

PXRD INVESTIGATION OF CHANGES IN DEHYDRATION BEHAVIOR OF GSK241572 HYDRATE AFTER MICRONIZATION

Feirong Kang

Physical Properties Group - Pharmaceutical Development GlaxoSmithKline King of Prussia, PA



This document was presented at PPXRD -Pharmaceutical Powder X-ray Diffraction Symposium

Sponsored by The International Centre for Diffraction Data

This presentation is provided by the International Centre for Diffraction Data in cooperation with the authors and presenters of the PPXRD symposia for the express purpose of educating the scientific community.

All copyrights for the presentation are retained by the original authors.

The ICDD has received permission from the authors to post this material on our website and make the material available for viewing. Usage is restricted for the purposes of education and scientific research.



PPXRD Website – <u>www.icdd.com/ppxrd</u>

ICDD Website - www.icdd.com

Outlines

Background and Objectives

Methodologies on Dehydration Analysis

- Thermal analysis
 - DSC/TGA
 - Hot-stage microscopy
- Gravimetric vapour Sorption
 - Isothermal dehydration
 - Kinetics modelling
- PXRD analysis
 - Reflection vs. transmission
 - Variable temperature PXRD
 - Rieveltd refinement

Conclusions

Background

- Many pharmaceuticals exist in both hydrated and anhydrous forms.
- The physicochemical, mechanical, processing, and biological properties of hydrates may differ significantly from those of the corresponding anhydrates.
- Understanding of dehydration behavior is required for process control and predicting stability of drug substance and drug product.
- Variable hydrates present relative complex dehydration behavior compared to stochiometrirc hydrates.

Compound in Study



- Form B, a hydrated form of this free base was chosen for development.
- Theoretical monohydrate 4.6%wt
- Classified as a variable, or nonstoichiometric hydrate
- Displays complex dehydration behavior pre and post size reduction

7-methoxy-1-methyl-5-(4-(trifluoromethyl)phenyl)-[1,2,4]triazolo[4,3-a]quinolin-4amine

(GSK241572)

Objectives

Characterize dehydration of the system pre- and post-micronization

- Various techniques were utilized including DSC, TGA, HSM, GVS, VT-PXRD etc.
- Study dehydration mechanism by the following approaches
 - Solid state kinetics modeling
 - Rietveld analysis
 - Crystal structural analysis

Scanning Electron Microscopy









HV Date Det Mag Spot WD GSK241572 4 KV 10/20/09, 11 53 ETD 77568 x 3 10.2 mm -500 nmDSC/TGA

Pre-micronization

Post-micronization



Distinct dehydration processes observed between non-micronized and micronized samples.

Hot-Stage Microscopy

Pre-micronization Post-micronization



Start recrystallization to needle-like crystals (anhydrous form)

Converted to anhydrous form



Start recrystallization to needle-like crystals (anhydrous form)

Converted to anhydrous form



Isothermal Dehydration



Drastically different behavior displayed with regards to the loss of water content over time. The rate of loss of non-micronized material is slower by ~ two orders of magnitude.

GVS Isotherms



GVS isotherms (25°C) for the interaction of water with micronized and non-micronized FormB.

Kinetic Modeling – Non-micronized

NETZSCH Thermokinetics 241572 unmic dehydration by GVS



Kinetic Modeling – Micronized

NETZSCH Thermokinetics 241572 mic dehydration by GVS



Summary so far....

- DSC, TGA, HSM, and GVS data showed distinct dehydration process between non-micronized and micronized materials.
- Kinetics modeling on dehydration showed
 - The non-micronized material exhibits a two-step dehydration process where a diffusion step is followed by a 2nd order step. The diffusion step is the rate limiting step based on that its dehydration activation energy is ~22 times higher than the 2nd step.
 - The micronized material follows a simple one-step process (2nd order)
- PXRD analysis will be discussed next to rationalize results.

Crystallographic Data

| Moiety formula | $C_{19}H_{15}F_3N_4O\cdot H_2O$ | |
|----------------------|---------------------------------|-----------------------------|
| Crystal system | Triclinic | |
| Temperature | 150(2) K | |
| Space group | P1 | |
| Unit cell dimensions | a = 8.473(6) Å | α= 102.09(4)° |
| | b = 12.542(7) Å | β=102.70(5)° |
| | c = 17.476(9) Å | $\gamma = 94.86(8)^{\circ}$ |
| Volume | 1754.8(18) Å ³ | |
| Z | 4 | |
| Density (calculated) | 1.478 Mg/m ³ | |
| | | |





PXRD



Line intensity change is due to preferred orientation

No baseline increase and peak broadening \rightarrow no phase separated amorphous content or significant degradation of crystallite integrity

Reflection Analysis – Non-Micronized



Reflection Analysis – Non-Micronized



Reflection Analysis – Micronized



Transmission Analysis – Non-Micronized





VT-PXRD – Non-Micronized



Form change observed at 150°C Line shifting (up to 100°C) less than 0.1 °20

VT-PXRD – Micronized



Form change observed at 150°C

VT-PXRD – Micronized



Line shifting 0.1-0.3 °20 to both lower and higher d-spacing directions changes to the unit cell are anisotropic between the hydrated and dehydrated states

Crystal Lattice Change upon Thermal Dehydration

Rietveld Analysis (HighScore Plus)

| | a/Å | b/Å | c/Å | α/\circ | β/° | γ/° | $V/\text{\AA}^3$ | V%change |
|--------------|-------|--------|--------|----------------|--------|-------|------------------|----------|
| mic INT | 8.623 | 12.636 | 17.358 | 102.25 | 103.07 | 94.47 | 1785 | 0.00% |
| mic 50C | 8.714 | 12.658 | 17.138 | 103.14 | 10274 | 94.32 | 1780 | -0.26% |
| mic 70C | 8.750 | 12.667 | 17.036 | 103.98 | 10272 | 93.74 | 1773 | -0.63% |
| mic 90C | 8.781 | 12.678 | 17.004 | 104.25 | 102.87 | 93.52 | 1775 | -0.54% |
| mic 100C | 8.798 | 12.685 | 17.006 | 104.28 | 102.92 | 93.5 | 1779 | -0.31% |
| | | | | | | | | |
| non-mic INT | 8.508 | 12.658 | 17.346 | 101.93 | 102.46 | 95.34 | 1766 | 0.00% |
| non-mic 50C | 8.552 | 12.745 | 17.281 | 101.84 | 101.62 | 96.95 | 1776 | 0.57% |
| non-mic 70C | 8.589 | 12.817 | 17.256 | 102.00 | 101.44 | 97.37 | 1793 | 1.50% |
| non-mic 90C | 8.630 | 12.833 | 17.222 | 102.11 | 101.55 | 97.14 | 1800 | 1.90% |
| non-mic 100C | 8.583 | 12.838 | 17.207 | 102.09 | 101.68 | 97.30 | 1787 | 1.19% |

Change in Cell Volume upon Thermal Dehydration



Slip Planes



A. (0 0 1) face (E_{att} = -28.3 kcal/mol), showing lack of egress channels for water

B. (0 1 0) face (E_{att} = -75.4 kcal/mol), C. (1 0 -1) face (E_{att} = -77.6 kcal/mol), both showing egress channels for water







Conclusions

- The drastic change in dehydration behavior of the hydrate after micronization is mainly due to
 - Breakage of water channels through longest dimension (a-axis)
 →shorten water travel length
 - Cleavage of several crystal faces that have low attachment energy
 → create more channels for water to escape
- PXRD analyses provide insight into the change of dehydration behavior.

Acknowledgements

- Glenn Williams
- Fred Vogt
- Rachel Forcino
- Jeff Brum
- Roy Copley