

# WMS UPDATE

GROUP MEETING

27/3/2018






MUHAMMAD HASAN & ALICE FAYTER

WARWICK

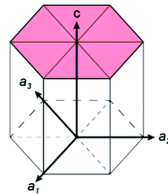
# OVERVIEW

- Introduction
- AFPIII
- AFPI
- Cryopreservation of bacterial cells
- Future work

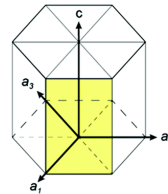
# INTRODUCTION: AFPs

	AFGP	Type I AFP	Type II AFP	Type III AFP	Insect AFP
Mass ( $\text{kg mol}^{-1}$ )	3-33	3-4.5	11-24	6.5	9
Representative structure					
PDB	-	1WFA	2PY2	1HG7	-
Binding plane	○ ● ○ ○	○ ○ ○ ● ○ ○ ● ○	○ ● ○ ○	○ ● ○ ○ ●	● ● ● ● ○
Origin	Antarctic notothenioids, northern cods	Right-eyed flounders, sculpins	Herring, sea raven, smelt	Eel pout, ocean pout, wolfish,	Darkling beetle, spruce budworm moth, midge, fly

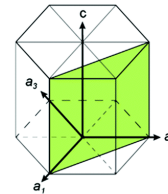
Crystal planes of hexagonal ice (Ih)



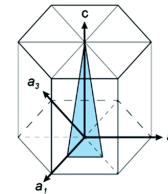
basal plane  
{0001}



primary prism  
plane {1010}



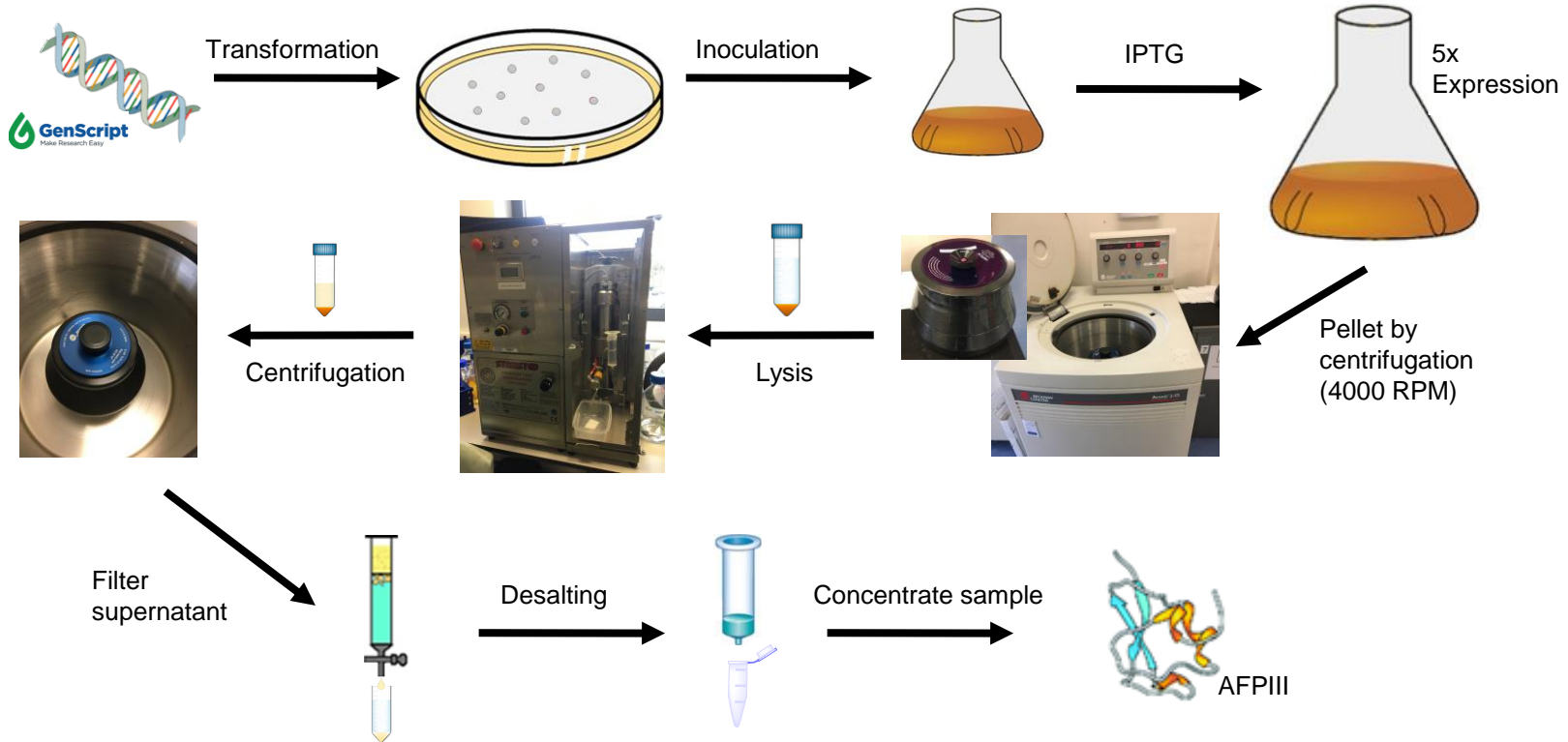
secondary prism  
plane {1120}



pyramidal plane  
{2021}

# AFPIII: Expression and purification

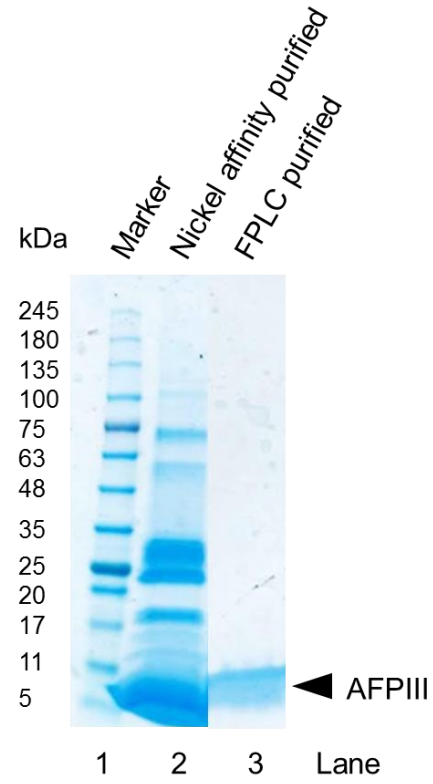
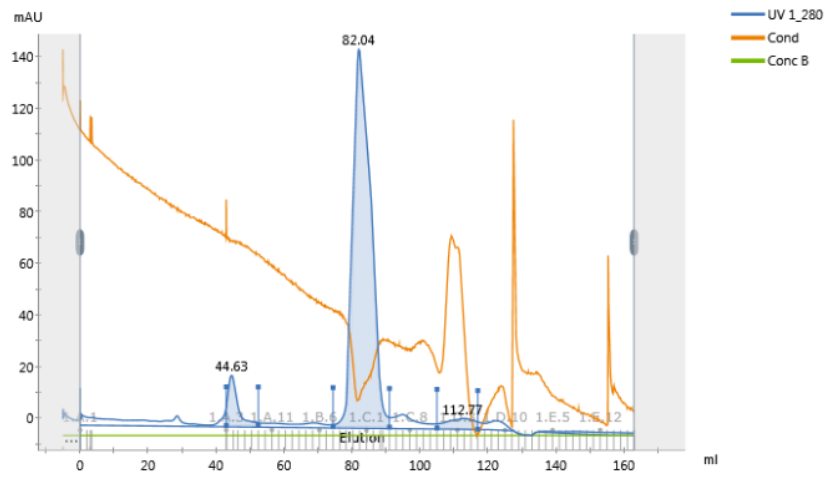
WARWICK



# AFPIII: FPLC/Gels

Concentration recorded by:

- BCA Assay
- Nanodrop (A280 absorbance) - Tyr & Trp
- Difficulties with FPLC originally



# bpAFPI: Construct

TRX-tag – His-tag – Thrombin – S.tag – rEK – TEV – AFPI

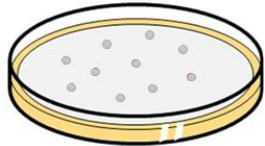
- Goal: Separate TEV protease, tag and leftover impurities from the AFPI peptide.

# bpAFPI: Expression

WARWICK



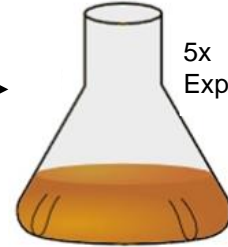
Transformation



Inoculation

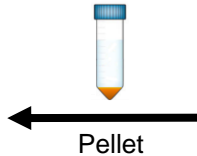


IPTG

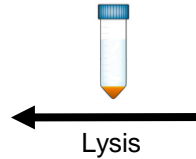


5x  
Expression

Pellet by  
centrifugation  
(4000 RPM)



Pellet

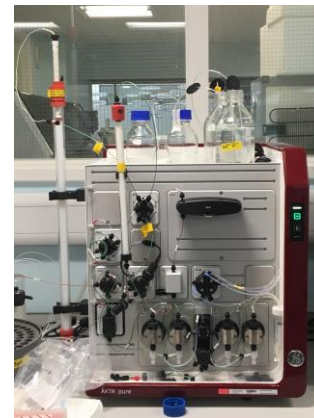
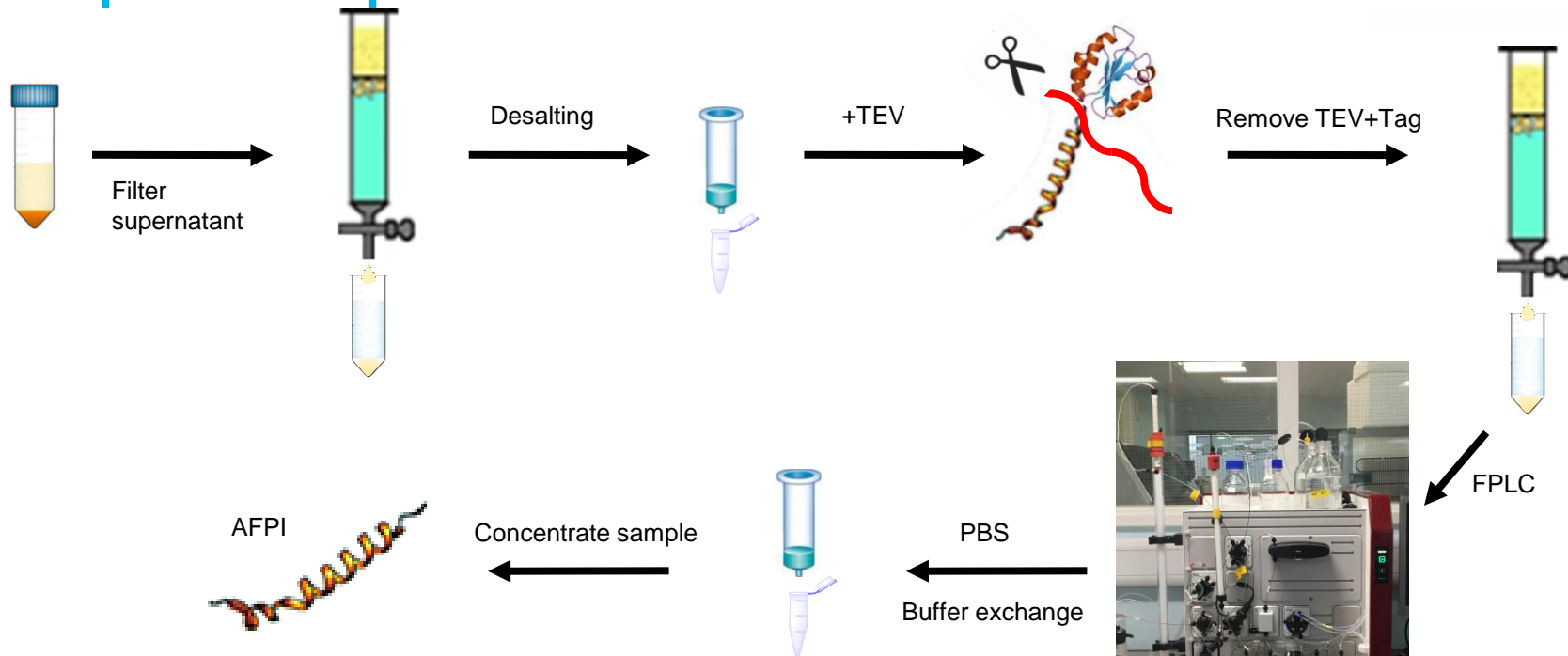


Lysis



# bpAFPI: purification

WARWICK



FPLC

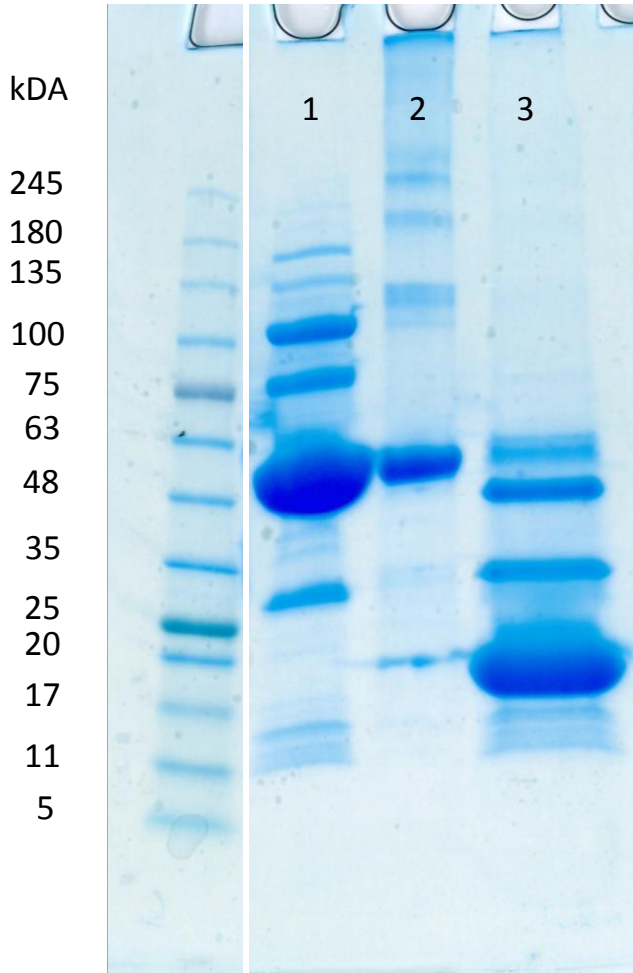


Concentrate sample

PBS  
Buffer exchange



# bpAFPI :Failed

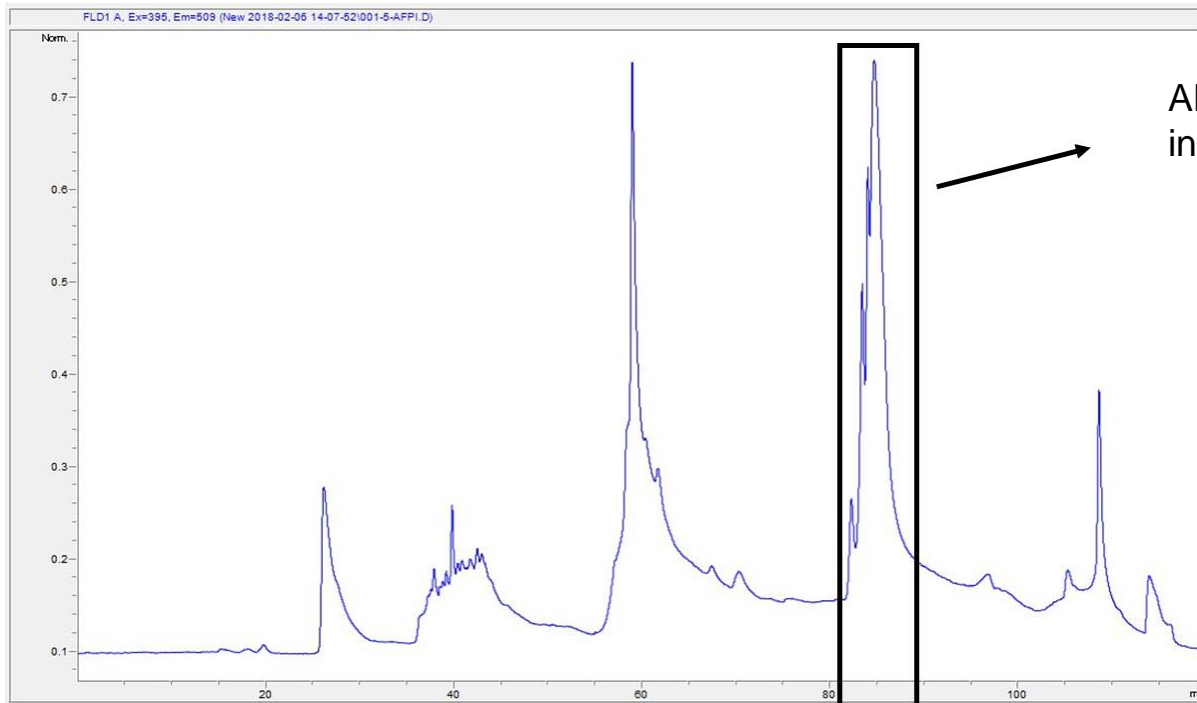


- 1 – AFPI after IMAC
- 2 – TEV after IMAC
- 3 – overnight incubation of AFPI-TEV

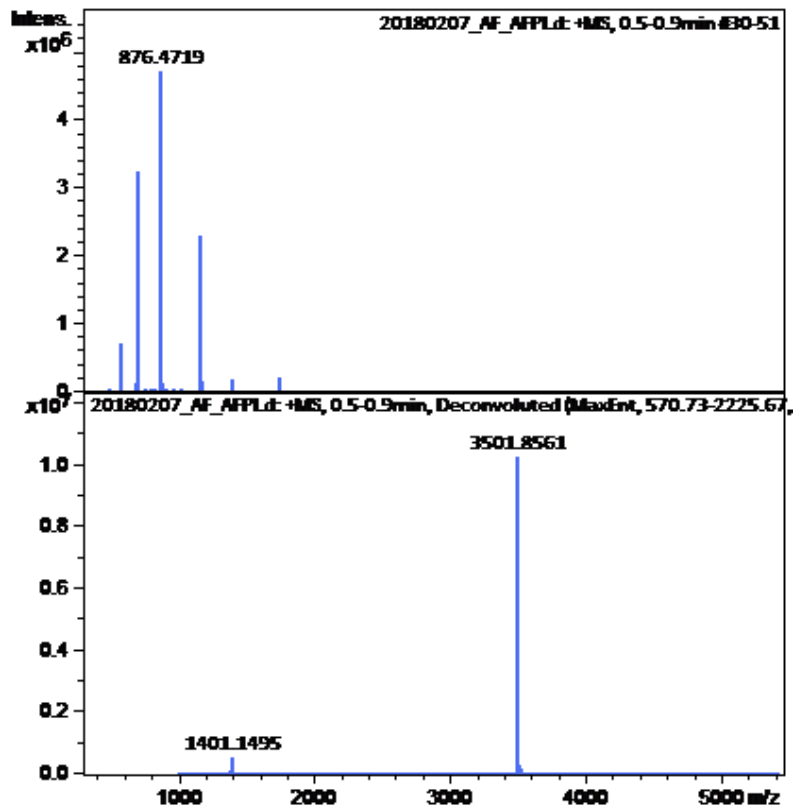
TRX-tag – His-tag – Thrombin – S.tag – rEK – TEV – AFPI



# bpAFPI: HPLC



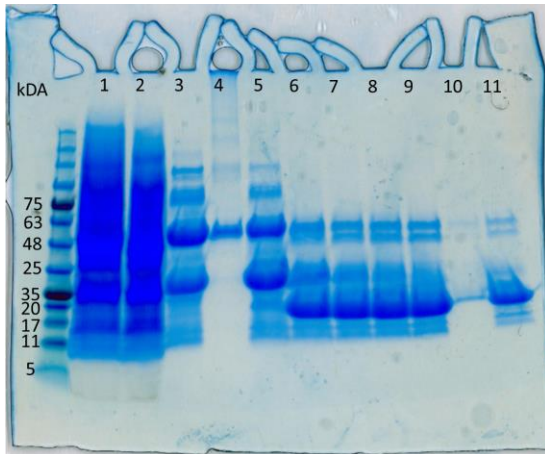
# bpAFPI: Mass Spec QToF



# bpAFPI: Success

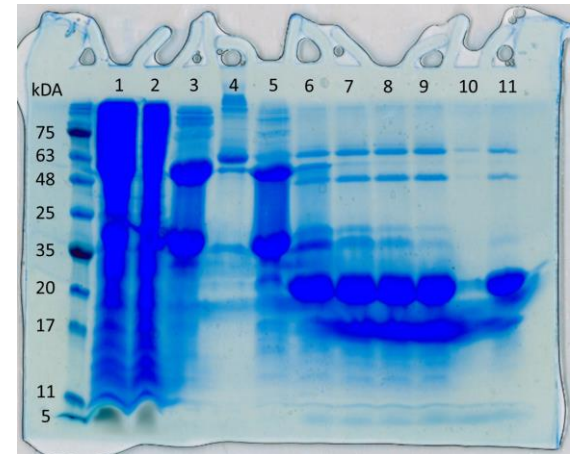
- The 40 AA peptide was expressed in fusion with a thioredoxin tag in pET32a vector (ref Davies paper).

**Tris-glycine gel**



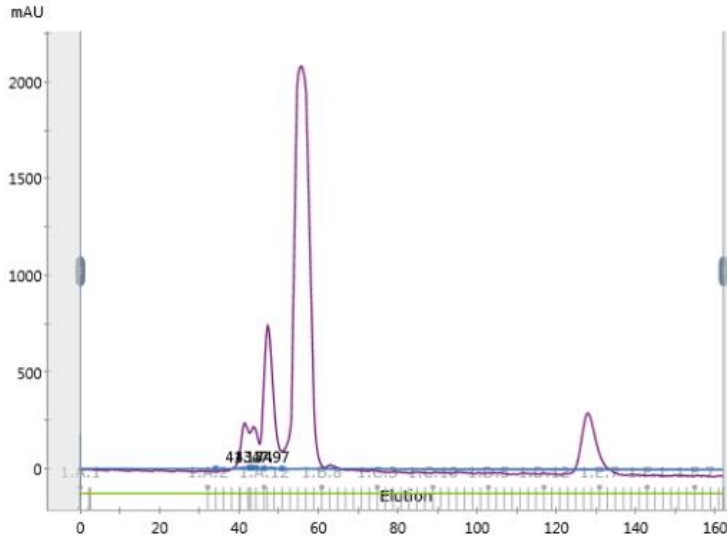
- 1 – Before IMAC
- 2 – Flow through
- 3 – Elution after PD10
- 4 – TEV
- 5 – 0 hrs after addition of TEV
- 6 – 1 hr
- 7 – 2 hrs
- 8 – 4 hrs
- 9 – Overnight
- 10 – FT from AFPI concentration
- 11 – left on the column after second IMAC

**Tris-tricine gel**

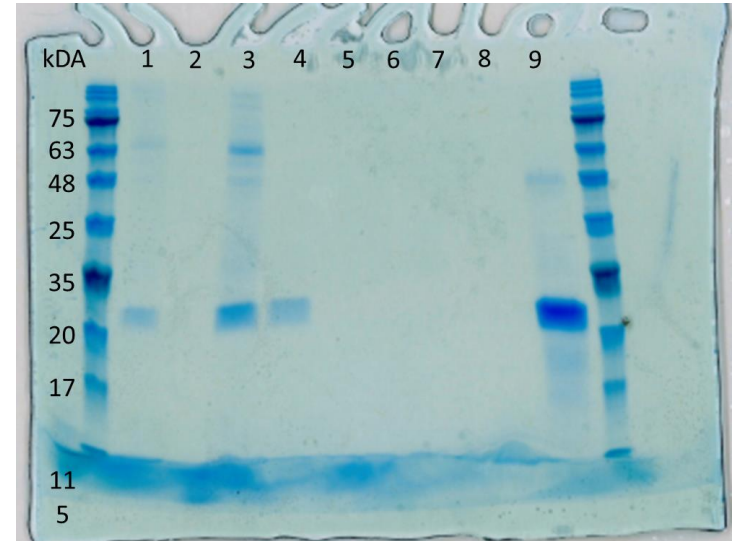


# bpAFPI: FPLC & Gel

Superdex 75 column = didn't work  
 Superdex 30 column = worked



- 1 – AFPI after IMAC
- 2 – FT from concentrator
- 3 – Concentrated AFPI
- 4 – B1
- 5 – B4
- 6 – B5
- 7 – B6
- 8 – E5
- 9 – old peptide sample



# bpAFPI: Proteomics Results

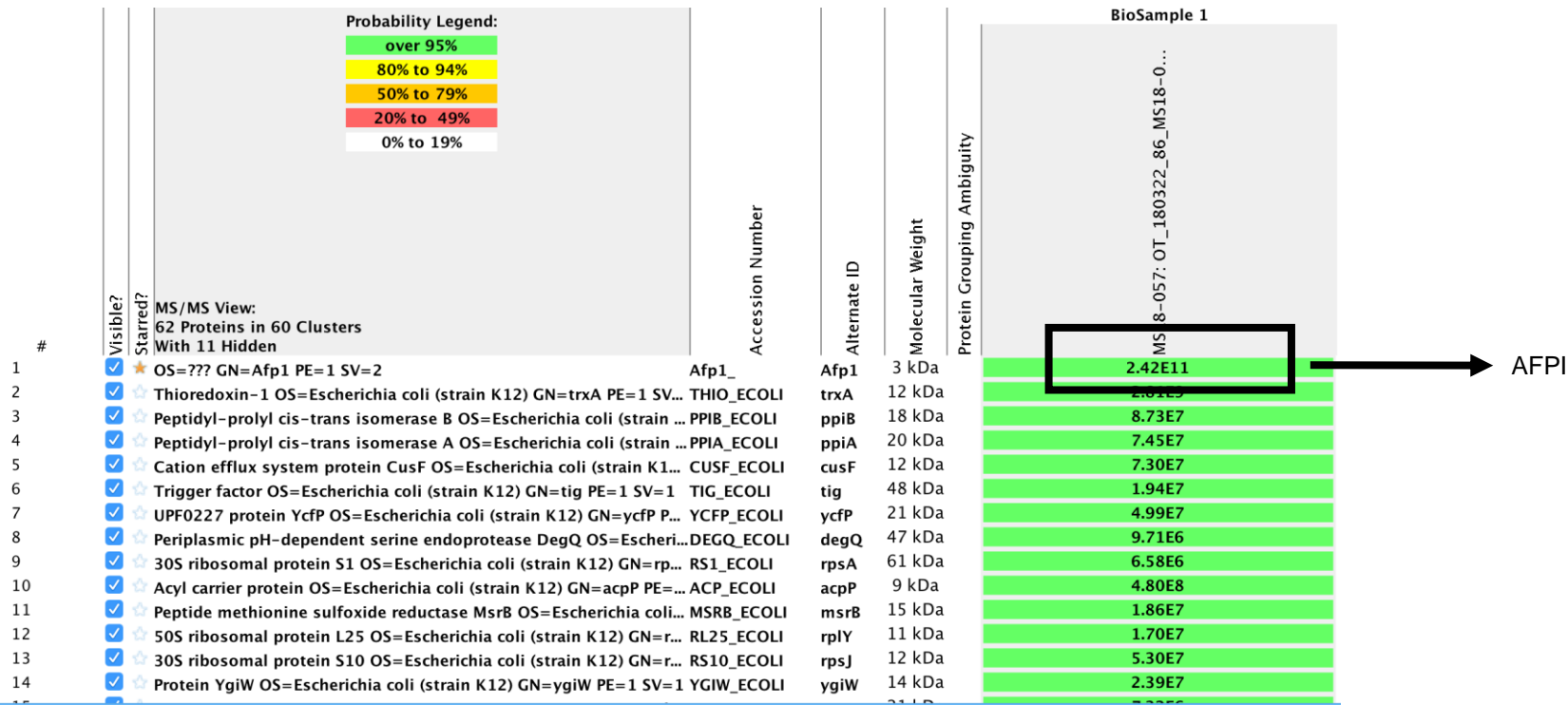
#	Visible?	Starred?	MS/MS View: 62 Proteins in 60 Clusters With 11 Hidden	Accession Number	Alternate ID	Molecular Weight	Protein Grouping Ambiguity
			<b>Probability Legend:</b> <span style="background-color: #90EE90; border: 1px solid black; display: inline-block; width: 15px; height: 10px;"></span> over 95% <span style="background-color: #FFFF00; border: 1px solid black; display: inline-block; width: 15px; height: 10px;"></span> 80% to 94% <span style="background-color: #FFD700; border: 1px solid black; display: inline-block; width: 15px; height: 10px;"></span> 50% to 79% <span style="background-color: #FF6347; border: 1px solid black; display: inline-block; width: 15px; height: 10px;"></span> 20% to 49% <span style="background-color: #F0F0F0; border: 1px solid black; display: inline-block; width: 15px; height: 10px;"></span> 0% to 19%				
1	✓	★	OS=??? GN=Afp1 PE=1 SV=2	Afp1_	Afp1	3 kDa	2.42E11
2	✓	☆	Thioredoxin-1 OS=Escherichia coli (strain K12) GN=trxA PE=1 SV=...	THIO_ECOLI	trxA	12 kDa	2.81E9
3	✓	☆	Peptidyl-prolyl cis-trans isomerase B OS=Escherichia coli (strain ...	PPIB_ECOLI	ppiB	18 kDa	8.73E7
4	✓	☆	Peptidyl-prolyl cis-trans isomerase A OS=Escherichia coli (strain ...	PPIA_ECOLI	ppiA	20 kDa	7.45E7
5	✓	☆	Cation efflux system protein CusF OS=Escherichia coli (strain K1...	CUSF_ECOLI	cusF	12 kDa	7.30E7
6	✓	☆	Trigger factor OS=Escherichia coli (strain K12) GN=tig PE=1 SV=1	TIG_ECOLI	tig	48 kDa	1.94E7
7	✓	☆	UPF0227 protein YcfP OS=Escherichia coli (strain K12) GN=ycfP P...	YCFP_ECOLI	ycfP	21 kDa	4.99E7
8	✓	☆	Periplasmic pH-dependent serine endoprotease DegQ OS=Escheri...DEGQ_ECOLI	DEGQ_ECOLI	degQ	47 kDa	9.71E6
9	✓	☆	30S ribosomal protein S1 OS=Escherichia coli (strain K12) GN=rp...	RS1_ECOLI	rpsA	61 kDa	6.58E6
10	✓	☆	Acyl carrier protein OS=Escherichia coli (strain K12) GN=acpP PE=...	ACP_ECOLI	acpP	9 kDa	4.80E8
11	✓	☆	Peptide methionine sulfoxide reductase MsrB OS=Escherichia coli...	MSRB_ECOLI	msrB	15 kDa	1.86E7
12	✓	☆	50S ribosomal protein L25 OS=Escherichia coli (strain K12) GN=r...	RL25_ECOLI	rplY	11 kDa	1.70E7
13	✓	☆	30S ribosomal protein S10 OS=Escherichia coli (strain K12) GN=r...	RS10_ECOLI	rpsJ	12 kDa	5.30E7
14	✓	☆	Protein YgiW OS=Escherichia coli (strain K12) GN=ygiW PE=1 SV=1	YGIW_ECOLI	ygiW	14 kDa	2.39E7

BioSample 1

MS18-057: OT\_180322\_86\_MS18-0...

# bpAFPI: TADA!

WARWICK



# Cryopreservation of bacterial cells

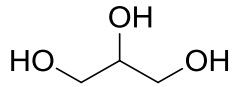
- Storage of lab strains
- Structure-activity relationship
- Industrial application in light of microbiome research



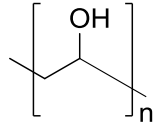


# THE CANDIDATES

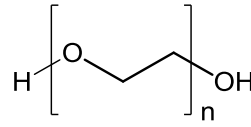
WARWICK



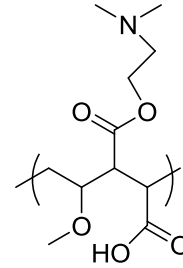
Glycerol



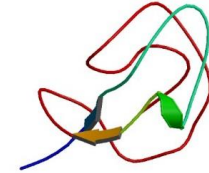
PVA



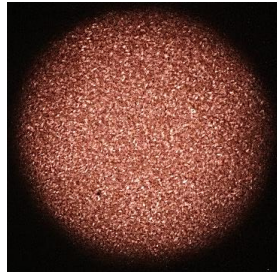
PEG



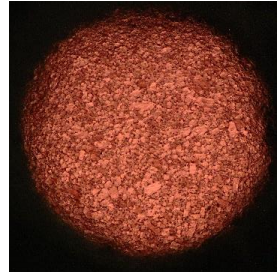
Poly(ampholyte)



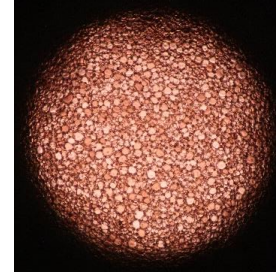
AFPIII



PEG/PVA



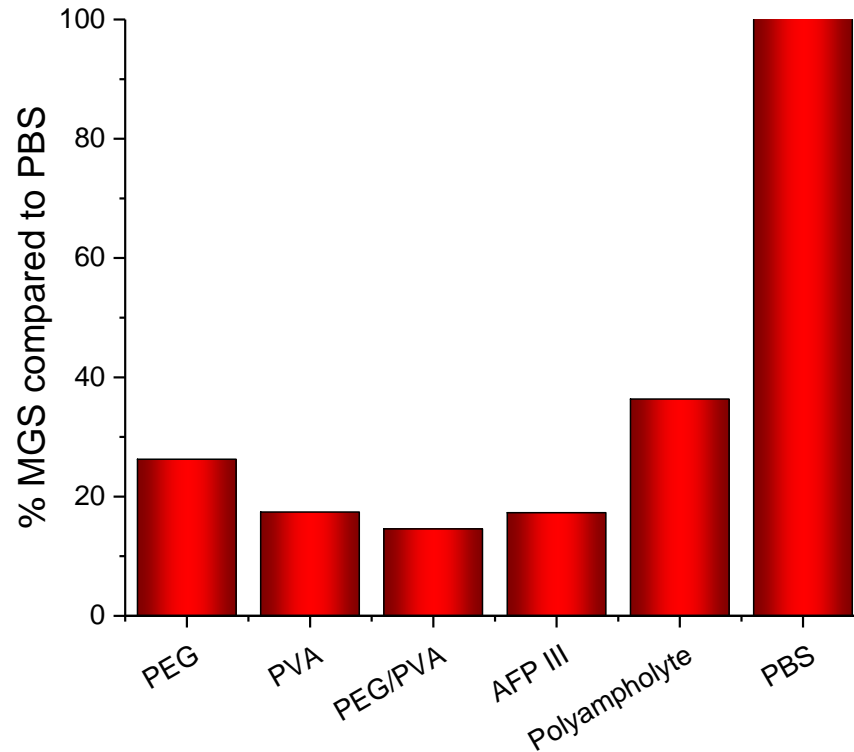
PVA



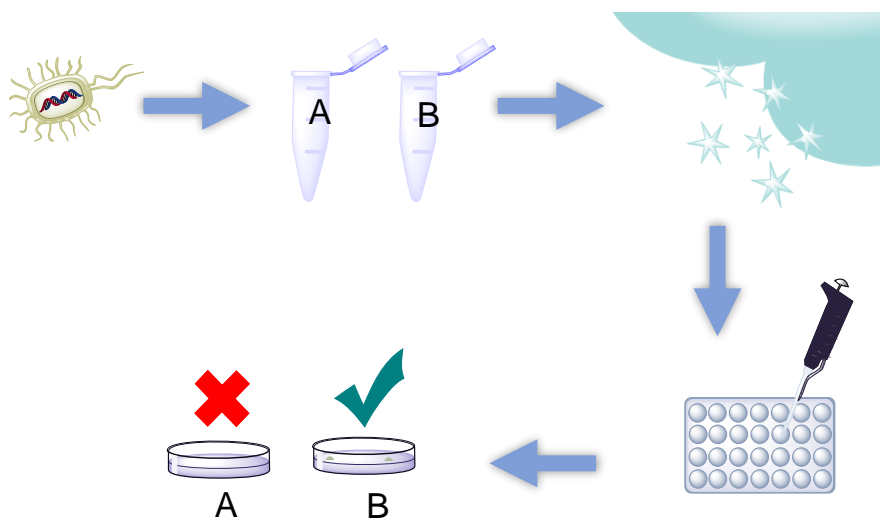
PEG

# Ice-recrystallization inhibition: results

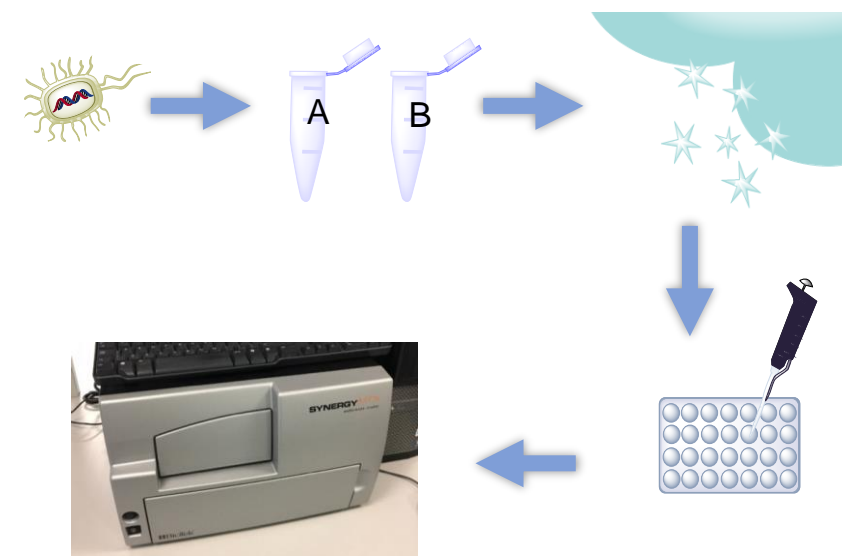
WARWICK



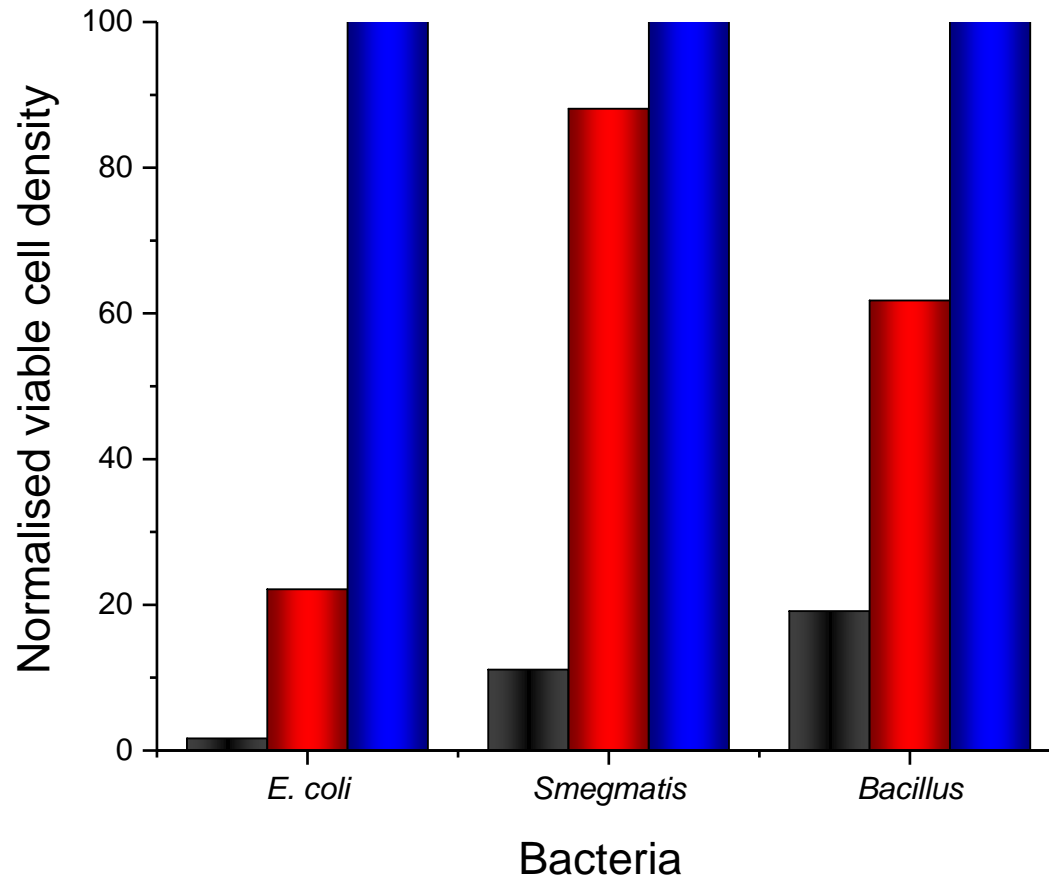
# Methods: Freeze-tolerance assays



Counting colonies

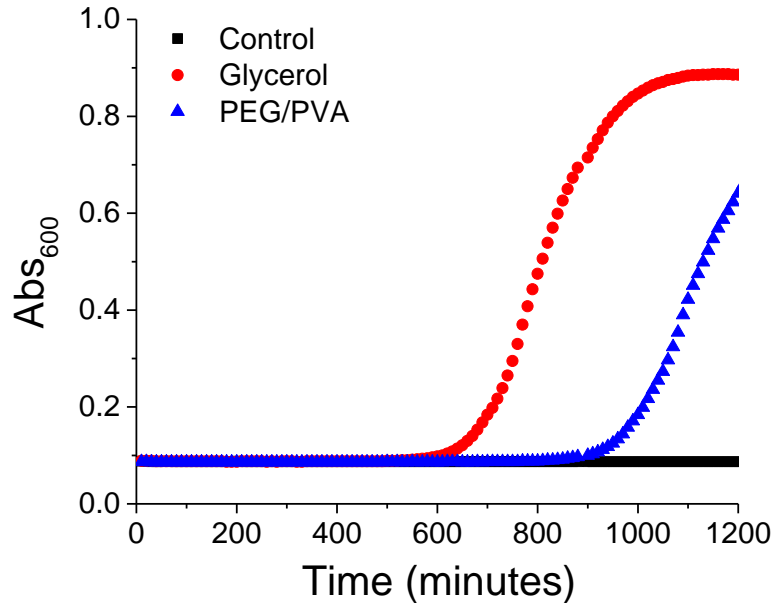


Monitoring growth profiles at OD600



**Control**  
**Glycerol**  
**PEG/PVA**

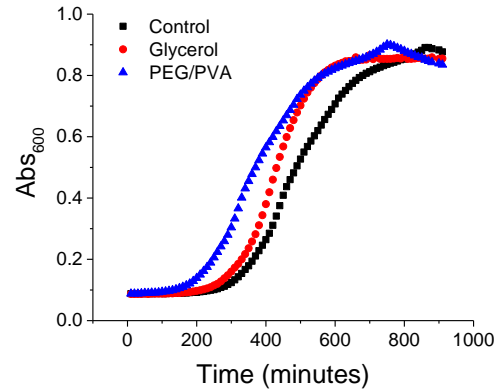
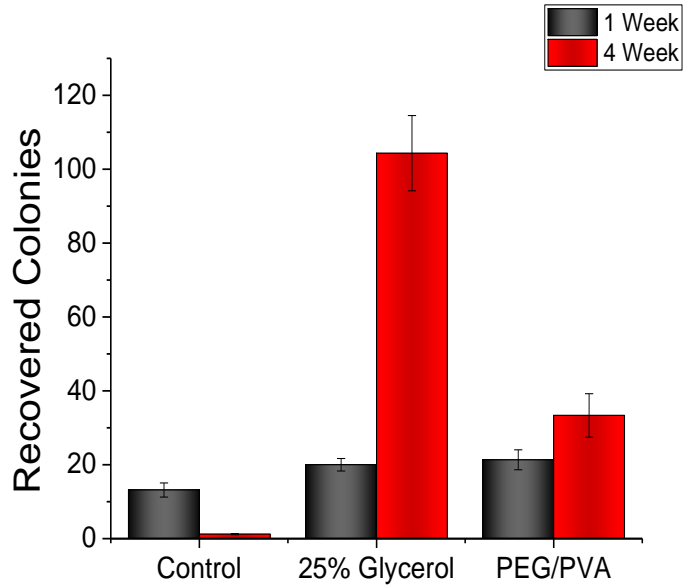
# *E. coli* growth profile post 7xFT with storage at -20°C



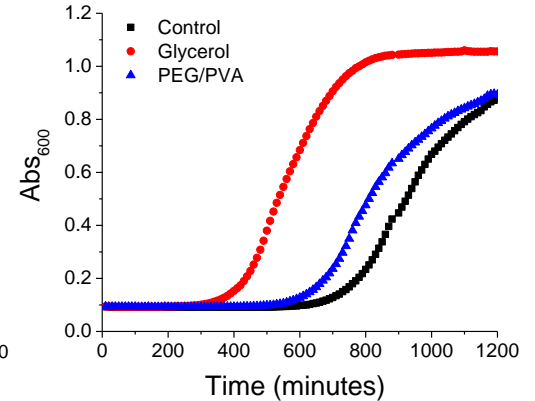
ALL



# Results after storage at -20 °C

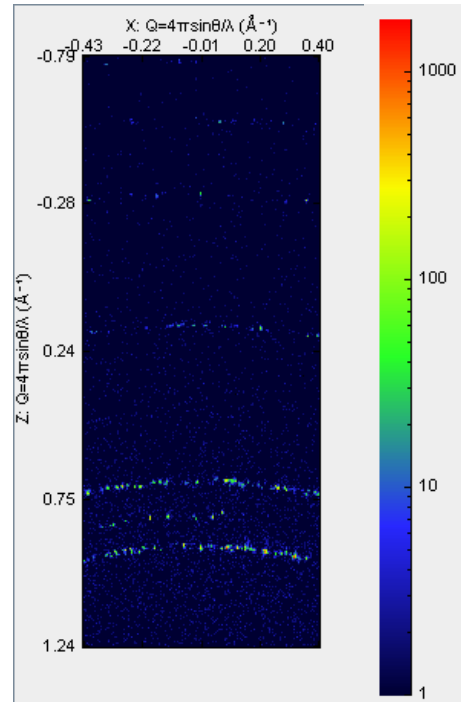
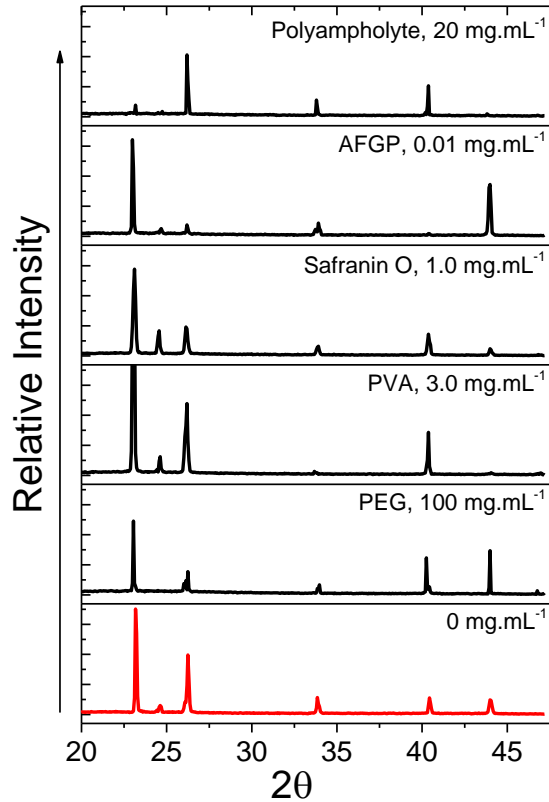


*E. Coli* after 1 freeze thaw cycle (Frozen in N<sub>2</sub> & left at -20 °C for 1 week)

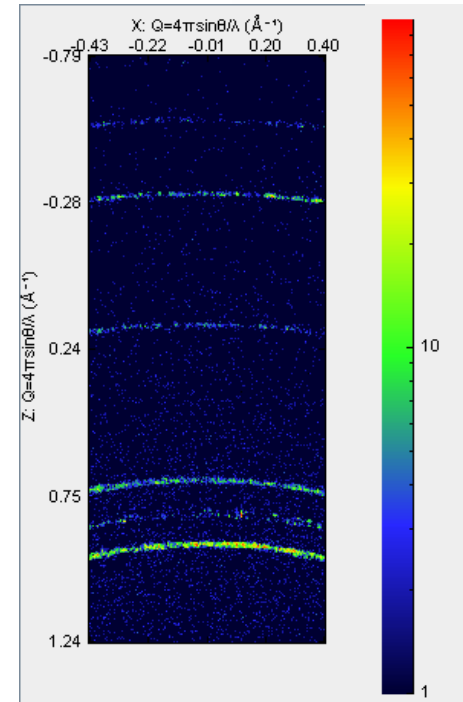


*E. Coli* after 1 freeze thaw cycle (Frozen in N<sub>2</sub> & left at -20 °C for 4 weeks)

# WAXS: Results & Hypothesis



Scattering pattern for Safranin O (0.1 mg.mL<sup>-1</sup>)



Scattering pattern for Safranin O (1 mg.mL<sup>-1</sup>)

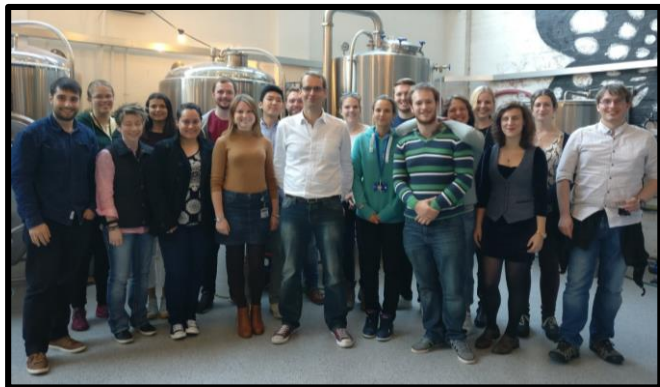
# Future Work



- LEA
  - Intrinsically disordered
  - Known to potentially provide cryoprotection
  - Distinct to IRI
  - Stabilisation?
- INP
  - Good to have one
- CRP
  - Cryoprotective protein
  - 29 kDa
  - Have seen it is 1000x more active than BSA
  - Just arrived!
  - Plan to: get genome & identify sequence/structure of protein
  - SFR



# Acknowledgements – GibsonGroup 2018



## Post-docs

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

## Undegrad Students

- Maelle Bottin
- Segun Wahab

## Phd Students

- 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

Funders £\$€



European Research Council

Established by the European Commission



EPSRC

Pioneering research  
and skills



The Leverhulme Trust

Noreen Murray Philanthropy  
Award

Friends of the University of  
Warwick Trust



THE UNIVERSITY OF  
WARWICK



Iceni  
Diagnostics



www.advantagewm.co.uk

[www.warwick.ac.uk/go/gibsongroup](http://www.warwick.ac.uk/go/gibsongroup)



@LabGibson