

**Better Nanoparticle Characterization** 

Improved Visualization, Counting and Sizing of Polydisperse Nanoparticle Colloids using ViewSizer<sup>®</sup> 3000

Jan "Kuba" Tatarkiewicz, PhD

**VP Engineering** 

**MANTA Instruments, Inc.** 



- Flow Cytometry (FC)
- Transmission Electron Microscopy (TEM)
- Static Light Scattering (SLS)
- Dynamic Light Scattering (DLS)
- conventional Nanoparticle Tracking Analysis (cNTA)



- Accurate & reproducible measurement of:
  - particle number concentration
  - particle size distribution
  - particle kinetic processes
- and visualization of highly <u>polydispersed</u> colloids

#### **Visualization of Brownian motion**





#### **Problem**





445 nm laser on polystyrene beads in water

#### Six orders of magnitude



- DLS large particles skew results (small ones not detected)
- DLS experimental complications that users overlook (concentration-dependent results)
- cNTA different sized particles can't be seen simultaneously (highly irregular images for large particles, dim for smallest)
- cNTA interrogated volume depends on particles sizes and their refractive indices (similar to FC problem when sizing)



# INTERNATIONALISOSTANDARD19430

## Particle size analysis — Particle tracking analysis (PTA) method

"Sample polydispersity affects the ability to track and therefore analyze different size fractions in the particle number-size distribution. [...] In a polydisperse sample large particles scatter a lot more than small particles making it difficult to detect or track small size particles."

#### MANTA solution US patent 9645070



(Multispectral Advanced Nanoparticle Tracking Analysis)



#### Sample of three color video





#### Size determination (Einstein 1905, Langevin 1908)



• Mean Squared Distance (*MSD* in 2D, *N* frames, jumps of *n*):

$$MSD(n) = \frac{1}{N-n} \sum_{i=1}^{N-n} (x_{i+n} - x_i)^2 + (y_{i+n} - y_i)^2$$

• Diffusion coefficient D (optimized least-square fit of MSD vs. n):

$$MSD(n) = (4 \cdot \Delta t \cdot D) \cdot n$$

• Hence diameter:

$$d = \frac{k_B T}{3\pi\eta D}$$

#### **Light sheet thickness**





#### **Concentration** (counts per volume)



- Observed volume depends on intensity of scattered light
- Calibration of interrogated volume is done using standards (various sizes and refractive indices) → *lookup table*
- Volume factor is calculated from average intensity of scattered light for each tracked particle (*takes laser power, camera exposure & gain into account*)
- Density of particle size distribution (PSD) is calculated with variable volume factor for each size bin

#### **Volume factor** US patent 9857283





### ViewSizer<sup>®</sup> 3000







cuvette w/insert US patent 9541490

#### **Specifications**



Range of particle sizes measured*	10 nm to 15 μm
Minimum sample volume	0.4 mL
Typical sample concentration	5 x 10 <sup>6</sup> to 1 x 10 <sup>8</sup> particles/mL
Sample temperature range (controlled)	10 °C to 50 °C, ± 0.1 °C (-15 °C to 110 °C available)
Dimensions	55 cm W x 66 cm D x 35 cm H
Weight	27 kg
Operational Environment	15 °C to 30 °C with < 85% RH

\*Sample material dependent

#### **NIST exploratory poly-standard**





#### Gold mixes: DLS vs. MANTA







#### TEM, DLS & cNTA vs. MANTA

*α-lactalbumin* nanoparticlesmade as per Arroyo-Maya et al.*J. Dairy Sci.* (2012) **95**, 6204-6214





#### Whole vs. fat-free milk





How to compare two distributions with unknown shapes (no theory)? Use the so called nonparametric tests like **Kolmogorov-Smirnow statistics**:

$D_{A,B}$	alpha	$D_{A,B,\alpha}$	Reject?
0.2335	0.050	0.0338	yes

d<sub>av</sub>=256 nm, SD=145 nm, CV=0.57 d<sub>av</sub>=163 nm, SD=68 nm, CV=0.42

### **Micellisation of a polymer**





If number of particles retained, diameter/volume increases -> growth by adding polymer

#### **Neat proteins**





#### Sample 1

Sample 2

#### **Viscosity of proteins**



7.0e+5 in water 6.0e+5 in protein 1 in protein 2 Density of PSD [counts/mL/nm] 5.0e+5 d<sub>in protein</sub> 4.0e+5  $\eta_{protein} =$  $*\,\eta_{water}$ d<sub>in water</sub> 3.0e+5 2.0e+5 1.0e+5 0.0 0 500 1000 1500 2000 Diameter [nm]

203 nm PSL in water and proteins

#### **Neat proteins PSD**





#### Lysozyme heated to 60 °C





### Fluorescence



Mix of three types of carboxylate fluorescent beads (all nominally 500 nm diameter, stained with Fluoresbrite®)



#### **Vesicles stained with fluorophore**



1,1'-Dioctadecyl-3,3,3',3'-Tetramethylindocarbocyanine-5,5'-Disulfonic Acid



Diameter [nm]



#### **Vesicles processed differently**



Vesicles stained with Dil<sub>c</sub>18(3)



$D_{A,B}$	alpha	$D_{A.B.\alpha}$	Reject?
0.0647	0.050	0.0297	yes

d<sub>av</sub>=278 nm, SD=166 nm, CV=0.60 d<sub>av</sub>=264 nm, SD=169 nm, CV=0.64

#### **Nanoparticles dissolution rate**





#### **Nanoparticles dissolution rate II**





Time [minutes]

#### **Settling rate for large particles**





$$d = \sqrt{\frac{18 * v * \eta}{g * (\rho - \rho_0)}}$$



#### **Protein crystallization rate**





Time [minutes]

Wittbold & Tatarkiewicz 2017, BioProcess International 15 (3)



polystyrene, also with PEG coating, silica, silver, gold, 316L stainless, sand/dirt, clay, CaO, YAG, SiO<sub>2</sub>, carbon, PMMA, LiMnO

sea water, fresh water, rain water, tap water, acetone, wine, urine, blood plasma, milk, ammonia, jet A-1 fuel

small molecule APIs, protein aggregates, silicon oil, protein crystals, liposomes, exosomes, vesicles, micelles, α-lactalbumin, rolled DNA, RNA, viruses, bacteriophages, emulsions, polymeric API carriers, self-adjuvanted proteins



- Individual particle method, not ensemble average
- Accurate density of PSD for polydisperse samples
- Concentration measured, not estimated
- Absolute method (no calibration w/standards needed)
- Particles and processes visualization

#### **Other benefits**



- Kinetic processes (time constants)
- Temperature range and ramp rates
- Real time agitation
- Real time reagent addition
- Multiple tests w/o changing sample

#### **Customers love ViewSizer® 3000**







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Thank you

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