Theory 3

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Freeze-Drying in Practice
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Martin Christ
Osterode am Harz, Germany





Analytical characterization

- Product attributes for designing lyophilization cycles
 - Differential scanning calorimetry: T_q, T_q, T_{eut}
 - Freeze drying microscopy: T_{collapse}
- Solid state characterization after lyophilization
 - Residual moisture (LOD, Karl Fischer, NIR, FMS)
 - Thermodynamic / Solid state (X-ray powder diffraction)
 - Specific surface area (BET)
 - Cake appearance at different levels (visual inspection, SEM, μ-CT)
 - Reconstitution





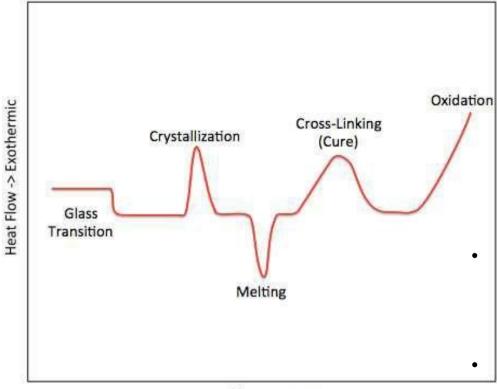
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Differential Scanning Calorimetry (e.g., Tg')

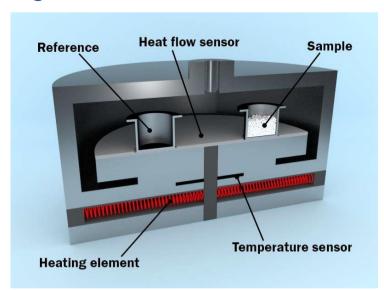


Temperature

Thermal analysis to detect physical transformation such as phase transitions (e.g. glass transition temperature $T_{g'}/T_g$, crystallization/melting point T_{eut} ...)

Measurement of the difference in the amount of heat required to increase the temperature of a sample compared to a reference with well-defined heat capacity as a function of temperature

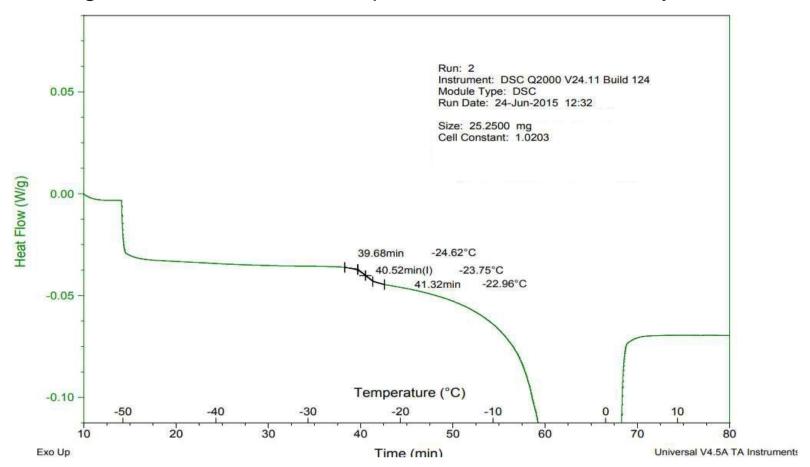
 Both the sample and reference are maintained at nearly the same temperature throughout the experiment





Differential Scanning Calorimetry (e.g., T_{g'})

Tg' = Glass transition temperature of the maximally freeze-concentrated solution

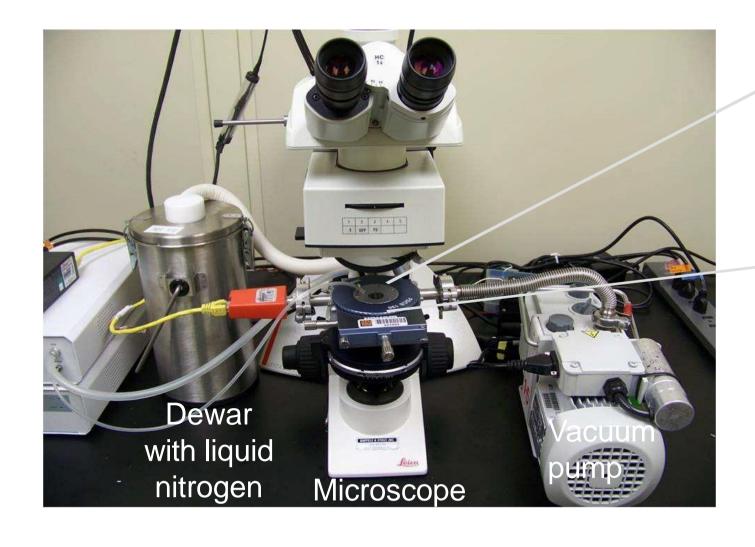


<u>Literature recommendation for design of DSC mesaurements:</u> Pansare SK, Patel SM. Practical Considerations for Determination of Glass Transition Temperature of a Maximally Freeze Concentrated Solution. AAPS PharmSciTech. 2016;17(4):805–19.





Freeze drying microscopy (T_{collapse})





Cryostage





Freeze drying microscopy (T_{collapse})



(Intact) frozen sample



Onset of collapse



Complete collapse

$$\rightarrow$$
 T_g' < T_{collapse} !!

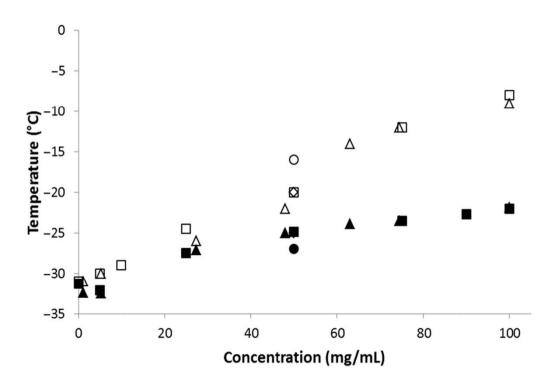
Rule of thumb: $T_{g'}$ ~2 °C lower than T_{c} (low protein conc., see side note)

For visualization: https://www.youtube.com/watch?v=SqM69VQboCl





Side note: differences in T_a' and T_c in dependence of protein concentration



Glass transition temperature of the maximally freeze-concentrated solution and T_c as a function of protein concentration. Values for mAb A (IgG1) are denoted by triangles, mAb B (IgG1) by circles, mAb C (IgG4) by squares, and Pro X (fusion protein, 90kDa) by diamonds. Closed symbols: $T_{g'}$; open symbols: T_{c} .

- The higher the protein concentration, the larger the deviation between T_c and T_d (up to ~14 °C at high protein conc.)
- Thus, Freeze drying microscopy analysis always more accurate than DSC only





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Residual moisture - Water content



balance

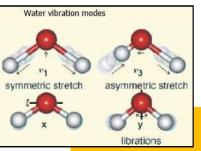
- component
- Destructive
- Loss on drying (LOD) may be challenging for water content low)



Quantitative water Karl-Fischer titration

CH₂OH + SO₂ + RN → (RNH)SO₂CH₂

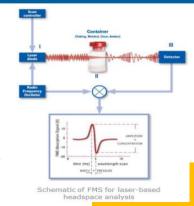
- Destructive
- Volumetric versus coulometric
- Extraction versus direct measurement



spectroscopy

Z R

- near infrared
- Non-destructive
- High throughput (can be
- multivariate calibration principal components and validation with KF-results



Headspace

laser light (1400 nm) and pressure analysis Non-destructive

- High throughput (can be automated)
- Vial format-specific calibration needed

More details in the excellent guest presentation by Derek Duncan

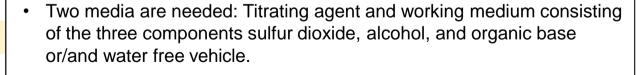
be translated into cake moisture via Karl Fischer

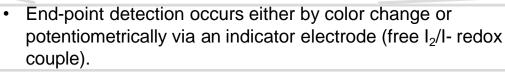
analysis **Gravimetric**



Karl-Fischer Titration

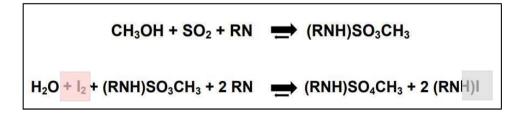
Volumetric Karl Fischer Titration







lodine is added by a burette during titration. Suitable for samples where water is present as a major component: 100 ppm - 100%









Coulometric Karl Fischer Analysis

lodine is generated electrochemically during titration. Suitable for samples where water is present in trace amounts: 1 ppm - 5%

- The working medium consists of the components sulfur dioxide, alcohol, and organic base or/and water free vehicle.
- Two electrodes are needed: One for lodine generation (anode), and one for potentiometric end-point detection via the indicator electrode (free I₂/Iredox couple).





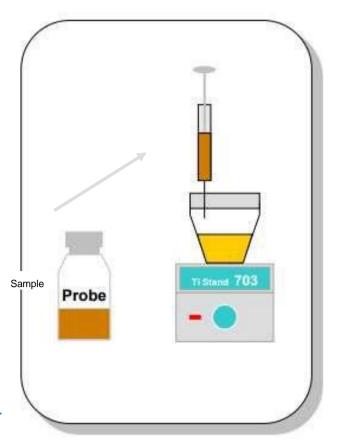


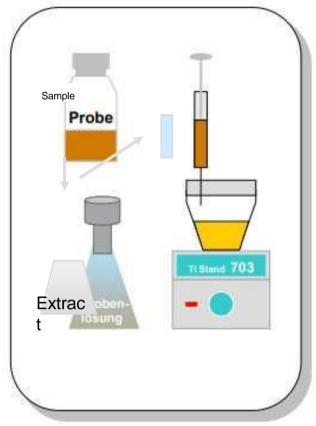
Karl-Fischer Titration

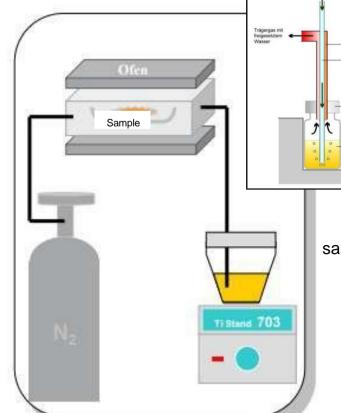
Direct Titration

Liquid Extraction

Evaporation*



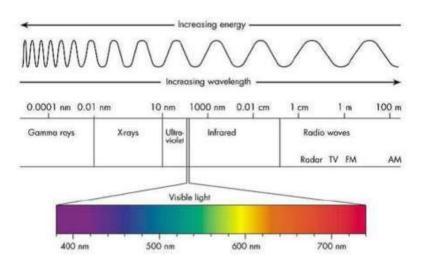


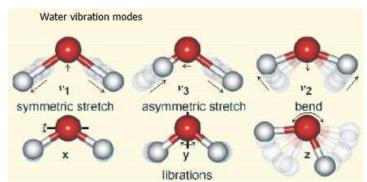


*Highly dependent on the sample and its heat sensitivity.

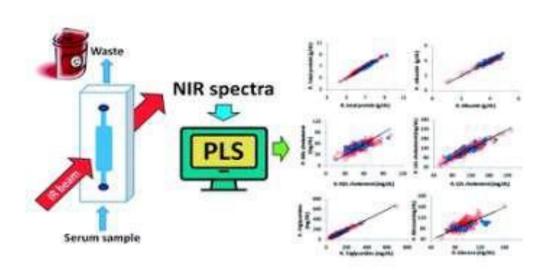


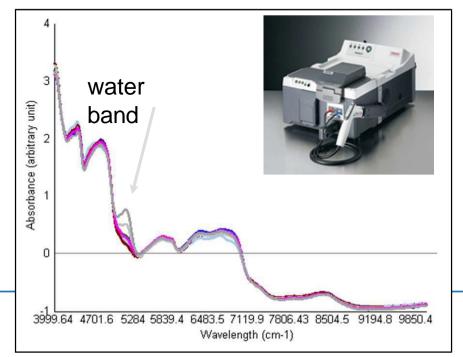
Residual moisture - NIR





- Molecule vibrations (overtone and combinations)
- Near infrared: ~760–2500 nm or 13.000–4.000 cm-1

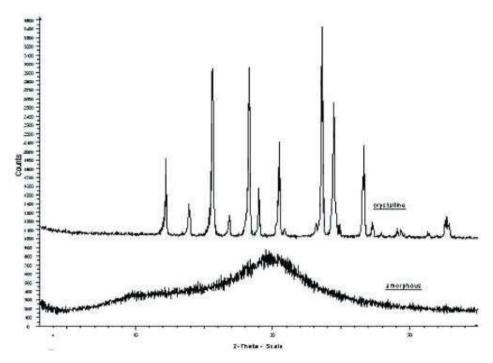




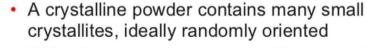




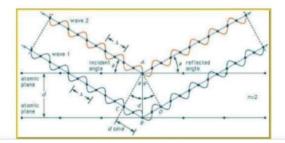
Xray powder diffraction - Morphology



The constructive and destructive interference can be measured as different intensities in the X-ray beam at given angles.



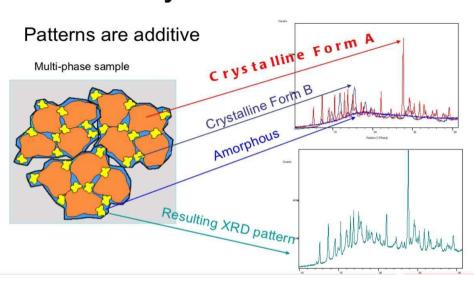
 Diffraction occurs when crystallites are oriented such that specific atomic planes are in the correct relationship with the incoming x-rays



Bragg's law: nλ=2dsinθ

Constructive interference is detected when the path-length difference is equal to an integer number of wavelengths

Mixture analysis





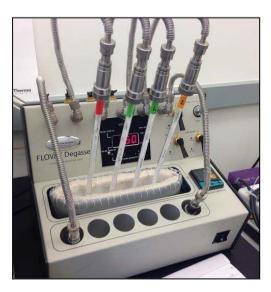


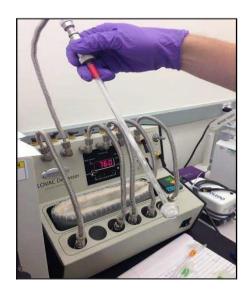
Specific surface area (BET)

S.Brunauer, P.Emmett, E.Teller Adsorption of Gases in Multimolecular Layers, J. Am. Chem. Soc., 1938, 60 (2), pp 309–319









- · Physical adsorption of a gas on the surface of the solid.
- Physical adsorption results from relatively weak forces (van der Waals forces)
 between the adsorbed gas molecules and the adsorbent surface area of the test
 powder. Thus, the determination is usually carried out at the temperature of liquid N2.
- Traditionally nitrogen or helium is used as adsorbate gas.
- Based on the BET theory, the amount of adsorbed gas corresponds to a monomolecular layer on the surface.
- The amount of adsorbed gas is correlated to the total surface area of the particles including accessible pores.









Cake appearance: visual inspection

Reprinted from: "Lyophilized Drug Product Cake Appearance: What Is Acceptable?" Patel S. Nail S. Pikal M. Geidobler R. Winter G. Hawe A. Davagnino J. Rambhatla Gupta S. 2017. J Pharm Sci. 106(7) Copyright [2017] © Elsevier B.V., its licensors, and contributors

Cosmetic defects versus impact on product quality?



Intact cake



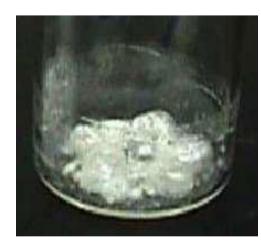
Shrinkage



Light collapse / melt-back



severe collapse / melt-back



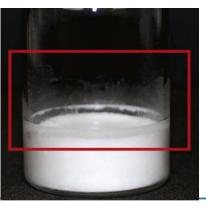
complete collapse collapse/melt-back



crack **COPYRIGHT © PDA**



dents



(minor) splashing

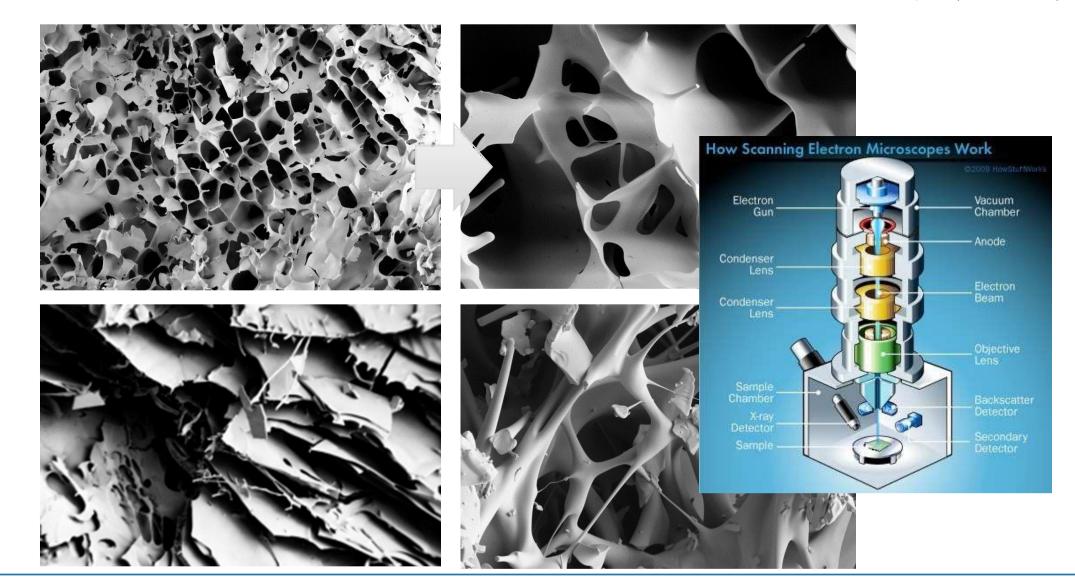


fogging





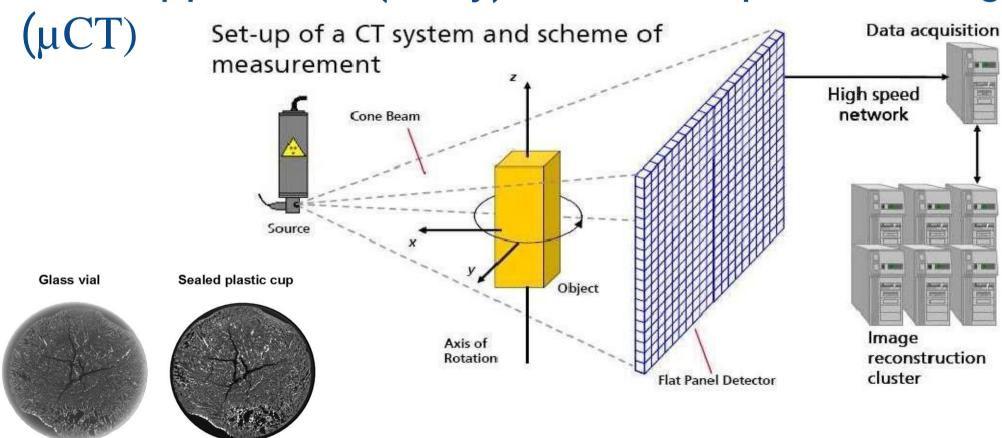
Cake appearance: Scanning electron microscopy (SEM)







Cake apperance: (X-ray) Micro-computed tomography



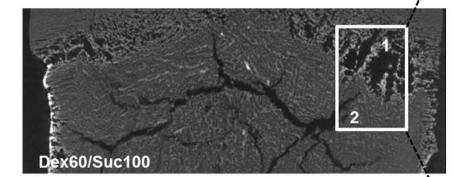
- A micro-focus x-ray source illuminates the object and a planar x-ray detector collects magnified projection images.
- Based on hundreds of angular views acquired while the object rotates, a computer synthesizes a stack of virtual cross section slices through the object.
- You can then scroll through the cross sections, interpolating sections along different planes, to inspect the internal structure.
- Selecting simple or complex volumes of interest, you can measure 3D morphometric parameters and create realistic visual models.



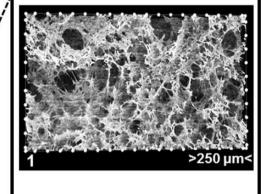


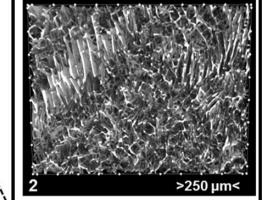
Cake apperance: Micro-computed tomography (µCT)

Dex0/Suc100









Microcollapsed structure

Intact sponge-like structure

Pros and cons and applicability of different imaging techniques summarized in and figures above reprinted from:





Determination of the reconstitution volume

Note: Reconstitution volume ≠ Filling volume (solid content needs to be considered)

Two practical approaches:

- 1. Measuring the loss on drying during freeze-drying (solution density unknown)
 - Weigh selected filled and semi-stoppered vials before and after freeze-drying
 - Mass difference can be accounted to water loss on drying
 - Mass loss can be converted to the reconstitution volume by division by density of water
- 2. Calculate the total amount of water that could be lost on drying (solution density known)
 - Determine the density of your formulated solution to be freeze-dried
 - Calculate the exact total solid content based on composition
 - Calculate the theoretically filled mass (by multiplication with formulation density)
 - Substract the total solid content from the theoretically filled mass
 - Calculated mass difference can be converted to the reconstitution volume by division by density of water





Reconstitution time







- → Water ideally flows along the side wall
- → Avoid foaming if samples contain surfactants
- → In case of long reconstitution times, gently swirling systems may be considered (no shaking!)

