

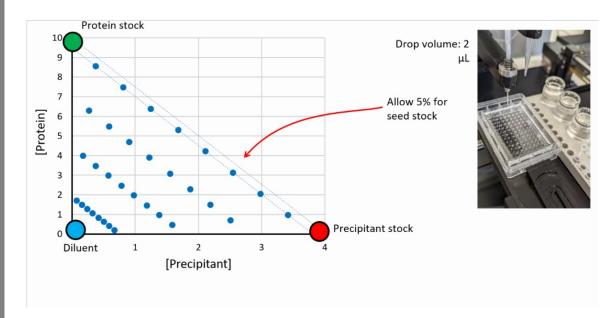
Dear Stefan

Explore your protein's phase diagram with our innovative and powerful new experiment script.

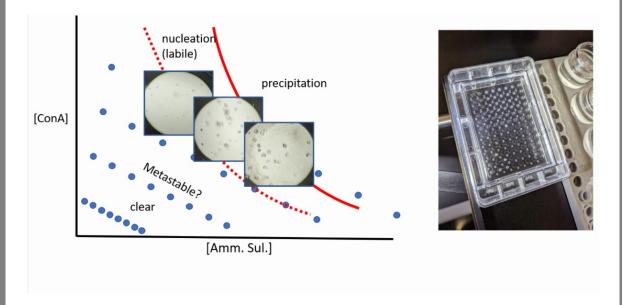
Douglas Instruments recommends a new approach to explore crystallization space systematically. The new experiment design samples a diagonal path across the phase diagram that is parallel to the expected solubility curve. This can give greater coverage of crystallization space than a typical protein-precipitant 2D concentration gradient, without requiring additional protein.

Case study using concanavalin A:

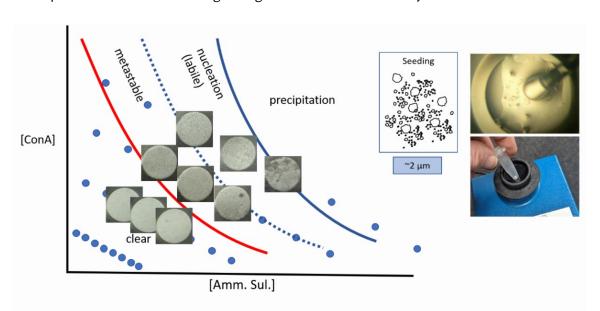
Experiment design: phase diagram experiment using concentrated protein (green), precipitant cocktail (red) and diluent (blue)



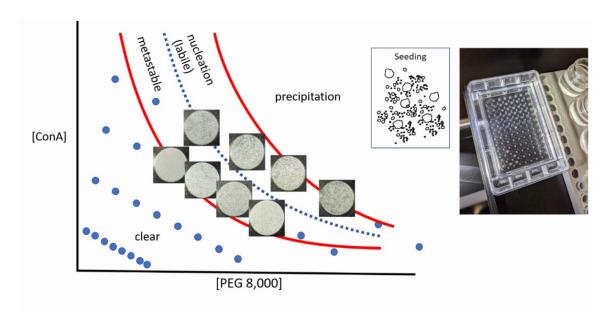
Experiment 1: three drops had crystals without seeding. This showed us the rough upper and lower boundaries of the nucleation zone



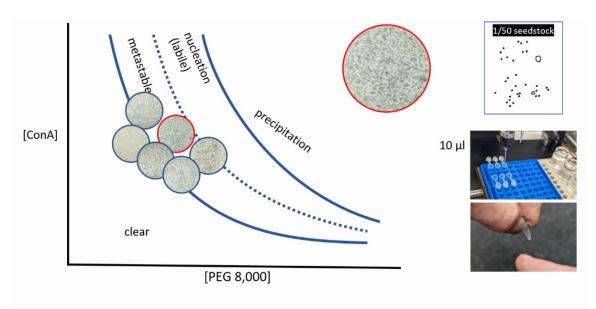
Experiment 2: we used one of the drops from experiment 1 to make a seedstock, and repeated the experiment with microseeding. This gave us the lower boundary of the metastable zone



Experiment 3: we repeated the same experiment with **50% PEG 8,000** replacing ammonium sulfate



Scaling up: using an Oryx8 we dispensed larger volumes to six PCR tubes with a 1/50 dilution of the seed-stock, at slightly lower PEG concentration. The PCR tubes were flicked, not vortexed.



Key benefits of new experiment:

- 1. **Expanded crystallization space**: Achieve broader testing with a systematic approach that can help identify optimal conditions for serial data collection, MicroED, or neutron diffraction.
- 2. **Simple and versatile**: this method is highly adaptable and often requires no changes between projects. Simply provide the protein and a pre-mixed precipitant cocktail at the desired maximum concentration, when requested by the software.
- Seeding: run experiments both with and without seeds to identify the metastable
 regions of the phase diagram. Seeding can also be utilized to control crystal size and
 density.

Conferences:

Douglas Instruments is attending the following meetings:

ECM 34, Padova

26 August - 30 August 2024

Rhine-Knee Regional Meeting, Konstanz

30 September - 2 October 2024

Visit our booth for:

Demonstrations
rMMS microseeding toolkits
Updates about our latest projects including crystallization methods for serial crystallography.

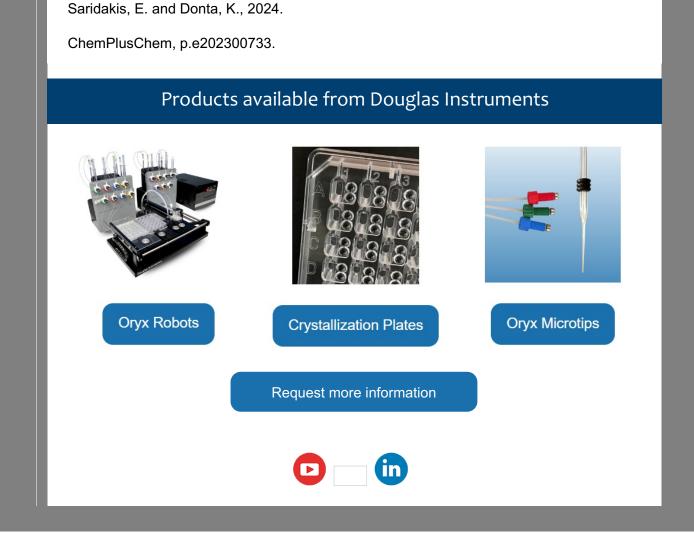
Recently published research using Oryx protein crystallization robots:

Rationally seeded computational protein design of **a**-helical barrels

Albanese, K.I., Petrenas, R., Pirro, F., Naudin, E.A., Borucu, U., Dawson, W.M., Scott, D.A., Leggett, G.J., Weiner, O.D., Oliver, T.A. and Woolfson, D.N., 2024.

Nature Chemical Biology, pp.1-9.

Protein Thermodynamic Properties, Crystallisation, and the Hofmeister Series



Douglas Instruments | Douglas House East Garston | Hungerford, RG177HD GB

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