Chemoinformatics – predicting the physicochemical properties of 'drug-like' molecules James F Blake

A few major advances have occurred in the area of physicochemical modeling of organic compounds during the past several years, spurred on by changes in the pharmaceutical industry. Recent advances include the ability to categorize and screen the overall physicochemical properties of potential drug candidates based entirely on their molecular structures and the ability to model the components that contribute to the oral absorption characteristics of potential drug candidates.

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Current Opinion in Biotechnology 2000, 11:104-107

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Abbreviations

NPSA	non-polar surface area
Pc	permeability coefficient
PSA	polar surface area

Introduction

It has been estimated that roughly 10% of the compounds that enter development eventually become marketed drugs and 40% of compounds fail due to poor pharmacokinetic properties [1]. The ability to predict so called ADME (absorption, distribution, metabolism, and excretion) properties from molecular structure would have a tremendous impact on the drug discovery process both in terms of cost and the amount of time required to bring a new compound to market. This review covers computational approaches to the prediction of various physicochemical properties of complex organic molecules from their molecular structures without the need of any experimentally derived parameters. The environment for the application of these computational approaches is the drug discovery and preclinical drug development settings. Specifically, we are interested in the prediction of physicochemical properties that play a critical role in determining the oral absorption characteristics of therapeutic agents, namely, aqueous thermodynamic solubility and permeability across biological membranes. Dressman et al. [2] have related this dependence to the absorption potential (AP) of a compound via:

$$AP = \log \left(P \cdot F_{\text{non}} \cdot S_{0} \cdot V_{L} \cdot X_{0}^{-1} \right)$$
(1)

Where *P* is the octanol-water partition coefficient, F_{non} is the fraction in nonionized form at pH 6.5, S_0 is the intrinsic solubility, V_L is the volume of the lumenal contents, and X_0 is the dose administered. Johnson *et al.* [3] formulated the maximum absorbable dose (MAD) of a compound as follows:

$$MAD = K_a \cdot S \cdot SIWV \cdot SITT \tag{2}$$

Where K_a is related to rat intestinal perfusion rate, S is the aqueous solubility at pH 6.5, and SIWV and SITT are the small intestine (SI) water volume and residency time, respectively. Both equations (1) and (2) describe the quantity of drug that could be absorbed under ideal conditions. The fundamental importance of $\log P$ (as a measure of lipophilicity/hydrophobicity) in the distribution of compounds in various biological systems and a number of computational methods for the estimation of $\log P$ from the molecular structure have been surveyed by Buchwald and Bodor [4] and will not be discussed herein. From equation (2), it is clear that maximum absorption occurs for highly soluble compounds with high permeability characteristics. It is also clear that in order to provide guidance to medicinal chemists, one must address the relative contributions of both solubility and permeability in absorption modeling. Additionally, one must be able to make reasonably accurate predictions for complex molecules before they are synthesized in order to reduce discovery time, which excludes the use of any experimentally derived parameters. Curatolo [5•] has recently reviewed the relationship between the physical chemical parameters of drug candidates and how they affect oral delivery. Readers interested in the complex relationships among the various ADME parameters and how they influence the overall human pharmacokinetic characteristics are referred to a review by Obach et al. [6]. The ultimate goal of modeling studies is the ability to predict the oral absorption properties of a compound based entirely on its molecular structure. This is a very challenging problem and the ability to predict the individual components of equation (2) would also be of great value.

Prediction of oral absorption

We begin our review with computational methods that address the overall oral absorption characteristics of potential drug candidates. The goal is to develop models and rules that will allow medicinal chemists to design compounds, or combinatorial libraries, that are heavily biased in favor of good oral absorption characteristics. In order to accomplish this, one must first define, or categorize, compounds that possess the desired properties. To this end, Lipinski et al. [7] were the first to conduct a systematic comparison of the physicochemical properties of compounds that are orally absorbed versus those compounds not expected to be well absorbed. The latter set were chosen to increase the size of the data set for analysis (i.e. there are not very many data points on poorly absorbed compounds). From a survey of compounds that entered Phase II clinical studies, Lipinski et al. found that poor absorption or permeability was probable for a compound when the molecular weight was above 500 amu (atomic mass unit), the calculated logP (ClogP; [8]) was

above five, and there are more than five hydrogen bond donors (sum of OH and NH groups), or ten hydrogen bond acceptors (sum of N and O atoms) present. If any two of these properties or features are present, the compound is likely to experience poor oral absorption characteristics. This is the so-called 'rule of 5'. The terms it employs are straightforward and are easily implemented by the practicing medicinal chemist. This work is important because it demonstrates that simple rules can be employed to categorize the properties of more favored compounds, and has proved very useful in the design and selection of compounds from virtual libraries. Of the top 100 best selling drugs in 1998 (74 are administered orally), only four fell outside the 'rule of 5' (CA Lipinski, personal communication). It should be mentioned that the 'rule of 5' addresses passive absorption and does not consider whether a compound might be involved in active transport mechanisms [9]. Clarithromycin is an example of a compound that has been shown to be actively transported but also violates the 'rule of 5'.

The 'rule of 5' provides an alert for potential absorption or permeation difficulties, but does not provide a quantitative estimate of a compound's absorption profile. The first true quantitative structure-property relationship (QSPR) study on a large (86 compounds) and diverse set of drug and 'drug-like' compounds was conducted by Wessel and coworkers [10**]. In this study, the measured human intestinal absorption (HIA) was correlated with six descriptors computed from the three-dimensional molecular structure (three charged partial surface area [CPSA] descriptors [11] and three topological and geometric descriptors). The CPSA descriptors are related to the physicochemical properties of molecules and have been successfully used in a variety of property prediction applications [10**,11-15]. In their study, a nonlinear model was used to fit %HIA with root-mean-square errors of 9.4% for the training set and 16.0% for an external prediction set. Given the diversity of the molecules under study and the complex nature of oral absorption, these results are very encouraging. As mentioned above, absorption depends on both the permeability and solubility of a compound; focused studies on each of these properties are described in the sections that follow.

Prediction of permeability

As it is difficult to measure intestinal permeability *in vivo* in humans, a number of models have been developed to provide a measure of the local absorption rate. The effectiveness of using Caco-2 monolayers [16–18], which are derived from a human cell line, in the modeling of human intestinal permeability has been reviewed recently by Lennernäs [19]. From an experimental standpoint, Pade and Stavchansky [20] have demonstrated that permeability coefficients derived from Caco-2 cells coupled with solubility measurements are reasonable predictors of the fraction of drug absorbed. Accordingly, there have been a number of recent efforts aimed at predicting the apparent

permeability coefficient (P_c) from Caco-2 membrane permeability studies. While log*P* has been employed in a number of studies involving biological membranes (as a measure of lipophilicity/hydrophobicity of a compound) [21], another popular quantity known as the polar surface area (PSA) of a molecule has been used in a variety of applications [22,23^{••},24–26]. The PSA is generally defined as the sum of the van der Waals (or solvent-accessible) surface areas of oxygen and nitrogen atoms including attached hydrogens [22,23^{••}]. The PSA of a compound can be related to its hydrogen bond accepting and donating ability.

van de Waterbeemd et al. [22], Palm et al. [23**], and Krarup et al. [24] have all incorporated PSA calculations in various forms to correlate the P_c (or log P_c) from Caco-2 studies with the PSA of a molecule, after screening a number of possible descriptors. In each of these studies, good correlations were generally obtained when the PSA was used alone or in conjunction with a limited number of descriptors. For example, Palm *et al.* [23^{••}] demonstrated that for a series of nine β receptor antagonists, the Caco-2 $\log P_c$ were highly correlated (cross-validation correlation coefficient q, $q^2 = 0.98$) to the dynamic PSA (i.e. multiple conformations of the compounds were averaged in the PSA calculations). Krarup et al. [24] also used dynamic averaging (via molecular dynamics simulations) of the PSA defined by the solvent-accessible surface to predict Caco-2 permeabilities for a series of six β -receptor antagonists and five ester prodrugs (correlation coefficient r, $r^2 = 0.98$). Both groups of researchers have noted that a balance between the nonpolar surface area (NPSA) and PSA may be important in permeability modeling. The incorporation of contributions from the NPSA was investigated by Stenberg et al. [26] for a series of 19 oligopeptide derivatives using Boltzmann weighted contributions from the PSA and NPSA fit to a sigmoidal function. For this system, the equation developed from the surface area terms outperformed similar equations that employed experimentally derived parameters, such as octanol-water (P), heptane-water, iso-octane-water, and heptane-ethylene glycol partition coefficients. For all of these studies, increasing PSA diminished membrane permeability. No distinction was made between hydrogen bond donating or accepting PSA contributions, which may warrant further studies.

In all of these cases, however, relatively small (~20 compound) data sets of closely related structural analogs were used. As is the case with all regression models, the predictive capabilities for compounds outside the training sets is not known, and should be investigated further to look for more generally applicable models. Although predicting P_c is important, Palm has suggested that PSA calculations could be used as a screening tool to weed out compounds with poor absorption properties, in a similar way to the 'rule of 5' [27]. From a study of 20 relatively diverse compounds, they concluded that structures with a PSA \geq 140 Å² would be poorly absorbed (< 10% fractional absorption), whereas those compounds with a PSA \leq 60 Å² would be well

absorbed (>90%). This was recently investigated further by Clark [25], wherein the data set from Wessel *et al.* [13] (74 drug and drug-like compounds) was used to demonstrate that compounds with a PSA \geq 140 Å² were indeed found to be poorly absorbed. The results of this study compare favorably with the predictions from 'rule of 5' alerts, and offers chemists an additional screening tool.

Prediction of aqueous solubility

Though aqueous solubility has been extensively studied, computational methods for the estimation of this highly important property are just beginning to demonstrate predictive capabilities for complex molecules. The difficulty lies in the fact that aqueous solubility is not an intrinsic property of the molecular structure per se, as the crystallographic environment of the solid must be considered as well. Aqueous solubility can be greatly affected by crystal polymorphism. For a recent review of the factors involved in crystal packing effects, see Dunitz and Gavezzotti [28]. We will restrict our discussion to those methods that have been applied to the prediction of aqueous solubility of relatively complex molecules that exist as solids at room temperature, without the need to incorporate experimentally derived quantities, such as the melting point [29•]. For smaller data sets with relatively simple molecules, a number of researchers have shown that reasonably accurate predictions of $\log S_{\rm w}$ (log of the aqueous solubility expressed as mol dm⁻³) are possible [12,15,30,31]. For a series of hydrocarbons and halogenated hydrocarbons, Huibers and Katrizky [31] were able to predict $\log S_{w}$ with r = 0.98, and standard error s = 0.4 from a three term equation based on volume, surface area, and topological descriptors. Similarly, Mitchell and Jurs [12] determined a nine-parameter model based on topological, CPSA, and electronic descriptors with an root-mean-square error of 0.394 log units for a collection of relatively simple organic compounds. More recent publications have looked at larger and more drug-like data sets.

Huuskonen et al. [32] assembled a data set of 211 drugs and related analogs for their solubility prediction experiments. By screening 101 descriptors, they were able to build an artificial neural network (ANN) model based on electrotopological state indices (E-state) [33]. Using a training set of 160 compounds and a test set of 51 compounds, they found 14 input parameters gave $r^2 = 0.86$ with s = 0.53 for $\log S_{\rm w}$ of the test set, and $r^2 = 0.9$ with s = 0.46 for the training set. Although this result is encouraging, care must be taken when using any topological indices in predictive studies because compounds or functional groups that are not well represented in the training set can lead to poorly predictive models for compounds outside the training set. A more general approach would be based on properties computed from the structures, such as the solvatochromic parameters of Abraham et al. In their most recent study, Abraham and Le [34.] determined a seven parameter equation with $r^2 = 0.92$ and s = 0.56 for a 659 compound training set. From the equation, Abraham

finds that increasing hydrogen bonding ability always leads to an increase in solubility, as does the polarizability, whereas an increase in volume leads to decreasing solubility. Interestingly, Abraham and Le concludes that for simple systems, standard errors as low as 0.3 log units are attainable, whereas more complicated drug-like data sets will have errors of ~0.5 log units.

Conclusions

Over the past several years, there has been a tremendous shift toward emphasis on optimization of ADME properties early in the life of drug discovery programs. This shift has been fueled to a large extent by an increase in the number of compounds with poor solubility and absorption properties entering development [7]. Research to date has provided a good conceptual framework for the prediction of properties of interest, namely, solubility and permeability, in connection with the overall goal of predicting human absorption [7,10^{••},23^{••},34^{••}]. Future research will probably focus on developing models with data sets that are larger and built around more diverse collections of compounds with a wide range of chemical functionalitites. Two strategies are likely to emerge in the area of physicochemical property prediction: those seeking to develop general rules in order to screen large numbers of compounds, and those attempting to provide increasing levels of accuracy for more diverse compounds.

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