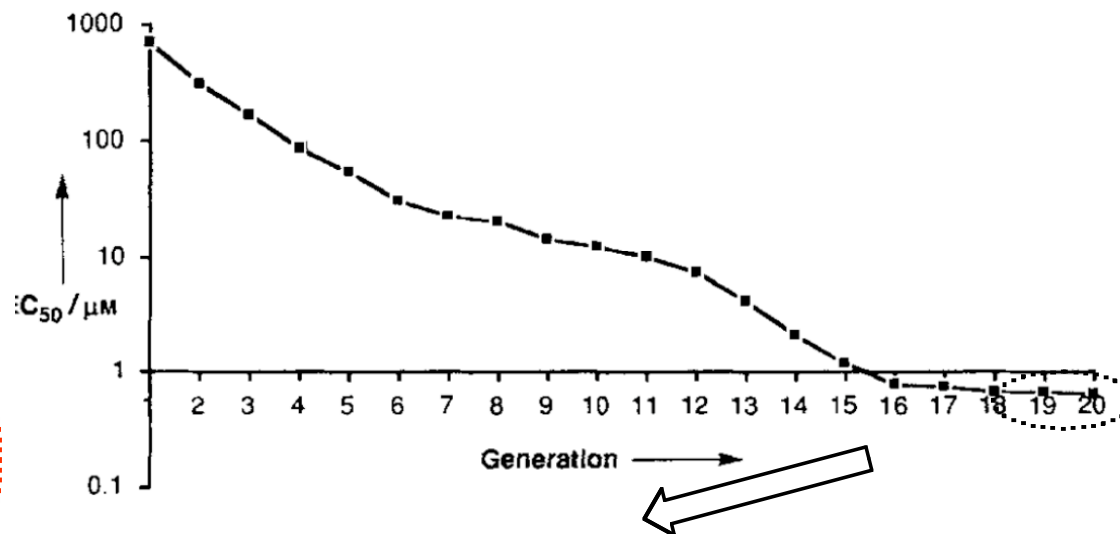
Replication

0010/ 01100/ 011/ 010011 \rightarrow 0010/ 01100/ 011/ 010011

Ambitious Application of GA (2)

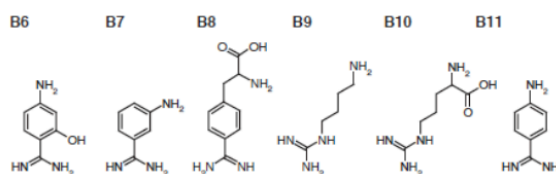
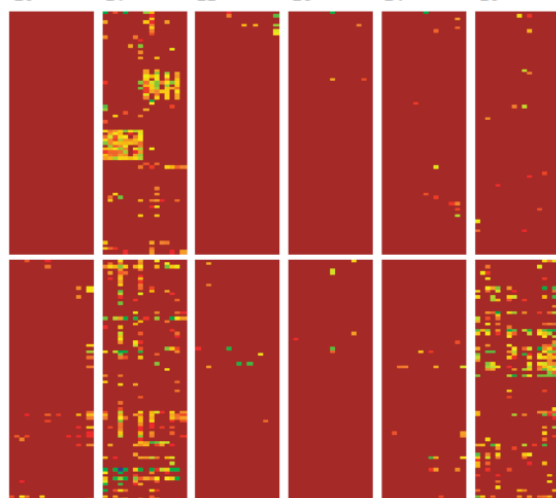
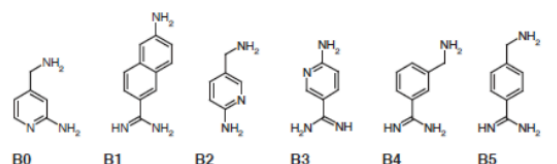
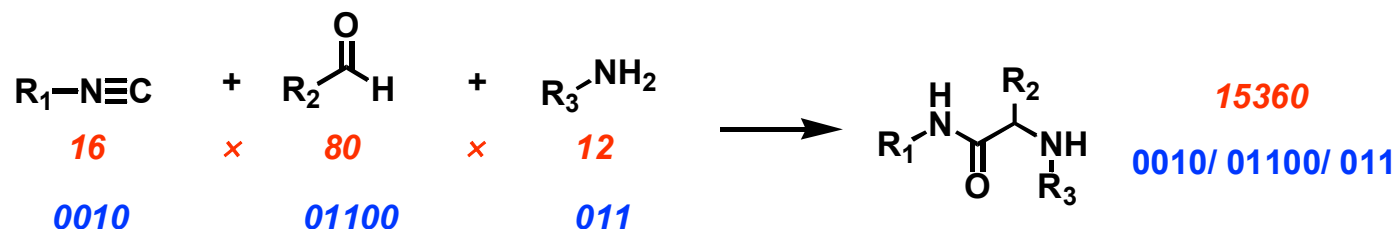
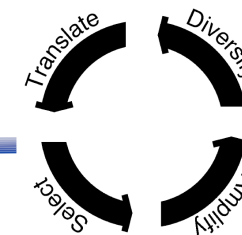


best 20 compounds × 20 generations



- > the best inhibitor exhibited $K_i < 1 \mu M$
- > **only 400 reactions** out of 160000 reactions
- > totally adjusting GA principals for targeting
- > but **nonpeptidic** (no structural biases)

Ambitious Application of GA (3)



> choosing the amines *with a structural bias* toward binding pocket (hydrogen-donor, hydrogen-acceptor etc)

⇒ in a manner analogous to *proteins or nucleotic acids*

> 15360 reactions were actually performed to provide library.

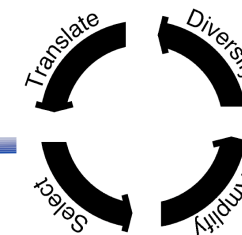
> IC₅₀ values were evaluated to be coded by spectral colors

> IC₅₀ < 1μM: 0.059 %, 1~10μM: 0.352 %, 10~100μM: 4.395 %

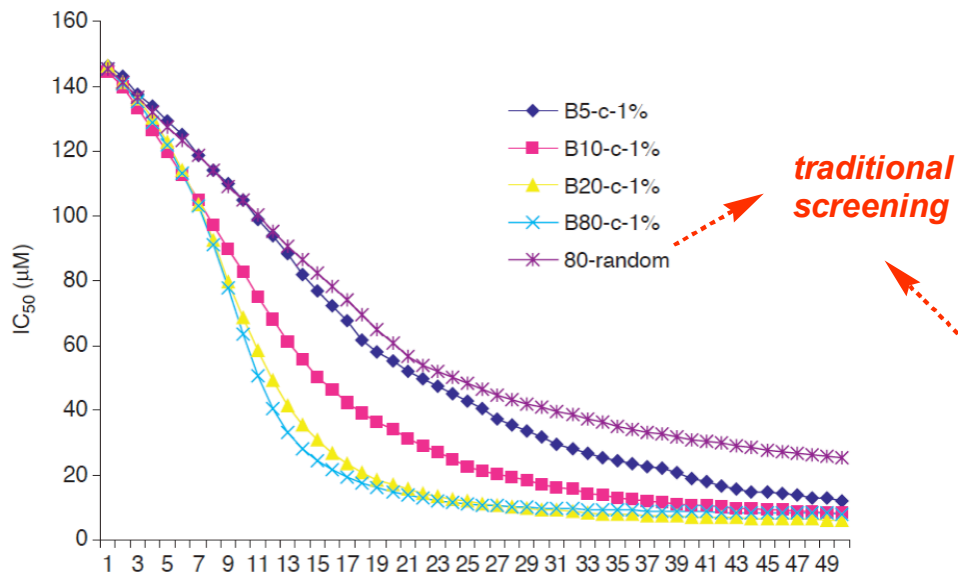


simulate GAs to *optimize the various parameters*

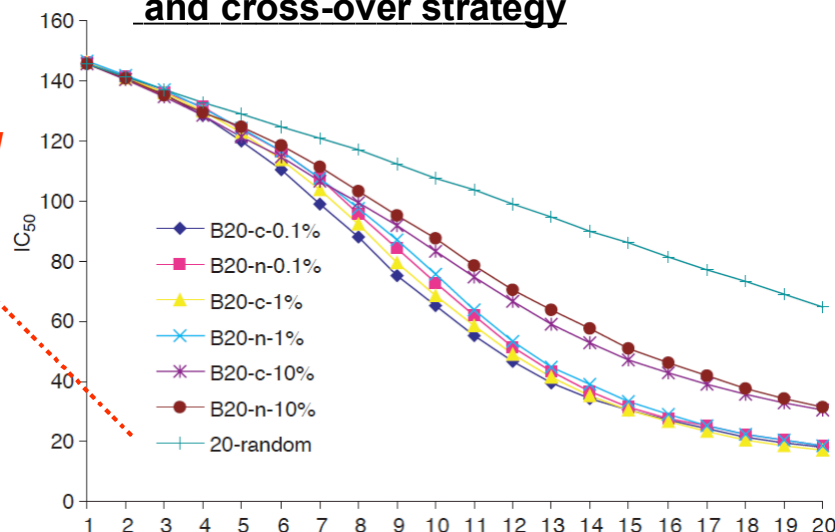
Ambitious Application of GA (4)



Optimization of generation size



Optimization of the mutation rates and cross-over strategy



> P (*the average activity*) = $(m_{GA}/N_{GA}) / (m_{random}/N_{random})$
 (m_{GA} , m_{random} = the slopes of the performance curves)

> $P = 23.6, 14.1, 10.3, 3.1$ ($N = 5, 10, 20, 80$)

> $P \times N = 118, 141, 206, 248$ ($N = 5, 10, 20, 80$)

> small N *learns more per individual*

> large N *learns faster*

> *judge the efficiency* of GAs or other methods

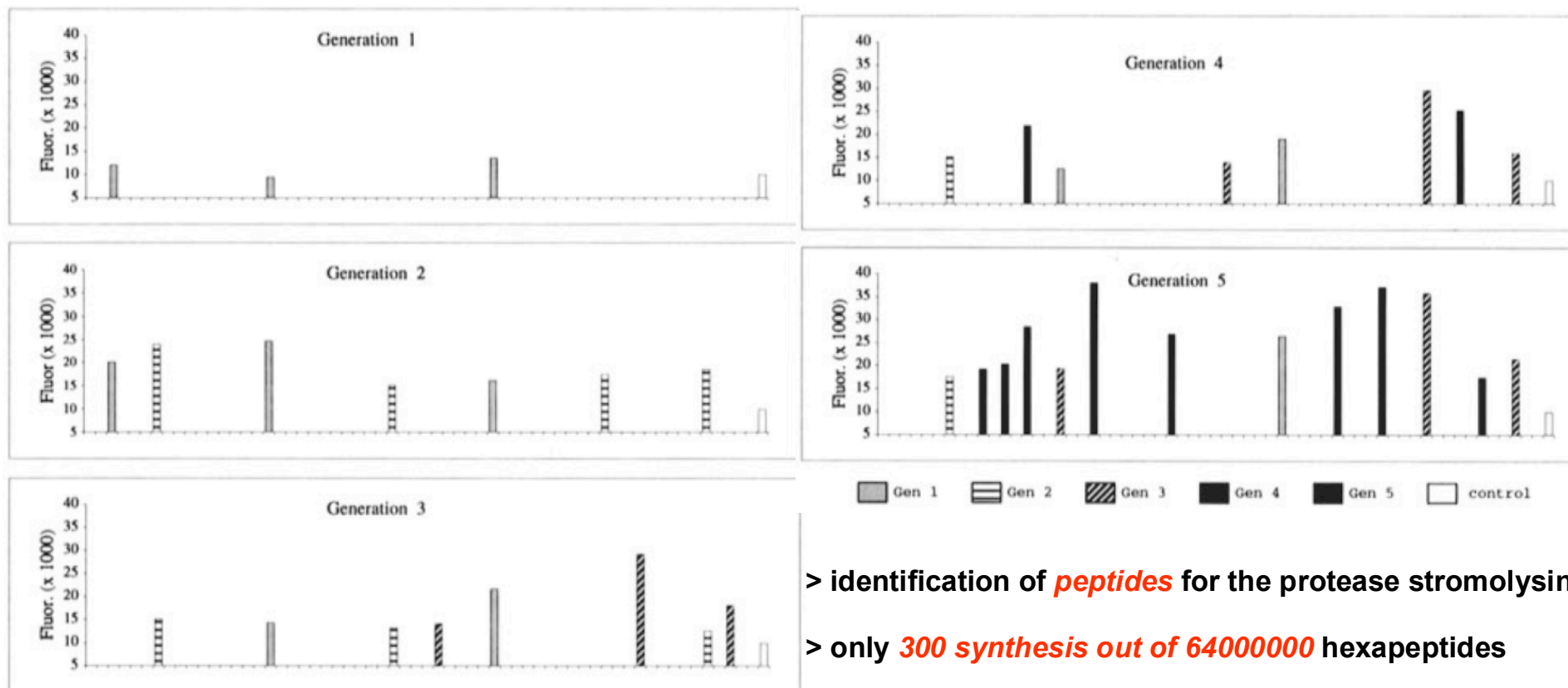
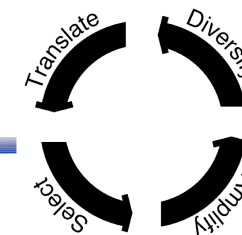
> c = crossover only between starting materials
 n = DNA-like crossover at any bit

> c *exerts a little better performance* despite of lower diversity

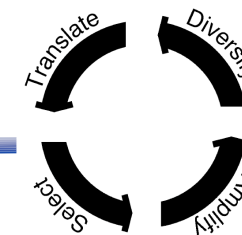
> *increasing the mutation rate deteriorated* the performance

→ *mutation destroyed* the acquired knowledges

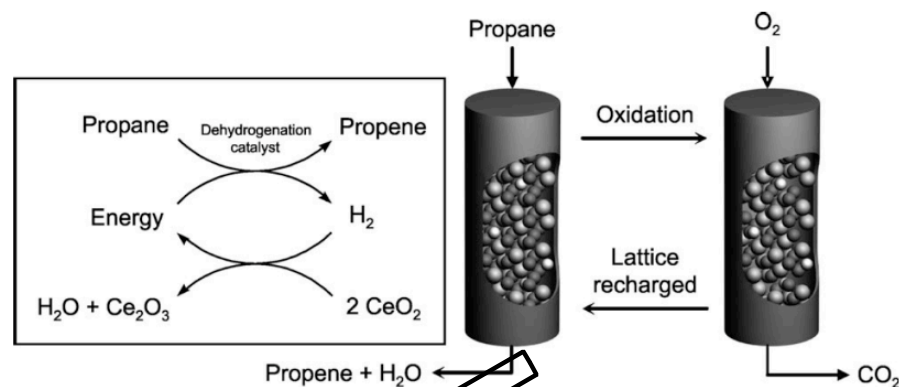
Various Applications of GAs (1)



Various Applications of GAs (2)

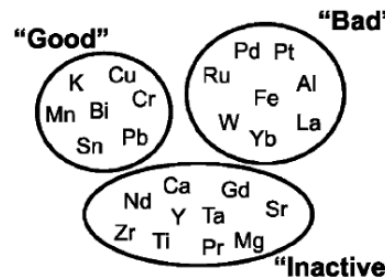
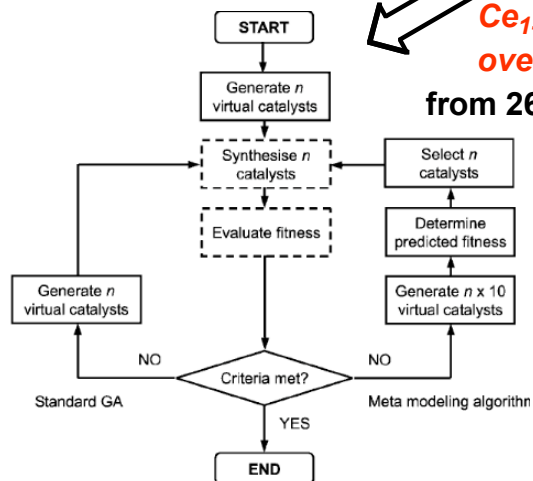


Selective Hydrogen Oxidation Catalysts via GAs

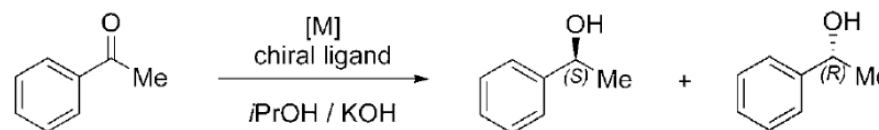


$Ce_{1-x-y}M^1_xM^2_yO_2$ dopants
over 17000 combinations

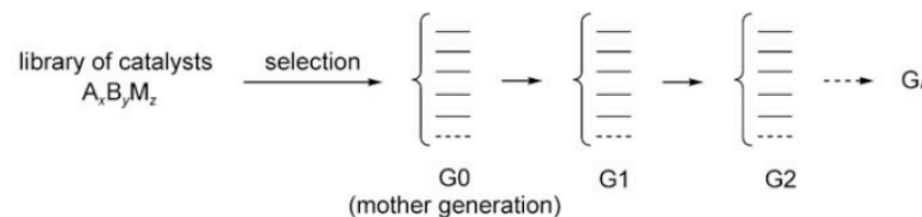
from 26 metals in 5 cat amounts



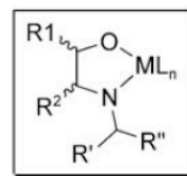
Asymmetric Hydrogenation Catalysts via GAs



A: chiral amino alcohol × 11 B: aldehyde × 30
C: metal (Ru, Ir) × 6

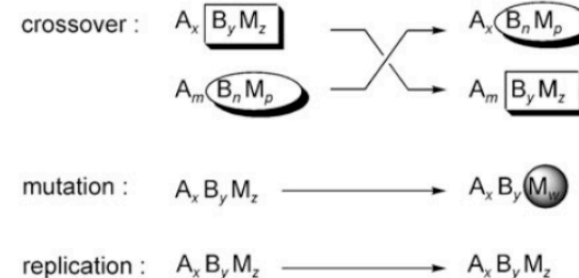


general catalyst structure



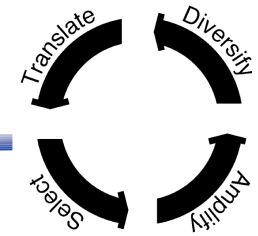
genetic code
 $A_x B_y M_z$

genetic operators



- > the best catalyst found by evaluating **only 10 % of the whole**
- > a **time reduced to 30 %** compared to the traditional methods
- > development of **double algorithm**

Features of GAs



Advantages

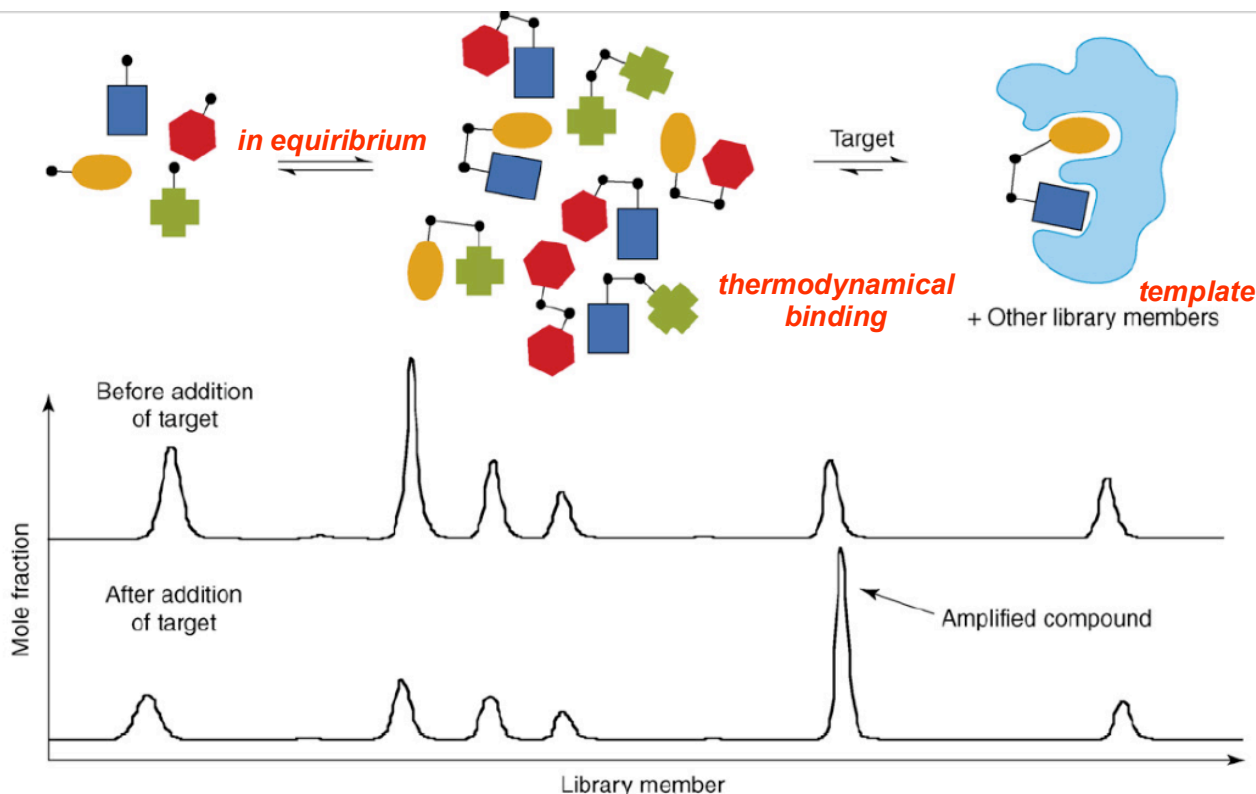
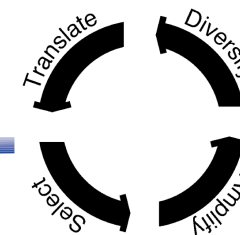
- > mimicking the evolutionary cycle in Nature
→ realizing *the diversity close to Nature*
- > a small fraction of the *'virtual library'*
→ not time-consuming
- > equally applied to *synthetic molecules* as to DNA or proteins

Disadvantages

- > the need to *synthesize and assay individual molecules*
→ What's the difference from traditional methods?
- > *insufficient* size and complexity
- > beneficial only for *the known developed systems*
→ known things of which *parameters set in advance*

2. Dynamic Combinatorial Chemistry

Dynamic Combinatorial Chemistry



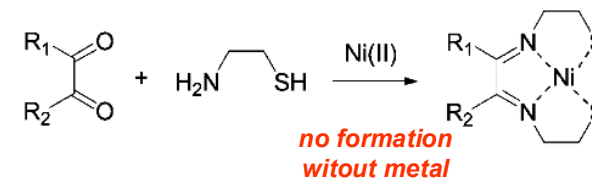
@ Fischer's study on carbohydrates in 19th century



@ Watson & Crick's DNA helix templates in 1950s



@ Busch's first example in 1960's



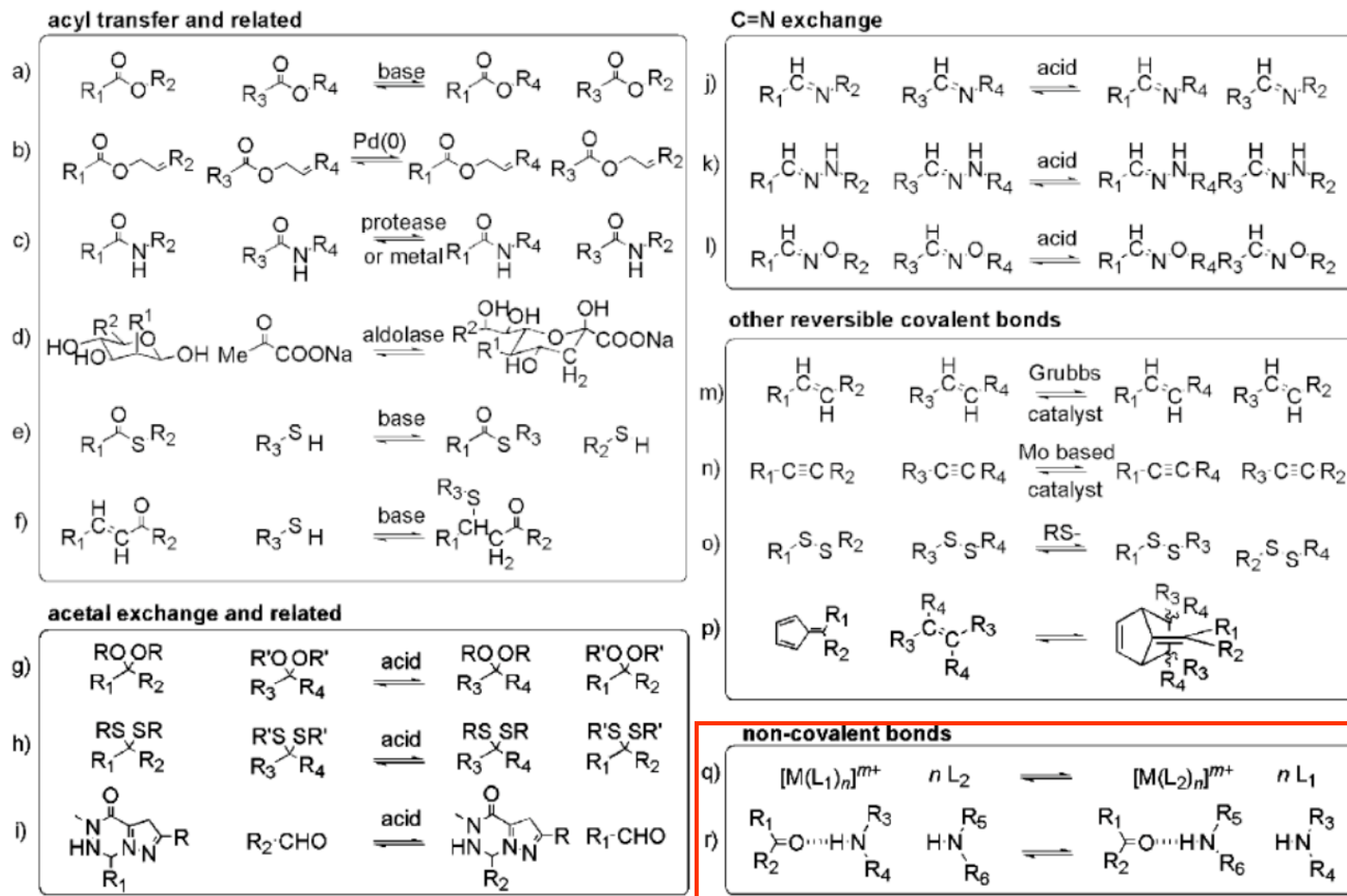
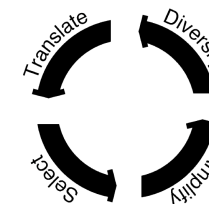
D. H. Busch et al. JACS. 1962, 84, 1762

- > the composition is determined by the **thermodynamic stability, in equilibrium**.
- > upon the addition of **templates**, desired molecules can form the stable complex.
- > not screen, but selection
- evaluated **in mixtures**, not indivisually
- > for the discovery of **small molecules, catalysts, and materials**

J.K.M. Sanders, S. Otto et al. Chem Rev., 2006, 106, 3652

D.R.Liu et al. Curr. Opin. Chem. Biol., 2007, 11, 259

Reversible Reactions Used for DCC

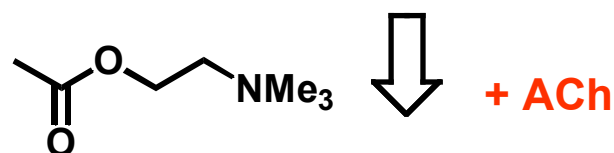
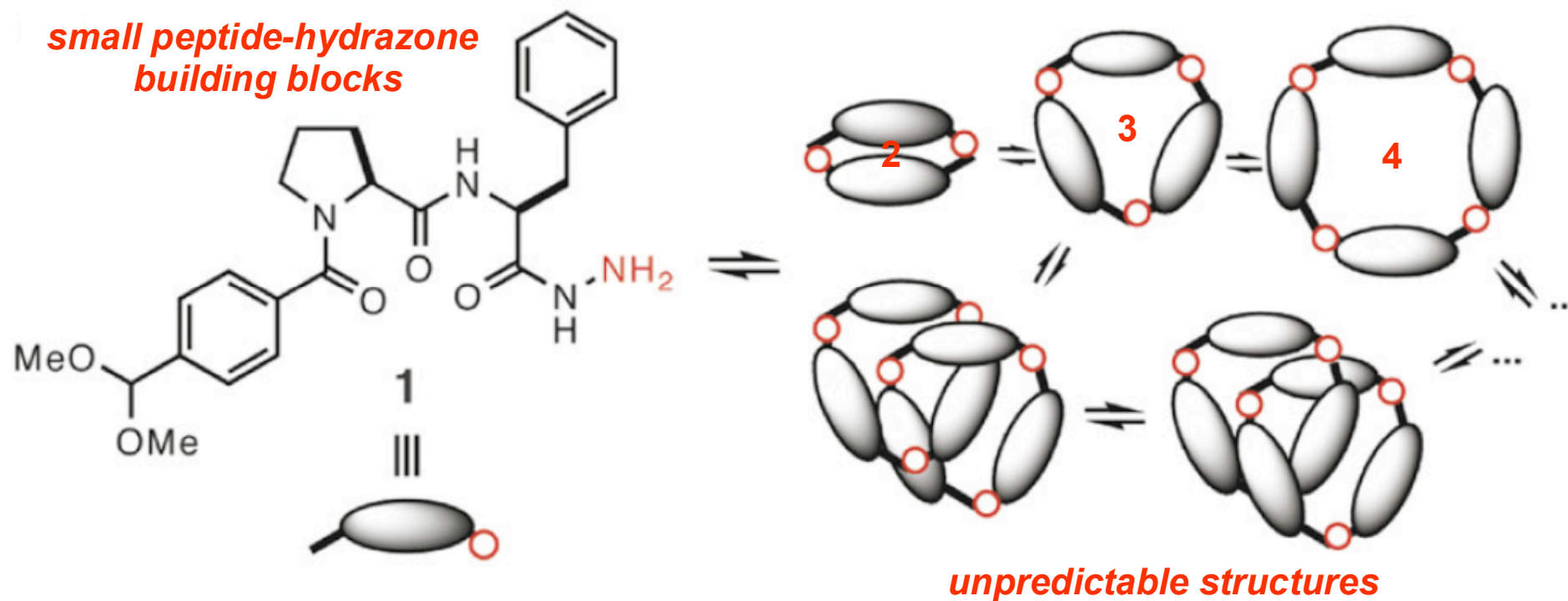
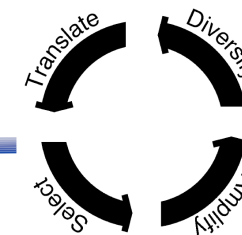


> weak and labile noncovalent bonds to achieve **rapid equilibrium**

VS

covalent bonds to ensure **thermodynamic stability**

DCC for Amplification of Molecules (1)

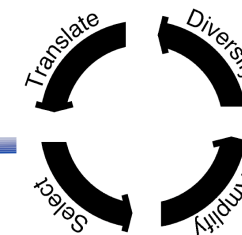


J. K. M. Sanders et al. *Org. Biomol. Chem.* **2003**, *1*, 1625

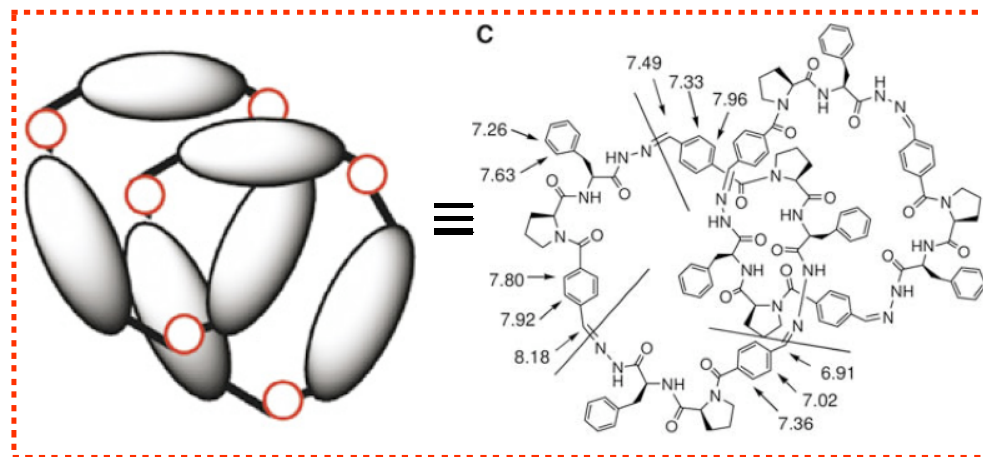
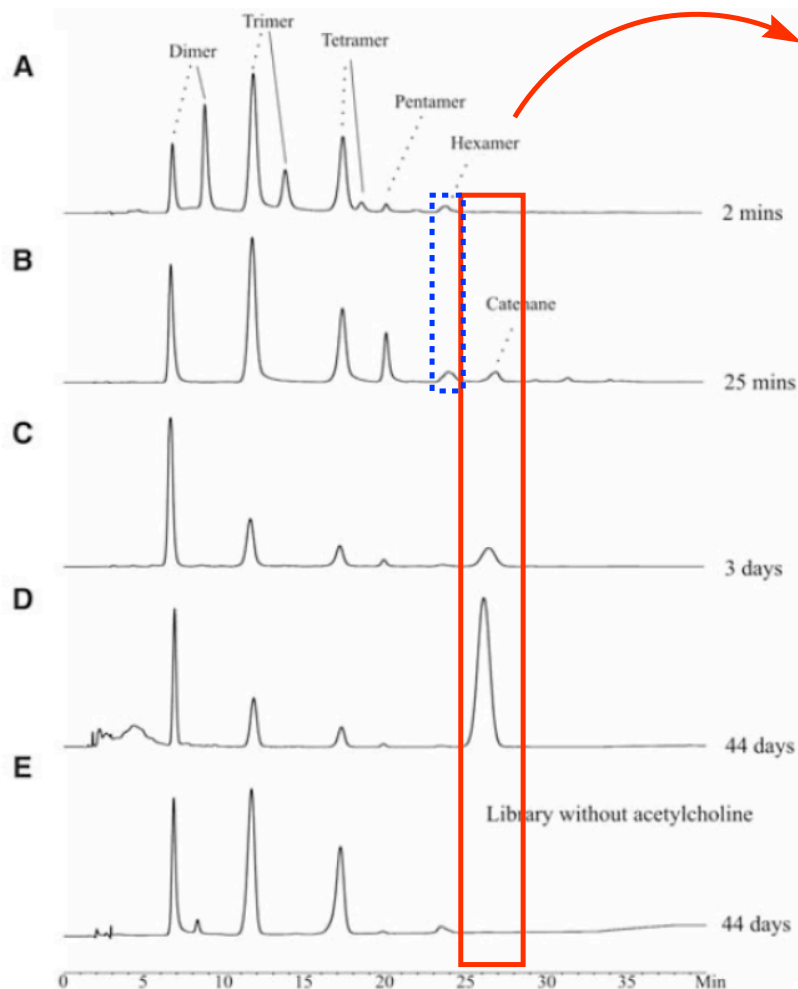
@ With the addition of ACh, accessing unpredictable structures ???

J.K.M. Sanders, S. Otto et al. *Science.*, **2005**, *308*, 667

DCC for Amplification of Molecules (2)

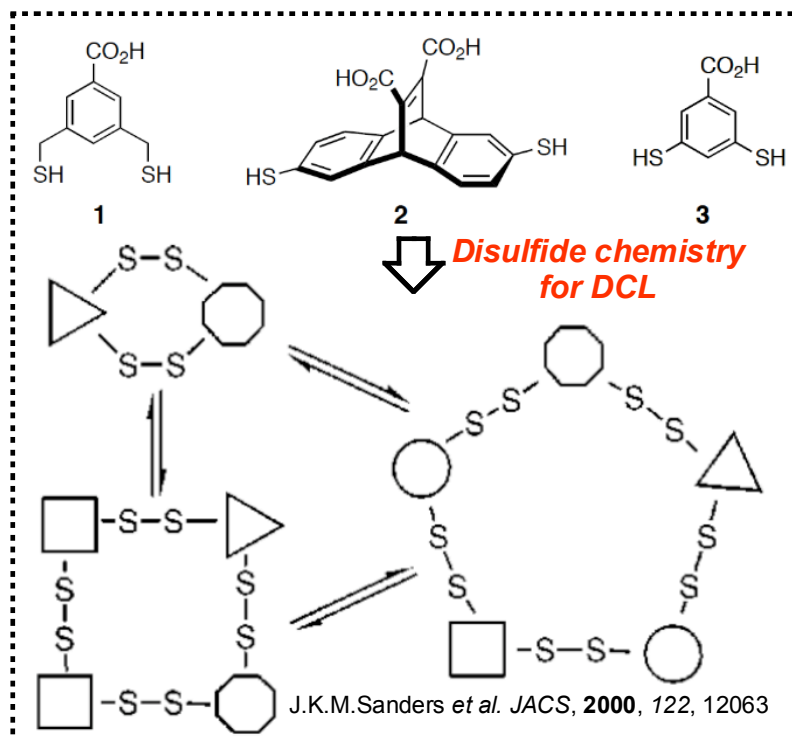
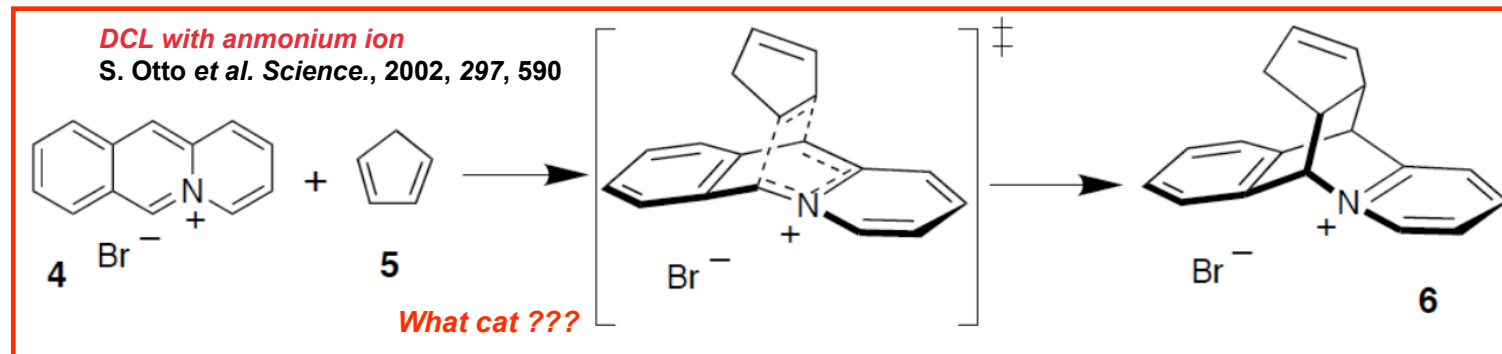
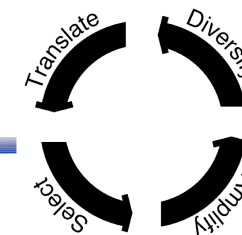


HPLC traces



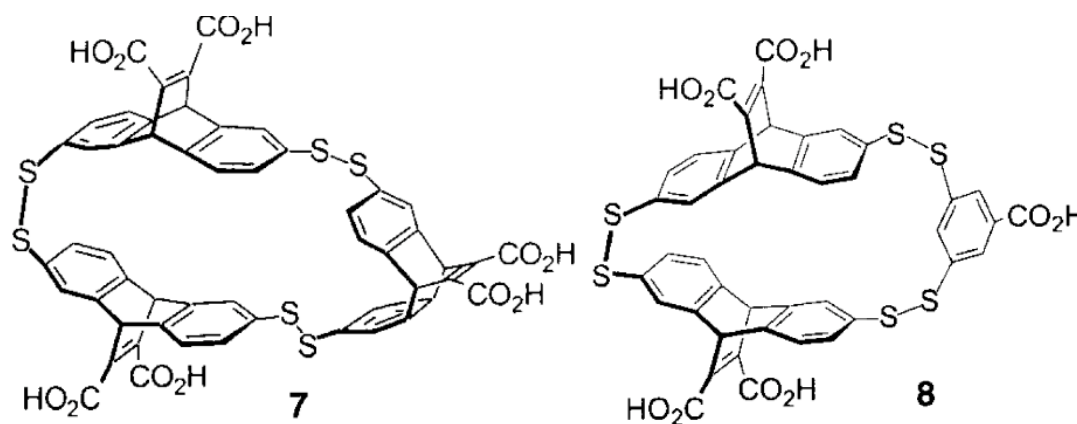
- > first **kinetically favoured product** was observed. (blue)
- > afterwards **thermodynamically favoured product** was observed. (red)
→ [2]catenane
- > [2]catenane was **amplified** at the expense of all the other materials.
- > determined also by NMR analysis and MS analysis
- > **only one diastereoisomer** in 67 % yield
- > **complex functions** from small and simple SMs

DCC for Catalyst Discovery (1)



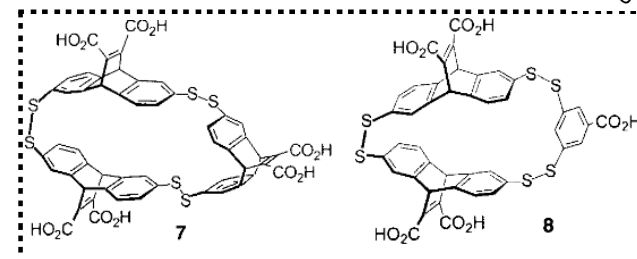
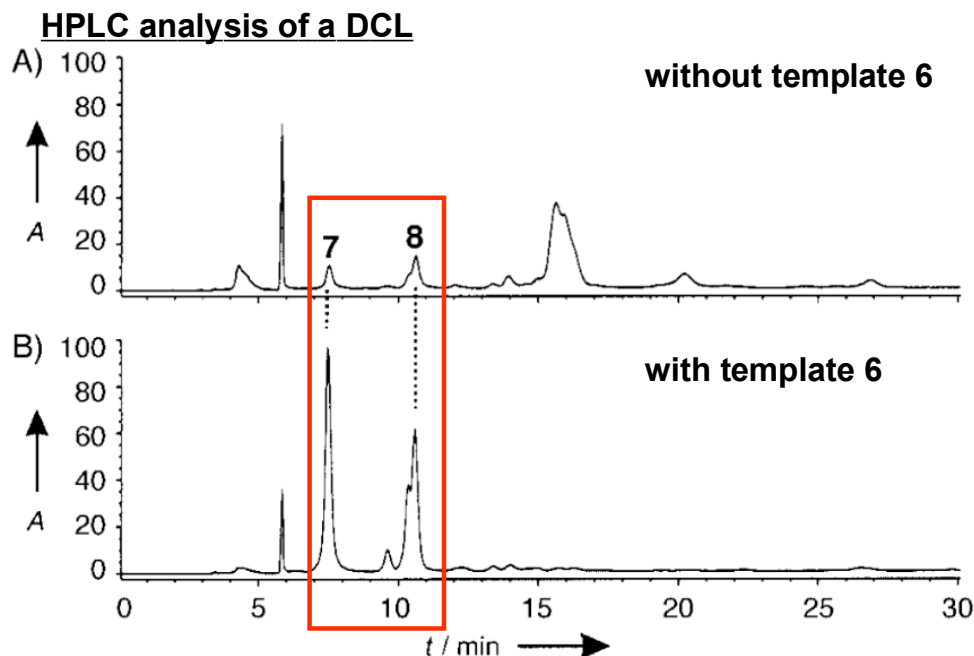
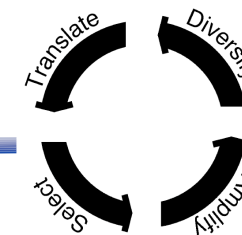
> **disulfide chemistry** for DCLs of macrocycles

> with high affinity for cationic hydrophobic molecules
 → the **Diels-Alder reaction**



S. Otto et al. *ACIE*, 2003, 42, 1270

DCC for Catalyst Discovery (2)



> 1, 2 and 3 mixed in water, under O₂

> **With template**, macrocycles 7 and 8 were obtained in the mixture.

ITC analysis of the thermodynamic datas

		7	8	
SM diene	4	K_1 [M ⁻¹]	1.3×10^5	6.4×10^5
		ΔG° [kJ mol ⁻¹]	-29.1	-33.1
		ΔH° [kJ mol ⁻¹]	-23.7	-40.6
		$T\Delta S^\circ$ [kJ mol ⁻¹]	5.4	-7.5
TM	6	K_1 [M ⁻¹]	2.4×10^5	3.9×10^5
		ΔG° [kJ mol ⁻¹]	-30.7	-31.9
		ΔH° [kJ mol ⁻¹]	-25.8	-38.5
		$T\Delta S^\circ$ [kJ mol ⁻¹]	4.9	-6.6

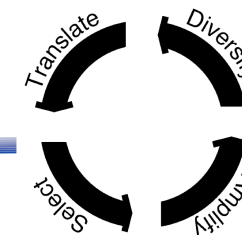
> 8 binds more strongly with 4

> 7 binds more strongly with 6

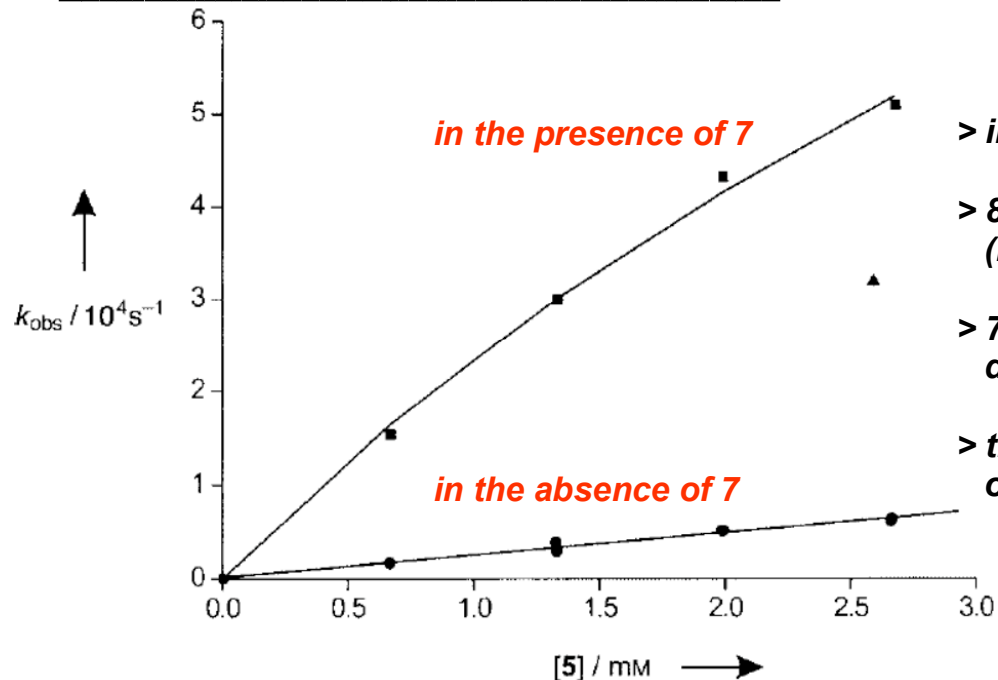
→ **incorporation of 5 into 7•4 complex**

→ 7 would be **catalytically active**

DCC for Catalyst Discovery (3)



Observed rate constant for the Diels-Alder reaction

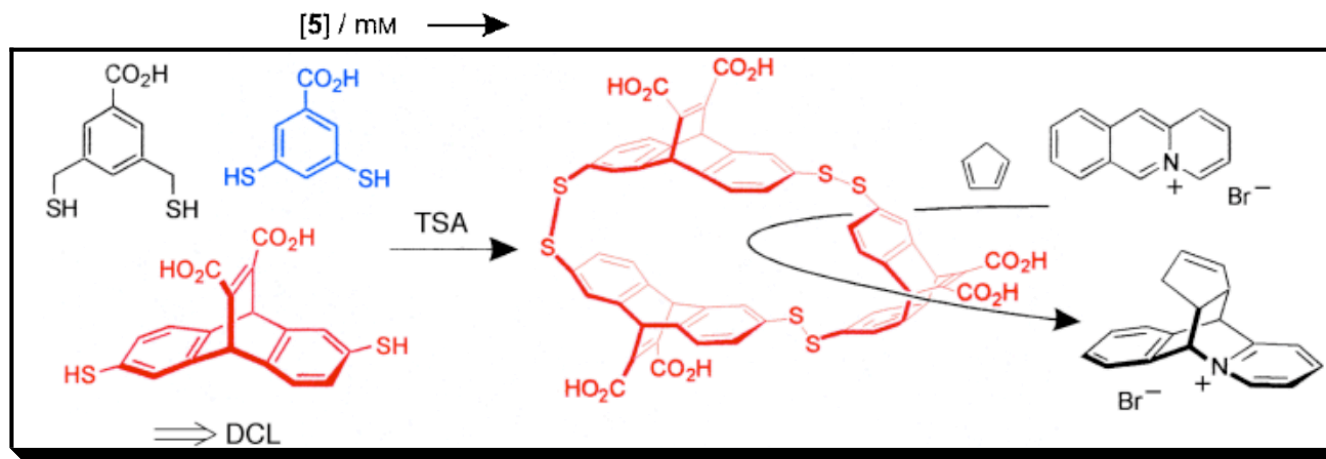


> indeed **7** induced a modest acceleration

> **8** turned out to be catalytically inactive (not shown)

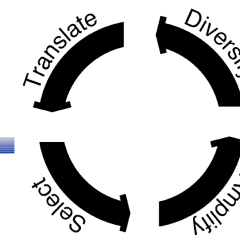
> **7** seemed to **leave enough space** to bind dienophile **5**

> the efficiency of **7** was moderate compared with other catalysts

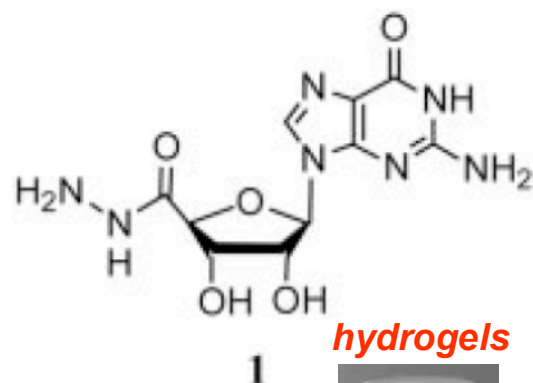


@ In the mixture, the reacton reversibly discovered the efficient catalysts

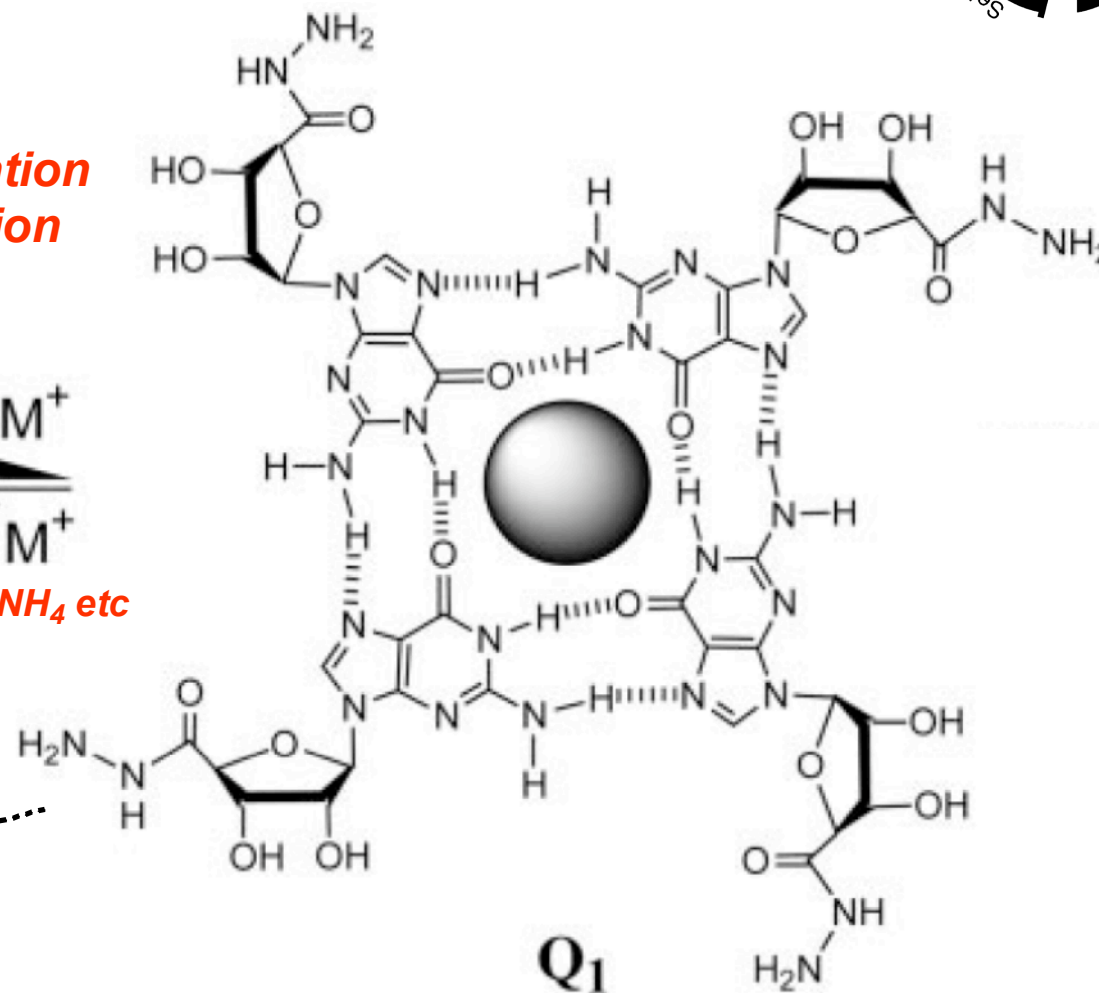
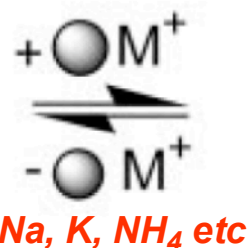
DCC for Functional Materials (1)



*Supramolecular formation
in non-covalent fashion*

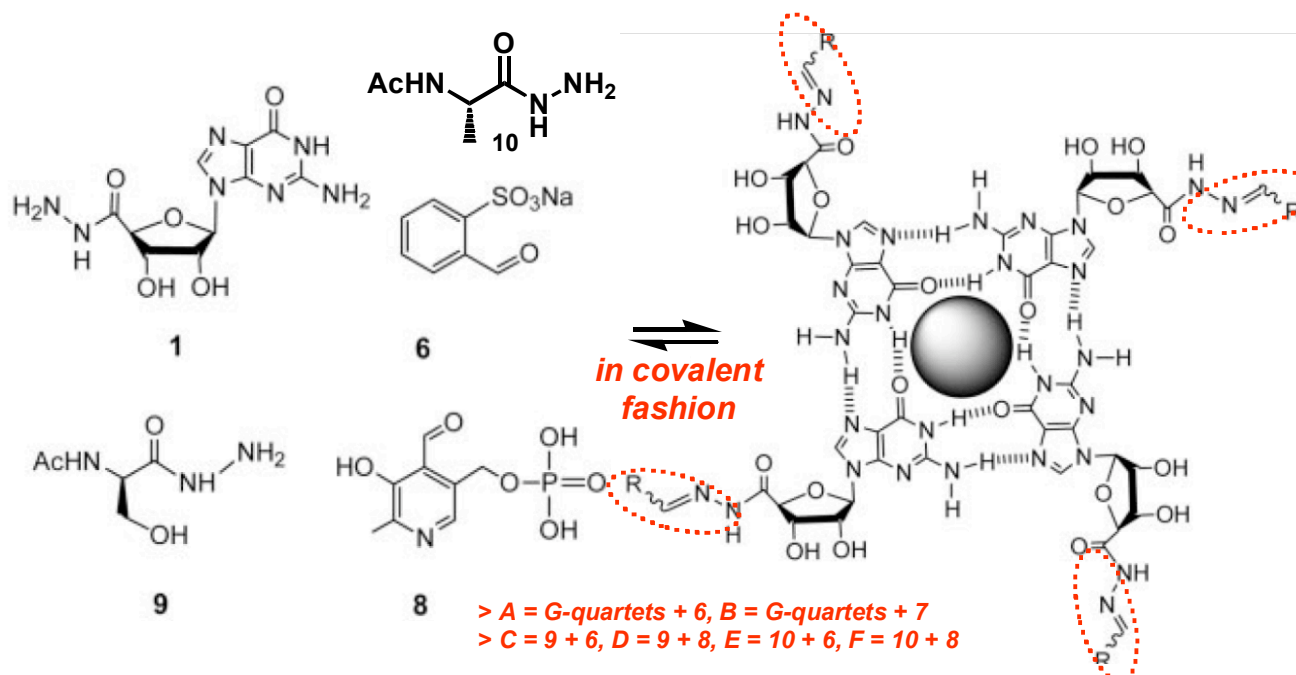
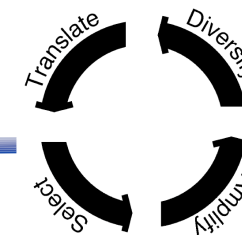


hydrogels



- > undergoing the association into *G-quartets* in the presence of *cations*
- > with *hydrogel formations*

DCC for Functional Materials (2)



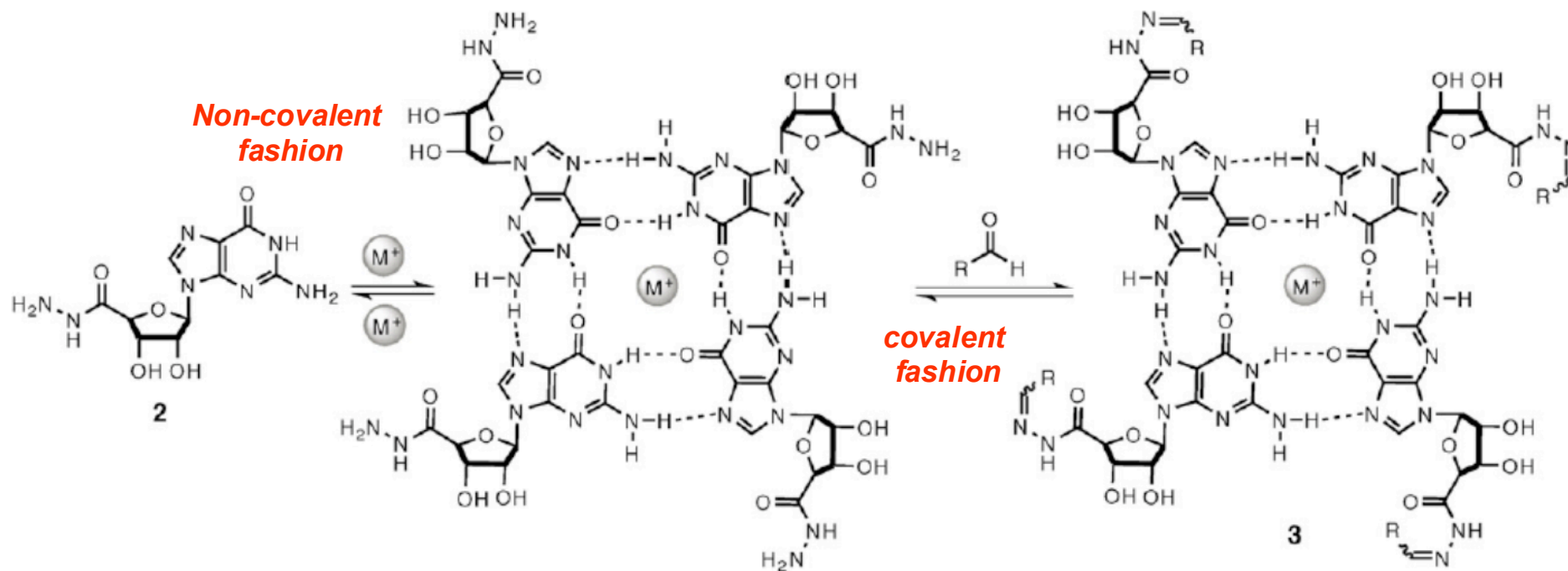
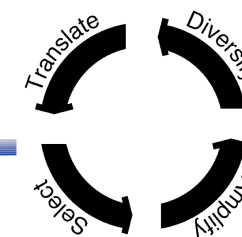
Entry	Hydrazides			Aldehydes		Acylhydrazones at the equilibrium, %					
	1	9	10	6	8	A	B	C	D	E	F
1	1	1	0	1	1	8	39	42	11	—	—
2	1	0	1	1	1	9	37	—	—	40	12
3	0	1	0	1	1	—	—	15	85	—	—
4	0	2	0	1	1	—	—	50	50	—	—
5	1	1	0	0	1	—	87	—	13	—	—
6	1	0	1	0	1	—	96	—	—	—	4
7	1	1	0	1	0	48	—	52	—	—	—
8	1	0	1	1	0	51	—	—	—	49	—
9	0	1	1	1	1	—	—	25	25	25	25

> ¹H NMR determined the equilibrium

> At higher temp (~ 80 °C), the selectivity got worse due to the *melting of hydrogels*

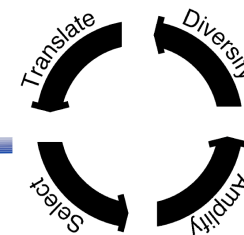
> with the *thermodynamic gelation* event, 1 preferentially scavenged 8.

DCC for Functional Materials (3)



- > the lability of **noncovalent interactions** + **reversible covalent** bonds
→ multistep self-assembly for DCLs
- > with **themoreversible hydrogels**
- > demonstrating DCC can be applied to a **supramolecular** scale
→ discovery of **materials** with desired bulk properties

Features of DCC



Advantages

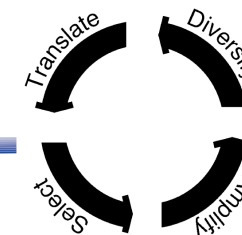
- > ***reversible and thermodynamical*** process in equilibrium
- > desired = amplified, undesired = removed
→ ***'automatically' selection + amplification***
- > applications to small molecules, catalysts, materials

Disadvantages

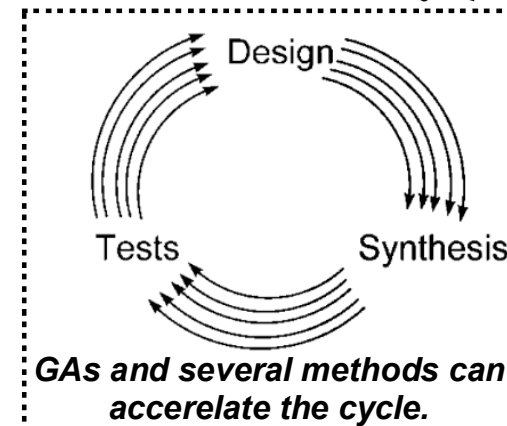
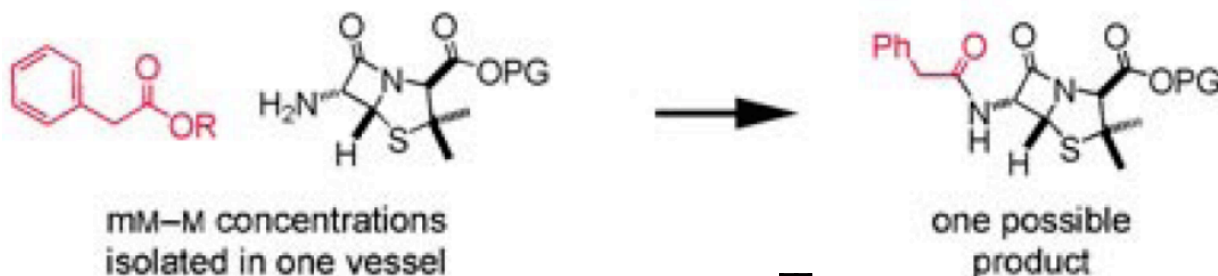
- > ***limited diversity*** of available building blocks for appropriate targets
- > difficulty of deconvoluting the complex mixtures
- > Life can be thought of as '***a kinetic state of matter***'
→ ***systems biology and systems chemistry ???***

3. DNA-templated Synthesis

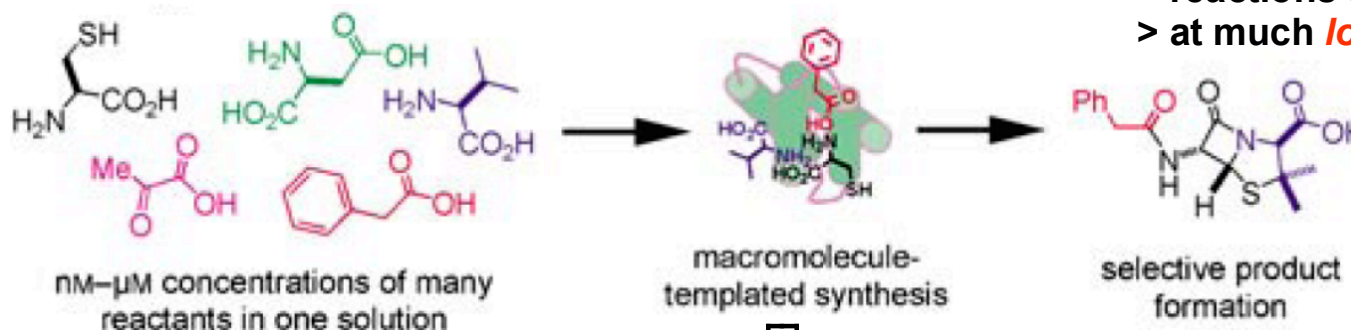
DNA for Evolutional Approaches



Chemists' approach to control reactivity



Nature's approach to control reactivity



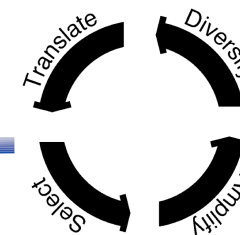
- > in **one-pot** fashion
- > reactions are **randomly** performed
- > at much **lower concentrations**

- > ideally suited to perform the iterative operation (**translation, selection, amplification and diversification**)
- power of **PCR** method

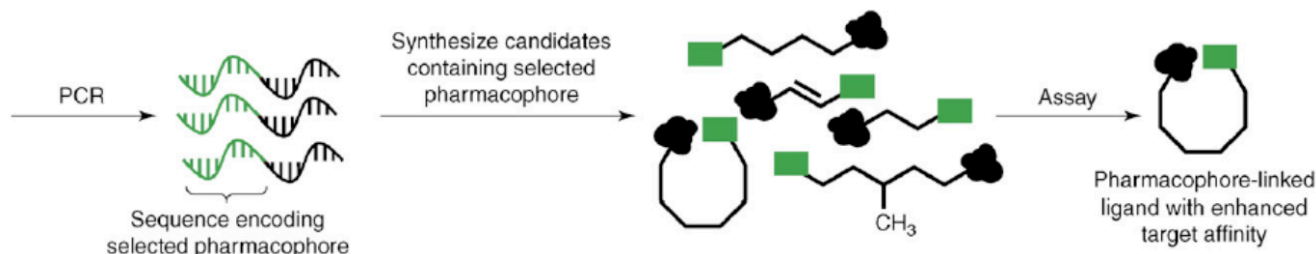
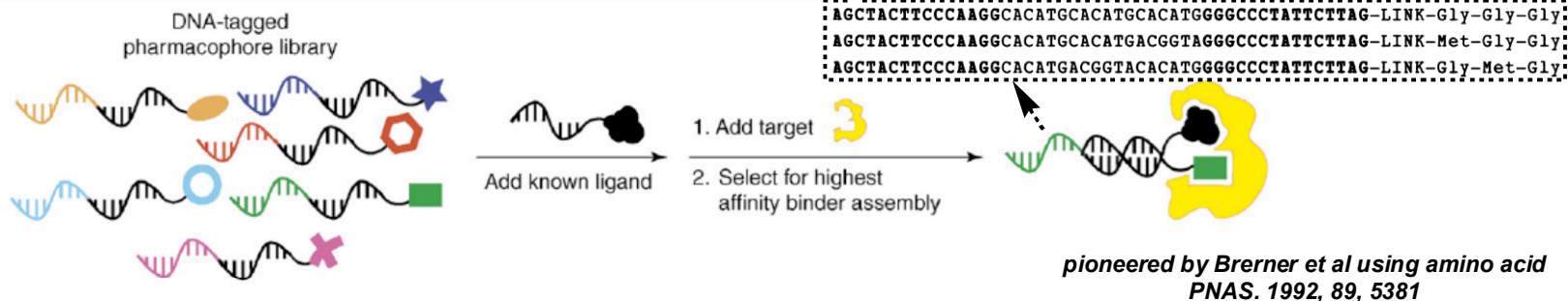
DNA as a tool for evolutional approaches

- > with **high information density**
- > with **chemical stability**
- > with the **ease of manipulation**

Several Methods

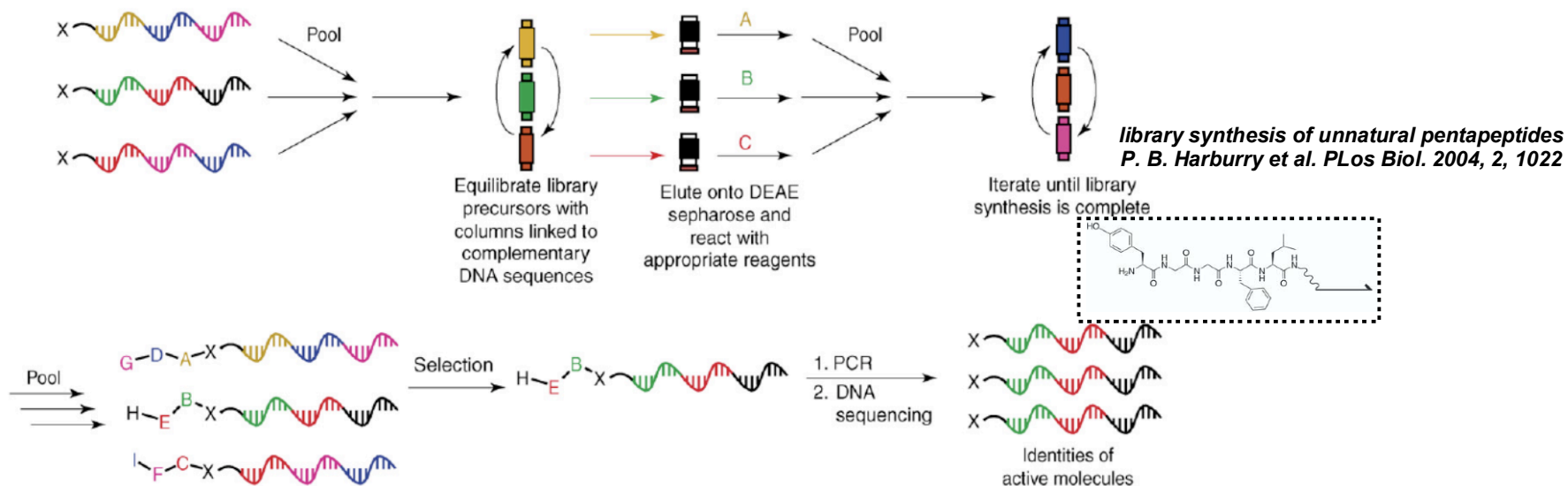


Encoding Chemical Libraries Using DNA > *identification of DNA-linked small molecules*



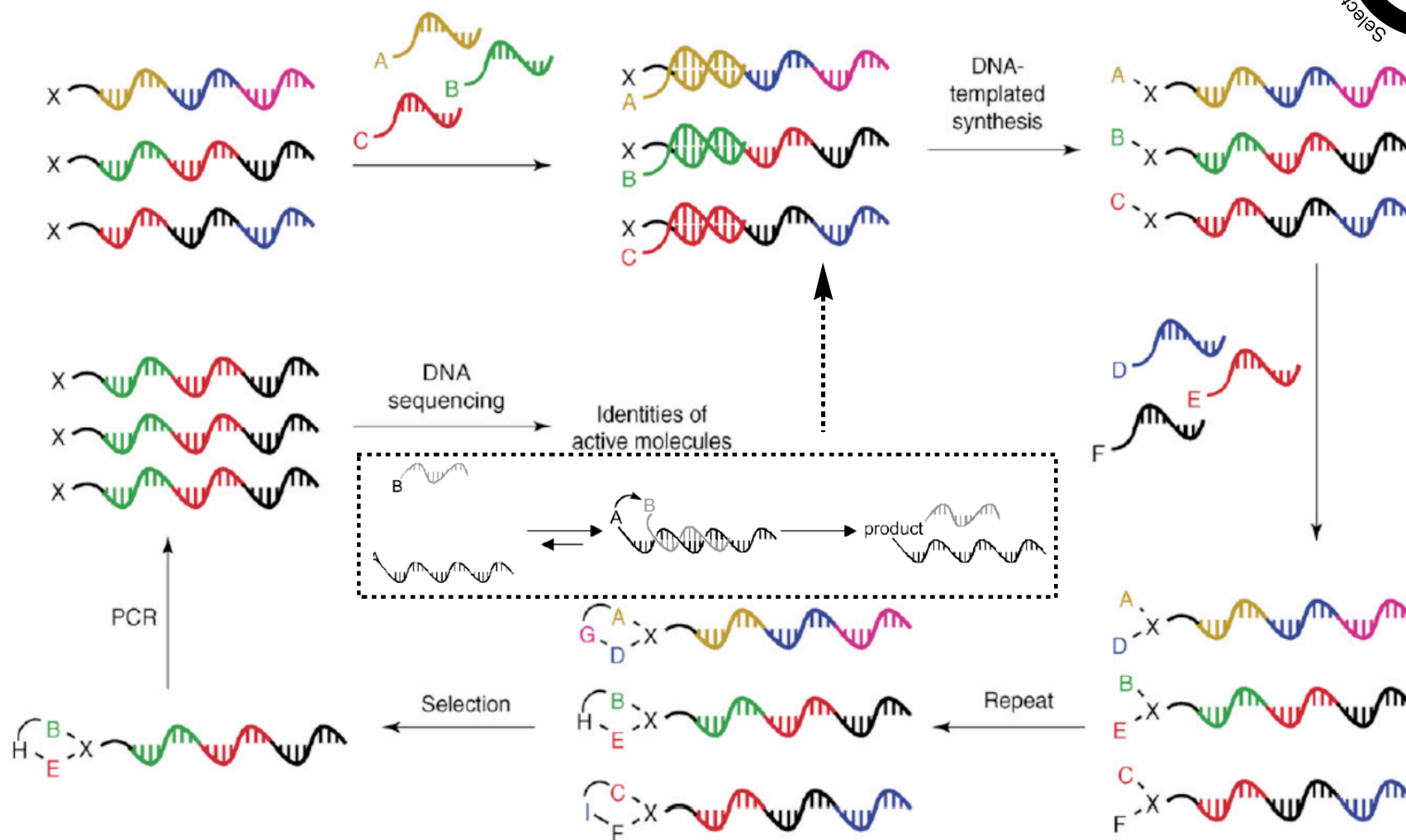
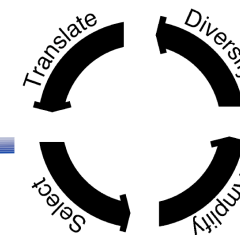
Also see:
 D. Neri et al. *Nat. Biotechnol.*, 2004, 22, 568

DNA Library > *reaction of DNA-linked small molecules individually*



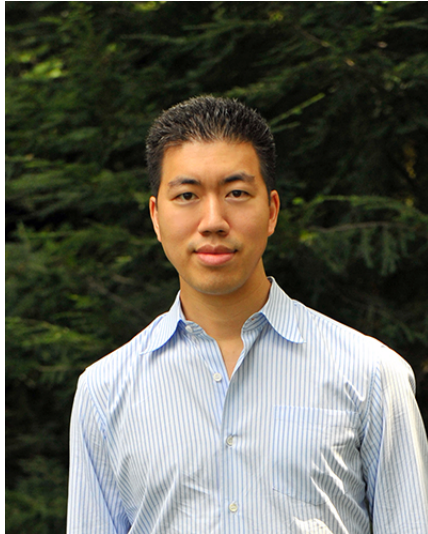
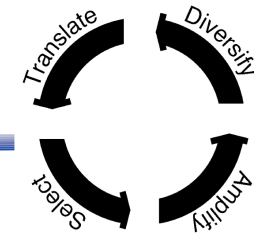
D.R.Liu et al. *Curr. Opin. Chem. Biol.*, 2007, 11, 259

DNA-Templated Synthesis



- > *spatial separation is not required*
- > *DNA annealing controls reactivity*
- *all templates and reagents coexist in a single solution*
- > Using complementary DNA-linked reagents *via Watson-Crick base-pairing*

David R. Liu as a Key Person



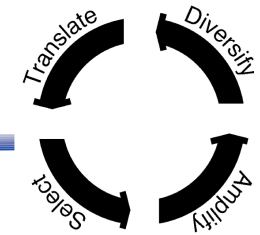
TRAINING

- 1991-1994
(B.A. research) Conducted research in synthetic organic chemistry, biochemistry, and molecular biology on 2,3-oxidosqualene cyclase (a key enzyme in steroid biosynthesis) under the guidance of Professor E. J. Corey at Harvard University.
- 1994-1999
(Ph.D. research) Probed the mechanism of chorismate mutases using site-directed natural and unnatural amino acid mutagenesis; designed new tRNAs for unnatural amino acid mutagenesis; initiated the engineering and evolution of proteins and nucleic acids for the site-specific incorporation of unnatural amino acids into proteins in living cells. Research conducted under the guidance of Professor Peter G. Schultz at the University of California, Berkeley.

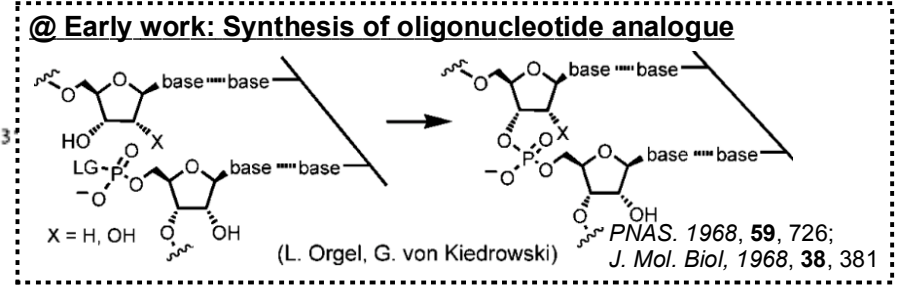
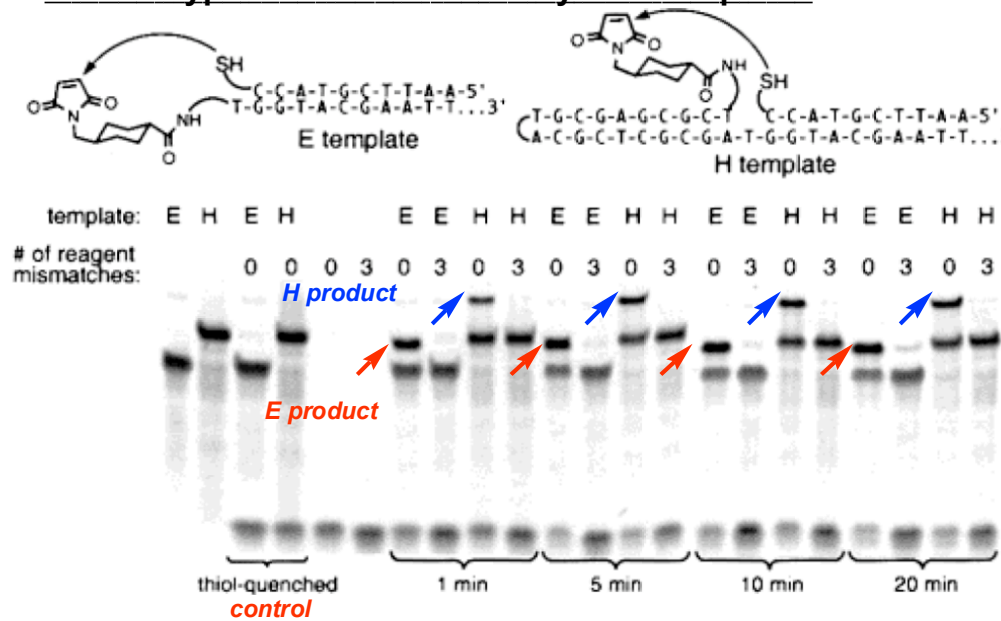
POSITIONS

- 1999-2003 Assistant Professor of Chemistry and Chemical Biology, Harvard University
- 2003-2004 John L. Loeb Associate Professor of the Natural Sciences and Associate Professor of Chemistry and Chemical Biology, Harvard University
- 2007-2010 Harvard College Professor, Harvard University
- 2005-present Senior Associate Member of the Broad Institute of Harvard and MIT (2010-present); Associate Member of the Broad Institute of Harvard and MIT (2005-2010)
- 2005-present Professor of Chemistry and Chemical Biology, Harvard University and Investigator, Howard Hughes Medical Institute

Pioneering Work (1)

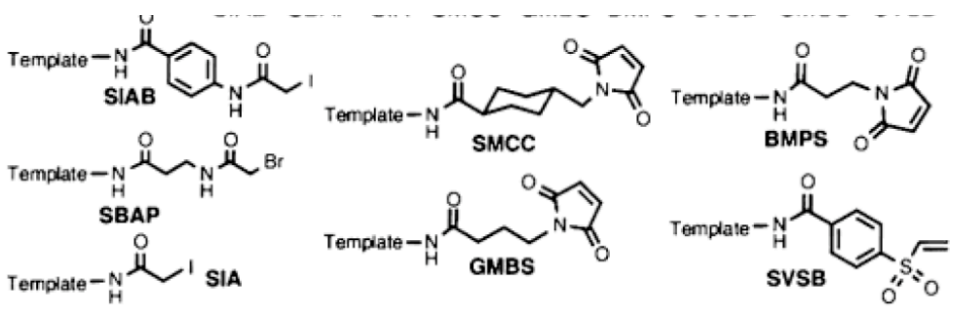
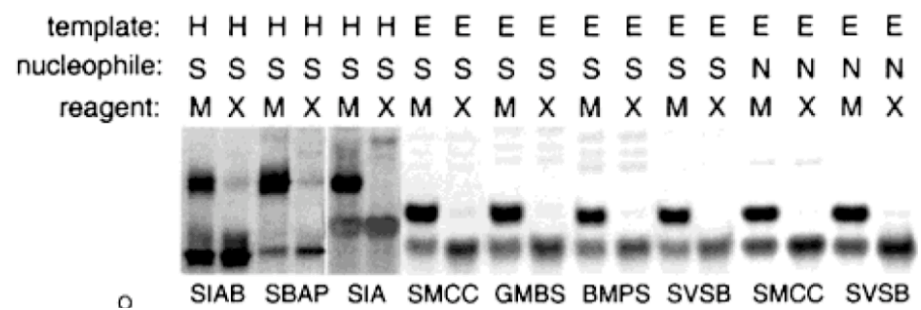


Michael-type reactions directed by DNA templates



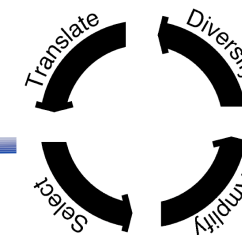
- > H and E DNA architectures exhibited **similar reactivities**.
- > no products were observed by using **mismatched DNAs**.
- > DNA-templated reactions proceeded even the products **differed from the natural backbones**.

Various reactions directed by DNA templates

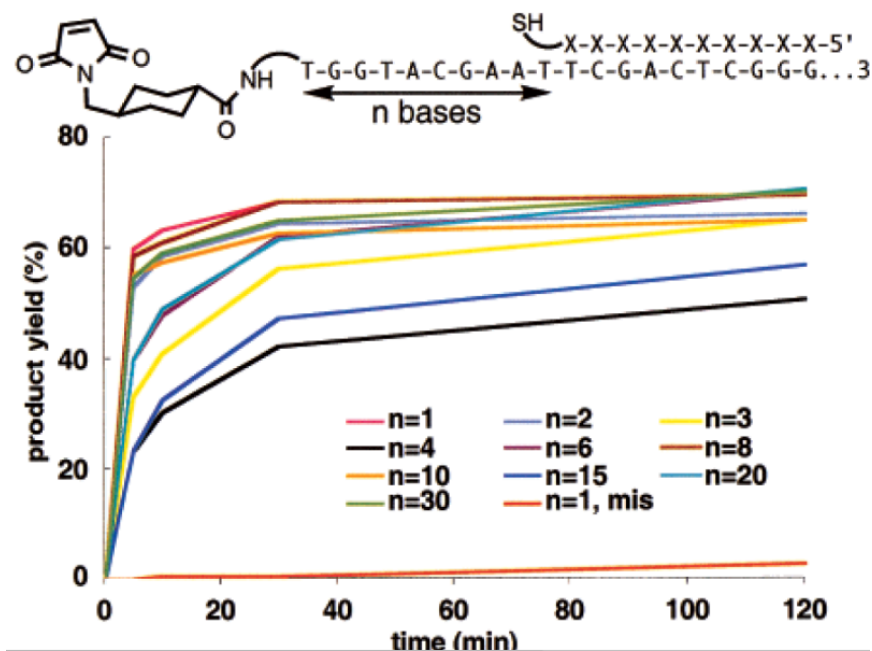


> **Other reaction systems** could be applied for (SN_2 substitutions, additions to $\alpha\beta$ -unsaturated carbonils etc).

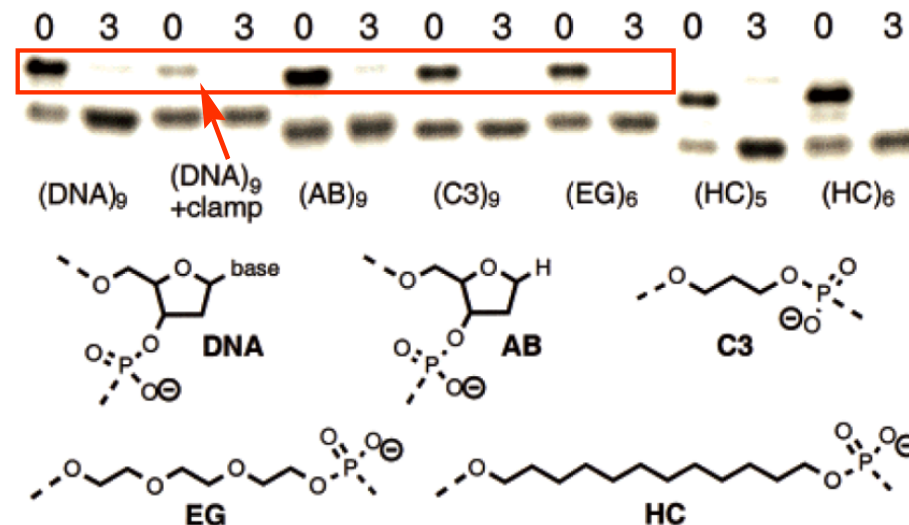
Pioneering Work (2)



Distant-independent (DTS) DNA-templated synthesis



Evaluation of possible contribution

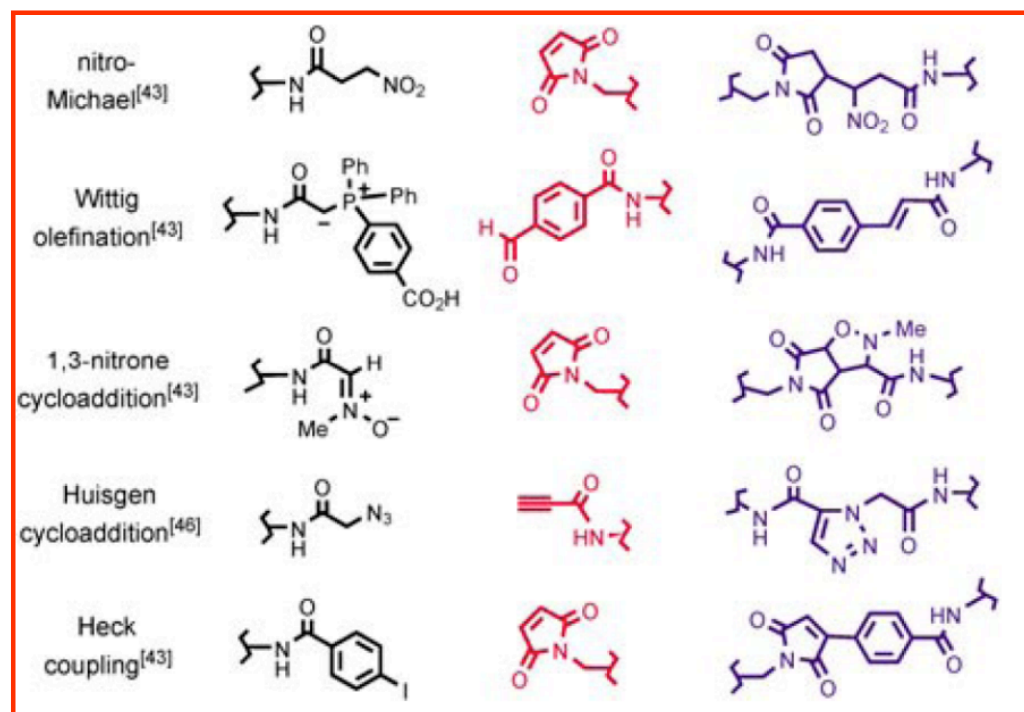
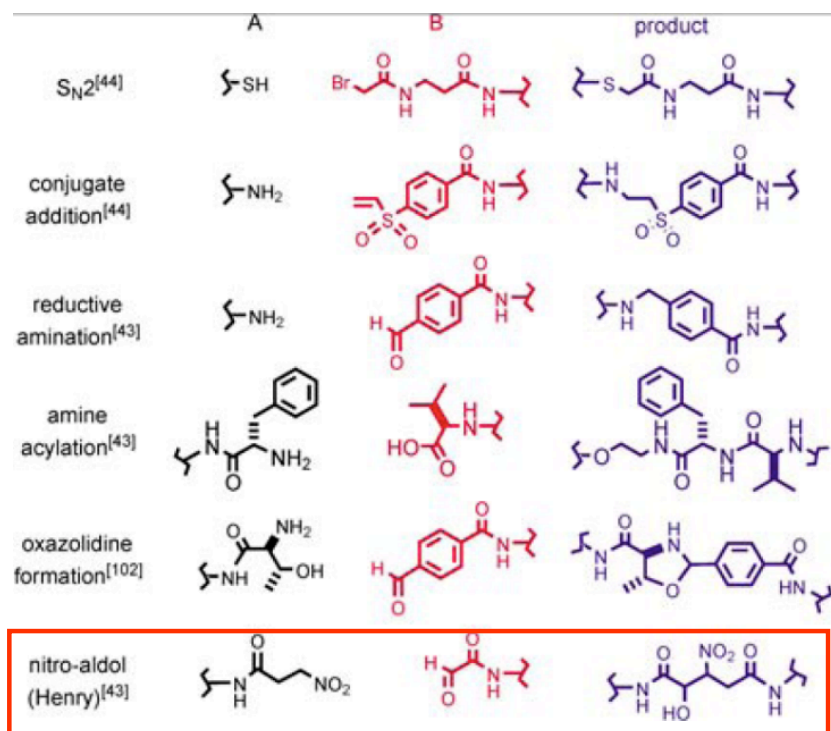
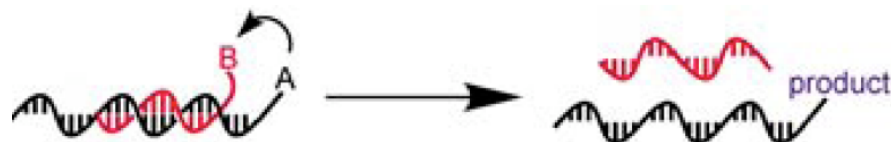
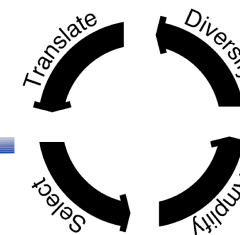


- > demonstrating remarkable **distance independence**.
- > Even in **30 bases distances**, efficiently reacted. (through transition state **200 membered ring !!!**)
- > **sharply contrast** with traditional organic synthesis

- > possible contributions were evaluated. (**conformational preferences, interbase interactions, backbone hydrophilicity etc**)
- > the backbone structural elements were **not responsible**.
- > **complementary DNAs to intervening region** significantly reduced the products. → **the flexibility of the region** is likely to be crucial.

@ The fact that **DNA annealing became the RDS** would explain the distant independence. (decreasing the conc. resulted in a marked reduce in the reaction rate)

Expansion to the Reaction Generality

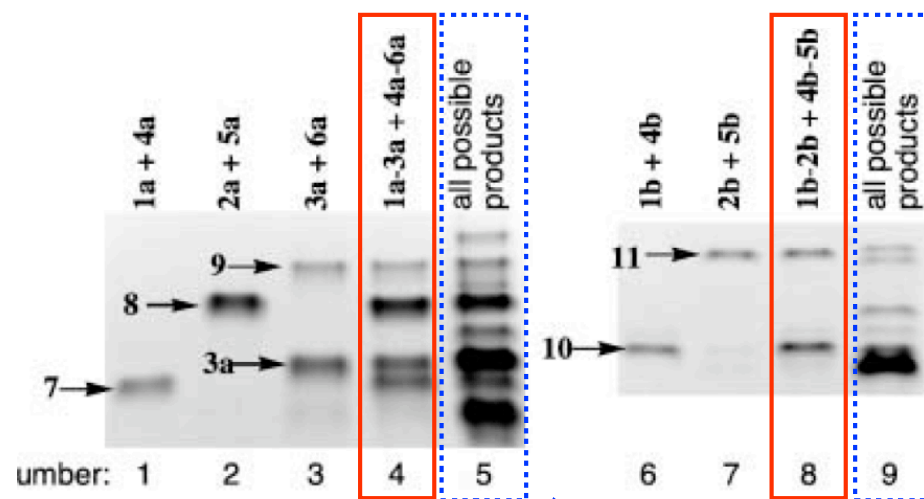
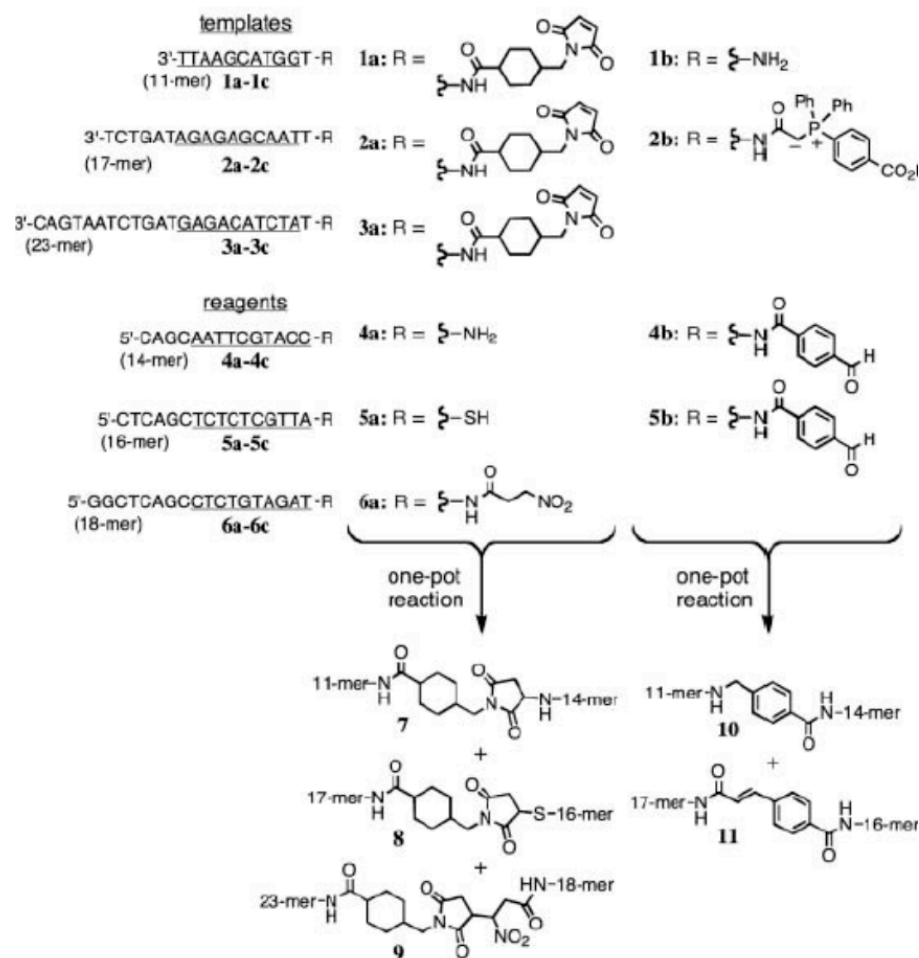
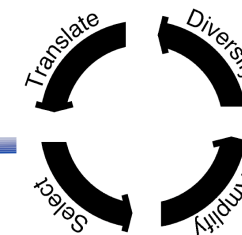


> the first useful **C-C bond forming reactions** (nitro-aldol, Wittig, Heck coupling etc)

D.R.Liu *et al.* *ACIE.*, 2002, 41, 1796

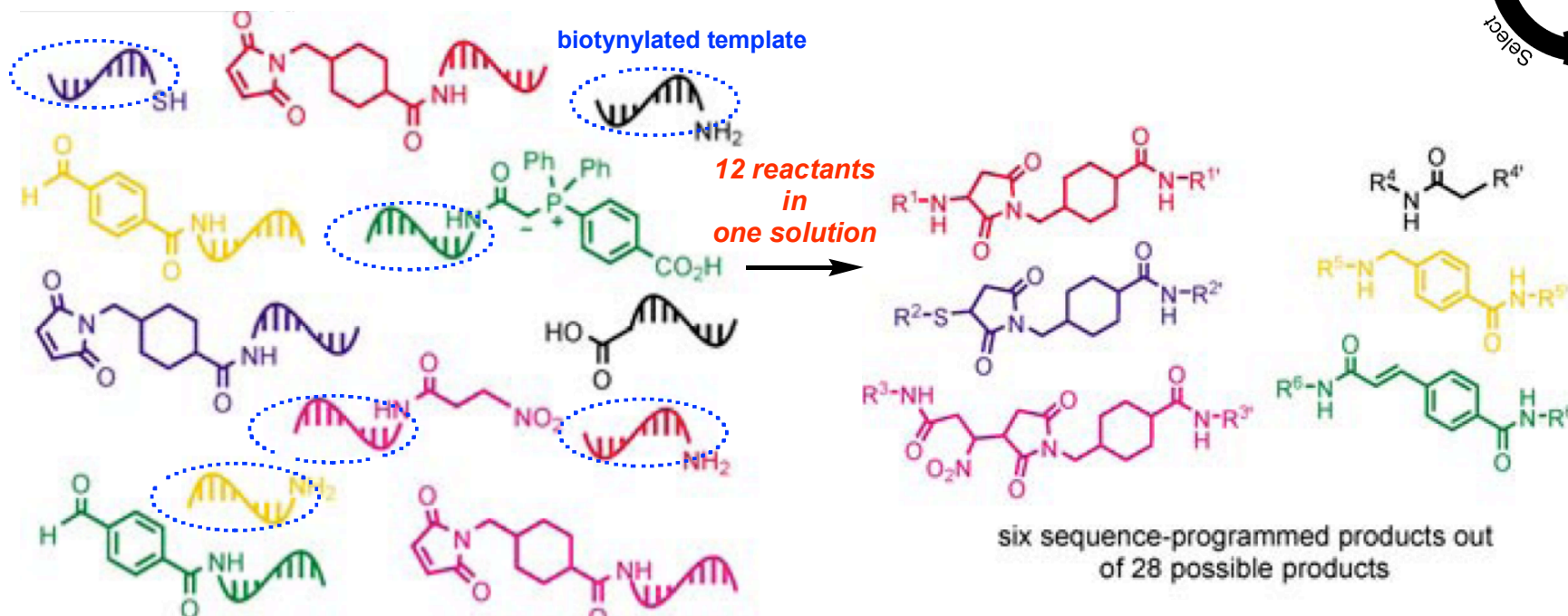
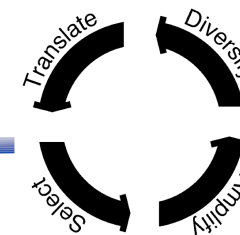
D.R.Liu *et al.* *ACIE.*, 2003, 42, 1370

New Modes of Controlling Reactivity (1)

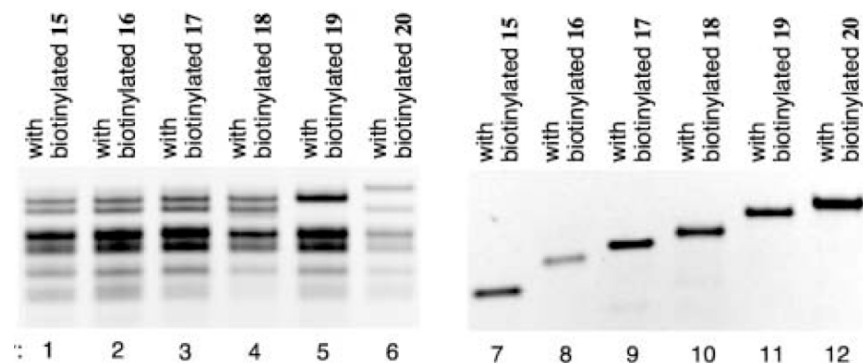


- > different type of reactions were conducted **in one solution**.
- > **the different length** of DNAs enabled easier detection.
- > **selective 3 reactions** took place to afford 7, 8, 9 exclusively (from 9 possible reactions)
- > other possible products were obtained in **< 5 %**.

New Modes of Controlling Reactivity (2)



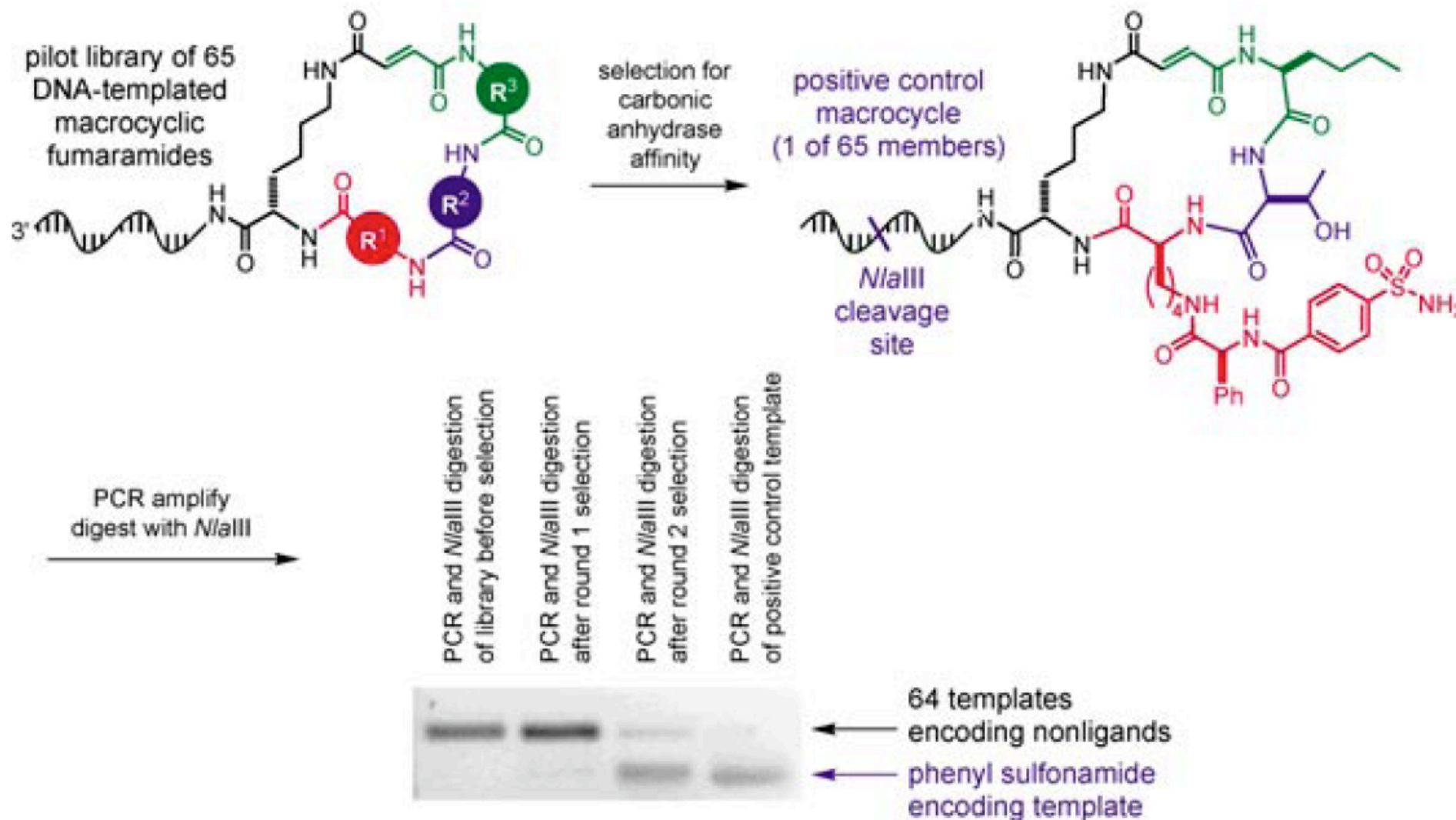
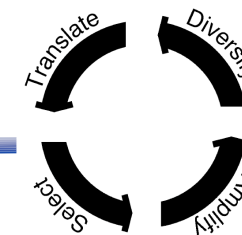
before purification with beads **after purification with beads**



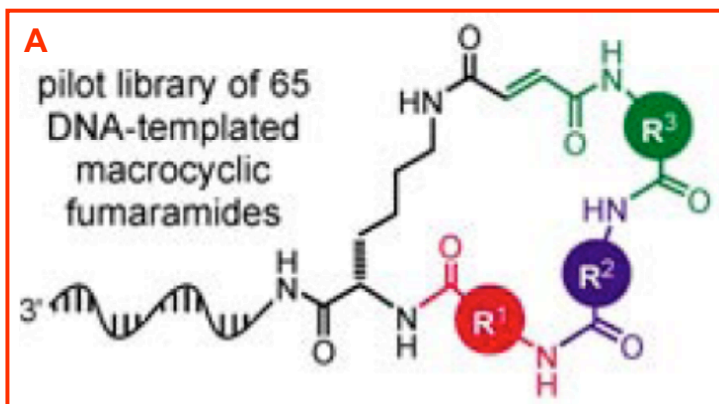
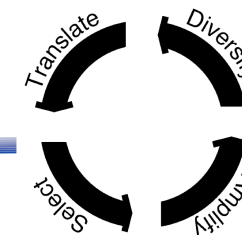
- > the reactions were performed containing 6 templates and 6 reactants **in one solution**.
- > 6 templates with **biotinylation**
- > biotinylated templates **could be purified** with magnetic beads.
- > **only the single product** was obtained programmed by DNA sequences.

@ DTS exhibited the ability to direct different types of reactions in one solution, incompatible under traditional reaction conditions.

Applications of DTS ~ Selection of a Library (1)

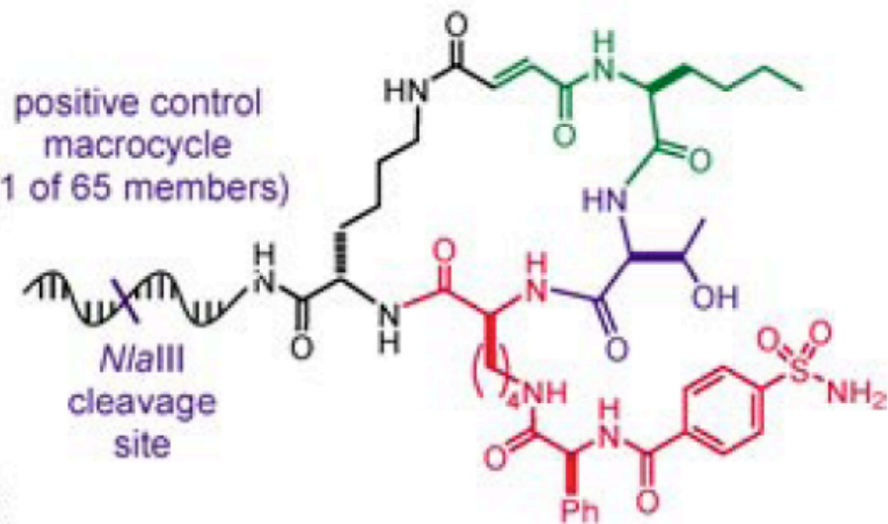


Applications of DTS ~ Selection of a Library (1)



selection for carbonic anhydrase affinity

positive control macrocycle (1 of 65 members)



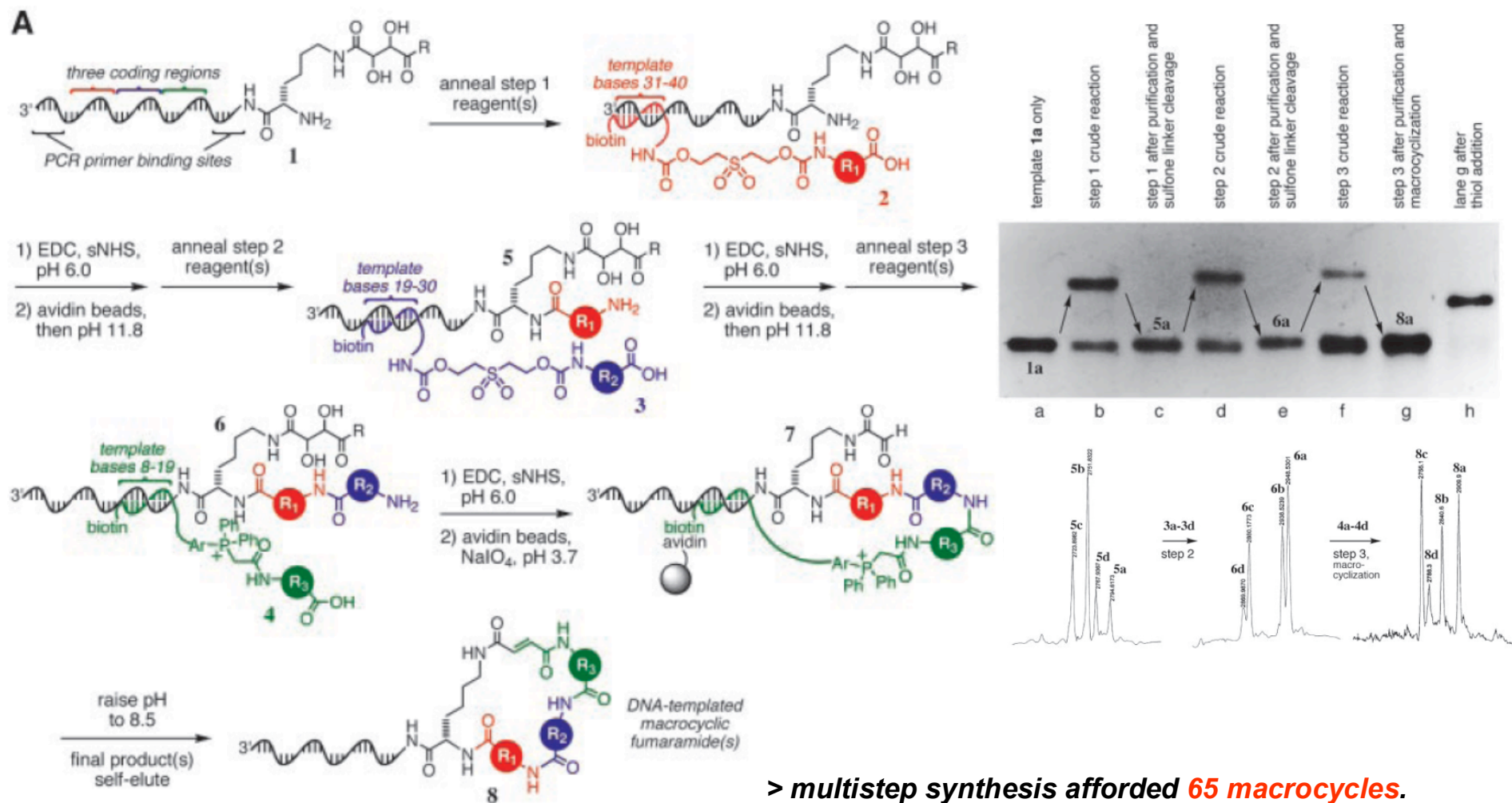
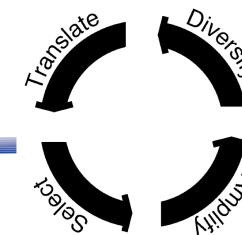
PCR amplify
digest with *NalIII*

PCR and *NalIII* digestion of library before selection
PCR and *NalIII* digestion after round 1 selection
PCR and *NalIII* digestion after round 2 selection
PCR and *NalIII* digestion of positive control template



← 64 templates encoding nonligands
← phenyl sulfonamide encoding template

Applications of DTS ~ Selection of a Library (2)



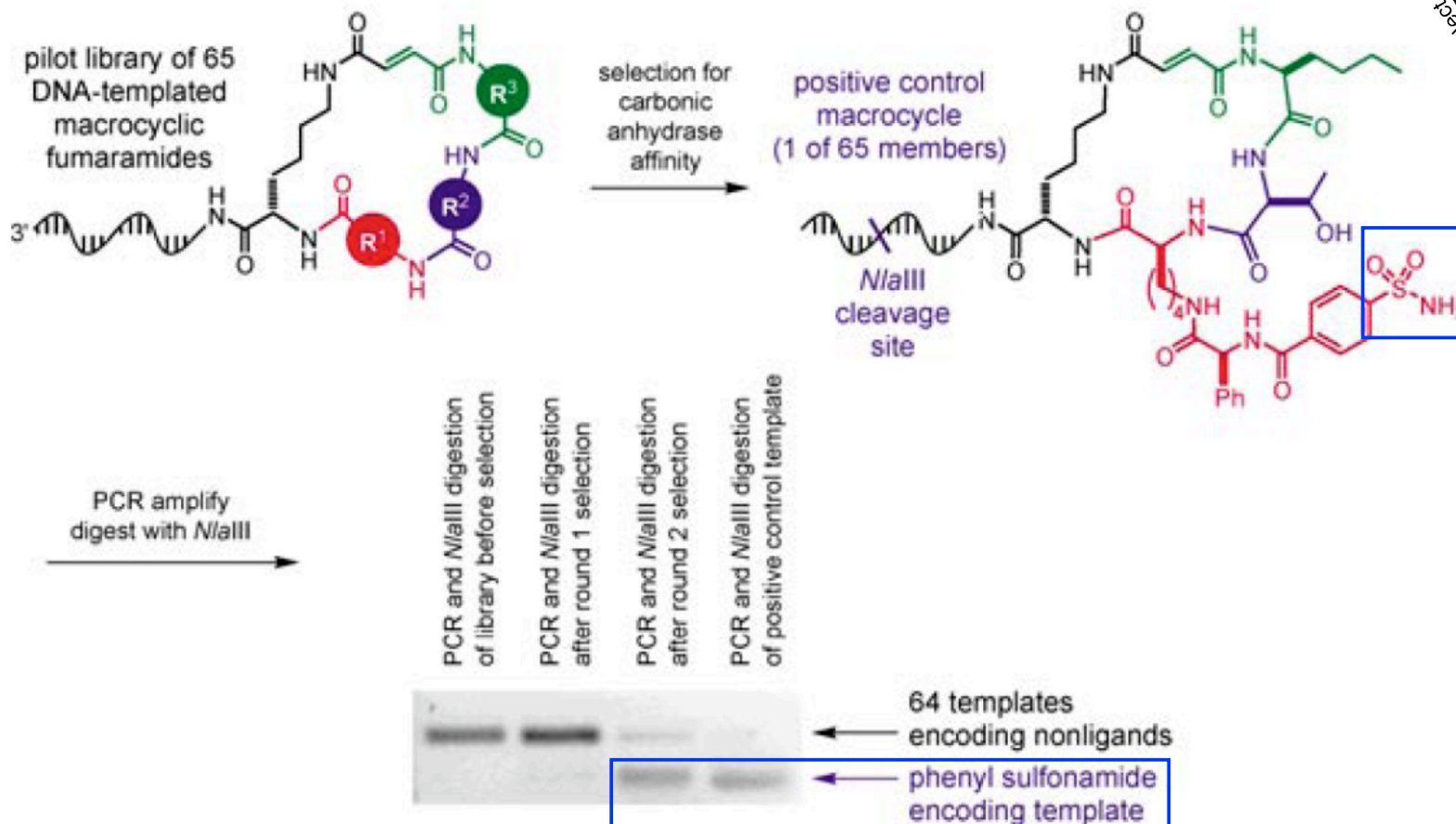
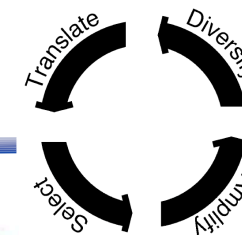
> multistep synthesis afforded **65 macrocycles**.

> each step was monitored by **PAGE** and **TOF-MS**

D.R.Liu *et al.* *JACS.*, 2002, 124, 10304

D.R.Liu *et al.* *Science.*, 2004, 305, 1601

Applications of DTS ~ Selection of a Library (3)

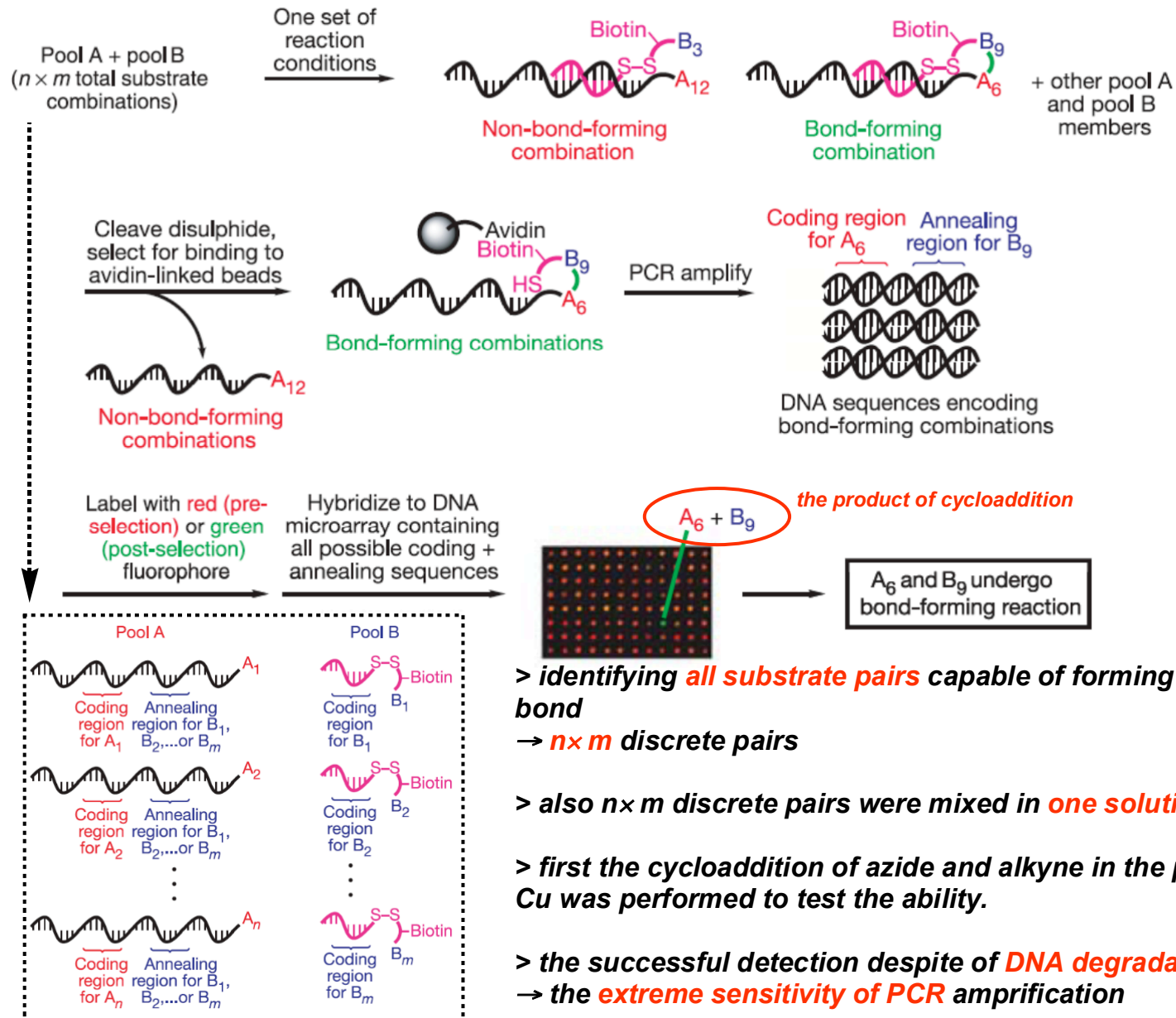
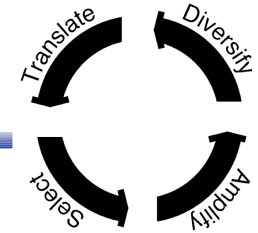


- > a pilot library of 65 macrocycles was utilized for **selective protein binding affinity** (carbonic anhydrase).
- > 2 selection rounds **enriched the single molecule** containing sulfonamide group.

@ DTS coupled with *in vitro* selection enables the translation, selection, amplification of DNA encoding **synthetic small molecules not biological molecules**.

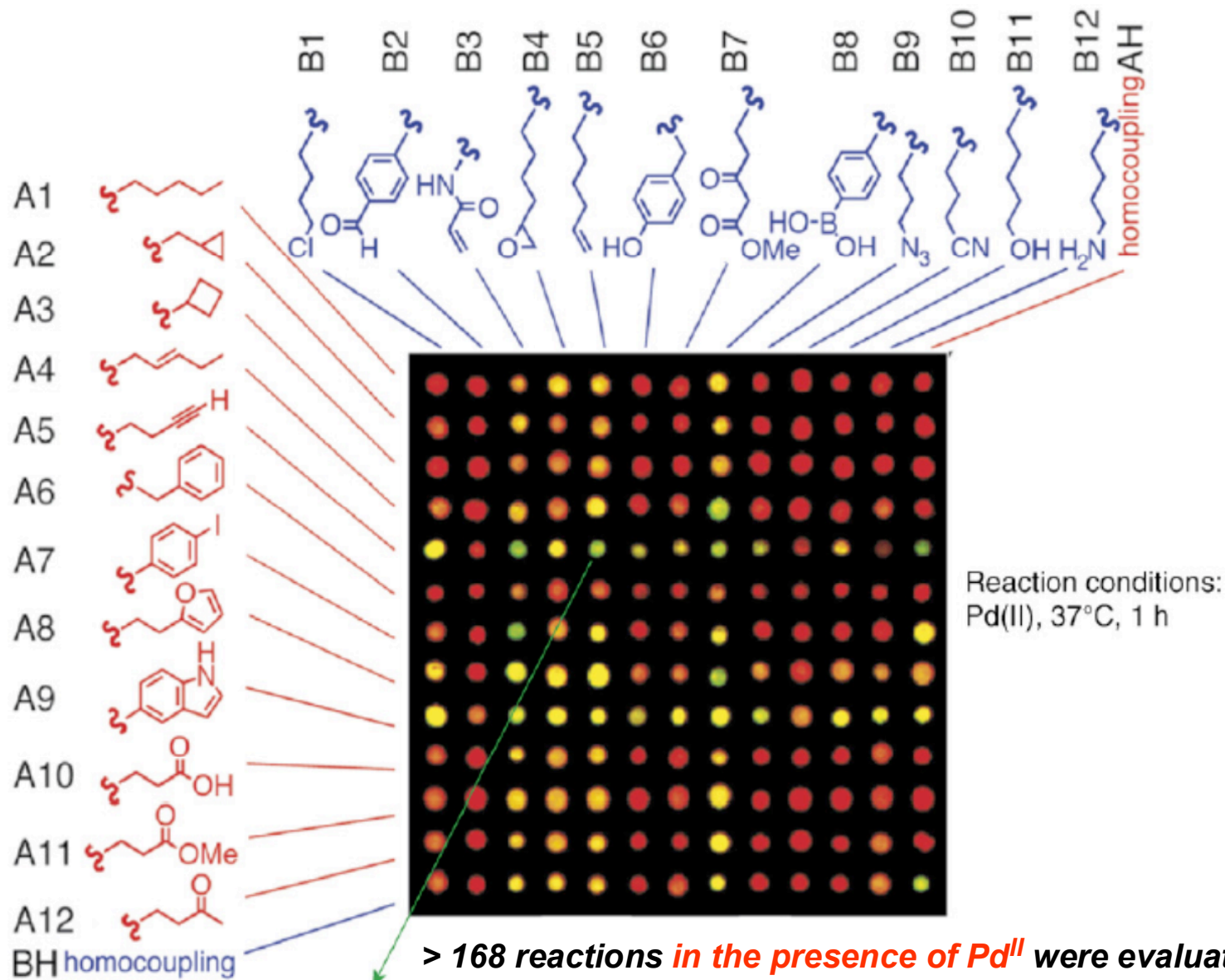
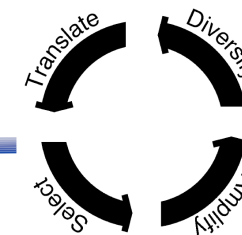
→ **the combined ability of synthetic chemistry and PCR method !!!**

Applications of DTS ~ Reaction Discovery (1)

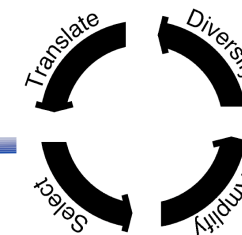


- > identifying **all substrate pairs** capable of forming a covalent bond
- $n \times m$ discrete pairs
- > also $n \times m$ discrete pairs were mixed in **one solution**
- > first the cycloaddition of azide and alkyne in the presence of Cu was performed to test the ability.
- > the successful detection despite of **DNA degradation by Cu**
- the **extreme sensitivity of PCR amplification**

Applications of DTS ~ Reaction Discovery (2)

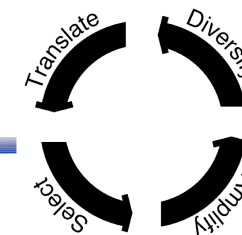


Applications of DTS ~ Reaction Discovery (3)

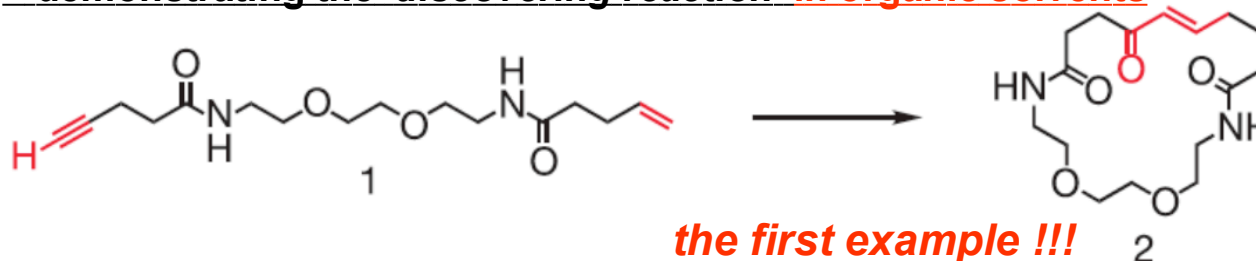


> characterization of array positives		Green/red fluorescence ratios		DNA-templated yields (%)		Product consistent with observed mass
Substrates		37°C	25°C	37°C	25°C	
		2.7	3.7	35	31	 R-CH ₂ -CH ₂ -C(=O)-CH=CH-CH ₂ -R ???
		3.5	3.1	28	20	
		1.6	1.9	36	34	
		2.6	2.7	45	42	
		3.0	2.8	57	39	
		1.8	<1.2	30	10	
		1.8	<1.2	19	<10	
		3.6	<1.2	39	14	 <i>Heck</i>

Applications of DTS ~ Reaction Discovery (4)



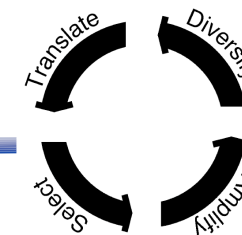
> demonstrating the '*discovering reaction*' in organic solvents



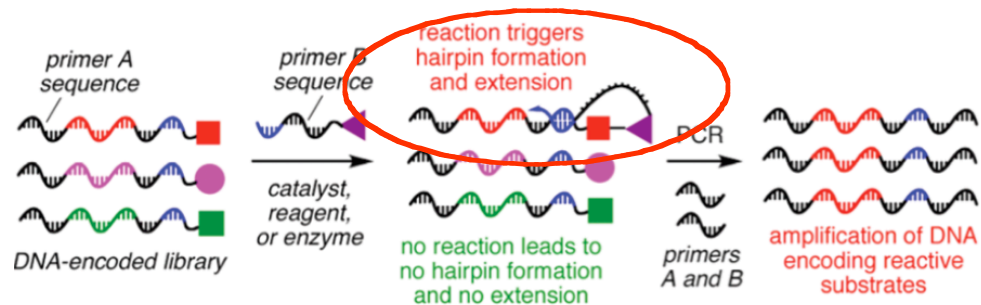
Entry	Metal(s)	Solvent	Conditions	Isolated yield
a	1 equiv. Na ₂ PdCl ₄	1 M NaCl in H ₂ O	25 °C, 15 h	86%
b	5 mol% Na ₂ PdCl ₄ 1 equiv. CuCl ₂	100 mM NaCl in H ₂ O	25 °C, 2 h	90%
c	5 mol% Na ₂ PdCl ₄ 1 equiv. CuCl ₂	9:1 THF: H ₂ O	25 °C, 4 h	91%
d	15 mol% Na ₂ PdCl ₄ 1 atm O ₂	9:1 THF: H ₂ O	25 °C, 14 h	73%
e	1 equiv. CuCl ₂	100 mM NaCl in H ₂ O	25 °C, 4 h	0%
f	1 equiv. CuCl	100 mM NaCl in H ₂ O	25 °C, 4 h	0%

@ the value of searching a large number of substrate combinations for unexpected reactions

Applications of DTS ~ Reaction Discovery (5)

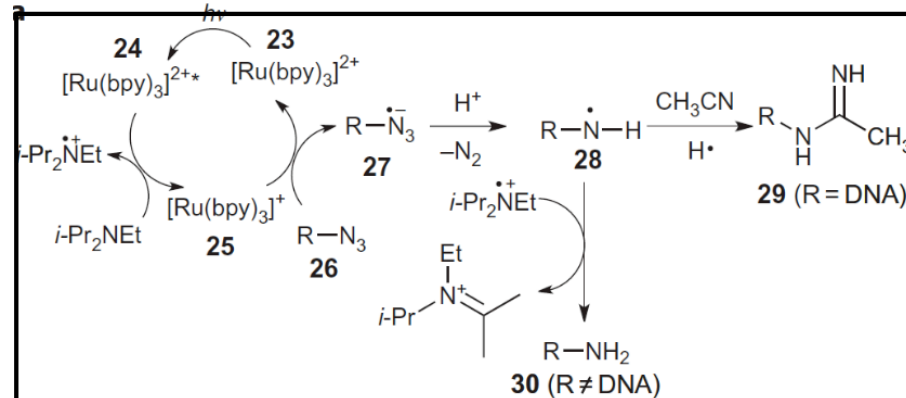


RDPCR (Reactivity-Dependent PCR)

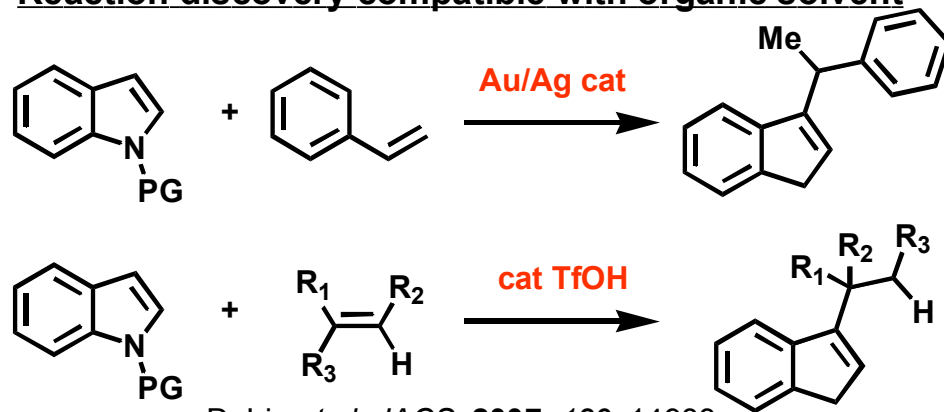


D, Liu *et al.* *JACS*. 2009, 131, 9189

Reaction discovery of mild azide reduction



Reaction discovery compatible with organic solvent



D, Liu *et al.* *JACS*. 2007, 129, 14933

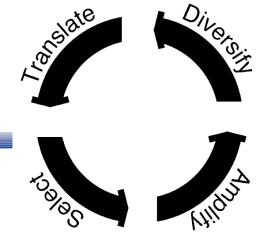
> in organic or aqueous solvent, open to air, at r.t

> compatible with various functional groups (alcohols, phenols, acids, alkenes, aldehydes etc)

> on oligonucleotides, oligo saccharides, protein

D, Liu *et al.* *Nat. Chem.* 2011, 3, 146

Features of DNA-Templated Synthesis



Advantages

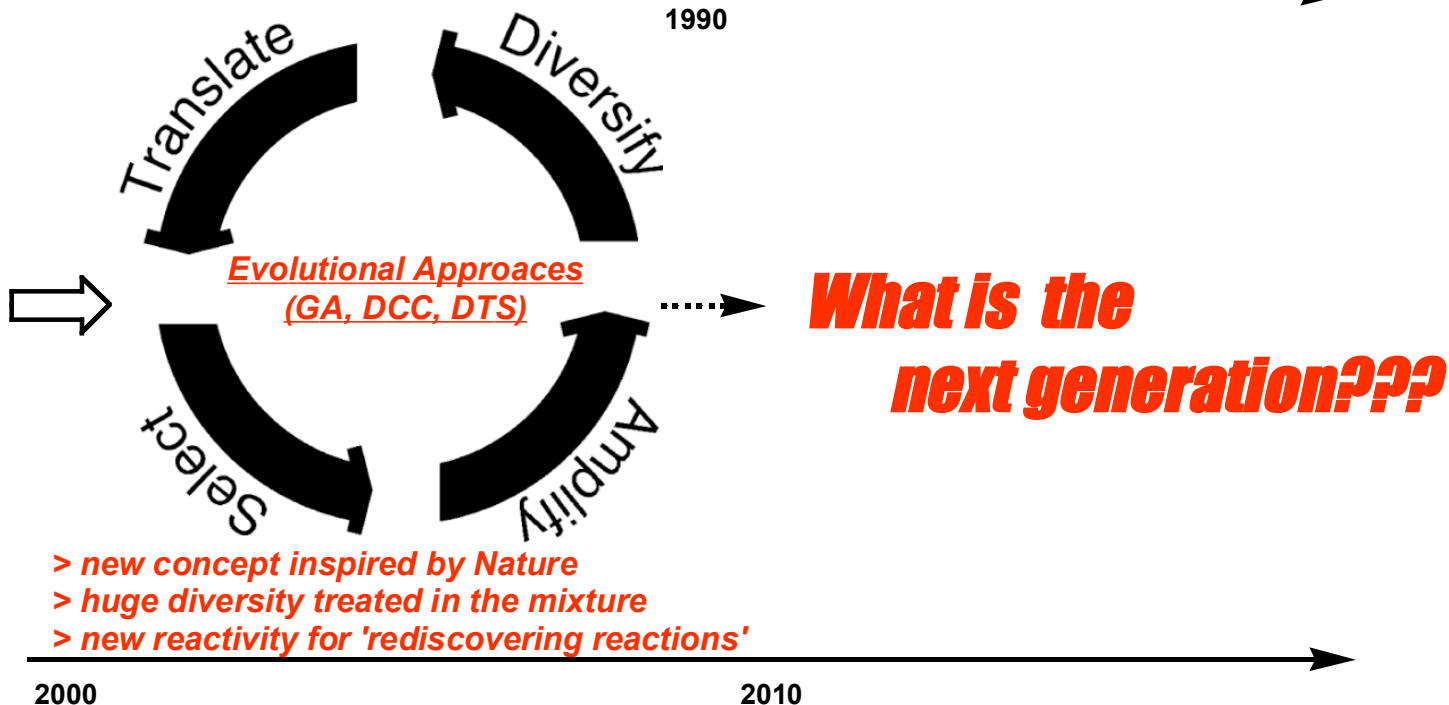
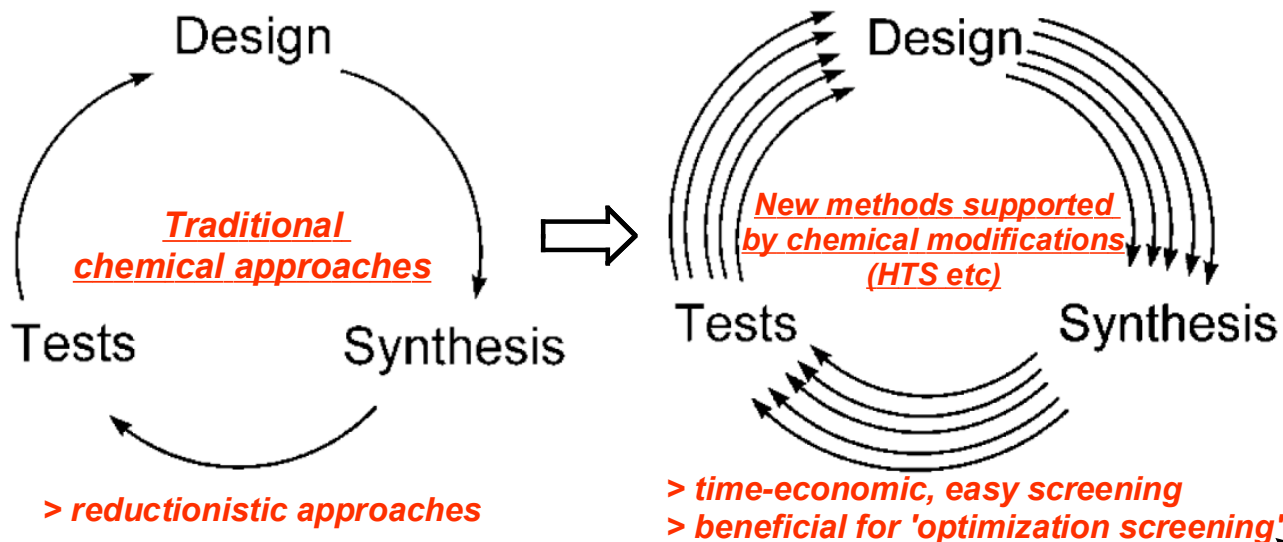
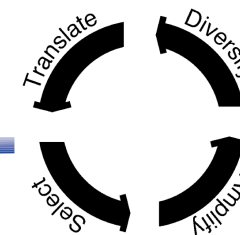
- > ***genuine evolutionary cycle*** (selection, translation, amplification)
- > with the ***synthetic molecules*** not amplifiable information (RNA etc)
- > in a single solution
 - ***the enormous magnitude of diversity, easiness, new reactivity***
- > ***reactivity controlled by DNA annealing***
 - ***selectivity***

Disadvantages

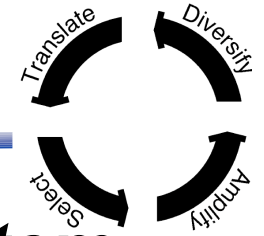
- > the need to prepare DNA-linked reagents
- > under conditions that ***support DNA hybridization***
(in the aqueous solvent, moderate temp., neutral pH etc)
- > just the combination of reactants in '***rediscovering reactions***'
 - the ability to develop ***conceptually new methodologies ???***

4. Future Direction

Toward Future Methodologies



Key Words

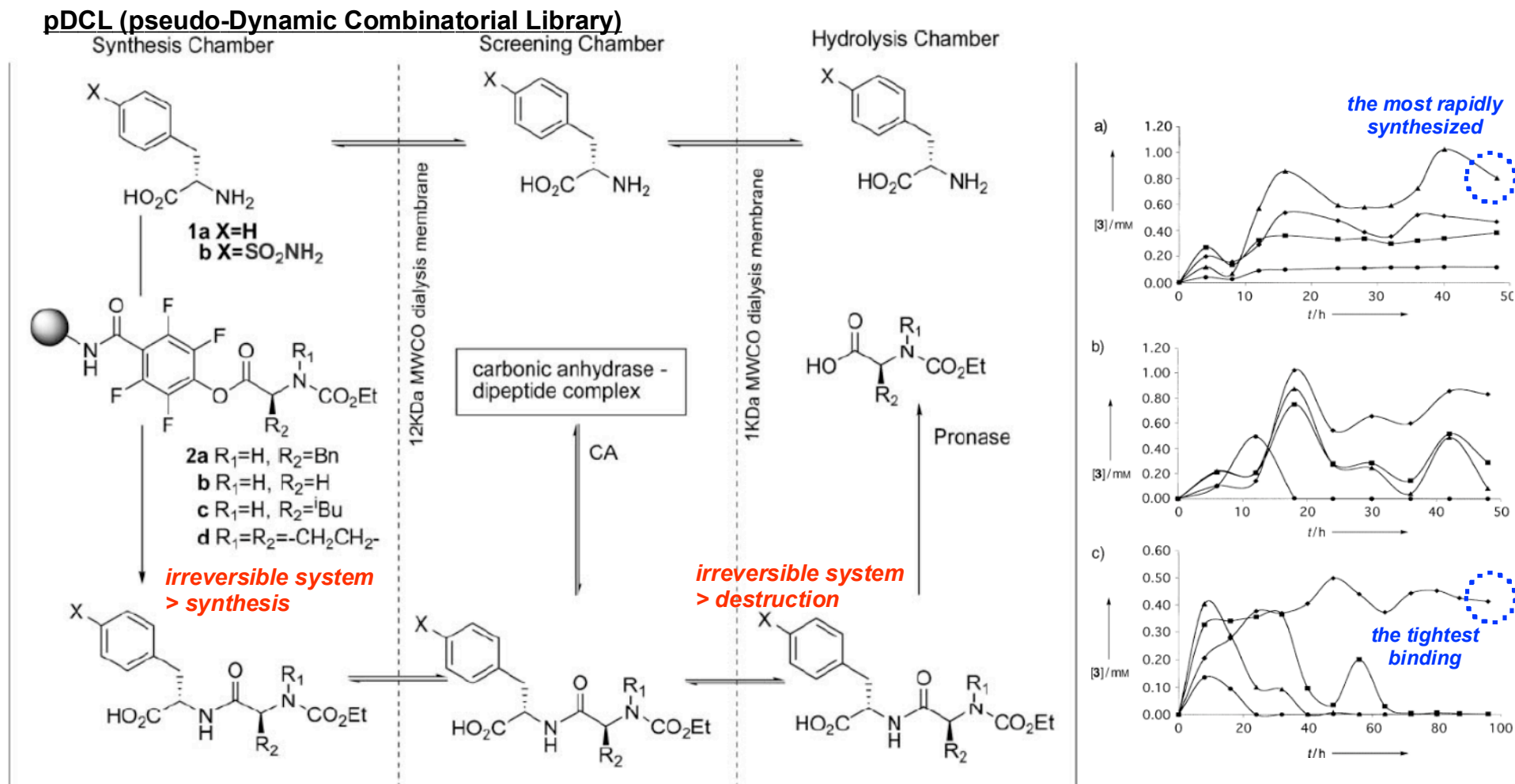
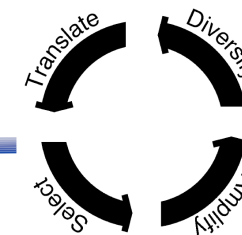


- > ***kinetic and thermodynamic combined system***
- > ***auto-catalytic***
- > ***self-replicating***
- > ***feedback***
- > ***oscillation***
- > ***systems chemistry***
- > ***application to medicine***

Also see: Mr. Sato's Lit. Seminar (B4)

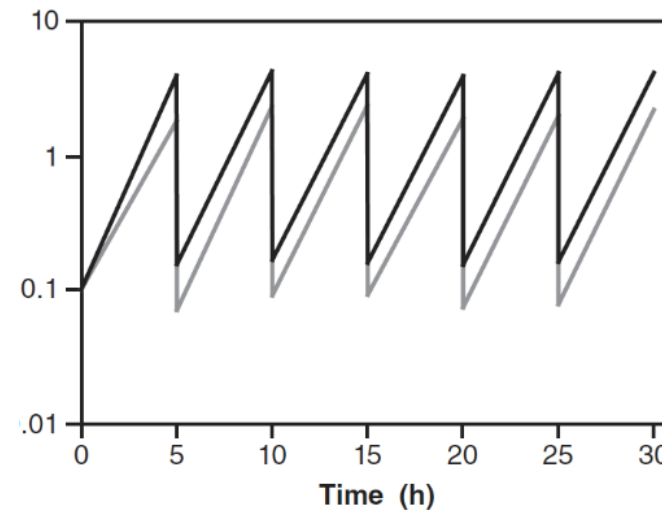
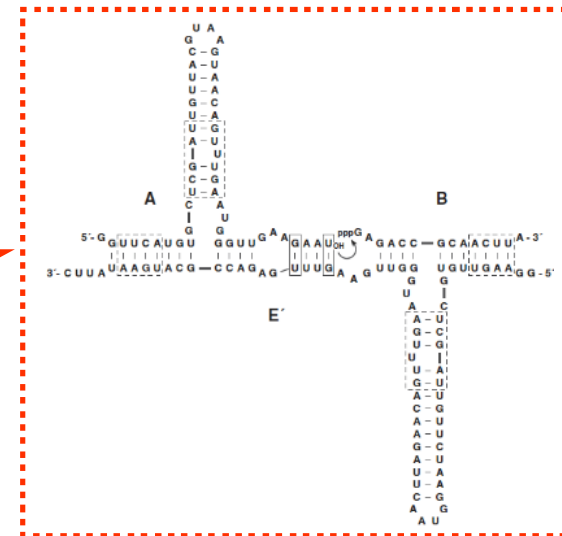
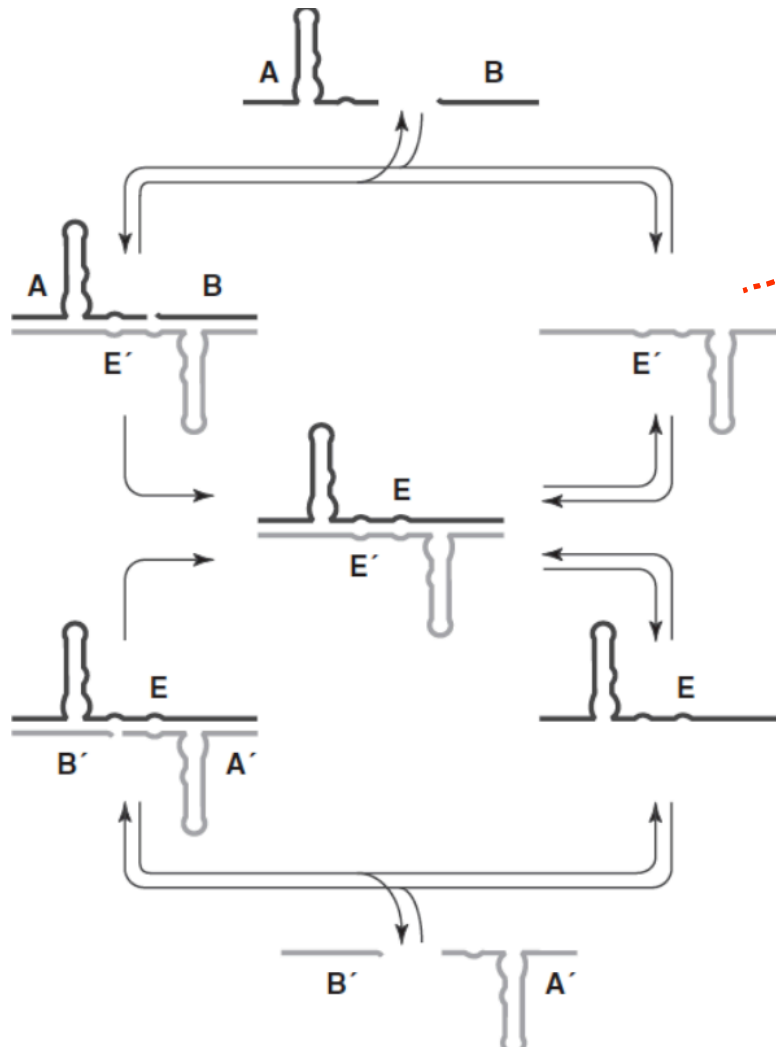
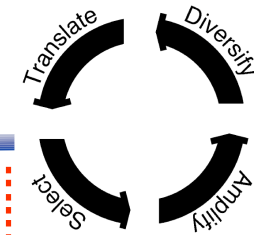
@ multiple component as a whole system

Molecular Networks under Kinetic Control



> **biology: balance between several processes that are kinetically controlled**
 → **drug discovery ???**

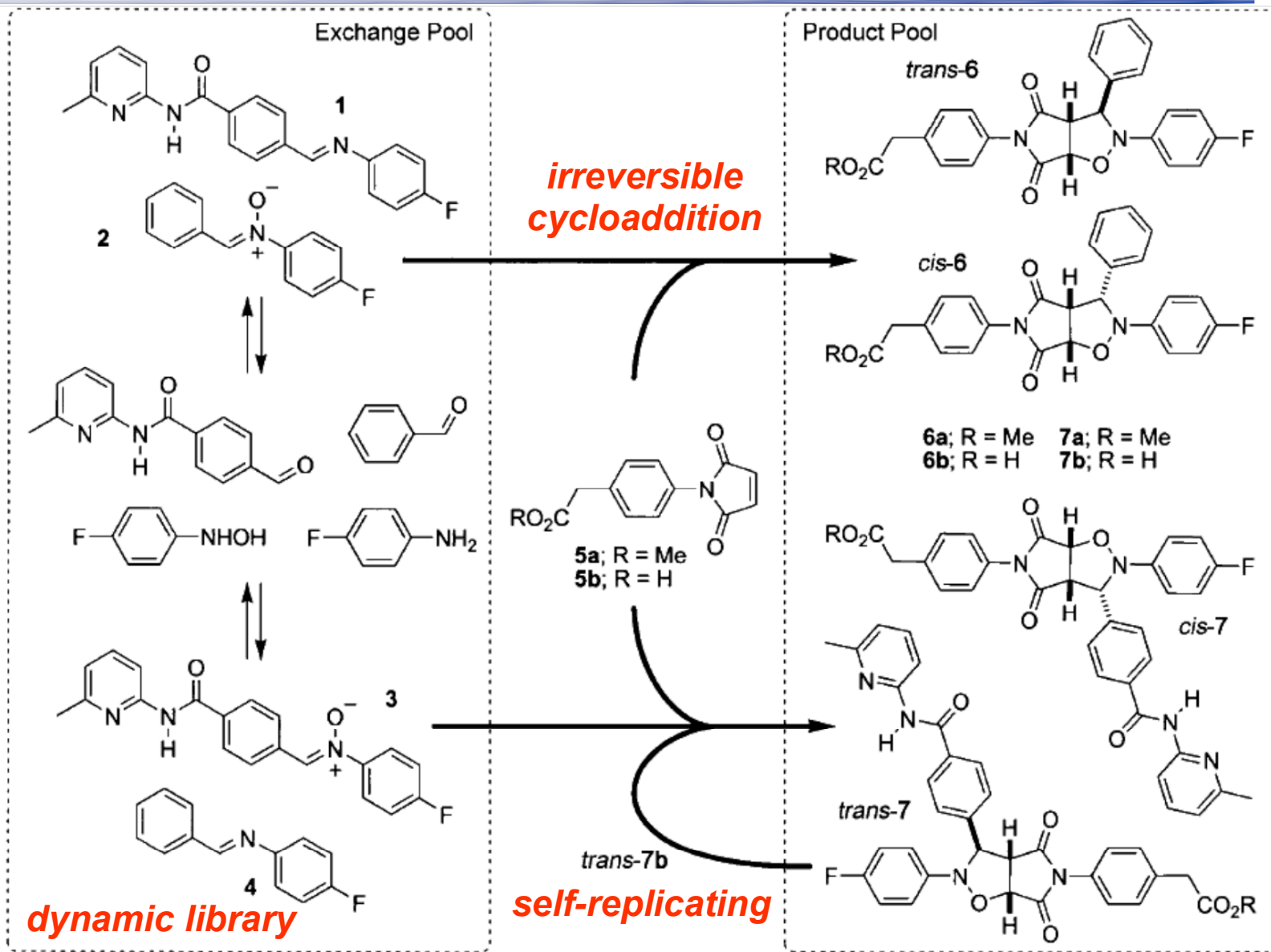
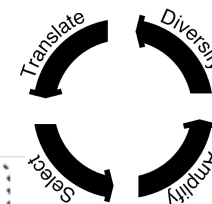
Self-Replicating (1)



- > **2 cross-replicating RNA enzymes**
- > **Self-sustained amplification without other proteins**

G. F. Joyce *et al.* *Science.*, 2009, 323, 1229
 Also see: Mr. Sato's Lit. Seminar (B4)

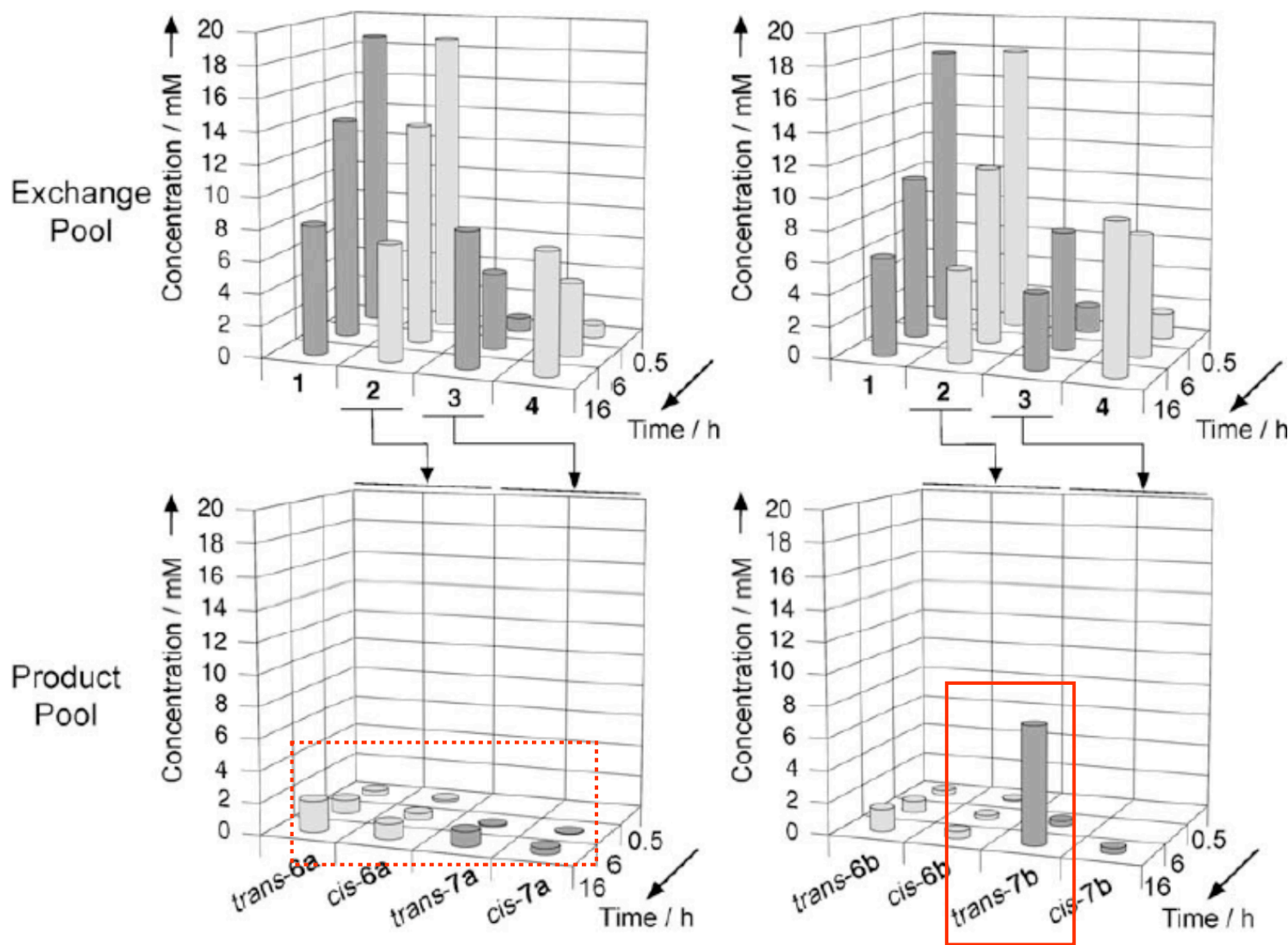
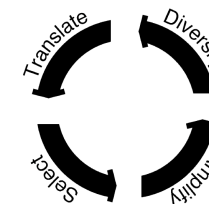
Self-Replicating (2)



D. Philip *et al.* *ACIE.*, 2006, 45, 6344

D. Philip *et al.* *ACIE.*, 2008, 47, 9965

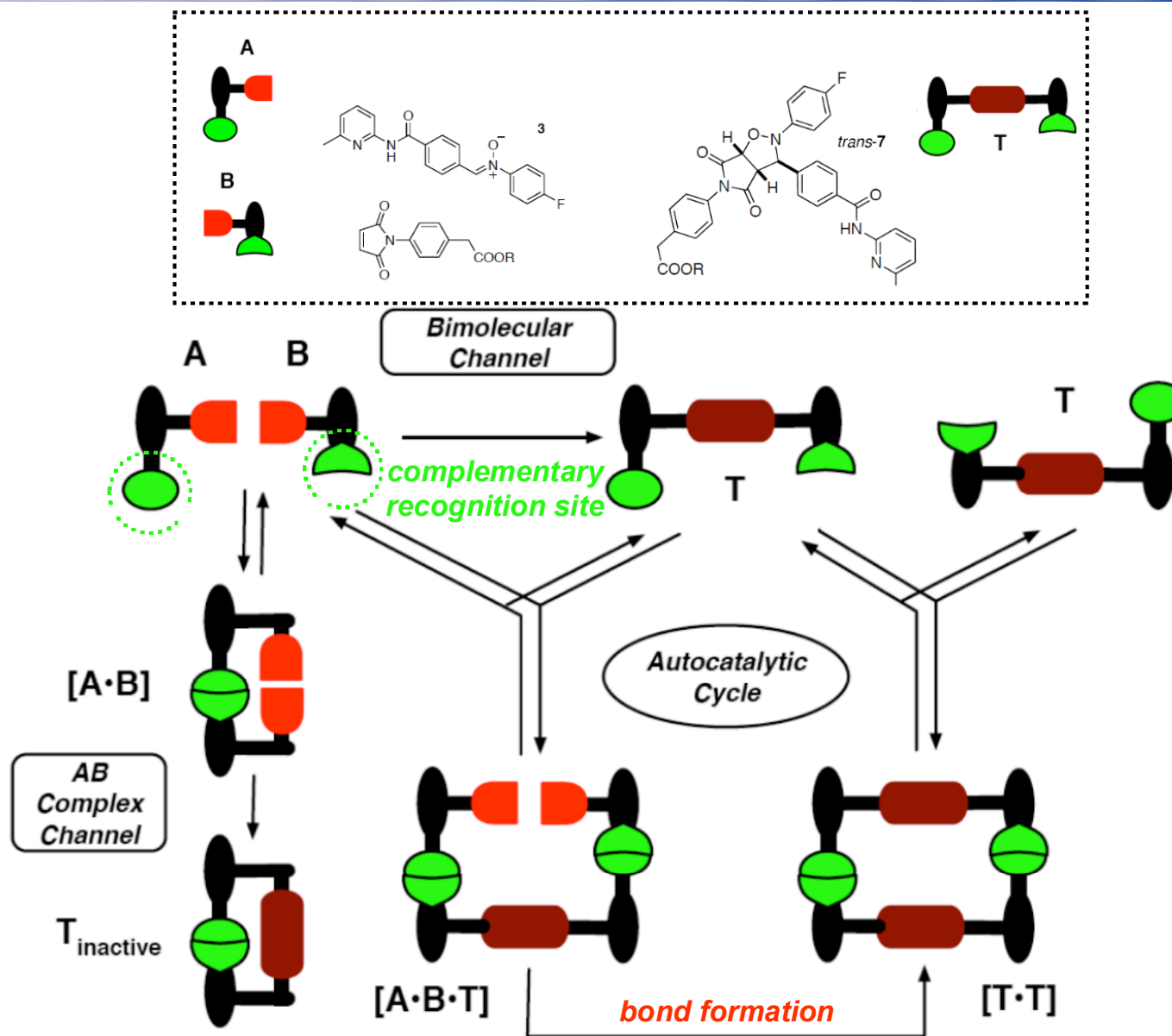
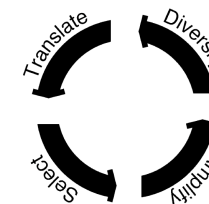
Self-Replicating (3)



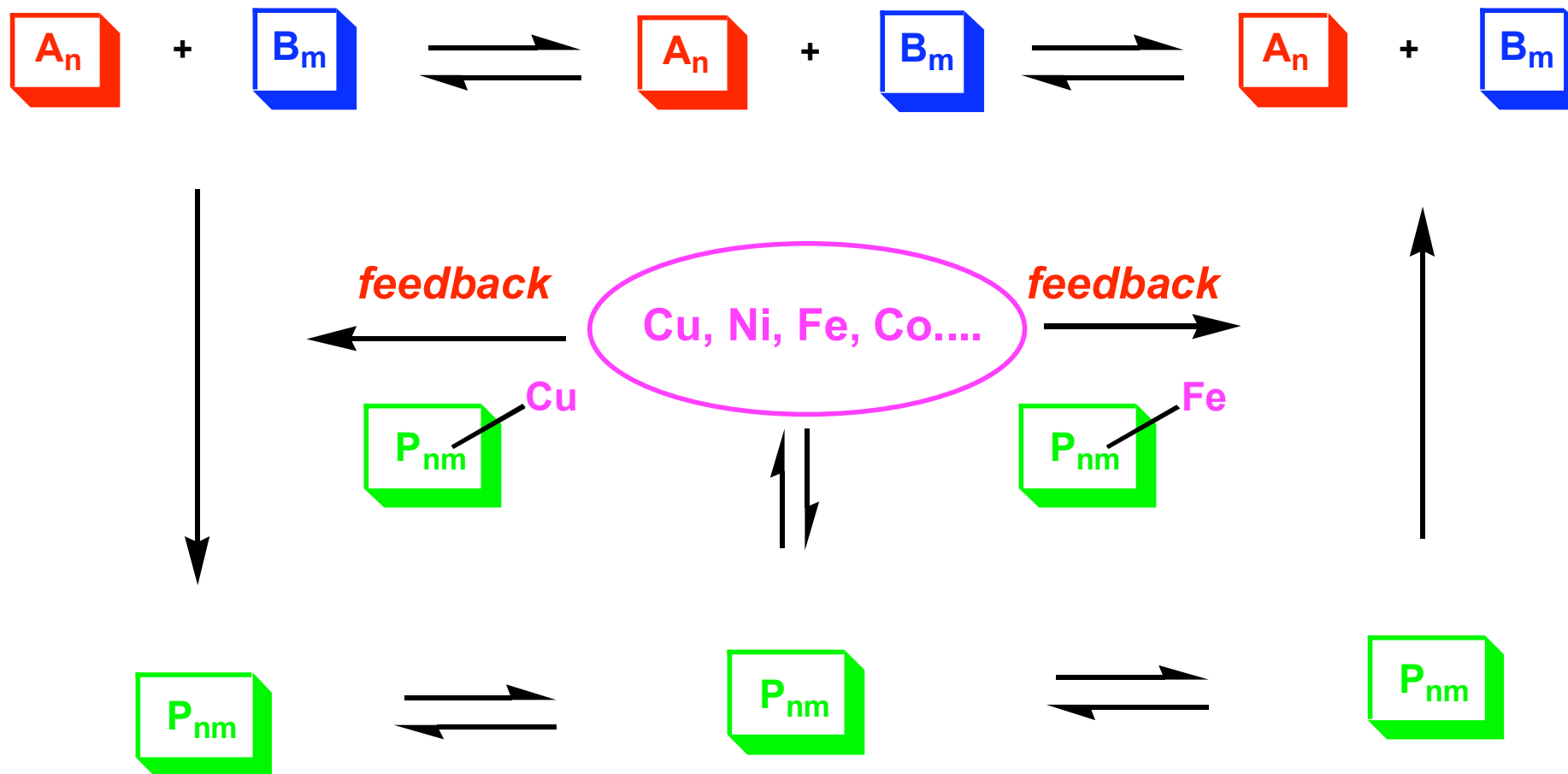
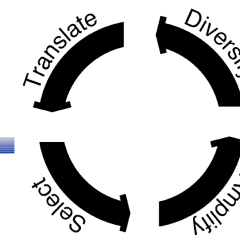
> the catalytic [3•5•trans 7] complex accelerated **own formation**

D. Philip et al. ACIE., 2006, 45, 6344 D. Philip et al. ACIE., 2008, 47, 9965

Self-Replicating (4)

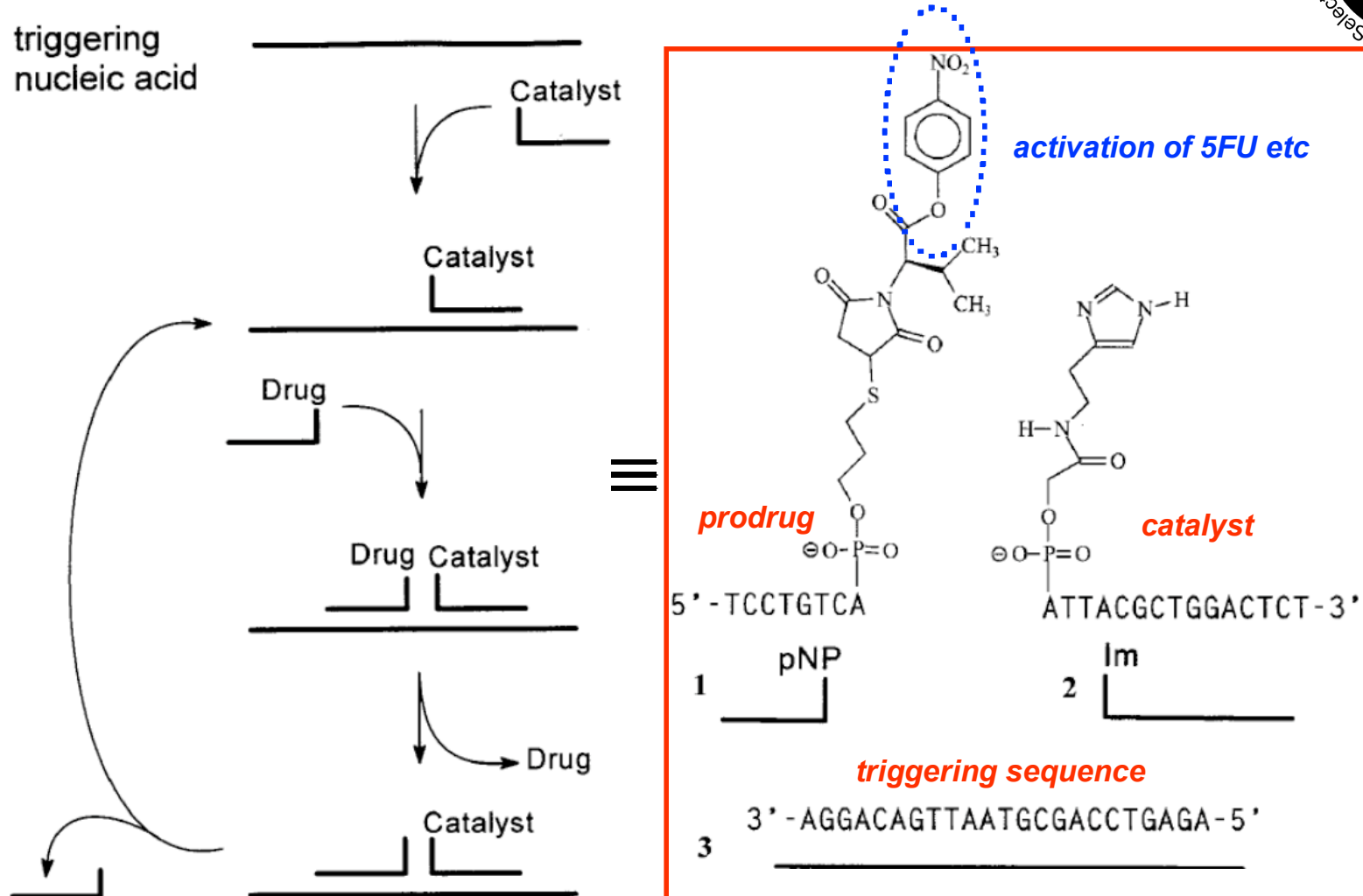
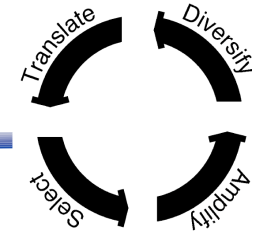


Ideal System



@ Can we construct the similar system as 'organic chemists of organometallics' ???

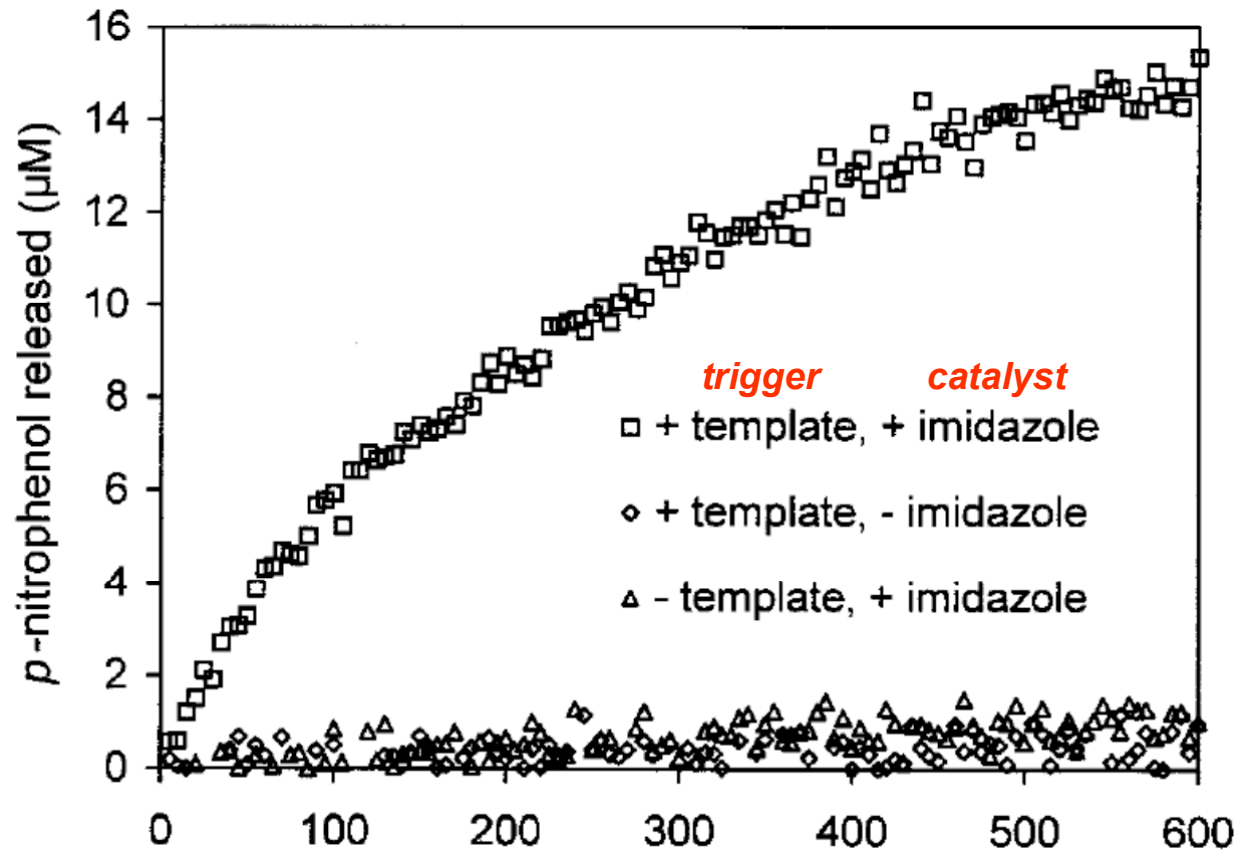
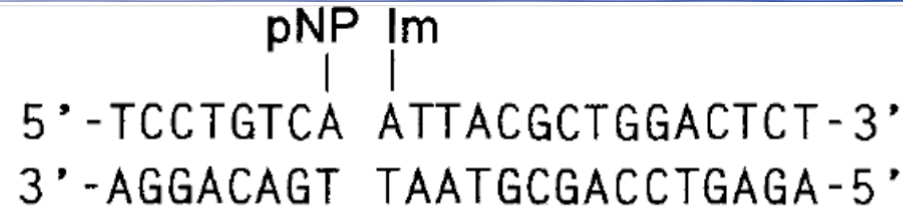
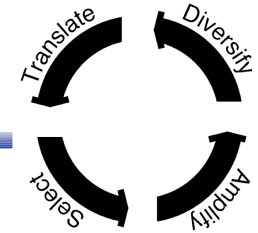
Application to Medicine (1)



> rational concept design for **'triggered catalytic drug release'**

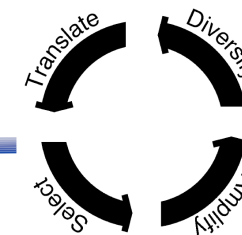
> the disease state (**binding by catalyst**) triggered the release of drugs

Application to Medicine (2)

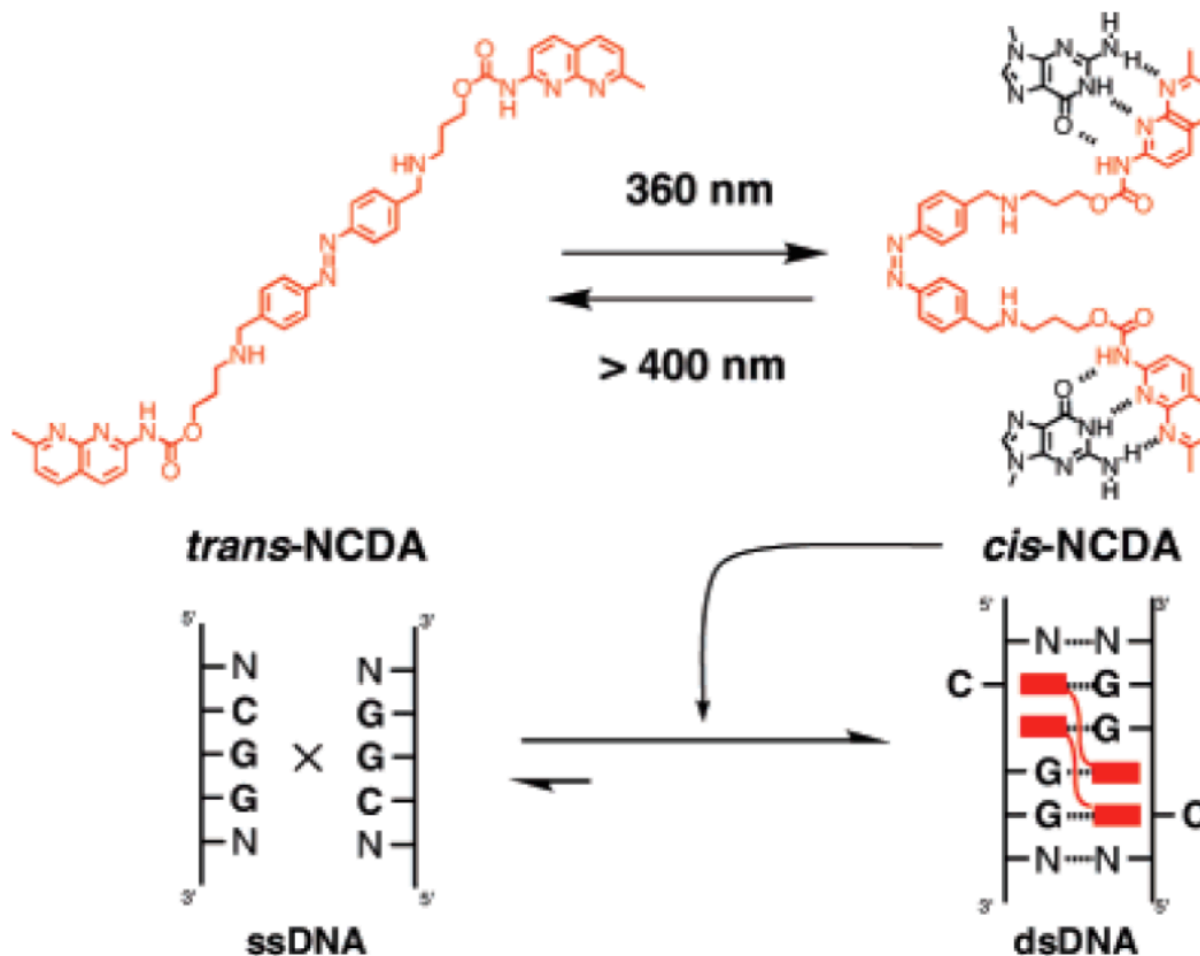


- > **only in the presence of a catalyst and a trigger**, drug release proceeded
- > **enzyme-like** behavior following Michaelis-menten kinetics
- > sensitive to **mismatch** sequences → **SNP** ???

Application to Medicine (2)



Phostoswithcable molecular glue for DNA



*@ Without using external switch, can we develop the original methodologies ???
→ New systems with 'oscillation' in time (rhythms) and space (patterns)*