Drug Repurposing: An Effective Tool in Modern Drug Discovery

V. S. Kulkarni*^a* **, V. Alagarsamy***a***, 1, V. R. Solomon***^a* **, P. A. Jose***^a* **, and S. Murugesan***^b*

a MNR College of Pharmacy, MNR Nagar, Fasalwadi, Sangareddy, Hyderabad, 502294 India b Department of Pharmacy, BITS Pilani, Pilani Campus, Pilani, 333031 India Received September 6, 2022; revised September 23, 2022; accepted September 25, 2022

Abstract—Drug repurposing is using an existing drug for a new treatment that was not indicated before. It has received immense attention during the COVID-19 pandemic emergency. Drug repurposing has become the need of time to fasten the drug discovery process and find quicker solutions to the over-exerted healthcare scenario and drug needs. Drug repurposing involves identifying the drug, evaluating its efficiency using preclinical models, and proceeding to phase II clinical trials. Identification of the drug candidate can be made through computational and experimental approaches. This approach usually utilizes public databases for drugs. Data from primary and translational research, clinical trials, anecdotal reports regarding off-label uses, and other published human data information available are included. Using artificial intelligence algorithms and other bioinformatics tools, investigators systematically try to identify the interaction between drugs and protein targets. It can be combined with genetic data, clinical analysis, structure (molecular docking), pathways, signatures, targets, phenotypes, binding assays, and artificial intelligence to get an optimum outcome in repurposing. This article describes the strategies involved in drug repurposing and enlists a series of repurposed drugs and their indications.

Keywords: drug repurposing, clinical trials, molecular docking, drug discovery, post-market safety **DOI:** 10.1134/S1068162023020139

INTRODUCTION STAGES OF DRUG REPURPOSING IMPORTANCE OF DRUG REPURPOSING AN IDEAL CANDIDATE FOR REPURPOSING STRATEGIES FOR DRUG REPURPOSING Phenotypic Screening Target-Based Methods Knowledge-Based Methods Signature-Based Methods Pathway- or Network-Based Methods Targeted Mechanism-Based Methods Molecular Docking Application of Drug repurposing in Drug Discovery Challenges **CONCLUSIONS REFERENCES**

INTRODUCTION

Drug repurposing is the technique of using an existing drug or drug candidate for a new treatment or medical condition for which it was not indicated before [1]. It was initially developed to treat a different

medical condition. It has been described as a serendipitous process that happens unexpectedly. In this process, the undesired side effects of drug molecules can also be a pointer to exploring the possibility of its effectiveness in an entirely different medical condition [2]. Usually, drugs with established safety in humans and tested and developed for efficacy in a particular disease other than the one for which they were developed [3]. This process brings the drugs directly to preclinical and clinical trials, skipping the drug development process, and thus reducing risk and costs [4].

STAGES OF DRUG REPURPOSING

The stages of repurposing are elucidated in Fig. 1 [17].

IMPORTANCE OF DRUG REPURPOSING

Repurposing can identify new compounds based on phenotypic benefits without explicitly defining the mechanism of action. This can be directly tested in preclinical animal models, and these results are more applicable to clinical applications and research. It may progress directly straight to Phase II clinical trials [5]. There is a minimum risk of failure with repurposed drugs [6]. The difference between traditional drug discovery and drug repurposing is described in Table 1.

¹ Corresponding author: e-mail: profvalagarsamy@gmail.com, drvalagarsamy@gmail.com.

Fig. 1. Stages of Repurposing.

AN IDEAL CANDIDATE FOR REPURPOSING

A drug has undergone clinical drug development and has been marketed as an ideal candidate for repurposing. A drug with well-established safety and toxicity studies in previous clinical trials, approved by the regulatory authorities, can skip clinical trials with sufficient data support and justification. The mechanism of action of the selected drug shall be established [3].

Those drugs that have gone through several stages of clinical development and have been unsuccessful for reasons other than safety are ideal candidates for repurposing. Some drug repurposing has occurred during clinical trials, like the well-known Viagra

Table 1. The main difference between Traditional Drug Discovery System and Drug Repurposing [6–13]

TRADITIONAL DRUG DISCOVERY	DRUG REPURPOSING
Include 5 stages:	Include 4 stages:
• Discovery and preclinical	• Compound identification
• Safety review	• Compound acquisition
• Clinical research	• Development
\cdot FDA review	• FDA post-market safety monitoring
• FDA post-market safety monitoring	
Generally, more time consuming	Less time consuming
High investment or cost	Lesser investment compared to traditional drug discovery
More risk of failure	Less risk of failure
Clinical efficacy and safety profile should be evaluated	Clinical efficacy and safety profiles already exist

Repurposing drugs have a significant advantage in decreasing the development cost and time to market over standard discovery. Data like pharmacokinetics, toxicology, and safety data from the standard discovery process [14–16].

Table 3. Methods of Repurposing [4]

Repurposing methods require various skills and knowledge of various factors, that is, it may be drug-oriented or disease-oriented, or treatment-oriented [4]

(sildenafil) indicated initially for hypertension and angina. The new role of treating erectile dysfunction was unravelled during the clinical trials.

There are examples of abandoned drugs with their toxicity resurfaced with different indications. Thalidomide, indicated for vomiting, was used to treat nausea in pregnant women and resulted in several congenital disabilities. Due to this tragic effect on fetal development, its use was banned or restricted in several countries. Later, the drug was repurposed for leprosy and multiple myeloma [18]. However, this type of repurposing has met objections from the scientific community [19]. The benefit-to–to-risk ratio can be considered during repurposing cases. A rational decision from the regulatory authorities is essential to address the concern "Is it worth the risk? Or can an existing therapy perform better than the repurposed drug?".

STRATEGIES FOR DRUG REPURPOSING

This approach usually utilizes public databases for drugs. Data from primary and translational research, clinical trials, anecdotal reports regarding off-label uses, and other published human data information available are included. Using artificial intelligence algorithms and other bioinformatics tools, investiga-

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tors systematically try to identify the interaction between drugs and protein targets. In-silico drug repositioning is a powerful technology with significant advantages, including speed and reduced costs [20]. There are usually three kinds of approaches; computational approach, biological experimental approach, and mixed approach, which are described in Table 2 [17].

Traditional phenotype-based screening methods do not need prior knowledge, and the repositioned drugs are just serendipitously tested. The integrated knowledge and elucidated drug action mechanisms increase with the complexity of modelling methods [4]. These methods are listed in Table 3.

Phenotypic Screening

The phenotypic screening method was used to discover molecules and biologics approved by the regulatory bodies. These methods do not include pharmaceutical or biological information and therefore are less likely to help elucidate any mechanisms of action of drugs. Most depend on serendipitous identification from tests aimed at specific diseases and medicines. The advantage of these methods (off-label use and phenotypic screening) is that they have a high chance of application to many drugs or conditions [4, 21].

Target-Based Methods

This method requires specific knowledge about the targets, such as 3D protein structures. Knowledgebased methods require knowledge about drugs or diseases, such as adverse effects, regulatory approval labels, records of clinical trials, and published disease biomarkers (potential targets) or disease pathways). These methods enable researchers to quickly screen any number of drug molecules with a known chemical structure (e.g., Simplified Molecular Input Line-Entry System SMILES). Target-based drug repositioning methods include.

• In-vitro and in vivo high-throughput and/or high-content screening (HTS/HCS) of drugs for a protein or biomarker of interest and

• In-silico screening of drugs or compounds from drug libraries, such as ligand-based screening or docking.

Compared to blinded methods, targeted-based methods improve the chances of drug discovery as targets are directly linked to the disease mechanism. Integrating target information and drug repurposing increases the possibility of finding therapeutically beneficial compounds [4, 21, 22].

Knowledge-Based Methods

These methods apply bioinformatics or cheminformatics approaches to include the available information of drugs, drug-target networks, chemical structures of targets and drugs, clinical trial information, FDA approval labels, signalling or metabolic pathways, and so on, into drug-repositioning studies. The information content of blinded and target-based methods may not be sufficient to identify new mechanisms beyond the known targets. Knowledge-based methods incorporate known information into predicting unknown mechanisms, such as novel drug targets, obscure drug-drug similarities, and new disease biomarkers. Knowledge-based methods include muchknown information into the drug repositioning process to improve prediction accuracy [4, 21].

Signature-Based Methods

Signature-based drug-repurposing methods use gene signatures derived from disease omics data with or without treatment to discover unknown off-target or disease mechanisms. Such genomics data can be assessed through publicly available databases, such as NCBI-GEO (http://www.ncbi.nlm.nih.gov/geo/), SRA Sequence Read Archive (http://www.ncbi. nlm.nih.gov/Traces/sra/), CMAP Connectivity Map and CCLE Cancer Cell Line Encyclopedia. Signature-based methods help uncover unknown mechanisms of action of molecules and drugs. Using computational approaches includes molecular-level mechanisms, such as significantly changed genes [4, 21].

Pathway- or Network-Based Methods

This method utilizes genetic disease data, available signalling or metabolic pathways, and protein interaction networks to reconstruct disease-specific pathways, thereby identifying the key target for repurposing drugs. They help narrow general signalling networks from a large number of proteins down to a specific network with a few proteins (or targets). These methods use pathway analysis or network biology methods to discover essential pathways from diseases' genetic, genomic, proteomic, and metabolic data to find new targets for repositioned drugs. Example: Signalling mechanisms of metastatic subtypes of breast cancer because the subtype signalling mechanisms are hard to elucidate from existing breast cancer pathways or the gene signatures [4, 21].

Targeted Mechanism-Based Methods

These methods integrate treatment omics (genetic) data, available signalling pathway information, and protein interaction networks to delineate the unknown mechanisms of the action of drugs. It aims to discover drug action mechanisms by identifying off-target or targeted pathways of treated drugs using drug omics data (before and after drug treatment). For example, drug resistance is an issue in cancer therapy. Although patients initially respond well to a drug, they often acquire resistance to that drug after a few months of treatment. Hence, successful drug treatment needs additional information about the mechanisms of action of drugs to find better drug targets. However, there are fewer studies on targeted mechanism-based methods that developed elegant computational models to predict the drug effects and related targeted pathways. This is because of the difficulties in deriving effective computational models [4, 21].

Molecular Docking

Molecular docking is a versatile tool used to predict the geometry and to score the interaction of a protein in a complex with a small-molecule ligand. Docking can be performed by docking a known drug into a large set of different target structures or a database of approved medications into one intended target. Molecular docking is a convenient and fast method to screen large libraries of ligands and targets, with a full range of sampling options [23]. 3D structures of the target shall be available through crystallography, nuclear magnetic resonance (NMR), or comparative models to carry out docking. Drawbacks and limitations include approximate scoring function and imperfect binding mode placement algorithms. However, these problems can be overcome by postprocessing docking results with more accurate scoring functions and other criteria [24].

Table 4. (Contd.)

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Application of Drug Repurposing in Drug Discovery

Besides developing new treatment options (such as immunotherapy and host-directed therapies), scientists worldwide are working to repurpose existing drugs against SARS-CoV-2 [27]. Table 2 gives a detailed list of repurposed drugs, and a few are repurposed for the

treatment of COVID-19. Another example is metformin, an antidiabetic drug that shows anticancer effects by decreasing the incidence of different cancers and inhibiting the proliferation and migration of cancer cells, activating apoptosis, and reducing EMT (epithelial-mesenchymal transition) and metastasis. Repurposing helps overcome antibiotic resistance. For

example, TB strains resistant to currently used drug combinations are found in all parts of the world. The product antibiotic, pyridomycin, discovered in the 1950s, is repurposed to treat TB, which takes the place of isoniazid [25, 26]. Table 4 contains a detailed list of repurposed drugs with their initial indication and repurposed use.

Challenges

Although screening efforts are relatively inexpensive, approved drug clinical trials are expensive. The advantage of drug repurposing is that the early stages of clinical development are complete; hence, the drugs can proceed to clinical studies. However, doing clinical studies directly without preclinical studies is a risk. Inaccurate identification of drugs may prove to have no significant impact on therapy or mortality rates, and it may result in a loss in terms of treatment and expense.

Identifying a new therapeutic indication for an existing drug is a significant challenge. Choosing the right therapeutic area for the drug under investigation, evaluating the clinical trials with respect to the new therapeutic use, and deciding which stage of the clinical study or preclinical study shall be restarted are a few challenges repurposing. New preclinical and clinical trials may be required to be carried out if the available data are not satisfactory and do not comply with the requirements of regulatory agencies.

Another critical issue is patent application and intellectual property rights (IPR). Patents or IPR can prevent some repurposed drugs from entering the market, and IP protection for drug repurposing is minimal. Hence, the regulatory constraints and the risk of abandoning repurposing projects due to unsatisfactory results can discourage the investment of money or resources towards drug repurposing. A spike in market demand can be an excellent motivator to shift researchers and investors into action.

CONCLUSIONS

There are numerous diseases for which good therapeutic options have not been developed. The concept of repurposing a drug enables exploring the hidden potential of many molecules and better utilization of therapeutic agents. For better drug repositioning, more in-depth understanding along with integrated approaches between computational and experimental methods may be required to ensure high success rates of repositioned drugs.

COMPLIANCE WITH ETHICAL STANDARDS

The authors declare that they have no conflicts of interest.

This article does not contain any studies involving human participants performed by any of the authors and does not contain any studies involving animals performed by any of the authors.

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