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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 method-ologies**. 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.

^{*}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.

^{*}Jorgensen WL., Science, 254: 954, 1991.

^{**}Oshiro CM et al.: J Comput Aided Mol Des., 9: 113, 1995.

^{***}Wlodek ST et al.: Protein Sc., 7, 573, 1998.

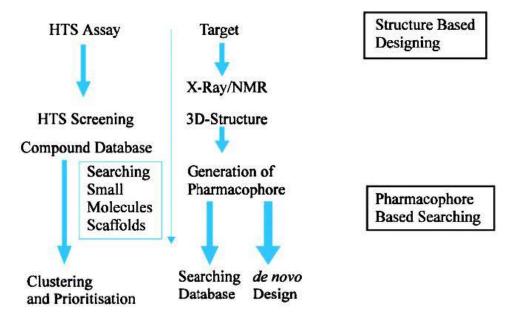
^{****}Giller VJ and Johnsen AP: In: Designing Bioactive Molecules: Three Dimensional Techniques and Applications, Martin YC and Willen P (Eds.): AM. Chem. Soc, Washington DC, 1997.

^{*****}Martin YC, J. Med Chem., 35: 2145, 1992.

^{******}Schneider G et al.: Angew Chem Int Ed Engl. 38, 2894, 1999.

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TARGET IDENTIFICATION SCHEME



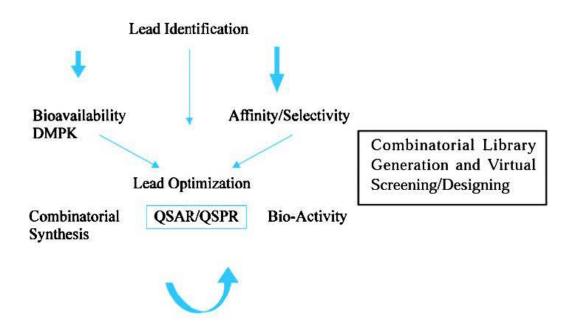


Fig. 3.11. Flow Chart of Rational Designing and Reverse
Designing plus in silico Techniques Used
in Drug Designing Method

[In Silico Techniques are shown in Boxes viz, structure-based designing; Pharmacophore-based designing; and combinatorial Library generation and virtual screening/designing]