# **Molecular Modelling Metered Dose Inhaler Suspension Formulations**



### **Reading Guide:**

Each slide is a different section Zoom into each panel Read along from left to right

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**Engineering and Physical Sciences Research Council** 





### **1. Introduction to Project**



#### **CONTEXT**

- Suspension formulations contain four key components
- Molecular modelling provides the ability to model specific surfaces of a crystalline particle
- Understanding interactions between different materials can potentially aid formulation development



#### **BACKGROUND**

- Molecules are represented as spherical atoms connected by springs
- Atoms are assigned point charges and force fields calculate potential energy
- In crystallography molecular positions are specified relative to their unit cell
- This is used to define planes in the bulk structure



#### **THIS POSTER**

- Phases must be simulated and validated individually before being combined
- This poster shows validation of fluticasone propionate's (FP) solid crystal structure
- Also, validation of liquid propellant HFA-134

### **2. Solid State of Fluticasone Propionate**



#### **METHOD**

- FP's crystal structure data came from the DAXYUX entry1
- Hydrogen positions were optimised and point charges assigned using AM1 method within MOPAC2
- Program habit $98<sup>3</sup>$  calculated interaction energies using generic force fields Dreiding4 and Tripos 5.25



#### **MORPHOLOGY VALIDATION**

- In this model, each face's growth rate was proportional to attachment energy
- Both force fields predicted a morphology that resembled the hexagonal profile of FP crystals from slow evaporation in methanol



#### **SURFACE CHEMISTRY**

- Morphology model shows the chemistry of each face
- (100) face exposes the hydrogen bond accepting O1 atom
- Large cavities between molecules

## **3. Liquid Propellant HFA 134a**



#### **METHOD**

- 463 HFA-134a molecules were simulated in a cubic box using molecular dynamics code DL\_Poly 4.096
- Atomic point charges came from the force fields being tested; OPLS<sup>7</sup> and PCFF<sup>8</sup>
- Volume could change with conditions of constant target temperature and pressure; ranging from 263 to 323 K and 5.6 atm, respectively
- Equilibration lasted 500 ps and sampling 1.2 ns



#### **VALIDATION**

- Thermal expansion was compared to physical values from the Peng-Robinson equation of state
- It showed a difference in density of  $+/-10$  % for the PCFF and OPLS force fields respectively



#### **STRUCTURE OF LIQUID**

- Radial distribution function (RDF) of PCFF simulations show higher peaks at lower temperatures
- It also resembled a previous Monte Carlo simulation<sup>9</sup> which further validates the results

### **4. Conclusion and Future Work 5. References**



- Molecular models of the solid and liquid phase were validated against physical values
- The different chemistry of FP's crystal surfaces was highlighted and structure of liquid HFA-134a was observed with an RDF plot
- The two phases will be combined to measure the free energy of wetting of different faces
- Then, further work will use this method to look at other materials in formulations

- 1. Cejka, J., Kratochvfl, B., Jegorov, A. & F, K. Crystal structure of fluticasone propionate, C25H31F3O5S. *Z Krist*. 220, 143–144 (2005).
- 2. Stewart JJP. MOPAC: A semiempirical molecular orbital program. *J Comput Aided Mol Des*. 4, 1–103 (1990).
- 3. Clydesdale, G., Roberts, K. J. & Docherty, R. HABIT95 A program for predicting the morphology of molecular crystals as a function of the growth environment. *J. Cryst. Growth* 166, 78–83 (1996).
- 4. Mayo, S. L., Olafson, B. D. & Goddard, W. A. DREIDING: A generic force field for molecular simulations. *J. Phys. Chem.* 94, 8897–8909 (1990).
- 5. Matthew, C., Richard III, D. C. & Nicole Van, O. Validation of the general purpose tripos 5.2 force field. *J. Comput. Chem.* 10, 982–1012 (1989).
- 6. Todorov IT, Smith W, Trachenko K, Dove MT. DL\_POLY\_3: New dimensions in molecular dynamics simulations via massive parallelism. *J Mater Chem.* 16, 1911–8 (2006).
- 7. Jorgensen WL, Maxwell DS, Tirado-Rives J. Development and testing of the OPLS all-atom force field on conformational energetics and properties of organic liquids. *J Am Chem Soc.* 118, 11225–36 (1996).
- 8. Sun H. Ab Initio Calculations and Force Field Development for Computer Simulation of Polysilanes. Macromolecules. 28, 701–12 (1995).
- 9. H. Do, R. J. Wheatley, & J. D. Hirst, Microscopic structure of liquid 1-1-1-2-tetrafluoroethane (R134a) from Monte Carlo simulation. *Physical Chemistry Chemical Physics*. 12, 13266–13272 (2010).

Thanks for viewing. Send questions to: **pm13vb@leeds.ac.uk**