



Protein Characterisation

mosquito

Automated Liquid Handling

- High throughout nanolitre dispensing
- True positive displacement pipetting
- · Genomics, crystallography, screening, assay development





iii ttplabtech



UNcle

Stability and Aggregation Analysis

- T_m & T₂₀₀, isothermal stability, thermal recovery, B₂₂ & k_n
- · Sizing, polydispersity & viscosity
- 48 samples at a time, 9 µL of each







dragonfly

Automated Liquid Handling

- Reliable low volume, non-contact dispensing
- · Rapid assay development, significant reagent & cost savings
- · Broad dynamic range





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Stunner

Only system to combine size & concentration

- · Concentration analysis
- Hydrodynamic size
- Polydispersity & protein aggregation







Trap

Dynamic Protein Interaction

- · Optical tweezers
- **DNA-RNA-protein interactions**
- · Protein folding and conformation











Lunatic

Batch Quantification of Protein, DNA & RNA

- Measure multiple 2µL samples at any concentration
- Samples stable in chip for up to 2hrs
- Actual concentration of DNA/RNA without impurities







XtaLAB Synergy

Structural Biology / Protein Crystallography

- · 4 fold higher flux PhotonJet sources
- Dual Wavelength sources (Mo/Cu/Ag)
- Leading edge HPC Detectors







HUNKY

Quantifying protein stability using ΔG

- ΔG and ΔG_{trend}
- Agg Path



LCHAINED



Big Kahuna

Cutomisable Automated Biologics Formulation

- · Automated buffer exchange in microplates
- Characterise formulations with limited material
- · Developability and stability screens







pUNk

Protein Sizing Platform (DLS)

- · Compact and fast
- Just 5µL sample (recoverable)
- · Protein size, per cent aggregation & polydispersity



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- Target Identification & Validation | cloning & expression | assay development
- Lead Identification | assay development | activity analysis
- Characterisation & Optimisation | biophysical | chemical
- Safety & Preclinical | formulation | stability
- Manufacturing & Regulatory | scale up | optimisation









- · Particle size, count and shape: Microscopy
- Particle identification: Raman spectroscopy and/or LIBS
- · Fully configurable to your requirements





