

Combinatorial libraries: strategies and methods for 'lead' discovery

Alan Spivey

Department of Chemistry
University of Sheffield

Key sources of information

- WWW:

- Diversity information pages [<http://www.5z.com/divinfo/>]
- *J. Combinatorial Chem.* [<http://acsinfo.acs.org/journals/jcchff/index.html>]
- *Combi. Chem. H.T.S.* [<http://www.bscipubl.demon.co.uk/cchts/index.html>]

- Books

- *Combinatorial peptide and nonpeptide libraries-a handbook*, Ed. G.Jung, VCH, Weinheim, **1996**.
- *Combinatorial chemistry-synthesis and application*, S.R.Wilson, A.W.Czarnik, Wiley, New York, **1997**.

- Reviews

- ‘Combinatorial chemistry’, *Chem. Rev.* **1997**, 97(2), special issue.
- ‘Combinatorial chemistry’, *Curr. Opin. Chem. Biol.* **1998**, 2(3) & **1999**, 3(3).
- ‘Combinatorial chemistry’, S. Borman, *Chem & Eng. News* **1997**, Feb24, 43.

Format and scope of lecture

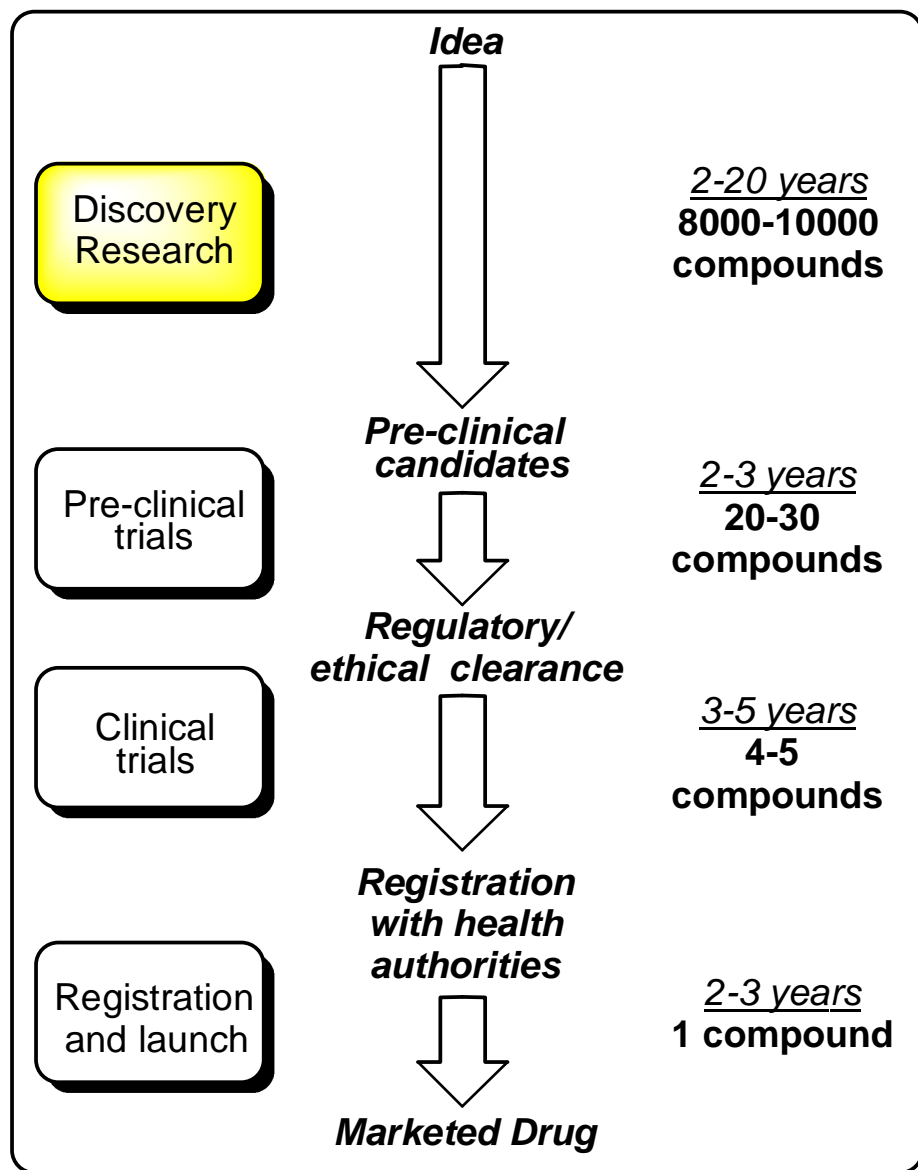
- What is combinatorial chemistry?
- The drug discovery process
- Approaches to combinatorial library synthesis:
 - mix and split synthesis
 - parallel synthesis
 - encoded tagging
- Library types:
 - oligomeric libraries
 - template based libraries
- Combinatorial drug discovery!

What is combinatorial chemistry?

Combinatorial chemistry is a useful tool for rapidly optimizing molecular properties, particularly ones that are difficult to design *a priori*...

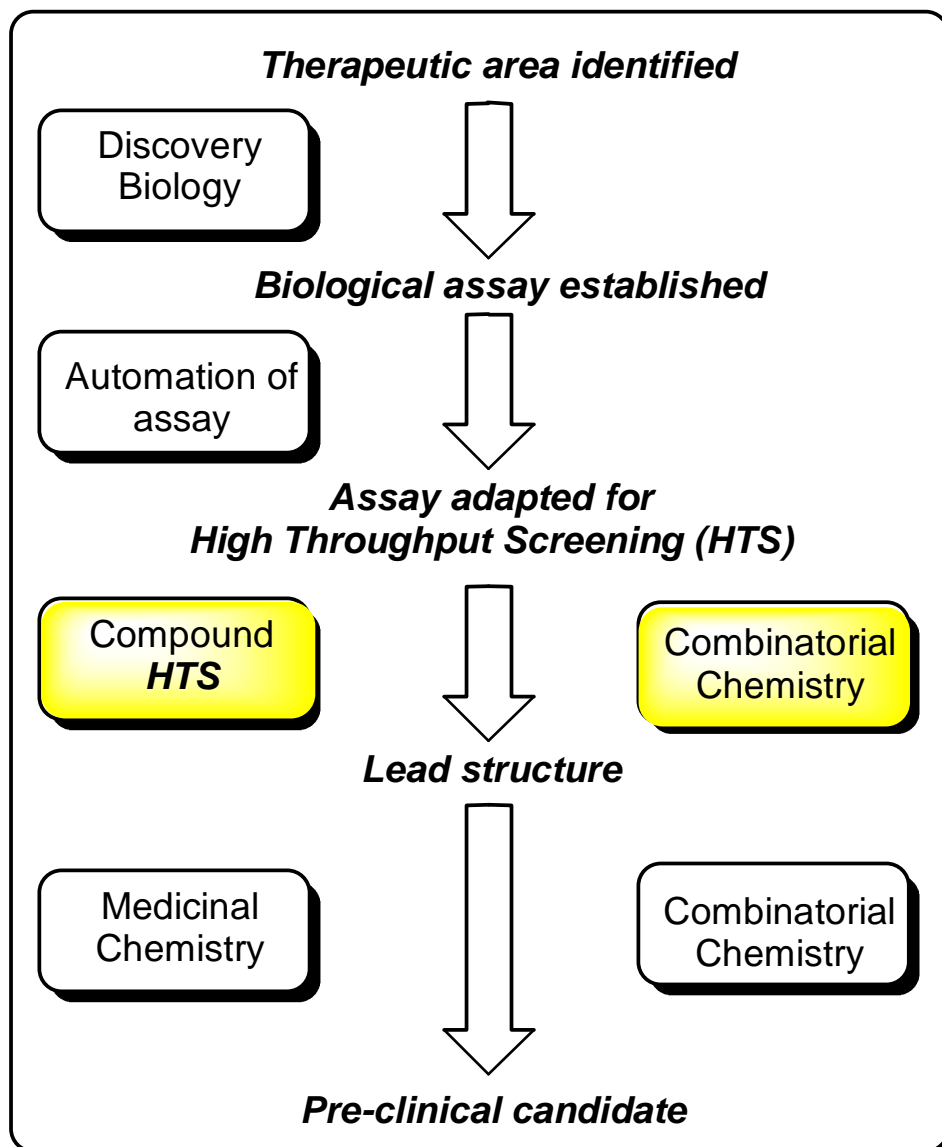
Nature uses a combinatorial approach to generate diverse functional macromolecules such as antibodies to recognize a vast array of antigens.

The drug discovery process



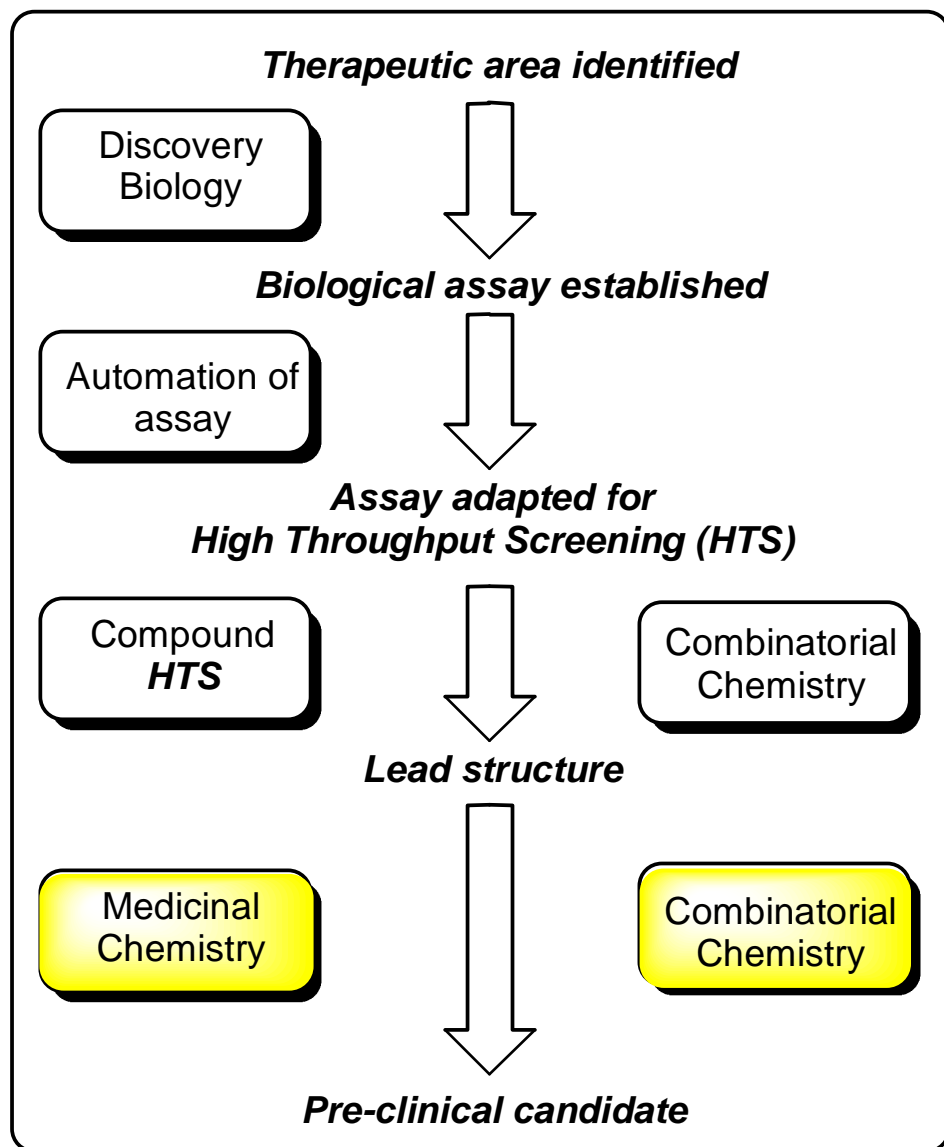
- The total cost of bringing a new drug to market is typically ~£250m (i.e. EXPENSIVE!)
- Of this, £170m is spent on DISCOVERY RESEARCH.
- This reflects the large amount of TIME involved in synthesising new compounds.
- A typical chemist can synthesise ~100 compounds a year using traditional techniques.
- **SOLID PHASE ORGANIC SYNTHESIS (SPOS)** and **COMBINATORIAL CHEMISTRY** are beginning to revolutionise this situation.

Discovery chemistry: stage 1



- **High Throughput Screening (HTS):**
 - Rapid, automated screening of compounds for specific biological activity.
- **Role of combinatorial chemistry:**
 - Very large libraries.
 - Maximum diversity libraries.
 - Mix and split libraries (& parallel synthesis).
 - Mixtures of compounds (& single compounds).

Discovery chemistry: stage 2



- **Medicinal chemistry:**
 - Systematic optimisation of molecular and physicochemical properties of lead compound
- **Role of combinatorial chemistry:**
 - Small libraries.
 - ‘Targeted/focussed’ libraries.
 - Parallel synthesis libraries.
 - Single compounds.

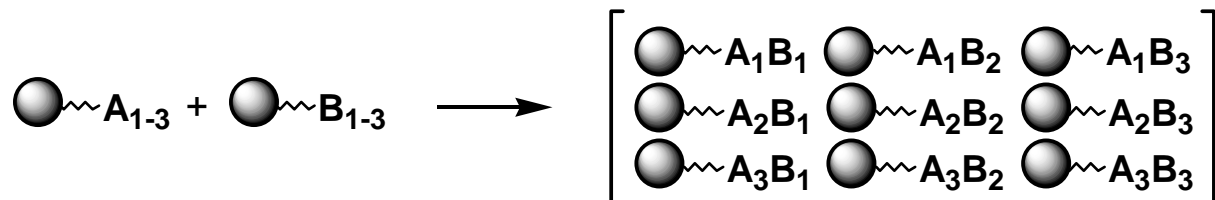
Traditional vs. combinatorial

- Traditional synthesis:



compounds prepared one at a time, characterised and screened

- Combinatorial synthesis:



reaction of 3 reagents A_x with 3 reagents B_y provides a library of 3^2 (i.e. 9) compounds A_xB_y

introduction of a third set of 3 reagents C_z increases the library size to 3^3 (i.e. 27) compounds $A_xB_yC_z$

Approaches to 'combinatorial' library synthesis

- ***In vivo* - biological methods:**

- Phage display, plasmids, polysomes etc.

- ***In vitro* - synthetic methods:**

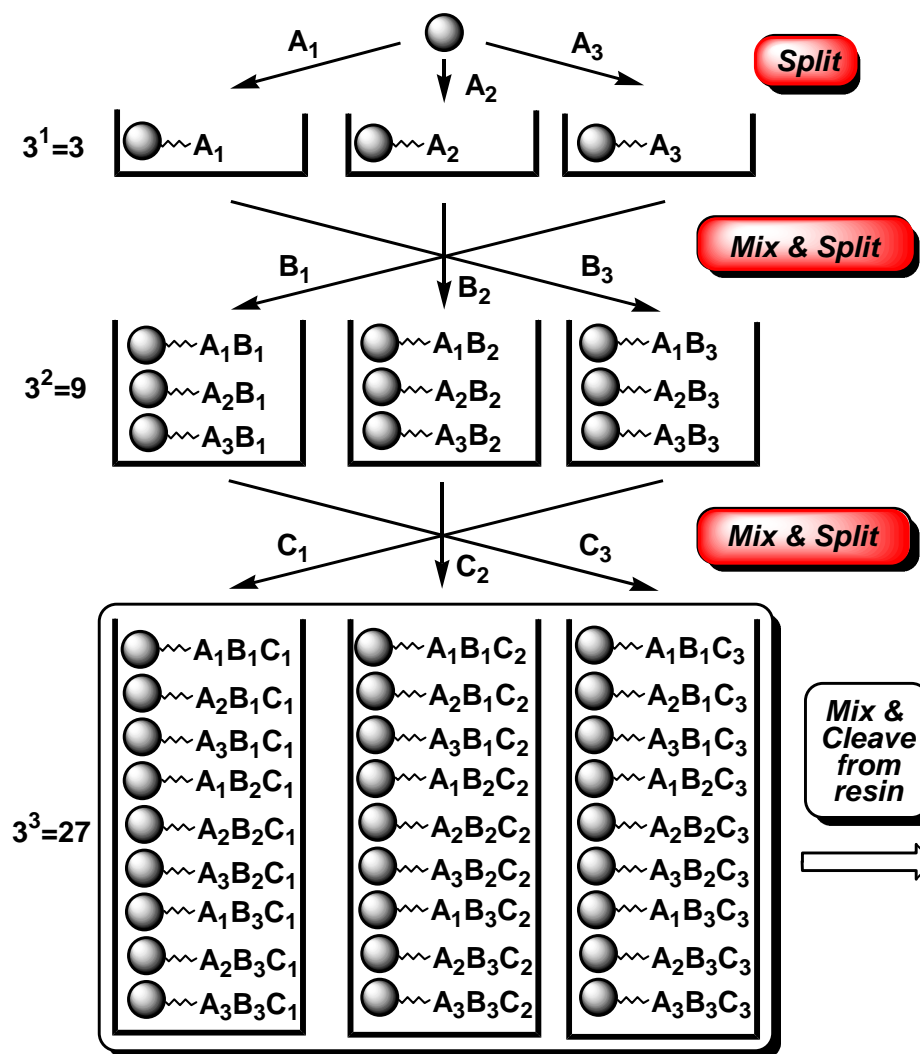
- **Mix and split using Solid Phase Organic Synthesis (SPOS).**

- Cleavage from the solid support following 'mix and split' results in complex mixtures (pools) of compounds. Screening of these mixtures yields 'hits' whose identity must be determined by 'deconvolution'.
- If screening can be performed 'on-bead' (i.e. 'one-bead one-compound' libraries) then deconvolution can be avoided.

- **Parallel synthesis using Solid Phase Organic Synthesis (SPOS).**

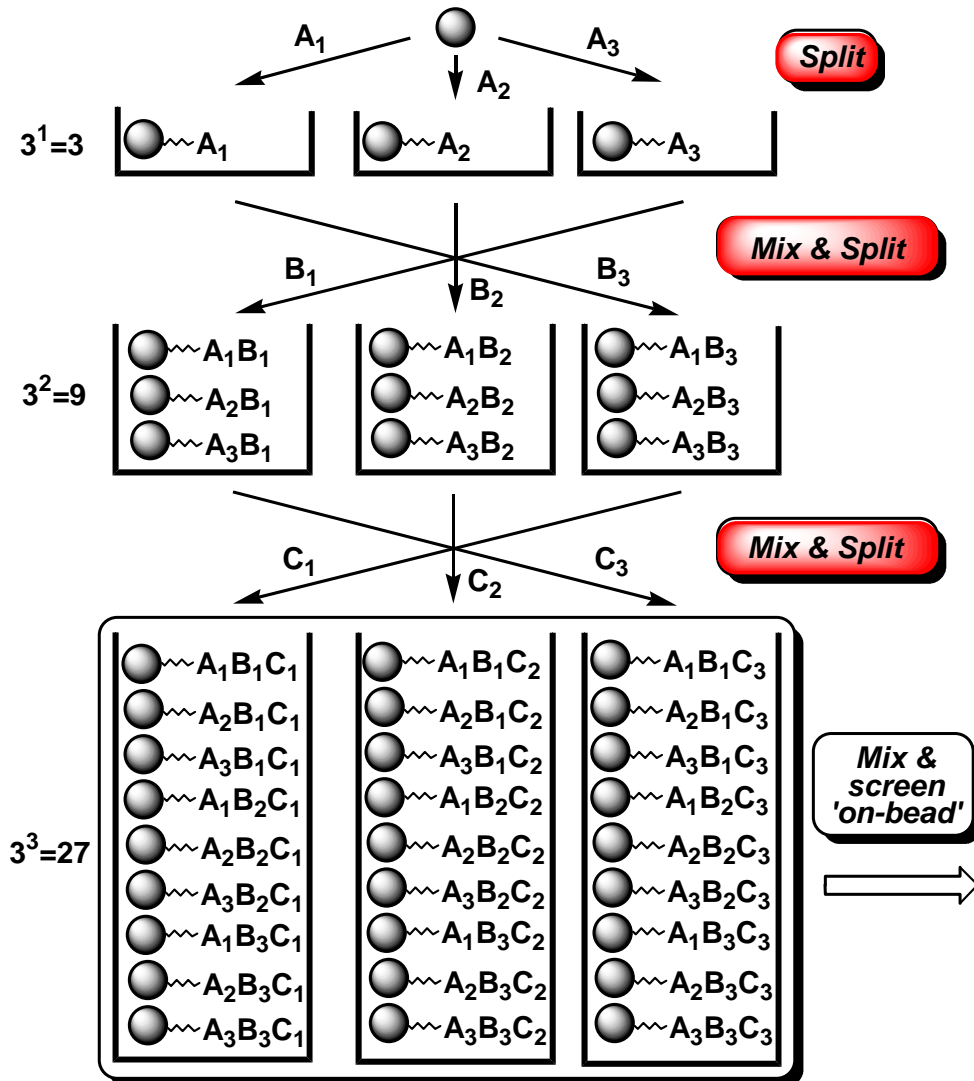
- Spatially separate synthesis of single compounds whose identity is uniquely defined by their location.

Mix & split synthesis: libraries of mixtures of compounds in solution



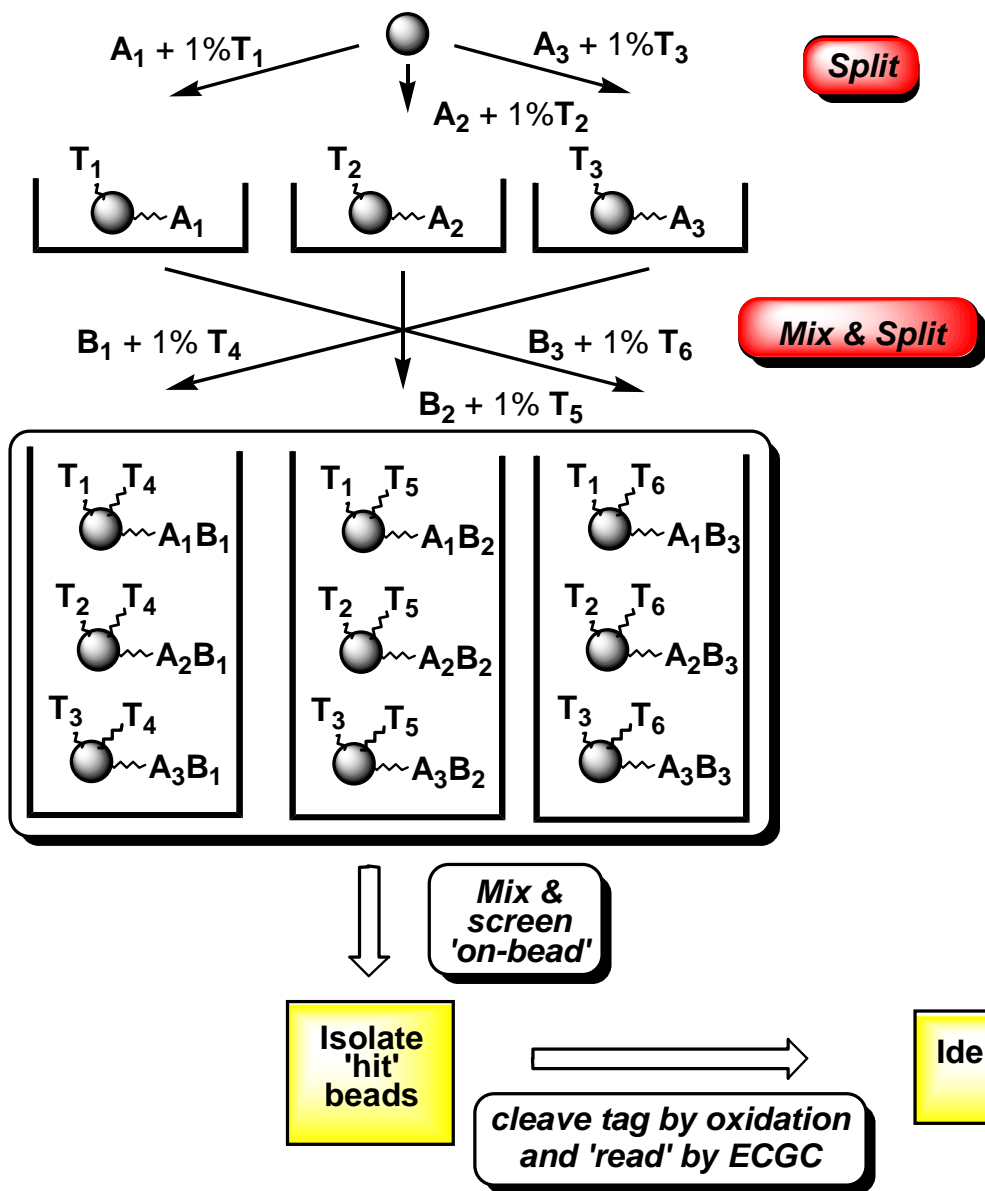
- Screening complex mixtures of compounds in solution can give false 'hits' due to synergistic effects.
- Identification of a compound within the mixture responsible for the 'hit' requires iterative **deconvolution**.
- Houghton *Nature*, 1991, 354, 84.

Mix & split synthesis: 'one-bead one-compound' libraries



- Requires a very sensitive screening protocol which can accommodate resin bound compounds.
- Identification of a 'hit' compound on (or from) a single bead ($\sim 100\text{pm}$) by:
 - analytical methods e.g. Edman sequencing of peptides, *MALDI-TOF MS*, single bead *NMR*...
 - reading '**encoding tags**' on beads.

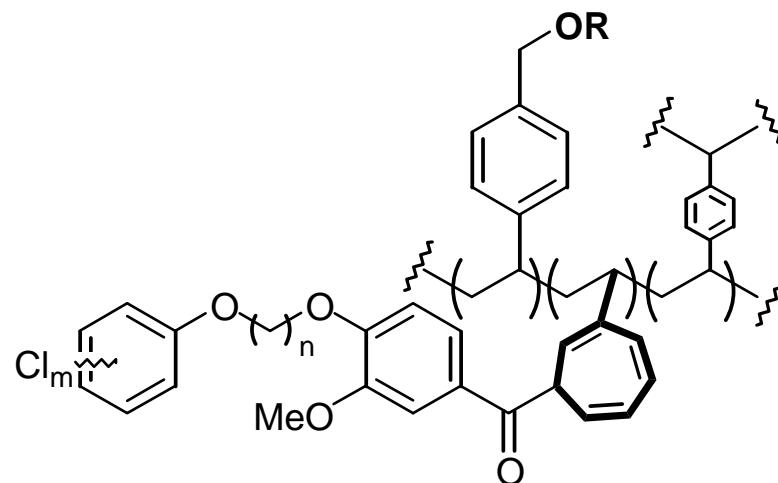
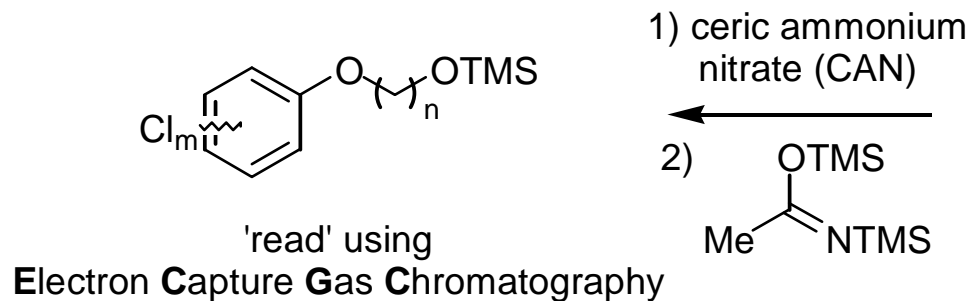
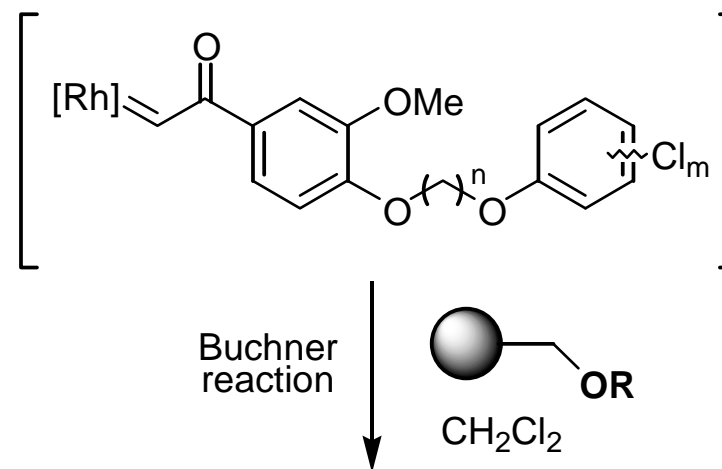
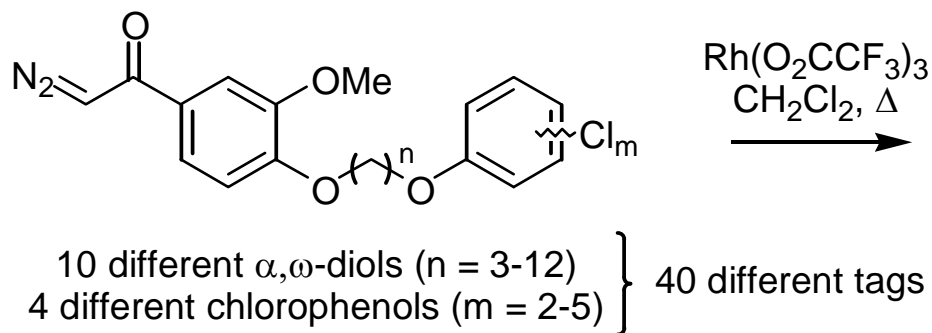
Clark Still's encoded tagging protocol



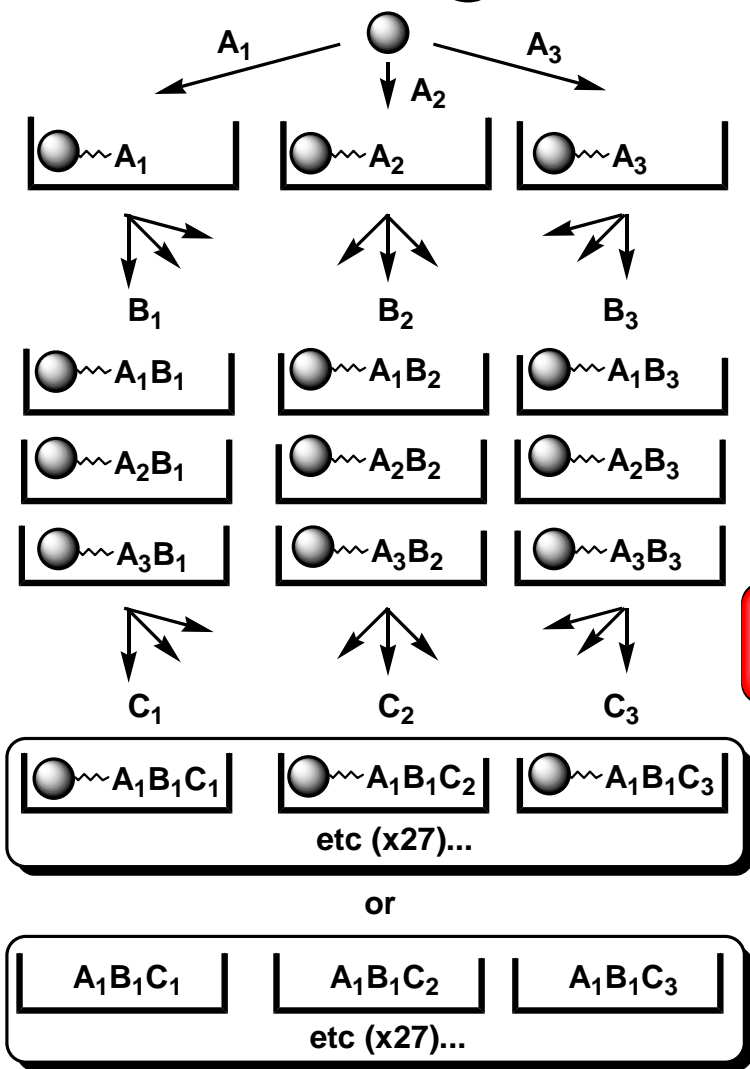
- Still *Acc. Chem. Res.* **1996**, 29, 155.
- Each 'monomer' used in the library synthesis has an associated encoded tag.
- The tags are chlorinated aromatic compound which can be analysed at sub-picomolar levels by **Electron Capture Gas Chromatography (ECGC)**.
- Allows for hit identification at one-bead fidelity for any type of library

Mechanism of Clark Still encoded tags

- Still *J. Org. Chem.* **1994**, *59*, 4723.



Parallel synthesis: spatial separation gives single compounds



Split

- Suitable for the synthesis of relatively small libraries as each compound requires its own reaction 'well'.

Split

- Each reaction 'well' may be anything from a small flask to a radio-frequency tagged 'tea bag' to an etched region on a silicon chip!
- Once screening has identified a hit no further work is required to deduce the identity of the active compound although it is routine practice to independently verify the structure.

**Split
± cleavage
from resin**

**Screen
each
'position'**

'Hit'

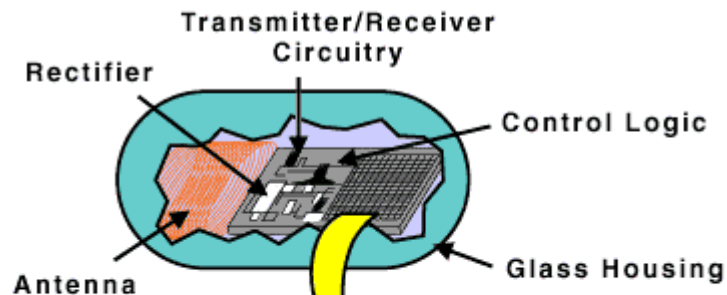
**Spatial location
of hit defines
its identity**

**Identity of
hit**

spatially separated libraries

Keeping track of 'tea-bag' parallel synthesis: Irori radio-frequency tagging

- <http://www.irori.com/>



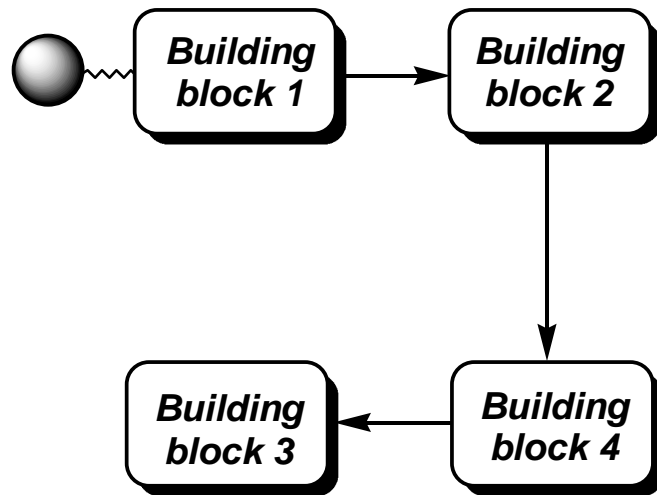
Synthesis Manager Data Base

1	2	3																	
		└─ 1-naphthoyl																	
		└─ nicotinoyl																	
		└─ 5-bromonicotinoyl																	



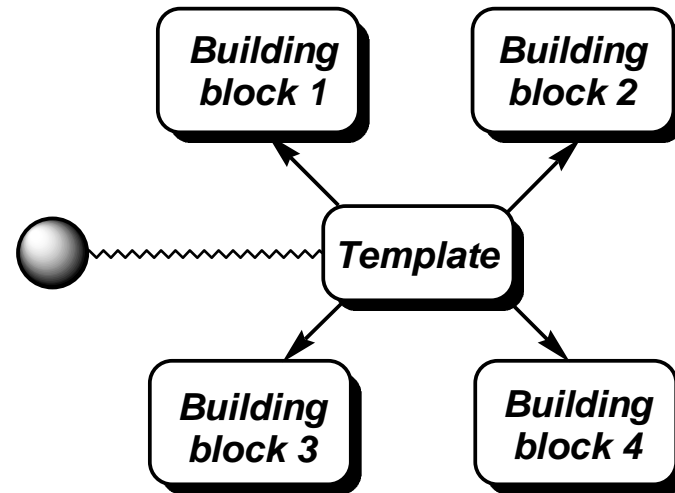
Library types

- There are essentially two strategically distinct types of library at the molecular level:



Oligomeric

peptides/peptoids
oligonucleotides
oligosaccharides
unnatural oligomers
polyaromatics

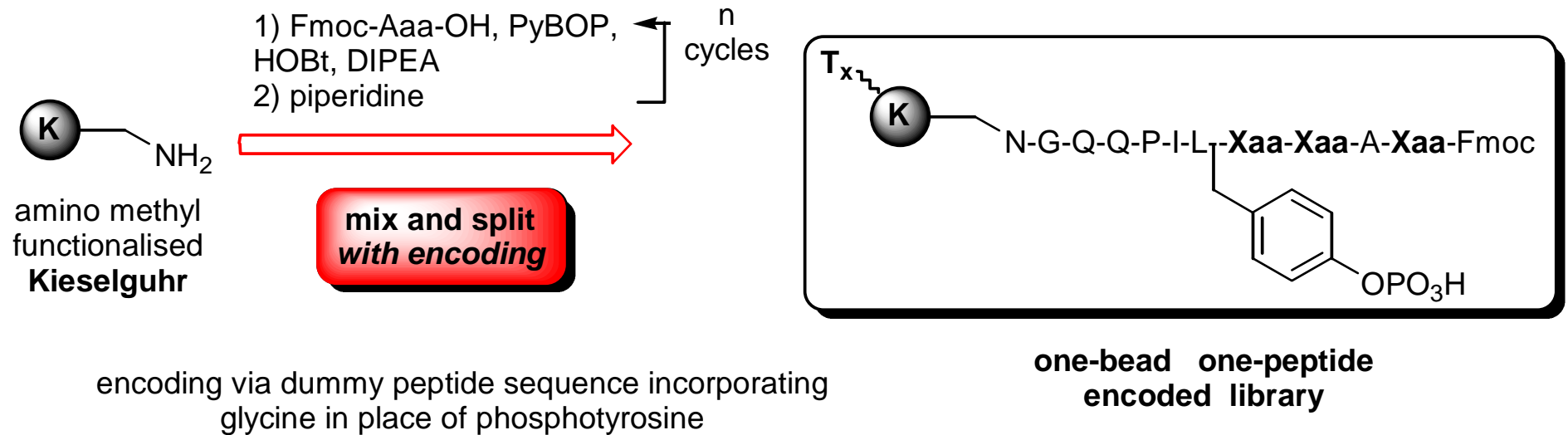


Template based

drug-like molecules
natural product-like molecules
heterocycle based molecules

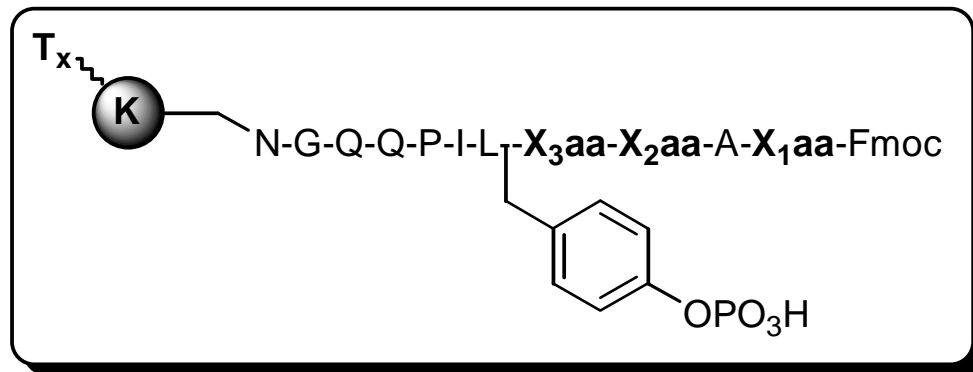
Balasubramanian's peptide library

- Protein tyrosine phosphatase substrate library (oligomeric).
- Balasubramanian *J. Am. Chem. Soc.* **1997**, *119*, 9568.
- **Library synthesis:**



Balasubramanian's peptide library

- Protein tyrosine phosphatase substrate library (oligomeric).
- Balasubramanian *J. Am. Chem. Soc.* **1997**, 119, 9568.
- **Library screening:**



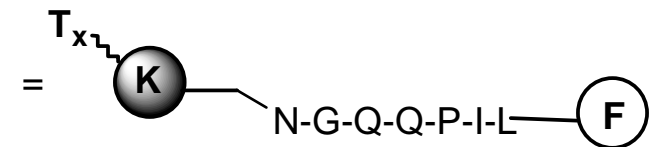
one-bead one-peptide
encoded library

- 1) leukocyte antigen receptor protein tyrosine phosphatase (*PTP*)
- 2) α -chymotrypsin
- 3) fluorescent labelling of *N*-terminus with carboxyfluorescein: **(F)**

Identity of PTP
substrate sequences

sequence
encoding tag peptide

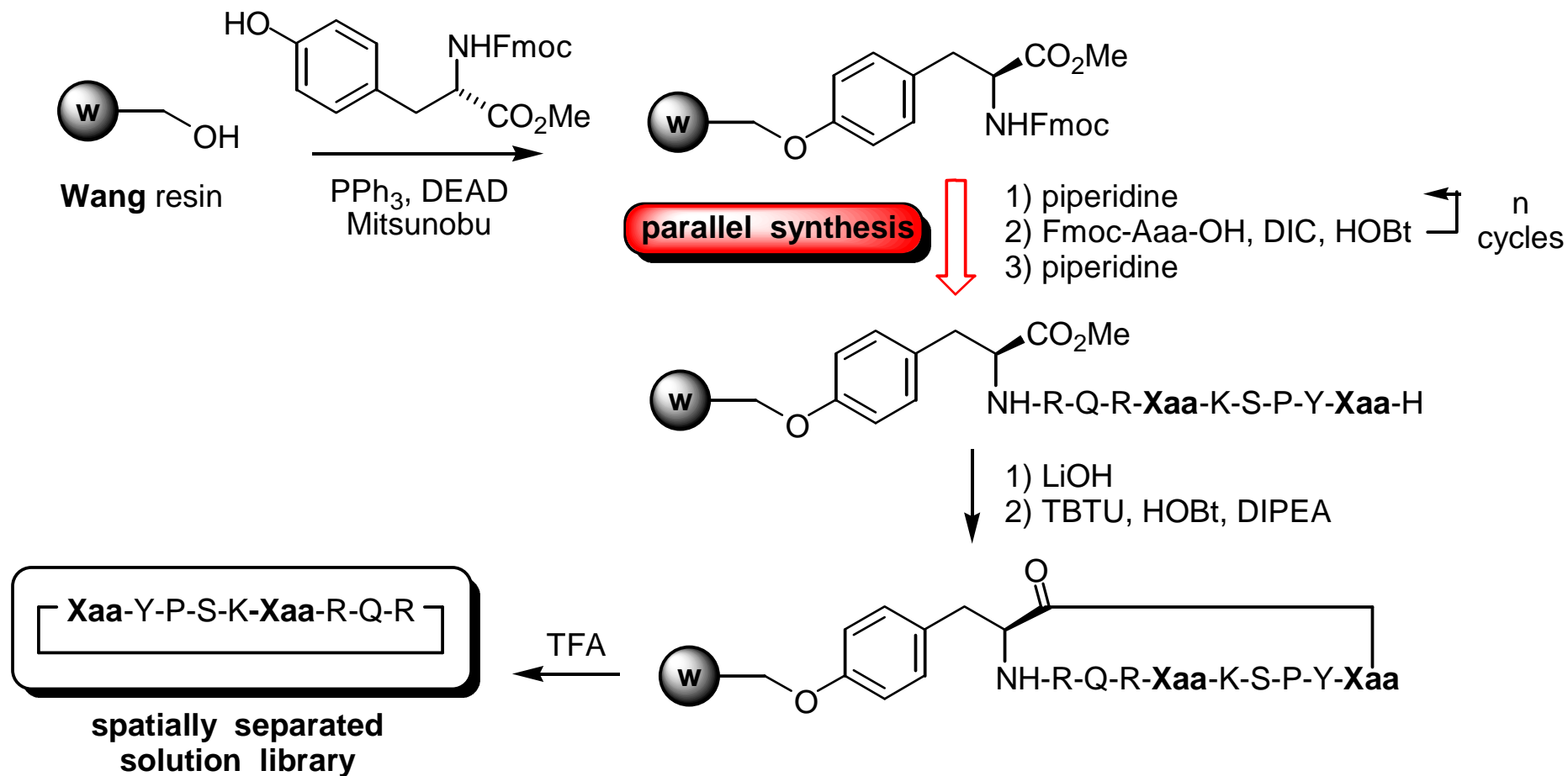
Hits



"preference for at least two acidic
residues in variable positions and
a glutamic acid residue at X_{1aa} "

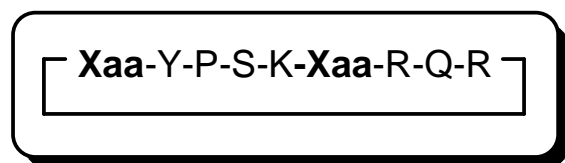
Beck-Sickinger's cyclic peptide library

- Neuropeptide Y analogue library (oligomeric).
- Beck-Sickinger *J. Org. Chem.* **1999**, *64*, 4353.
- **Library synthesis:**



Beck-Sickinger's cyclic peptide library

- Neuropeptide Y analogue library (oligomeric).
- Beck-Sickinger *J. Org. Chem.* **1999**, *64*, 4353.
- **Library screening:**



spatially separated
solution library

competitive binding assay in solution
with radiolabelled neuropeptide Y



Hits

identity defined by
spatial location

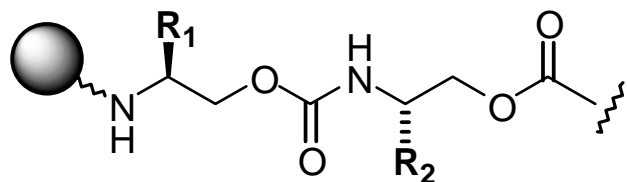
"weak competitive
binding
at μM level by range
of derivatives"

Oligonucleotide libraries

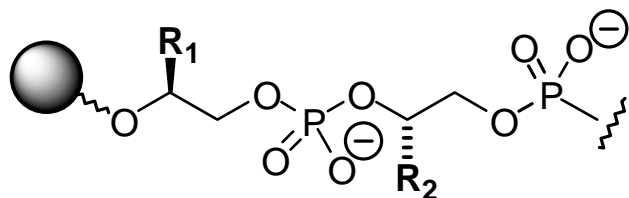
- These oligomeric libraries are generally prepared using *in-vivo* 'biological' methods and screened using **S**ystematic **E**volution of **L**igands by **E**xponential enrichment (SELEX) procedures.
- e.g. The discovery of very high-affinity RNA and DNA ligands to human IgE which inhibit binding to the Fc ϵ receptor I.
- Wiegand *J. Immunology* **1996**, 157, 221 (and references therein).

Unnatural backbone oligomer libraries

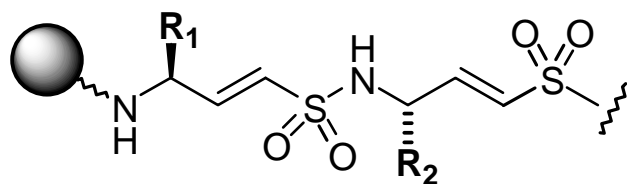
Backbone



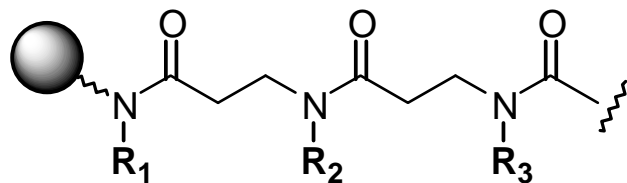
oligocarbamate



oligo-phosphodiester

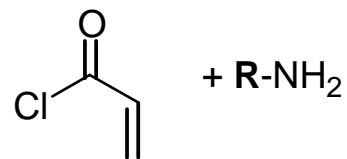
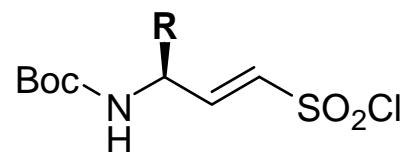
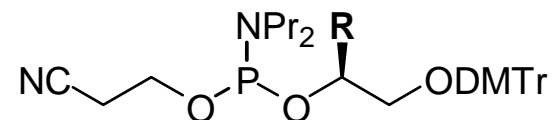
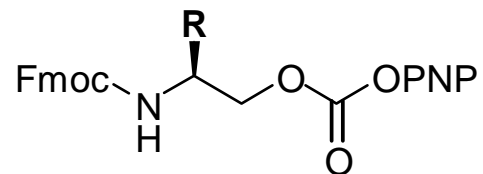


vinylogous sulfonamidepeptide



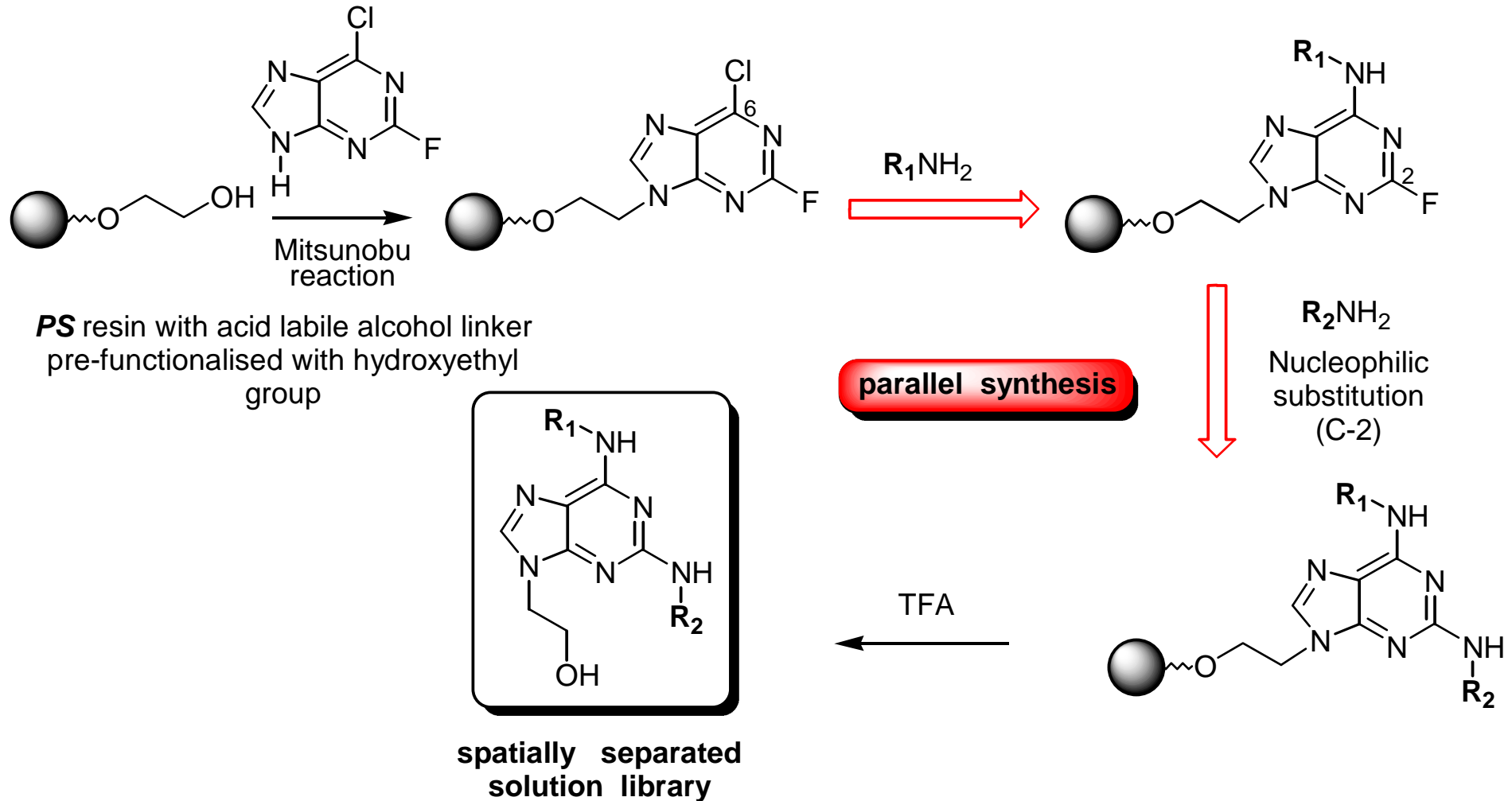
vinylogous sulfonamidepeptide

Monomers



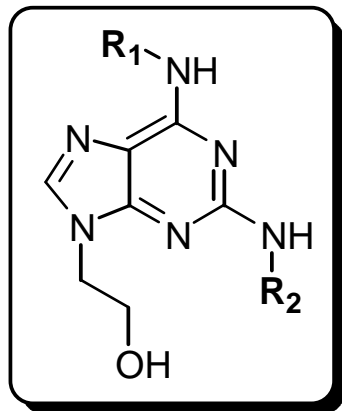
Schultz's purine library

- Kinase inhibitor library (template based).
- Schultz *Science* **1998**, 281, 533.
- **Library synthesis:**



Schultz's purine library

- Kinase inhibitor library (template based).
- Schultz *Science* **1998**, 281, 533.
- **Library screening:**



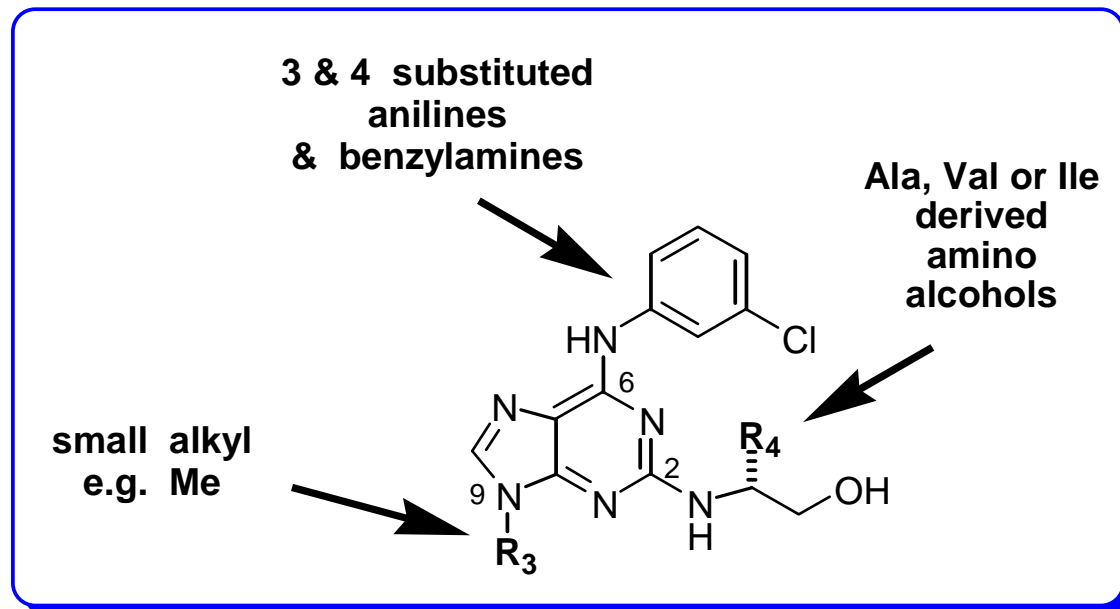
spatially separated
solution library

Screen each position
for inhibition of
h-CDK2-cyclin A kinase complex



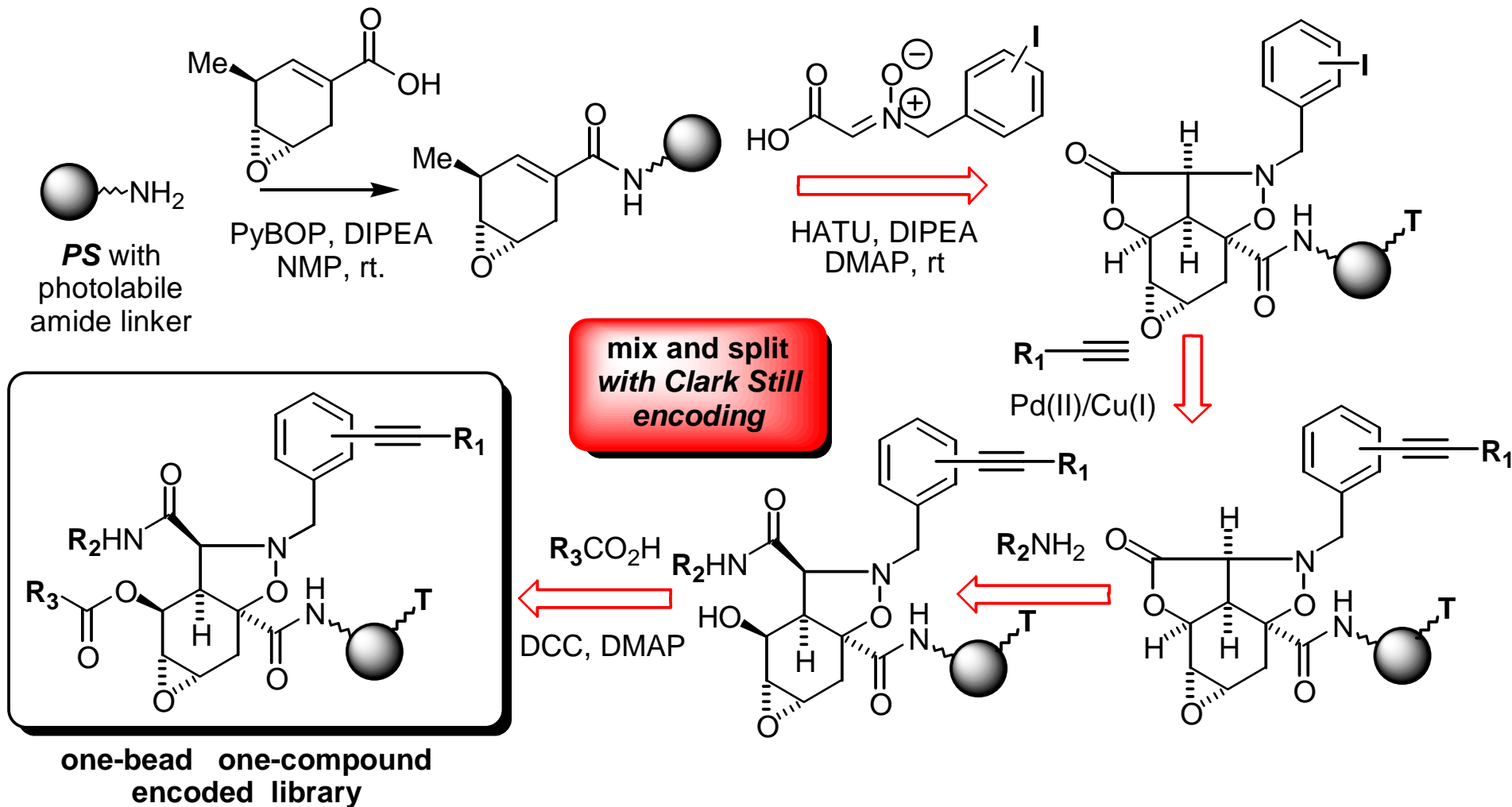
Hits

identity defined by
spatial location



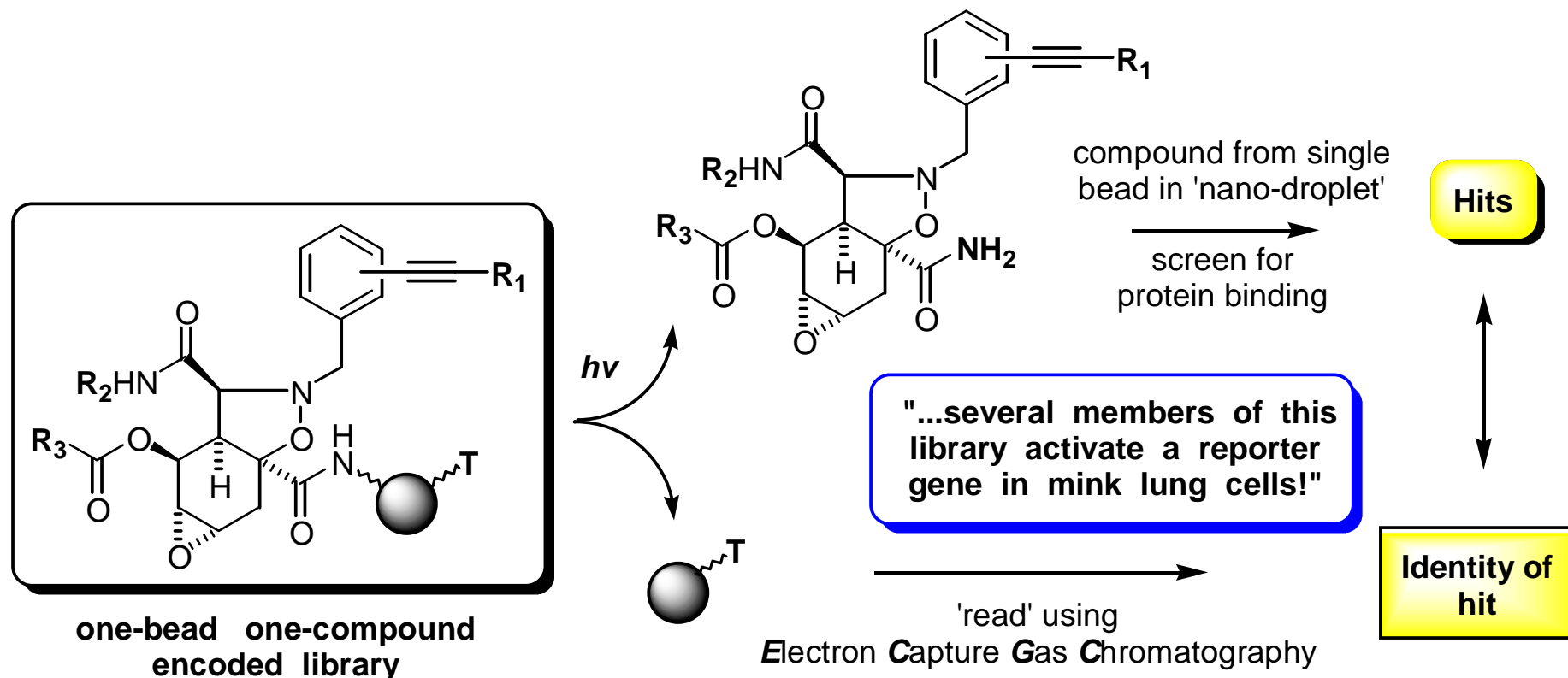
Schreiber's 'natural product' library

- Protein epitope binding library (template based).
- Schreiber *J. Am. Chem. Soc.* **1998**, *120*, 8565.
- **Library synthesis:**



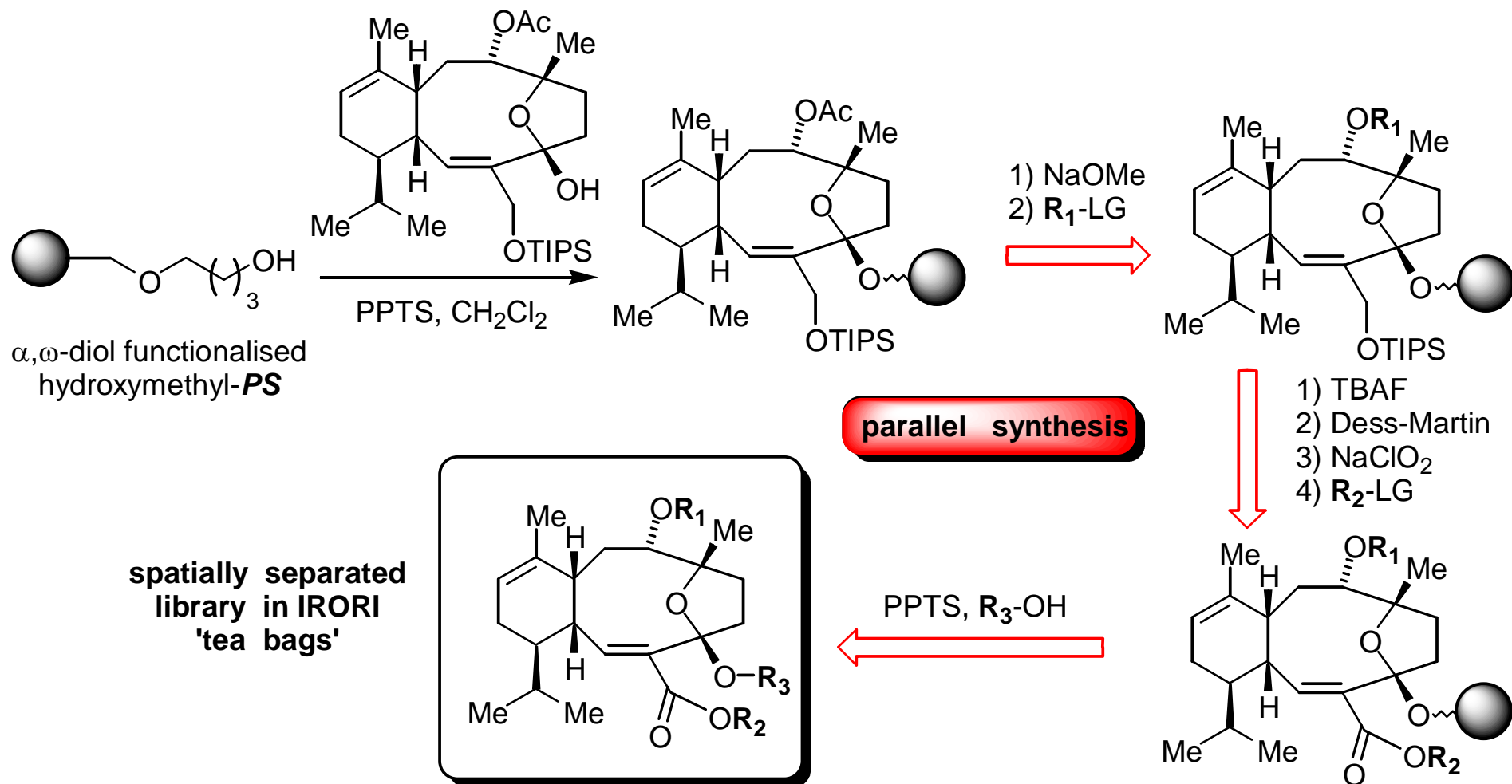
Schreiber's 'natural product' library

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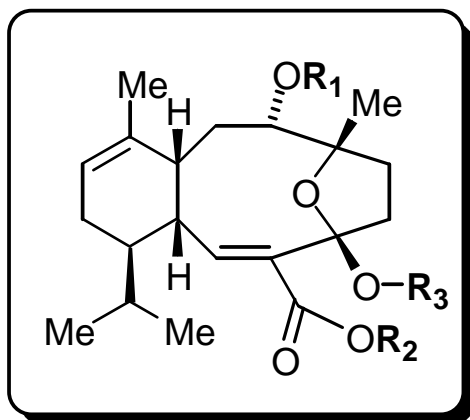
Nicolaou's sarcodictyin library

- Tubulin-microtubule disruptant library (template based).
- Nicolaou *J. Am. Chem. Soc.* **1998**, *120*, 10814.
- **Library synthesis:**



Nicolaou's sarcodictyin library

- Tubulin-microtubule disruptant library (template based).
- Nicolaou *J. Am. Chem. Soc.* **1998**, *120*, 10814.
- **Library screening:**

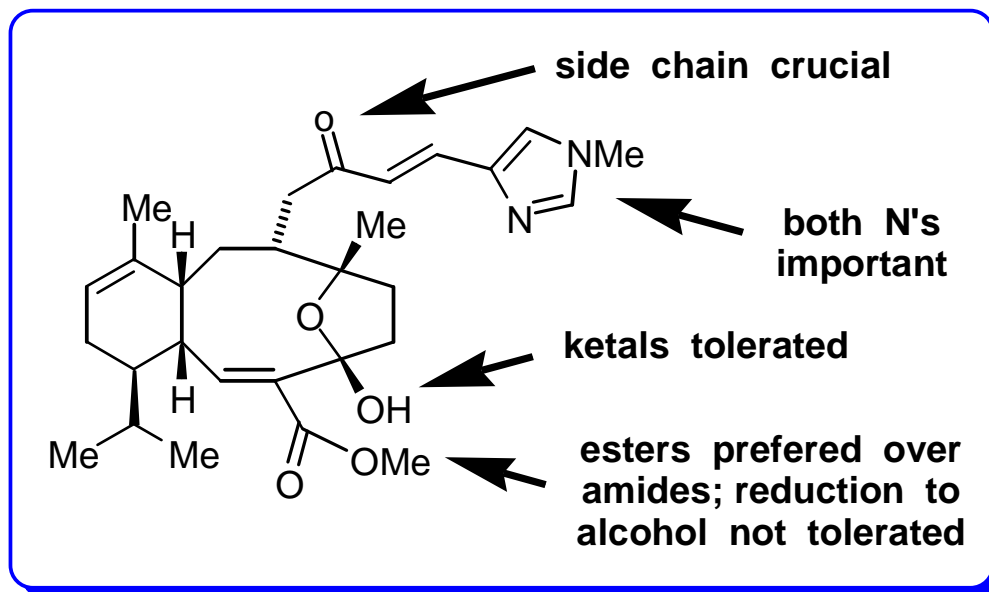


spatially separated
library in IRORI
'tea bags'

screen for induction of tubulin polymerisation
and
cytotoxicity with ovarian cancer cells

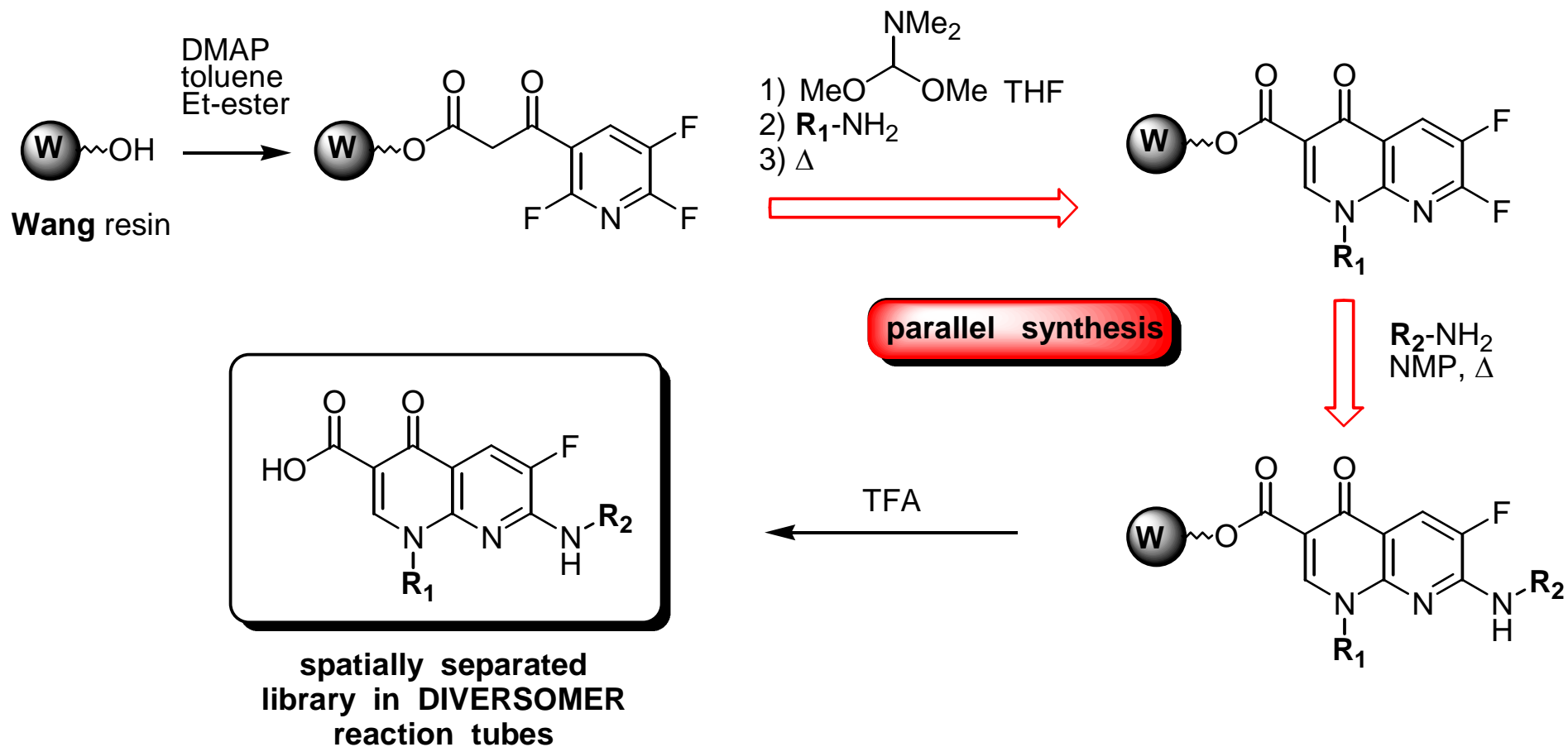
Hits

identity defined by
spatial location



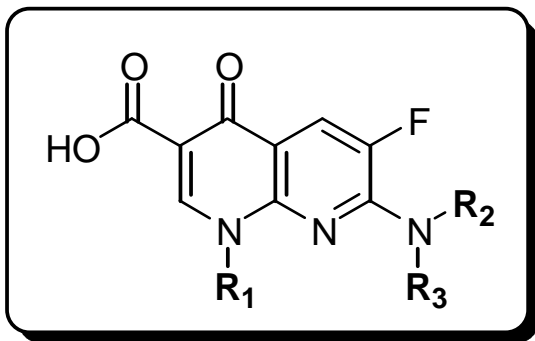
DeWitt's quinolone library

- Ciprofloxazin analogue library (template based).
- DeWitt *Tet. Lett.* **1996**, 37, 48115, and patent: WO 94/08711, **1994**.
- **Library synthesis:**



DeWitt's quinolone library

- Ciprofloxacin analogue library (template based).
- DeWitt *Tet. Lett.* **1996**, 37, 48115, and patent: WO 94/08711, **1994**.
- **Library screening:**

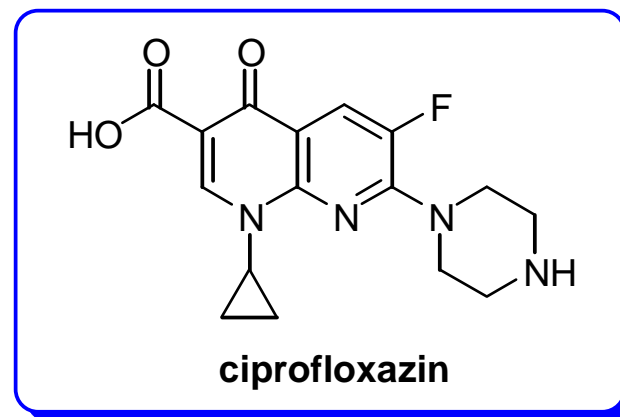


spatially separated
library in DIVERSOMER
reaction tubes

Screen for gyrase inhibition in solution

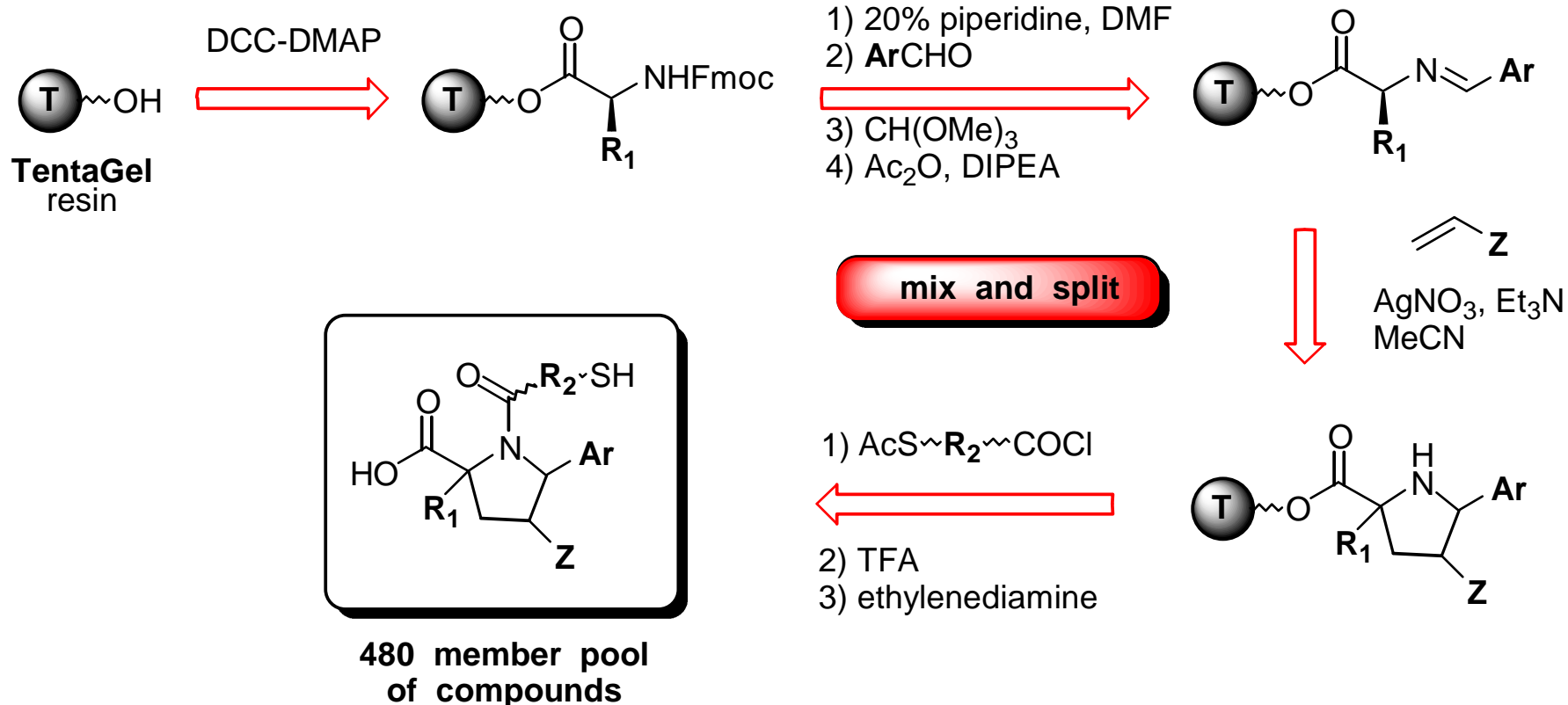


identity defined by
spatial location



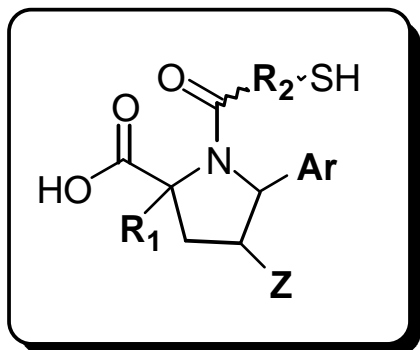
Gallop's mercaptoacyl proline library

- Angiotensin **C**onverting **E**nzyme (ACE) inhibitor library (template based).
- Gallop *J. Am. Chem. Soc.* **1995**, *117*, 7029.
- **Library synthesis:**



Gallop's mercaptoacyl proline library

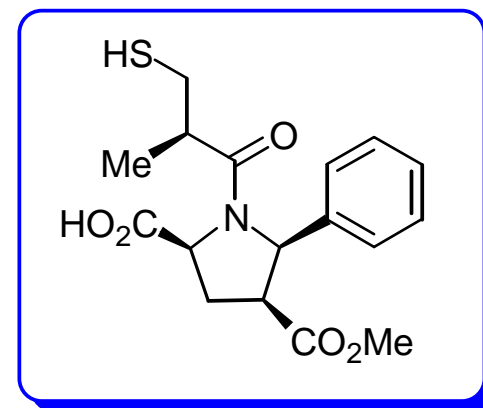
- Angiotensin **C**onverting **E**nzyme (ACE) inhibitor library (template based).
- Gallop *J. Am. Chem. Soc.* **1995**, *117*, 7029.
- **Library screening:**



480 member pool
of compounds

Screened for
ACE inhibition

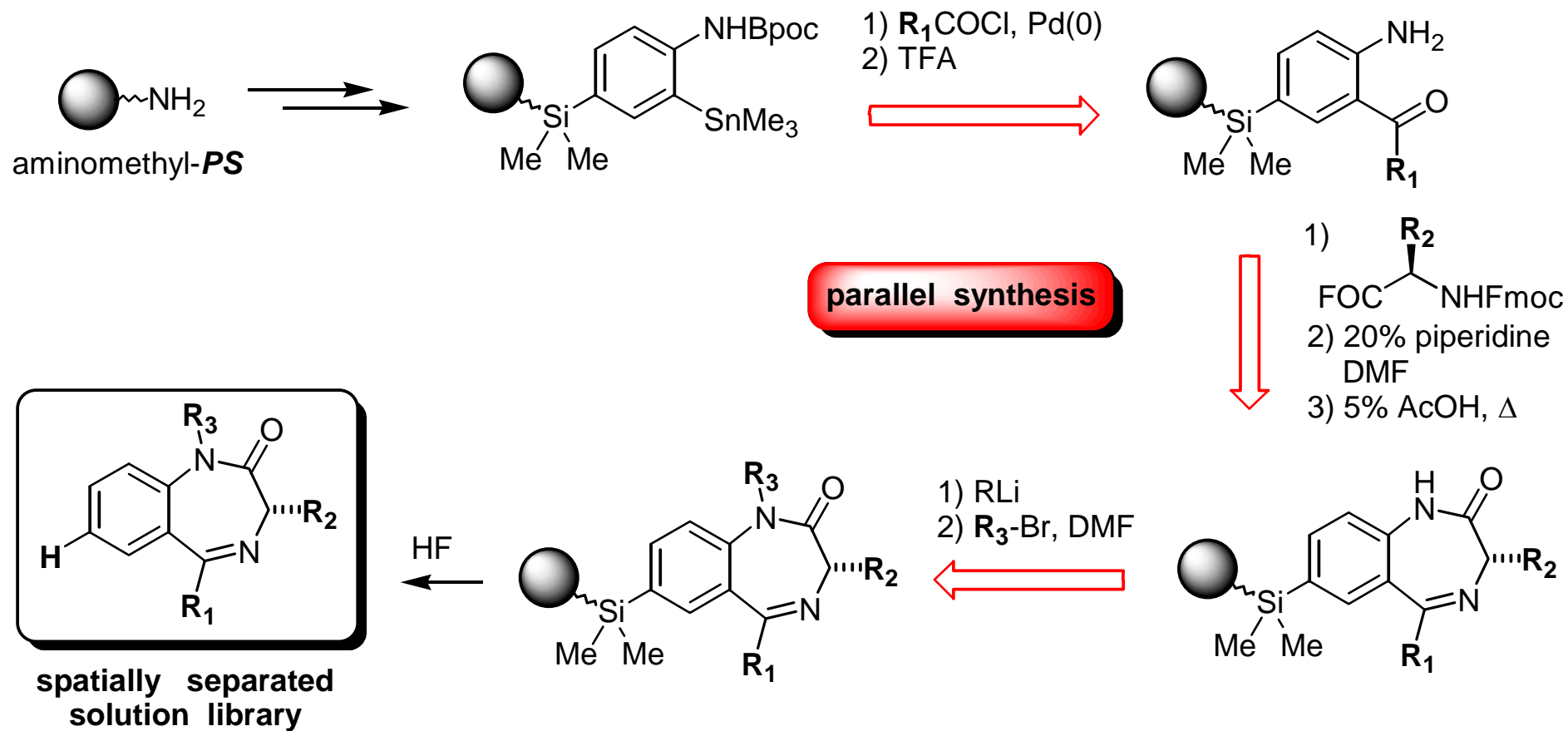
----->
DECONVOLUTION
VIA FOUR ITERATIONS OF
SUB-LIBRARY RE-SYNTHESIS
AND SCREENING



K_i ~ 160pM
3 x more potent than captopril

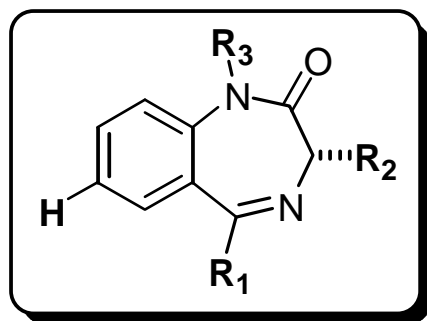
Ellman's benzodiazepine library

- Benzodiazepine library (template based).
- Ellman *J. Org. Chem.* **1997**, *62*, 2885.
- **Library synthesis:**



Ellman's benzodiazepine library

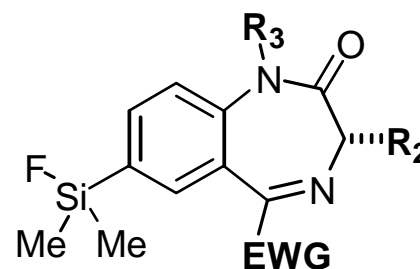
- Benzodiazepine library (template based).
- Ellman *J. Org. Chem.* **1997**, *62*, 2885.
- **Library screening:**



spatially separated
solution library



screening of this library was not reported because it pertained that some of the compounds in the library still contained silicon due to an anomolous cleavage mechanism which was particularly troublesome when R₁ was an electron withdrawing substituent



Summary

- What is combinatorial chemistry?
- The drug discovery process
- Approaches to combinatorial library synthesis:
 - mix and split synthesis
 - parallel synthesis
 - encoded tagging
- Library types:
 - oligomeric libraries
 - template based libraries
- Combinatorial drug discovery.