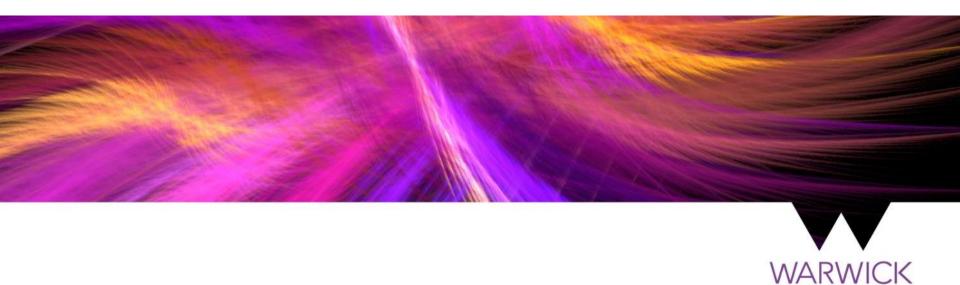
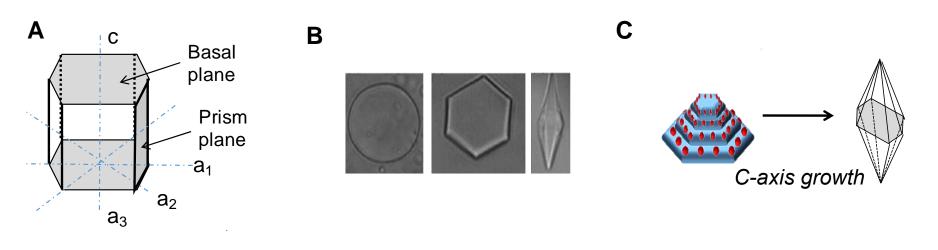
Work Update - December

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Introduction

- Ice formation and growth can be a serious problem
- Much research ongoing in many fields relating to ice
- Little understood about its mechanism
- This work investigates the ice growth process and mechanism of action of IRI active compounds; I have specifically been looking at dynamic ice shaping (modification of morphology of ice crystal) and ice recrystallisation inhibition
- As well as the dynamics of nuclei in frozen samples



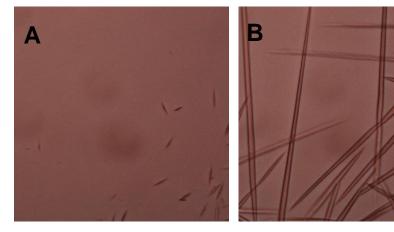
Projects & Aims

- Characterise the changes in ice structure upon addition of iceactive compounds (via microscopy, sucrose assays, X-ray scattering)
 - Antifreeze proteins
 - PVA
 - Safranin O
- Investigate what we can learn about the mobility of water and how it is affected by antifreezes using solid state NMR
 - Study relaxation rates
 - Compare motions in ice-active and inactive compounds, proteins
- Obtain further understanding of the mechanism of growth to improve techniques and designs for ice nucleation promoters and cryoprotectants.

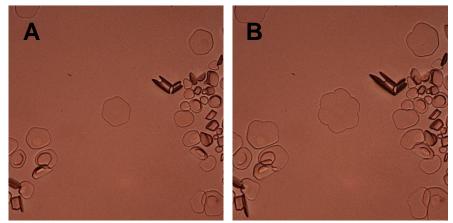
Sucrose Assay Results

- Used to study morphology of ice crystals
- Sample dissolved in 45 w% sucrose and PBS.
- Initial examples below for samples which ice shape and do not:
 - AFGP shows the expected needles as the crystals grow.
 - For the copolymer so far it appears to not ice shape (hexagonal crystal)

AFGP

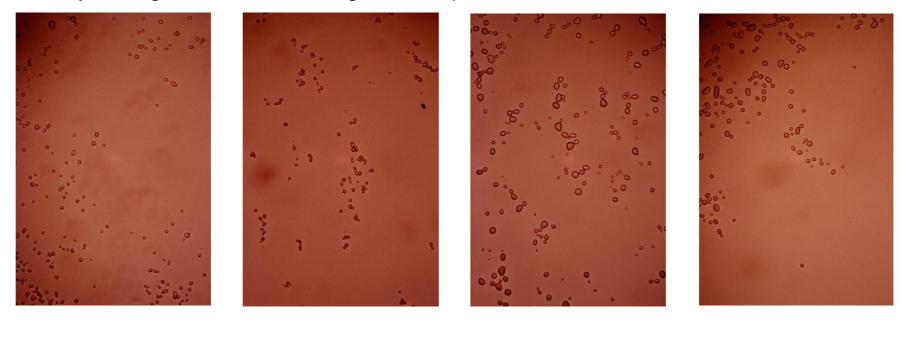


Ben G copolymer



Comparisons of crystal growth morphology

 Study the morphologies of ice upon addition of PVA and AFPIII-SNAP and if they change when bound to gold nanoparticles



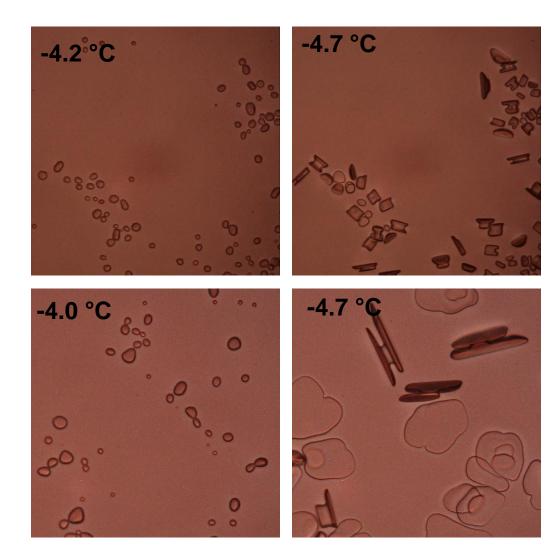
AFP-SNAP AFP-SNAP PVA-Gold -Gold

PVA

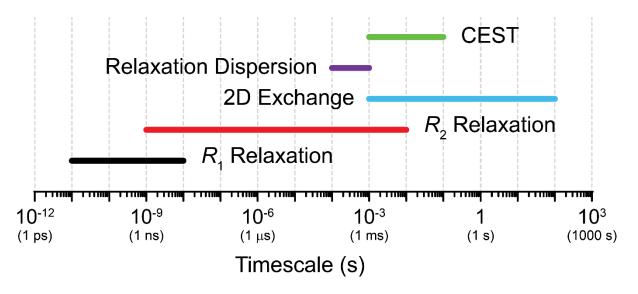
Close up on PVA comparison

PVA (0.1 mg.mL⁻¹)

PVA-Gold



Solid State NMR



- Performed T1, T2 and T1rho experiments on water, CPAs and negative controls using a MAS frequency of 10 KHz
- Measured relaxation rates to study mobility of water by looking at dynamics of protons
- Tried multiple experiments at different timescales (shown in figure above), but we saw the most useful data from relaxation dispersion (purple) and variable temperature R₂ relaxation studies (red).

Solid State NMR

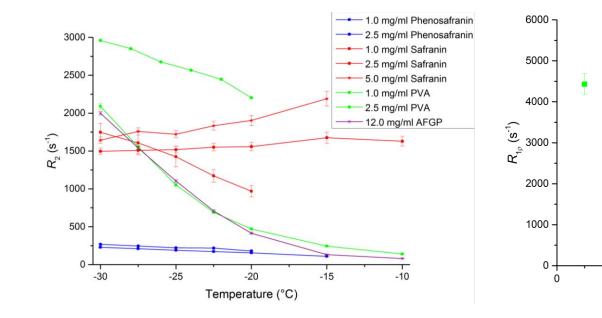


Fig 1. R_2 relaxation measurements between -30 and -10 $^{\circ}$ C

Fig 2. Relaxation dispersion data, R₁ against spinlock frequency

30

Spinlock (kHz)

20

10

1.0 mg/ml PVA

1.0 mg/ml Phenosafranin

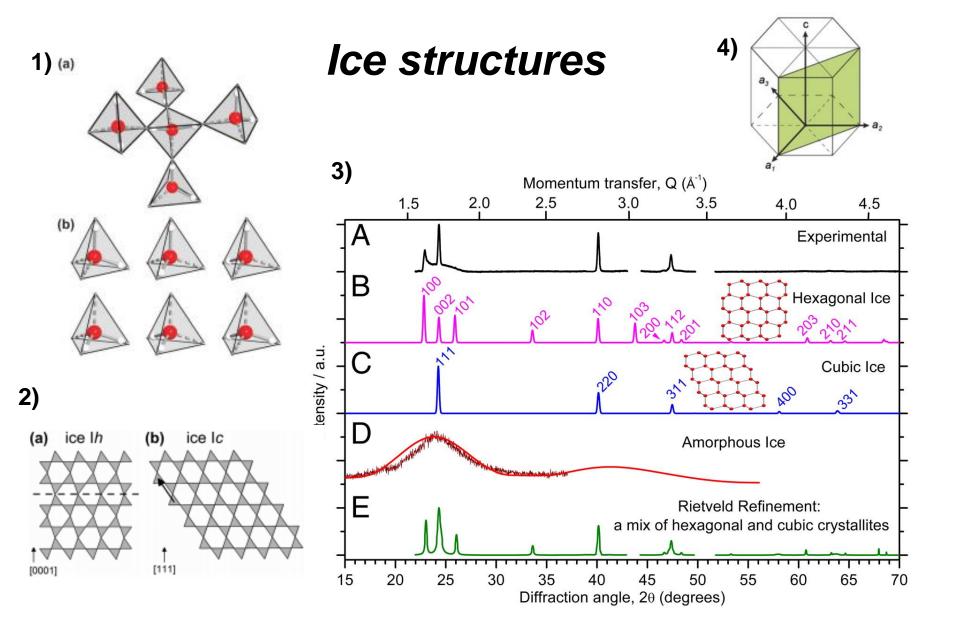
10.0 mg/ml Safranin

40

50

Solid State NMR

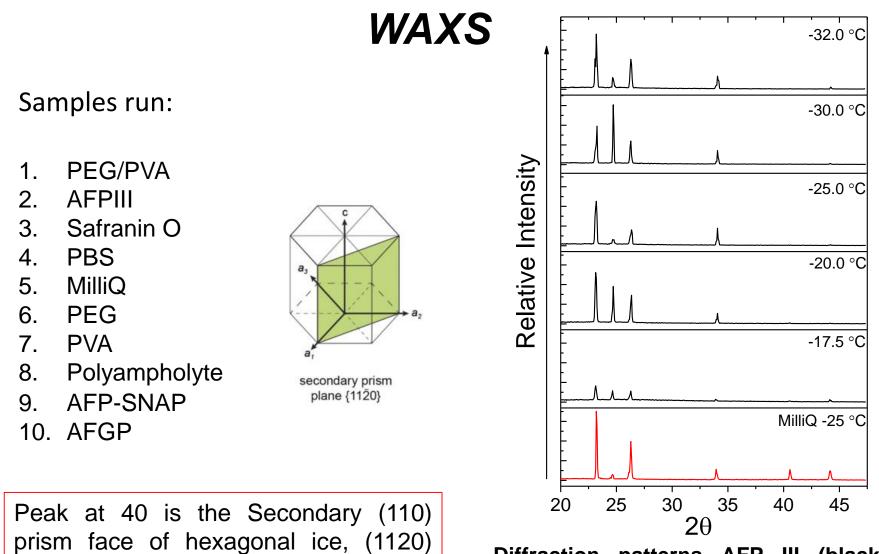
- Initially difficult to study ice
- We are hoping to compare macroscopic effects (eg. DIS, IRI) with the motions at different timescales.
- We so far have seen there is definitely a difference, just need to look at more samples eg AFGP.
- Perform viscosity measurements to compare to R2



Freezing points

Sample	Freezing point (°C)	Melting point	\overline{x} T _n (microscopy)
PBS:AFP III	-17.5	2.83	?
H ₂ O:Safranin	-23.66	4.33	
PBS	-24.03	3.39	-34.09
MilliQ	Ca26	Ca. 4	Ca40
H ₂ O:PEG	Ca24	Ca. 2.5	
H ₂ O:PVA	Ca22	Ca. 4	-37.38
PBS:Polyampholyte	-22.35	1.99	
PBS:AFP III-Snap	-21.5		-20.70



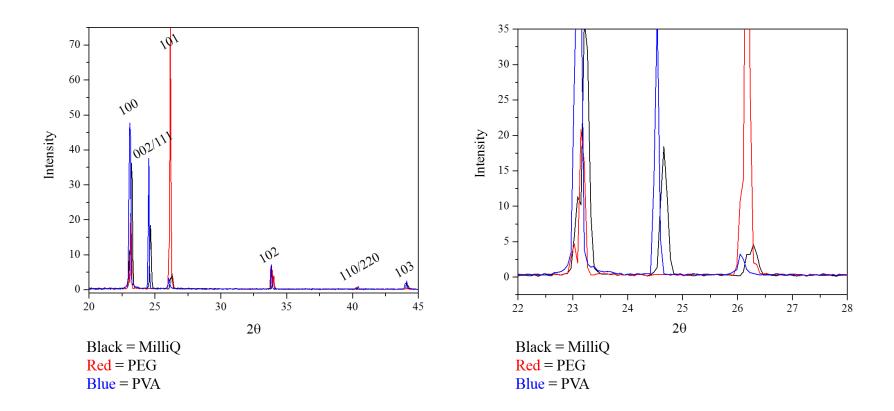


prism face of hexagonal ice, (1120) secondary prism plane Compared to

Diffraction patterns AFP III (black) compared to MilliQ (red) as the temperature decreases

WAXS

- WAXS analysis of water diffraction patterns from -10 to 10 degrees and the effect of PVA, PEG
- Peak splitting, slight shifts and changes in intensity observed



Labels for peaks based on Salzmann 2012

WAXS ideas

Possible reasons for peak splitting:

- Deformation of hexagonal ice by surface active components
- Potentially artefacts form
- Crystallisation of a solute but do the peaks not correspond to the known crystalline forms of solutes in systems studied
- Another ice polymorph ice IV/III?
- In structure changes due to pressure build-up due to volume expansion during water-ice transition.

Other results:

- Safranin O works differently to AFPIII, PVA and PEG.
- Support for PVA/PEG solutions working synergistically

To do:

- Need to analyse the latest data sets on AFGP
- Perform splat assay in x-ray



Protein Work & Cryopreservation

- AFPIII expression and purification to compare to PEG/PVA
- Cryopreservation of different bacteria strains

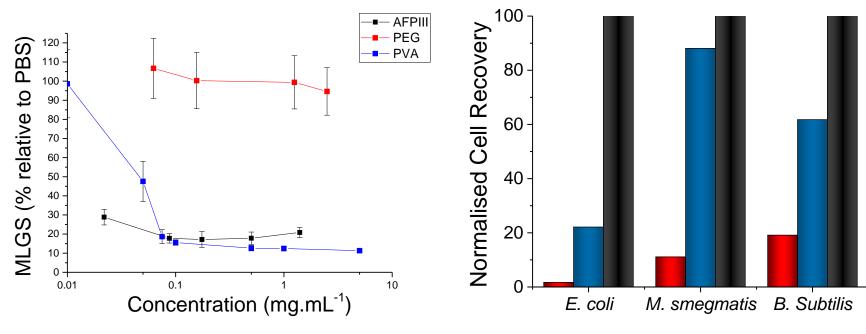


Fig 1. concentration dependence of IRI activity

Fig 2. Cell recovery after 7 FT with no CPA (red), glycerol (blue) and PEG/PVA (black)

What to do next

- Continue WAXS and then move on to SAXS to compare results directly with what we see from SPLAT assays
- Can also look at the gold nanoparticles in SAXS to first observe scattering of nanoparticles and then to see if they are coated by the proteins/how scattering is affected
- Analyse all the data to see what it means
- NMR of different DP PVAs
- Viscosity studies

Acknowledgements – GibsonGroup 2017



Post-docs

- Dr. Sarah-Jane Richards
- Dr. Caroline Biggs
- Dr Collette Guy
- Dr Lucienne Otten
- Dr Muhammed Hasan
- Dr Antonio Laezza

Undegrad Students

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 Adam Jones Segun Wahab

Phd Students

- Sang Ho Won
- Lewis Blackman
- Benjamin Martyn
- Joseph Healey
- Chris Stubbs
- Ben Graham
- Trisha Bailey
- Laura Wilkins
- Marie Grypioti
- Julia Lipecki
- Vinko Varas
- Gabriel Erni Cassola
- Robyn Wright
- Alice Fayter
- Alex Baker
- Ruben Tomas
- Iain Galpin



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Funders £\$€