

Uniting Cheminformatics and Chemical Theory to Predict the Intrinsic Aqueous Solubility of Crystalline Druglike Molecules



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Introduction

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- \geq Poor aqueous solubility remains a major cause of attrition in the drug development process.
- \geq In previous work, energy terms from a computed thermodynamic cycle had been used as descriptors in a multi-linear regression model for intrinsic solubility. Accuracy much better than from direct computation and comparable to leading informatics approaches was achieved. > In recent work, we have shown that accurate first principles calculation is now becoming possible – provided that both the crystalline and solution phases are described by accurate theoretical models.^[1] > Sophisticated machine learning techniques have been applied to many problems in the chemical sciences.

Methods

- \succ The free energy terms are partitioned into physically meaningful terms. The relevant solute – solute, solute – solvent and solvent – solvent enthalpy and entropy terms are computed.
- > DMACRYS is used to calculate the crystal lattice energies. Gaussian 09 (G09) is used to calculate the gaseous and solution phase energies.
- > The SMD solvation model was used; gas and solution phases were calculated at the HF/6-31G(d,p) and M062X/6-31G(d,p) levels of theory.^[1] \geq To understand the quantitative results, we have performed machine leaning methods such as: * Partial Least Square Regression (PLS): $Y = XB + \varepsilon$ where $B = [(X^T X)]^{(-1)} X^T Y$
 - Random Forest Regression (RF): ensemble of many decision trees.
 - Support Vector Regression (SVR): $f(x, \omega) = \sum_{i=1}^{m} \omega_i g_i(x) + b$
 - ***** Radial SVR: $e(-\gamma |x \gamma|^2)$

Chya II

Informatics Section

Gaseous

Cheminformatics Theoretical energies Data (Pre-processed)

XXXXXXX

11 C + 222

AS sub

A sup

Machine Learning Methods

Optimising the parameters for different



Results and Discussion

- \blacktriangleright We have used 100 drug-like molecules for this study. Where possible, SMILES were taken from a single source (ChemSpider) due to the variability in the interpretation of non-canonical SMILES strings.
- > The descriptors were calculated from SMILES strings in the Chemistry Development Kit (CDK). These descriptors include Molecular Weight, XLogP, Freely rotatable bonds, number of H-bond acceptors etc.
- For further analysis aimed at improving the accuracy, and to evaluate different descriptor sets, we used various machine learning approaches and compared the performance of such methods.
- Figure 2: boxplots represent the distribution of RMSE of various machine learning methods using 10 fold CV for different sets of descriptor. Here, the red dot represents the average performance (RMSE) of different models.

- Figure 1: represents the correlation between the HF/6-31G(d,p) theoretical calculation, which outperformed the M062X/6-31G(d,p) calculation, and the experimental LogS values.
- \succ Table 1 summarises the results of a linear regression analysis of the theoretical prediction against the experimental results.
- > Our initial results, a linear regression analysis, suggested an error of almost 3 LogS units, too large for a useful model.

importance are ongoing.

- Table 2 (A and B): reports the average over 10 fold CV of RMSE and R squared scores of different models and descriptor sets. These results suggest that RF performed slightly better than other predictive models when fitted with all sets of descriptors.
- ➢ Overall, machine learning methods significantly improved upon the linear M062X and Descriptors Only 30 relationship, suggesting that ÷. the theoretically calculated data could be explained by 25 non-linear QSPR models. **RMSE** 1.20 Our results also suggest that SE 1.10 RM ⁰⁸ the prediction is significantly improved if theoretical ~ energies are combined with cheminformatics descriptors. 1.10 learning > Machine based SVR PLS RF solely on computed energy terms and a study of variable

RMSE 1.10

08

PLS

RMSE PLS	Descriptors	Descriptors	Only
PLS			
	1.093	1.110	1.174
RF	1.086	1.107	1.134
SVR	1.119	1.111	1.132
<u> Table 2(B)</u>			
	M062X &	HF &	Descriptors
R Squared	Descriptors	Descriptors	Only
PLS	0.595	0.594	0.559
RF	0.602	0.583	0.559
SVR	0.575	0.576	0.559
	RF SVR Table 2(B) R Squared PLS RF SVR	RF 1.086 SVR 1.119 Table 2(B) M062X & R Squared Descriptors PLS 0.595 RF 0.602 SVR 0.575	RF 1.086 1.107 SVR 1.119 1.111 Table 2(B) M062X & HF & Descriptors R Squared Descriptors Descriptors PLS 0.595 0.594 RF 0.602 0.583 SVR 0.575 0.576

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References

SVR

RF

structures in DMACRYS.

1. Palmer, D. S.; McDonagh, J. L.; Mitchell, J. B. O.; van Mourik, T.; Fedorov, M. V., Journal of Chemical Theory and Computation, 8, 3322 (2012).