

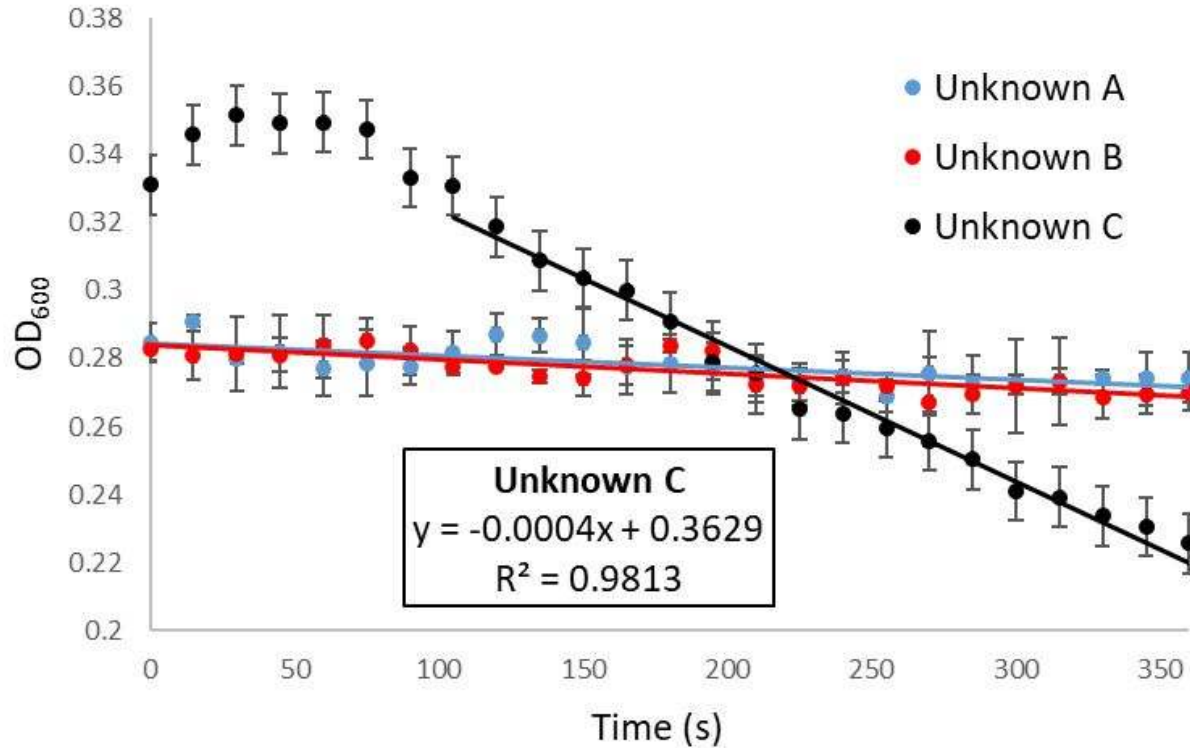


pUCK - 19 Lab Challenge

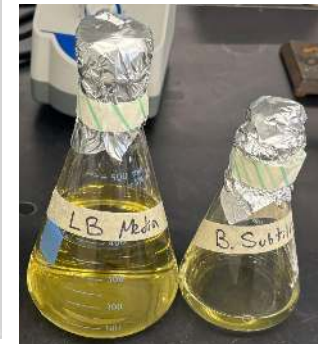
BIOC4001



Enzymatic Assay

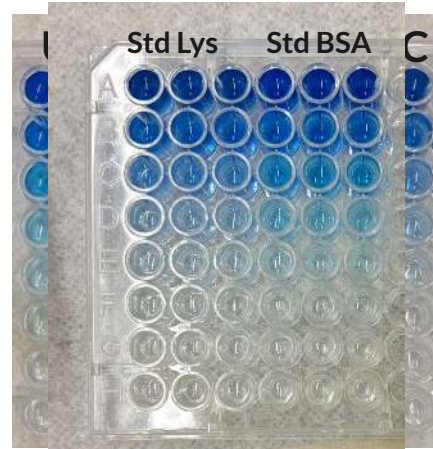
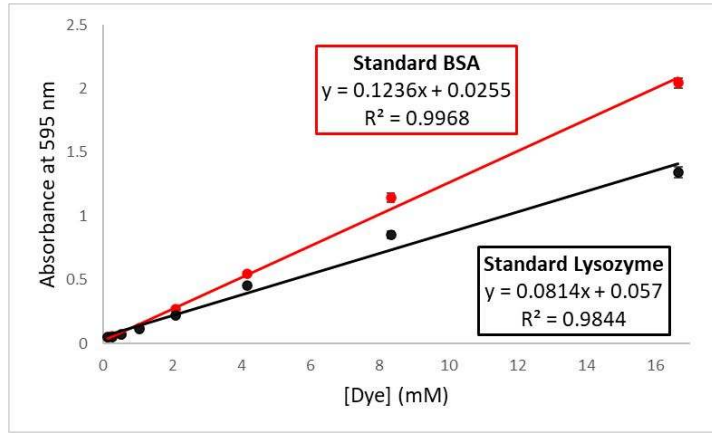


Yeast

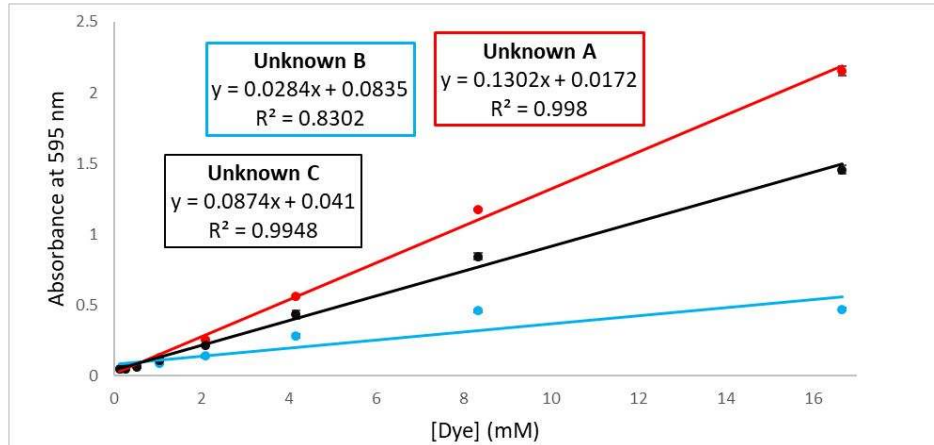


B. Subtilis

Dye Ligand Binding Assay and E1% Measurements



Fraction (0.1 mg/mL)	E1% (280 nm) $\left(\frac{L}{g \cdot cm}\right)$
Std Lys	21.0 ± 2.2
Std BSA	6.63 ± 1.00
Unk A	6.27 ± 0.25
Unk B	-1.63 ± 0.67
Unk C	21.8 ± 0.92



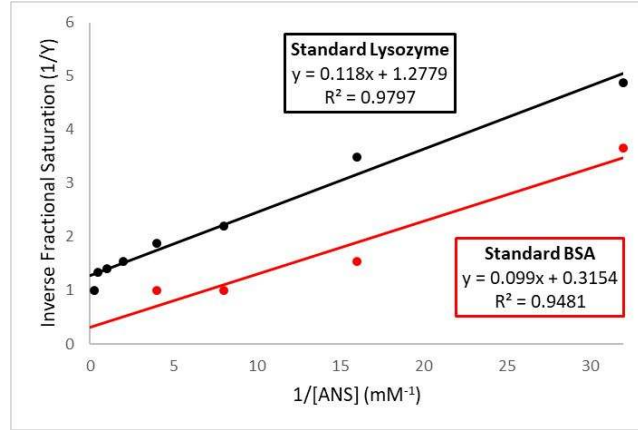
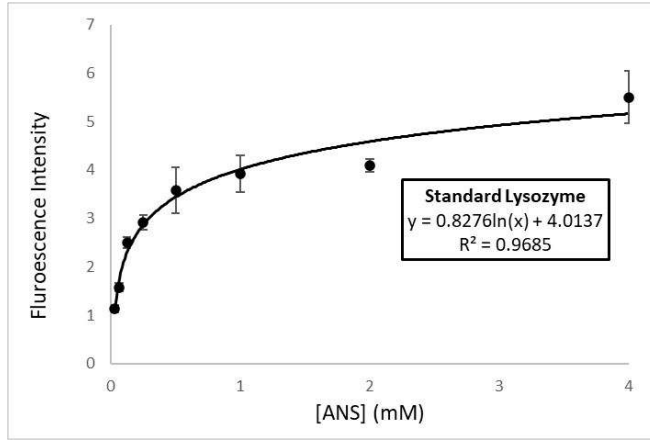
E1% is absorbance of 10 mg/mL solution

Lysozyme has a higher ratio of tyrosine and tryptophan in its structure than BSA

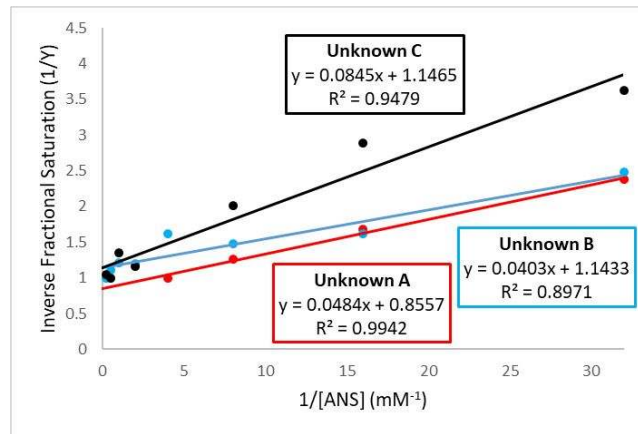
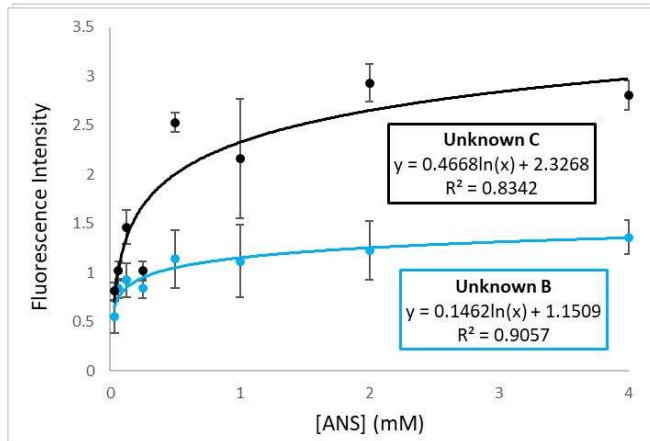
Unknown B had a negative absorption at 280 nm, indicating likely absence of protein, however had small absorbance at 260 nm range

ANS Fluorescence Binding Assay

$$\frac{1}{Y} = \frac{K}{n} \cdot \frac{1}{[L]} + \frac{1}{n}$$



	Dissociation Constant, K (mol/L)	Binding Sites (n)
Std BSA	0.31383	3.171
Std Lys	0.09234	0.7826
Unk A	0.05656	1.1686
Unk B	0.03522	0.8747
Unk C	0.07372	0.8722



MW BSA = 66.4 kDa

MW Lys = 14.4 kDa

Correction = 4.61

	Corrected Dissociation Constant, K (mol/L)
Std BSA	0.31383
Std Lys	0.02002

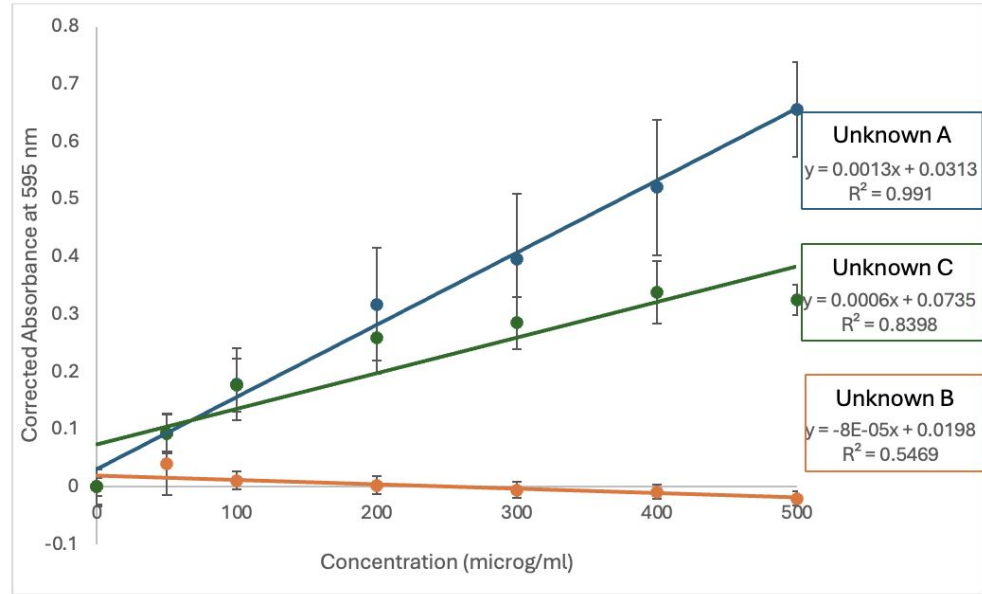
Bradford Assay

BSA would have a steeper slope and broader linear range:

-More arginine, lysine and histidine residues that strongly interact with the dye

Lysozyme may have a shorter linear range:

-high pI (11.4) makes it more prone to aggregation in the acidic conditions





Size-Exclusion Chromatography

The molecular weight cut-off (MWCO) for the spin column is 40 kDa

-BSA (MW=66 kDa) is expected to flow through the column and have a high amount of recovered protein

-Lysozyme (MW= 14.4 kDa) does not meet the cut-off

Sample	Amount of Protein (μg)
Unknown A	898.91 $\pm$ 599.77
Unknown B	86.739 $\pm$ 66.093
Unknown C	367.46 $\pm$ 245.43
BSA Standard	1034.5 $\pm$ 689.8

Semi-Qualitative (Organic Solvent Precipitate)

- Organic solvents exclude water from proteins and cause flocculation/aggregation.
- Lysozyme is Hydrophilic and interact more strongly with water
- BSA has hydrophobic surface and hydration shell is less stable
- BSA is expected to precipitate at lower organic solvent concentration
- pH precipitation did not work and made no use of salting out

Literature Precipitation Thresholds (%)

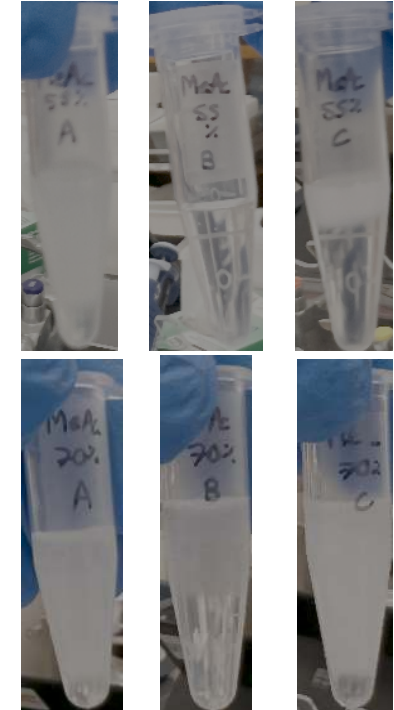
Solvent	Protein	Ppt		
		Starts	Major	Complete
EtOH	BSA	35	50	60
	Lys	50	60	70
MeAc	BSA	45	55	70
	Lys	60	70	80

Ppt Start Test



EtOH
Top 35%
Bottom 50%
A is BSA
C is Lys

Major Ppt Test

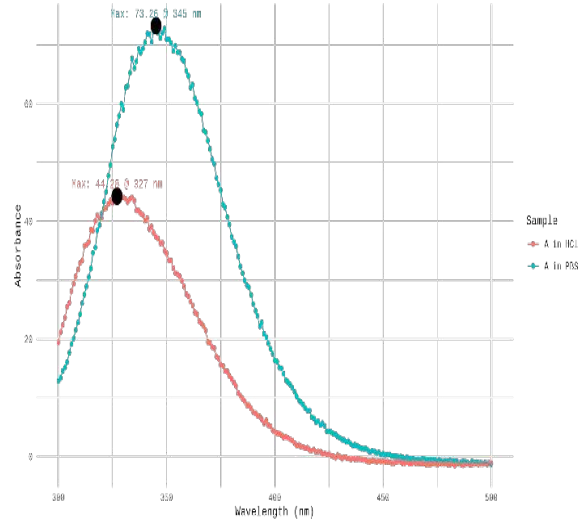


MeAc
Top 55%
Bottom 70%
A is BSA
C is Lys

Quantitative (UV-Vis)

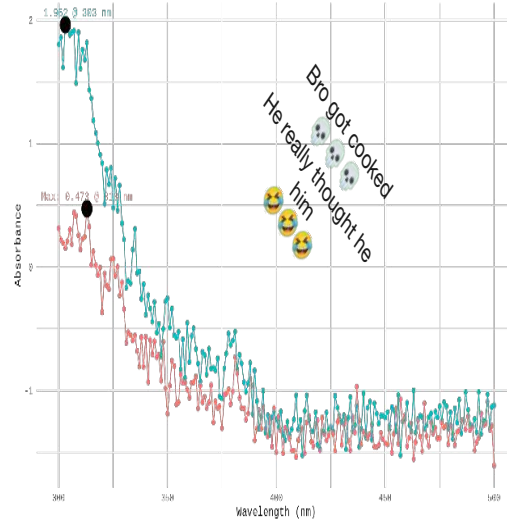
A is BSA

Absorbance vs. Wavelength (Samples A in PBS and A in HCl)



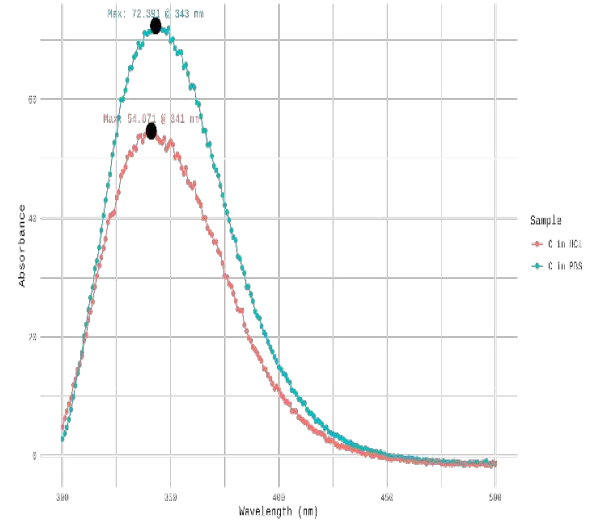
B is Unknown

Absorbance vs. Wavelength (Samples B in PBS and B in HCl)



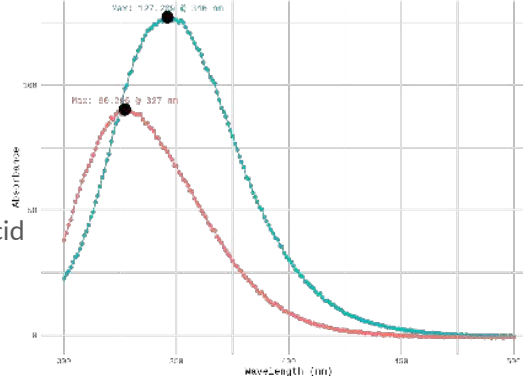
C is Lysozyme

Absorbance vs. Wavelength (Samples C in PBS and C in HCl)

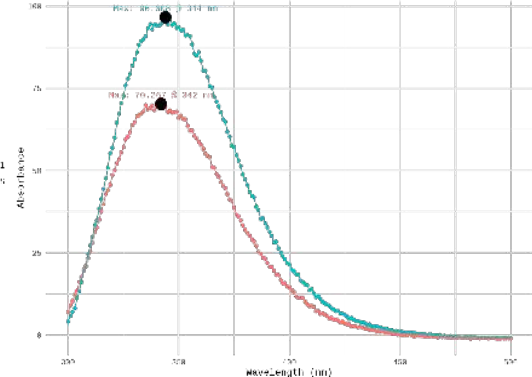


- PBS (pH 7.4) and HCl (pH 2)
- BSA is larger and more complex, held by ionic interactions
- Lysozyme is small and compact, held by covalent interactions
- Conformational change more important in BSA in Acid
- Tryptophan Fluorescence affected by environment
- Blue-Shift as Trp residues in BSA are more shielded from solvent.

Absorbance vs. Wavelength (Samples BSA Std in PBS and BSA Std in HCl)



Absorbance vs. Wavelength (Samples Lys Std in PBS and Lys Std in HCl)



Summary of Results of Each Fraction

Fraction
Std BSA
Std Lys
Unk A
Unk B
Unk C



References

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