

Compatibility Screening of Drug-Polymer Blends Using Computational Modelling



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Abstract Summary

- Many drugs present solubility challenges which adversely affect bioavailability
- Drug-polymer solid dispersions can improve solubility by maintaining the drug in an amorphous state
- This study investigated computational modelling for use in pre-formulation studies of drug-PVP solid dispersions
- Computational modelling was able to provide insight to temperature dependence of miscibility and to calculate key mixing parameters

Materials and Methods

Software

- Materials Studio (BIOVIA) [3]
- Blends module used for calculations and analysis
- Geometry optimisation in Forcite module
- **COMPASS II force field applied to operations**
- Ultra-fine calculation quality used (10,000,000 energy samples and 100,000 cluster samples with energy bin width of 0.02 kcal/mol)

Structures Studied

- **PVP screened against ibuprofen, carbamazepine and itraconazole**
- Monomer unit and drug structures drawn in Materials Studio before geometry optimisation
- **Degree of polymerisation set to 22 to mimic K-12 PVP**
- Head and tail atoms of PVP monomer unit set as non-contact to simulate full polymer chain

Outputs

- Binding energy distribution
- χ parameter
- Mixing energy
- Free energy of mixing
- Phase diagrams of drug binary mixtures with K-12 PVP

Conclusion

Pros:

- ✓ May save time and resources
- ✓ Ability to vary chain length
- ✓ Results may indicate probability of demixing and crystallisation
- ✓ Provides insight to temperature dependence of miscibility
- ✓ Calculates key mixing parameters
- ✓ Merit as a screening tool when used alongside established characterisation techniques

Cons:

- ✗ Rate of mixing could not be established
- ✗ Predictive power for crystallinity is unclear
- ✗ Unable to take moisture content into account
- ✗ Unable to take time-dependence of miscibility into account
- ✗ Not a reliable alternative to experimental study

References

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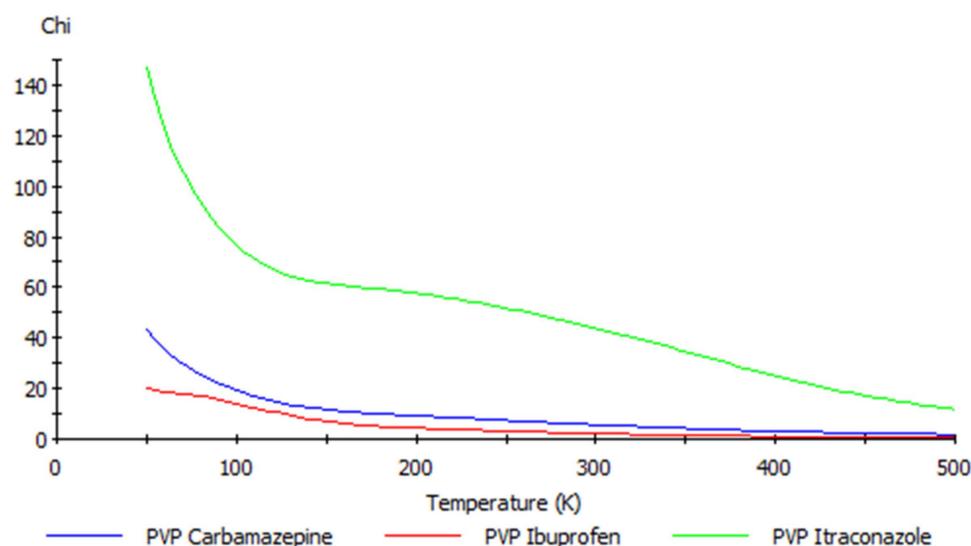
Introduction

Compatibility screening is an essential aspect of pre-formulation study [1]. **Miscibility is a key parameter defining physical stability [2].** This work sought to demonstrate how computational modelling may be used in pre-formulation studies to predict drug-polymer miscibility in solid dispersions. This could serve as a preliminary indicator of stability. Interactions in various binary mixtures were studied, with **PVP used as the model polymer screened against poorly water soluble model drugs.** Findings illustrate the merits and limitations of the method in determining miscibility and physicochemical properties, compared to experimental data and established techniques.

Results and Discussion

Computational modelling showed the model blends would not be stable. **This contradicts experimental findings.** Results may be predictive of long term physical instability. The rate of demixing was not provided, however the **magnitude of the χ parameter may indicate the probability of demixing and crystallisation at a given temperature (Figure 1).** The Blends module was unable to take moisture content into account, or the time-dependence of miscibility. Ability to vary chain length is a distinct advantage of the software, allowing for study of a very broad range of polymer grades. Information on binary mixtures is provided in a fraction of the time taken to obtain experimental data. Computational modelling may save time and resources, but **inaccurate results may mislead** when characterising potential formulations. **Results of this study did not correlate well with experimental data.** Computational modelling can be a useful complementary technique when used alongside, rather than in place of, more established techniques such as DSC, optical microscopy, and X-ray diffraction.

Figure 1 – Temperature dependence of χ parameter



Next Steps

- This research makes a contribution to the evaluation of potential approaches for studying drug and polymer chemistry
- Future work could investigate the influence of force field selection on the results
- COMPASS II was used in this study, but Materials Studio also offers Dreiding and Universal force fields
- A greater range of drugs could be studied, with differing physicochemical properties