

Fourth Editio.

# Medicinal Chemistry

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# 6. REVERSE DESIGNING

Interestingly, **reverse designing** has come into being by virtue of the introduction of *two* extremely important scientific discoveries, namely : (a) **High Throughput Screening**, and (b) **Combinatorial Chemistry**.

## 6.1. High Throughput Screening

Biological testing procedures may be adopted profusely and automated meticulously in an extremely innovated process known widely as **high throughput screening**, that could be able to carry out the normal, rapid, and precise testing scores of newly designed chemical structures all and sundry at a time. It is, however, pertinent to mention here that in several occasions it is absolutely feasible as well as possible to make use of the enormous fruitful advantages of various well-known **gene-cloning methodologies**. In this manner, one has to first and foremost **clone the desired receptor**, and subsequently measure the **binding phenomenon** of the **newly synthesized drug molecules** to the corresponding **cloned receptor**.

## 6.2. Combinatorial Chemistry

Even with the advent of a plethora of such widely accepted and practised methodologies as : Superb statistical methods, traditional synthetic techniques, and time-tested biological screening procedures, invariably prove to be very expensive and sometimes trun out to be non-productive in nature at the end. In fact, the tremendous stress and strain of this sort of cumbersome testing procedures ultimately led to the enterprising technique commonly known as **combinatorial chemistry**. Interestingly, it overwhelmingly makes use of comprehensive and extended libraries of chemical functional moieties which specifically interact either with a '**base molecule**' or with a '**parent molecule**' in a highly systematic small quantum of well-defined purely synthetic stepwise procedures.

Baum and Borman\* (1996) postulated that '**combinatorial chemistry**' refers to a particular sophisticated method of minimizing the effective cost of drug discovery whereby the following *three* cardinal objectives may be accomplished with utmost satisfaction and fruitful results, namely :

(a) to determine altogether 'new leads'

- (b) to find newer 'prototype drug molecules', and
- (c) to refine and optimize the QSAR.

Salient Features : The various *salient features* of 'combinatorial chemistry' are as enumerated under :

(1) **Chemical diversity of products** : Useful libraries of '**reactive**' chemical functional moieties invariably give rise to the **chemical diversity of products** which shall be duly screened for respective biological activity.

(2) Chemistry involved is not only graceful and stylish but also comparatively simple, whereby a few 'same reactions' could suffice in yielding thousands of drug molecules in a specific congeneric series.

<sup>\*</sup>Baum R and Borman S : Chem. Engg. News., 74 : 28, 1996.

(3) Invariably, one makes use of the **solid-state synthetic techniques** to allow the **desired growth of drug molecules** upon **polymer support**.

(4) The '**chemical reactions**' involved in (2) and (3) above must fulfil *three* vital and important criteria, such as : (*i*) clean reaction, (*ii*) reproducible reaction, and (*iii*) high yielding reaction.

(5) '**Robotics**' have been employed profusely to cut down the 'effective cost of synthesis' drastically.

In a rather broader perspective towards the ever increasing and eternal (never-ending) search for newer '**drug molecules**' *i.e.*, chemical entities that essentially requires specific and noteworthy biological characteristic feature do require *two* kinds of approaches, for instance :

(a) Rational Designing : It is considered to be the most 'popular technique' by virtue of the fact that it bears a direct relationship along with a methodical stepwise development of the creative genius and wisdom of a medicinal chemist to explore and exploit the Lock-and Key Model with respect to the ligand-receptor docking.

Rational designing usually encounters the following three obstacles and hinderances, such as :

- (i) limitations\*. e.g., conformational flexibilities for both ligand and receptor,
- (ii) conformers\*\*. i.e., binding with higher-energy conformers.
- (*iii*) active conformers\*\*\*. *i.e.*, influence of salt and water concentration upon the active conformers.

However, all the aforesaid *three* obstacles have been duly taken care of and adequately attended to, with an aim to design a '**drug**' that could mimick vividly the same in an *in vitro* model.

(b) Reverse Designing. Essentially involves the grouping together and searching\*\*\*\* of functionally and structurally identical chemical entities\*\*\*\* by making use of common and biologically effective motif, termed as **pharmacophore**\*\*\*\*\*, which is specifically found either in the **corporate** or **commercial** database. Importantly, at every articulated step carried out meticulously in the **intricate discovery phenomenon** one has to heavily depend upon the manipulative skill(s) related to **CADD**, which provides an extra mileage plus meritorious advantage in **data-processing** into several vital and relevant informations for future analysis in **drug design**.

The elaborated and comprehensive 'Flow Chart' has been dipicted in Figure 3.11 that evidently depicts the *two* above cited processes *viz.*, rational designing and reverse designing together with the latest well-known *in silico* techniques (in box) that are employed very commonly in various processes associated with drug design.

<sup>\*</sup> Jorgensen WL., Science, 254 : 954, 1991.

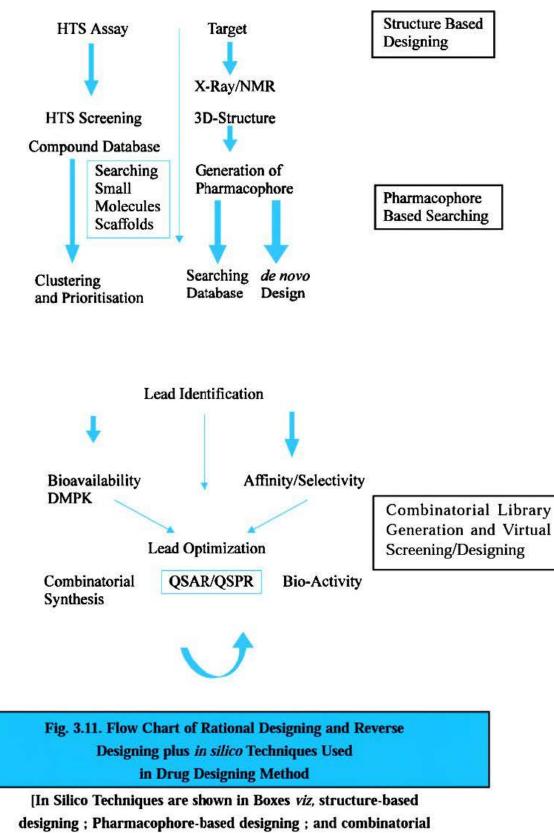
<sup>\*\*</sup>Oshiro CM et al. : J Comput Aided Mol Des., 9 : 113, 1995.

<sup>\*\*\*</sup>Wlodek ST et al. : Protein Sc., 7, 573, 1998.

<sup>\*\*\*\*</sup>Giller VJ and Johnsen AP : In : Designing Bioactive Molecules : Three Dimensional Techniques and Applications, Martin YC and Willen P (Eds.) : AM. Chem. Soc, Washington DC, 1997.

<sup>\*\*\*\*\*</sup>Martin YC, J. Med Chem., 35 : 2145, 1992.

<sup>\*\*\*\*\*\*</sup>Schneider G et al. : Angew Chem Int Ed Engl. 38, 2894, 1999.



### TARGET IDENTIFICATION SCHEME

Library generation and virtual screening/designing]