Principles of Combinatorial Chemistry

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1. Basic ideas & concepts

Basic idea of combinatorial chemistry:

- Preparation of a large number of different compounds at the same time
- ➤ High throughput- screening provides the most promising substances

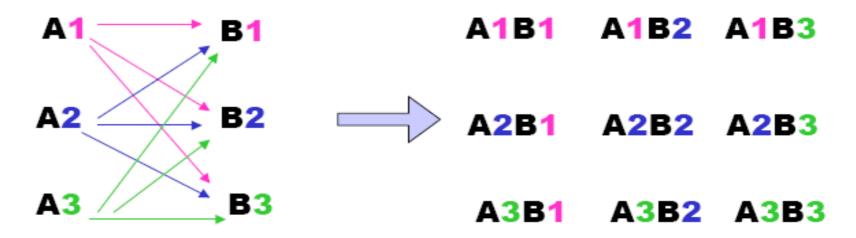
Conventional Reaction: $A + B \longrightarrow A-B$

Combinatorial Chemistry: $A(1-n) + B(1-n) \longrightarrow A(1-n) - B(1-n)$

Combinatorial Chemistry as a valuable tool in drug discovery and material science

Combinatorial Libraries:

- ➤ Def.: Collection of finally synthesized compounds
- Size: depends on the number of <u>building blocks</u> used per reaction and the number of reaction steps, in which a new building block is introduced
- ➤ Typical: 10² up to 10⁵ compounds



9 different compounds

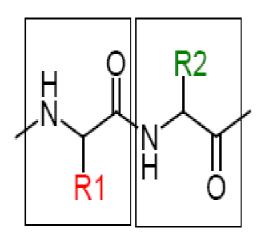
Distinction: Scaffold- based vs. Backbone- based libraries

1. Scaffold- based libraries:

<u>Definition:</u> Core- structure, which all compounds of the library have in common

 Scaffold can consist of several single building blocks (here: Aminoacid and Aminobenzophenone)

2. Backbone- based libraries:



Building block A Building Block B

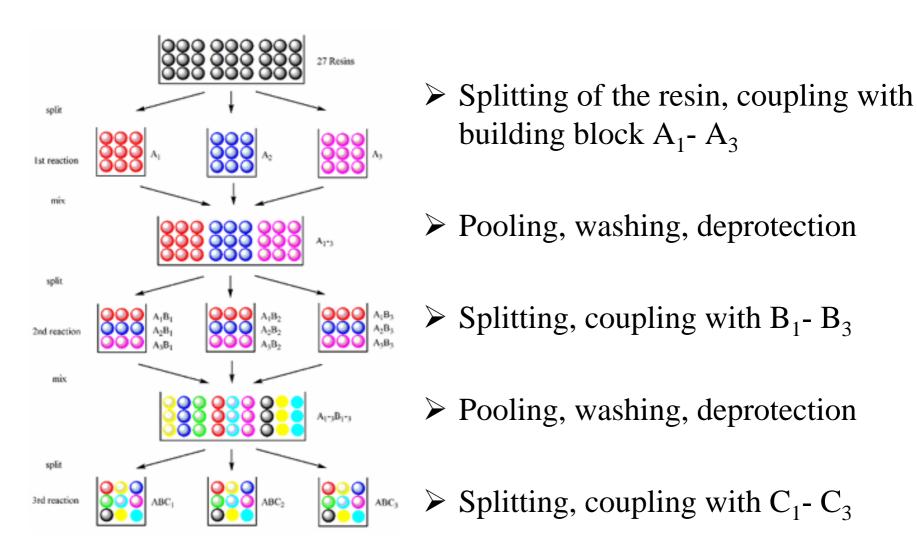
Further Examples: Nucleic Acids, Carbohydrates ➤ Main function: Lead identification & lead optimization e.g. in the drug discovery process

➤ Different approaches to generate new libraries:

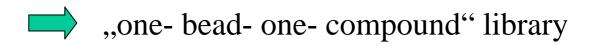
- "Random libraries"
- "Focused Libraries"

2. Synthetic methods & techniques

Split- Pool- Synthesis



➤ After a Split- Pool- synthesis: just one single compound is bound to each resin bead



> Split- Pool- Procedure requires a solid support

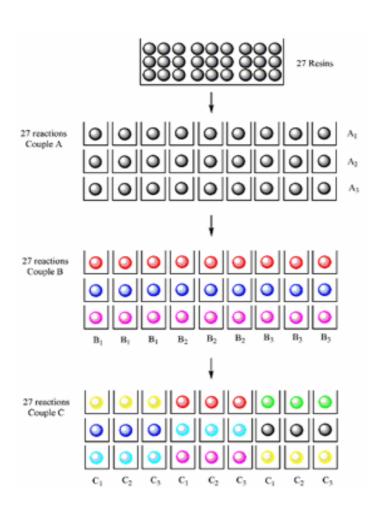
Advantages:

- Only few reaction vessels required
- Method of choice for large libraries (up to 10⁵ compounds)

Disadvantages:

- Threefold amount of resin beads necessary
- Only little amounts of the synthesized compounds available

Parallel Synthesis



- \triangleright Coupling with building block A_1 A_3 (1/3 of the resin beads for each building block), then washing, deprotection
- \triangleright Coupling with building block B_1 B_3 (1/3 of the resin beads for each building block), then washing, deprotection
- \triangleright Coupling with building block C_1 C_3 (1/3 of the resin beads for each building block), then washing, deprotection

- ➤ Concept: Compounds are synthesized in parallel using spatially separated compartments
- > ,,One vessel one compound"- philosophy
- > Solid supported as well as solution chemistry is possible

Advantages:

- Each compound is substantially ,,pure" in its location
- Defined location provides the structure of a certain compound
- Easier biological evaluation

Disadvantage:

• Applicable only for medium libraries (several thousand compounds)

Techniques:

> Conventional Technique: Using resin beads

Reaction vessels: 96 well Microtiterplate (MTP) or an array of chromatography tubes etc.

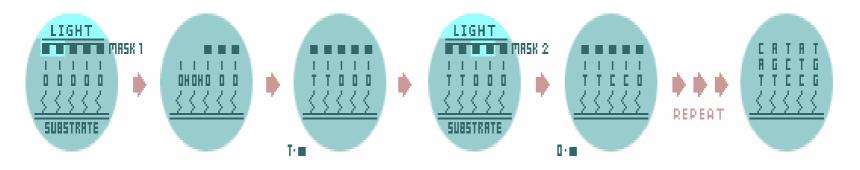
> Multipin-Technique:

Array of polimeric pins serves as solid support.

Polymer is functionalised with a variety of linkers



➤ Light- directed synthesis on silica wafers (based on semiconductor photolithography):

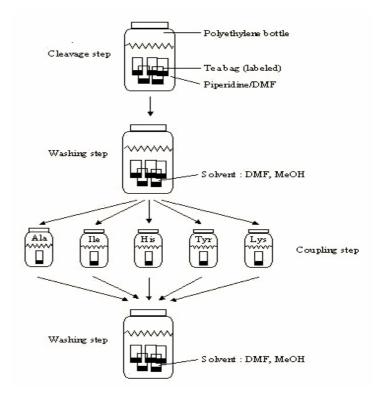


Key Points:

- ➤ Each set of building blocks contains a photolabile protecting group
- ➤ Only the building blocks which have been exposed to light can be coupled with another building block
- Pattern of masks and the sequence of protecting groups define the final structure of the compounds synthesized
- Each member of the library is synthesized at a specific location on a functionalised silica wafer
 - > Libraries: 50000 compounds located in a 50μm square site

Compromise between parallel- and split- pool- synthesis: Tea- Bag- Method (Houghten et al., 1984)

- "Tea Bag": 15 x 22 mm mesh packets filled with resin beads
- One single bead is
 (formally) replaced by a
 "macrobead" (= teabag),
 that contains many small
 ones
- Split- Pool- Protocol occurs batchwise



Advantages:

- •Greater Quantity of each compound is available at once (structural characterisation)
- •Labelling of the tea bags: Easier identification of each compound

Characterisation of combinatorial libraries

➤ Analytical characterisation

➤ Biological evaluation

Analytical characterisation:

Evaluating the success of a combinatorial synthesis by determining the yield and the purity of the compounds

Separate single compounds:

➤ Purification by conventional techniques (e.g. chromatography), determination of the yield by weighing the substances, confirmation of purity by elemental analysis or NMR-spectroscopy

Compound mixtures:

- ➤ Highly sensitive methods are required
 - Mass spectroscopy coupled with HPLC or capillary electrophoresis (CE)

Biological evaluation:

Determining the most (bio-)active substance of a mixture

Bioassays (ELISA, FIA,...)

Methods:

- 1. On bead screening:
- Compounds are still covalently attached to the solid support
- The solid-bound library is treated with a labelled biological target (receptor)
- Selection of the labelled beads (highly automated methods!) followed by structural characterisation
- Requirement: Solid support/ Linkers have to be watersoluble

Advantages:

- Useful for huge libraries (>10⁵ compounds)
- Time saving compared to other methods
- Re-use of the already assayed library

Disadvantage:

• Solid - bound compounds often show different bioactivity compared to free substances

2. <u>Deconvolution</u>

- Preparation of sets of sublibraries (Each of them contains compounds, where the identity of one single building block is known; the building blocks at the remaining positions contain all possible variations)
- Screening of the sublibraries provides the mixture with the highest bioactivity.
- Different proceedings: Iterative deconvolution/ deconvolution by positional scanning

Problems:

- Preparation of the sublibraries may be time-consuming
- The sublibrary with the highest bioactivity determined does not necessarily contain the most bioactive compound (synergistic effect of multiple compounds)

Sources:

- G. Jung: Combinatorial Chemistry, Synthesis, Analysis, Screening, p. 1- 34, Wiley VCH, Weinheim 1999 (Chemistry Library: 86/VK5500J95)
- Pictures: http://www.nvu.edu/classes/ytchang/book/c007.html