

Expression and Characterization of a Metalloprotein in Bacteroides Fragilis

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BF4112: Orphan protein BF4112 (from *B. fragilis*) was identified from the protein data bank as a possible Cys-Tyr containing protein. BF4112 is a metalloprotein and contains a cysteine residue (Cys98) in close proximity to a tyrosine residue (Tyr52), as well as a nearby metal binding site.

Figure 1: Crystal structure of apoBF4112.

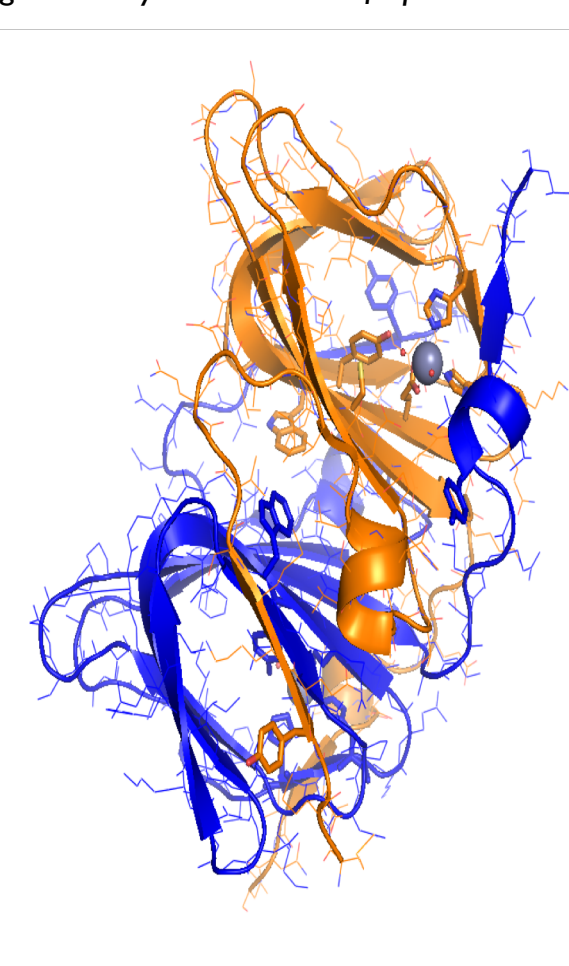
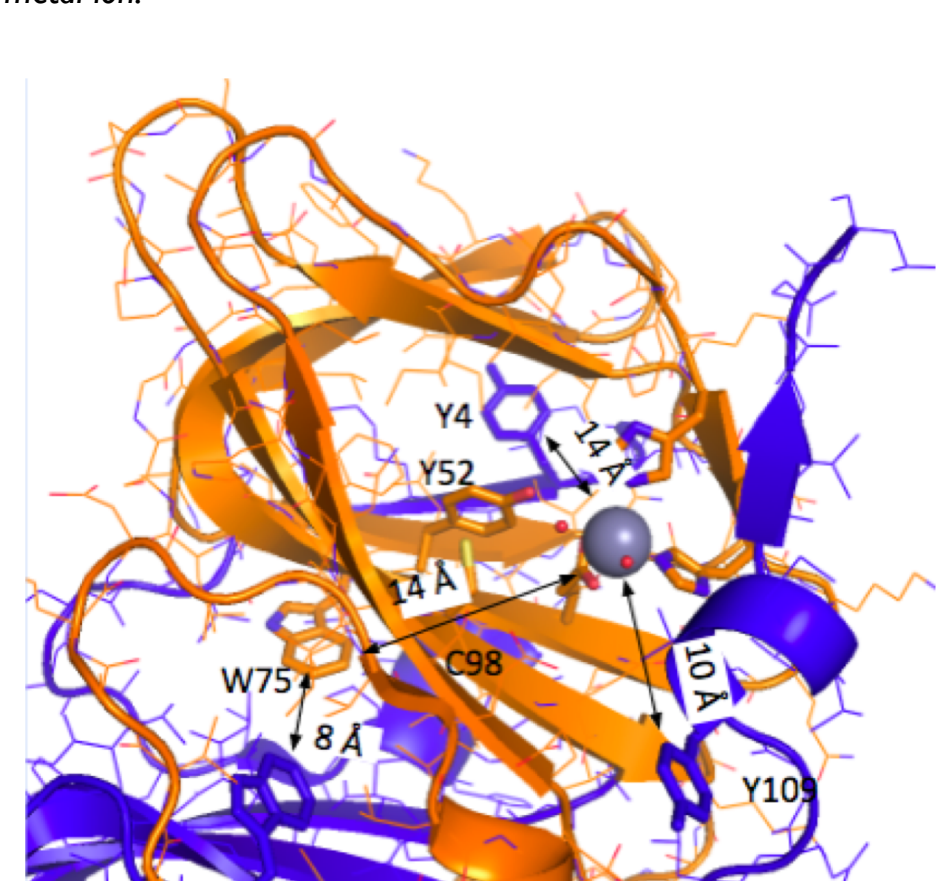


Figure 2: apoBF4112 active site. Distances between key residues shown. Cross-link forms between Y52 and C98. Sphere represents metal ion.



Cys-Tyr: Cysteine-Tyrosine modifications are a specific class of tyrosine PTM and remain relatively uncharacterized. We suggest that Cys-Tyr modifications may be more abundant than previously thought. The relevant thioether functional group forms at the 3' carbon of a tyrosine residue. Importantly, the few Cys-Tyr PTMs that have been identified are redox-active cofactors.

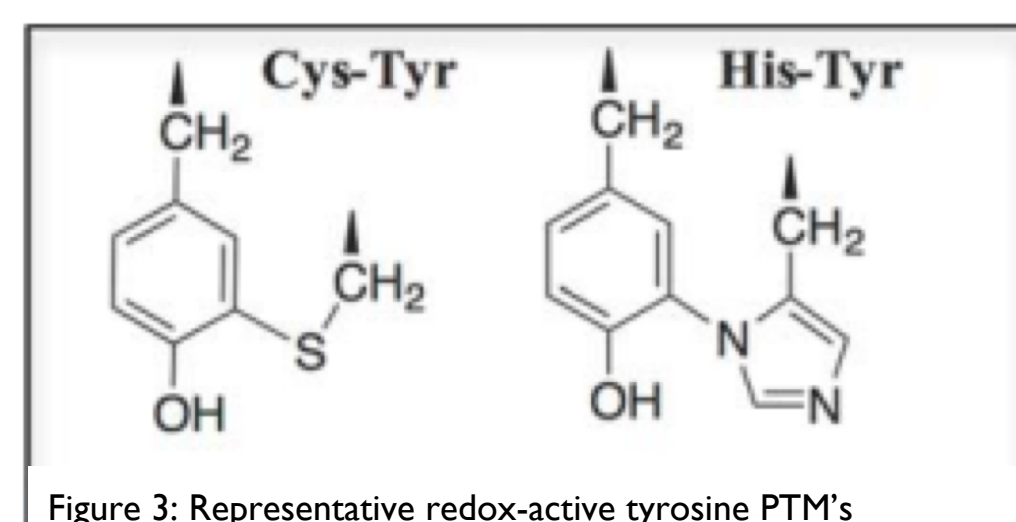
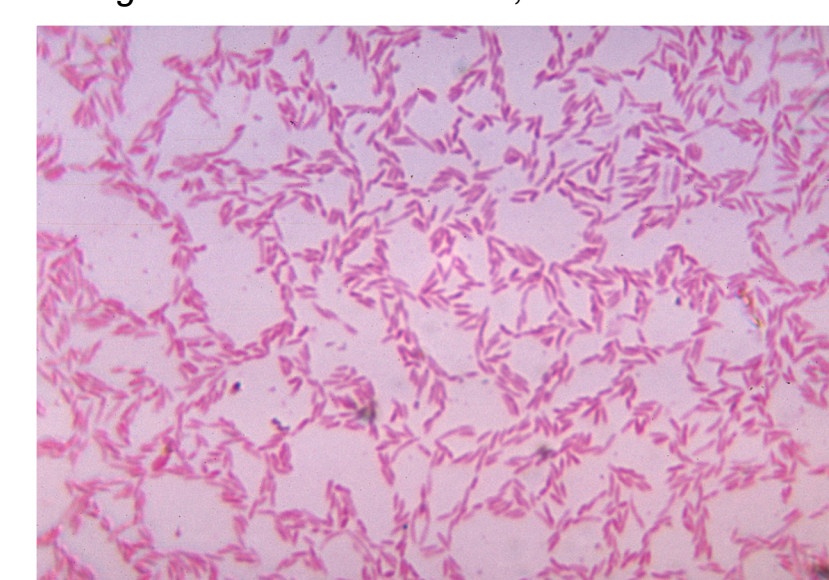


Figure 3: Representative redox-active tyrosine PTM's

Bacteroides fragilis: B.fragilis

Bacteroides fragilis is an anaerobic gut bacteria that is the most common isolate from pathogenic gastrointestinal infection. While *B. fragilis* is an anaerobe, its aerotolerance is well documented and likely contributes to its pathogenicity. While we have expressed BF4112 in *E. coli*, we sought to express BF4112 in *B. fragilis* to characterize the protein in its native environment. Characterization includes: whether Cys-Tyr forms, the native metal ion, and the function of the protein.

Figure 3: *Bacteroides fragilis* Image: CDC/Dr. V.R. Dowell, Jr.



Methods

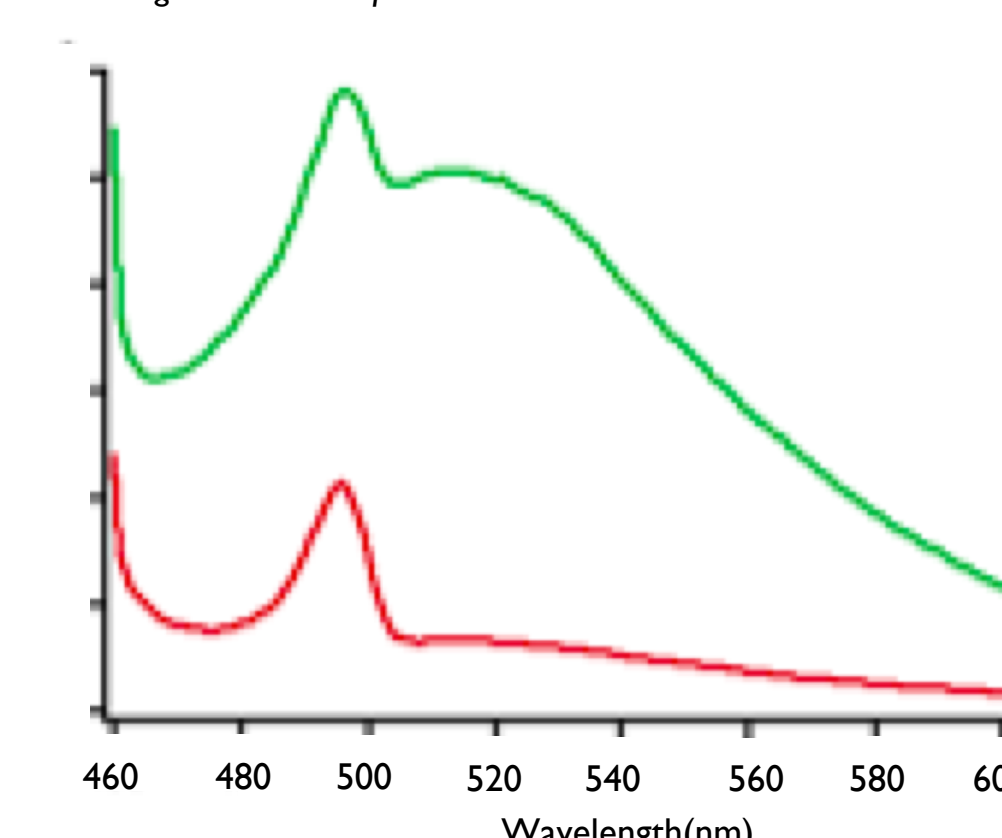
The following procedure was developed and optimized to express BF4112 in *Bacteroides fragilis*:

- Electrocompetent *B. Frag* cells were prepared and subsequently electroporated with His-BF4112 or pER 153 plasmid DNA.
- Transformants grew on TSA plates supplemented with hemin, cysteine, and erythromycin for 2 days.
- Single colonies were grown to 1 liter (or 25mL) anaerobically. Maltose was added to induce expression.
- BF4112 IL cultures were pelleted and suspended in Tris buffer before lysis using French Press.
- Cell lysate was purified using affinity chromatography
- Protein was concentrated, ran on SDS PAGE, and analyzed for iron content using the iron-binding agent ferrozine.
- pER 153 cultures (25mL) were lysed using sonication and analyzed using a fluorimeter.

Expression

The flavin-mononucleotide-fluorescent-based protein (FbFP) encoded by plasmid pER 153 (with maltose inducible promoter) was used to validate and optimize expression. pER 153 was subcloned to express His-BF4112. Experiments tested the effects of maltose addition and optimized the time period and amount of maltose for maximal protein expression in *B. fragilis*.

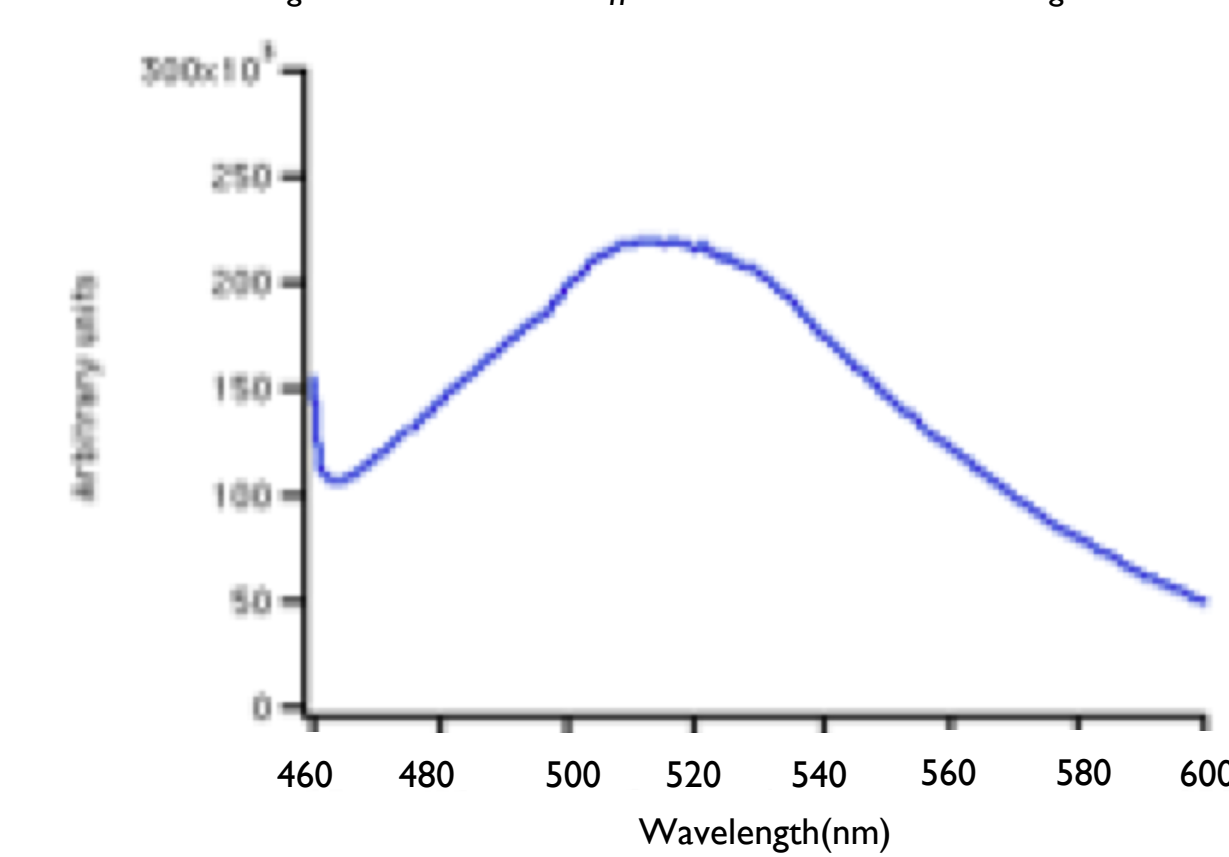
Figure 4: FbFP expression +/-maltose.



Legend: +Maltose (.5%) (green line), -Maltose (red line)

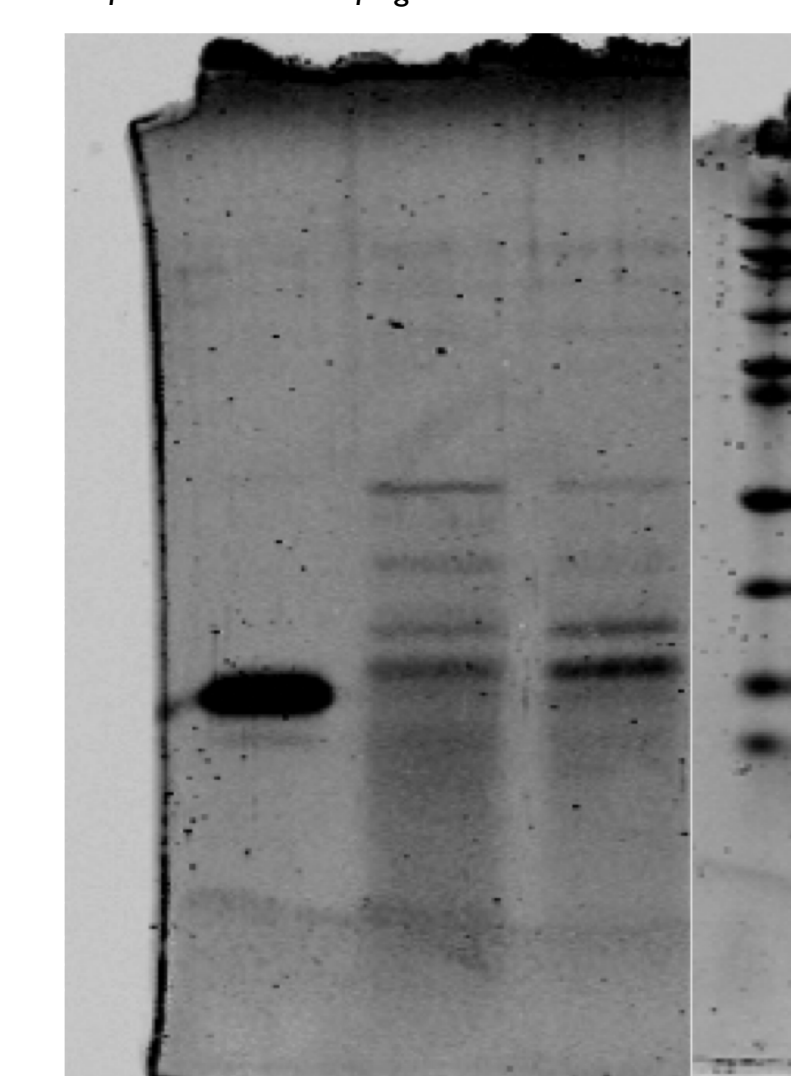
- Addition of maltose at .5% shows significant fluorescence increases with a maximum at 513nm
- Expression is relatively constant from 15-30 hours.

Figure 5: Fluorescence difference between two lines in Figure 5.

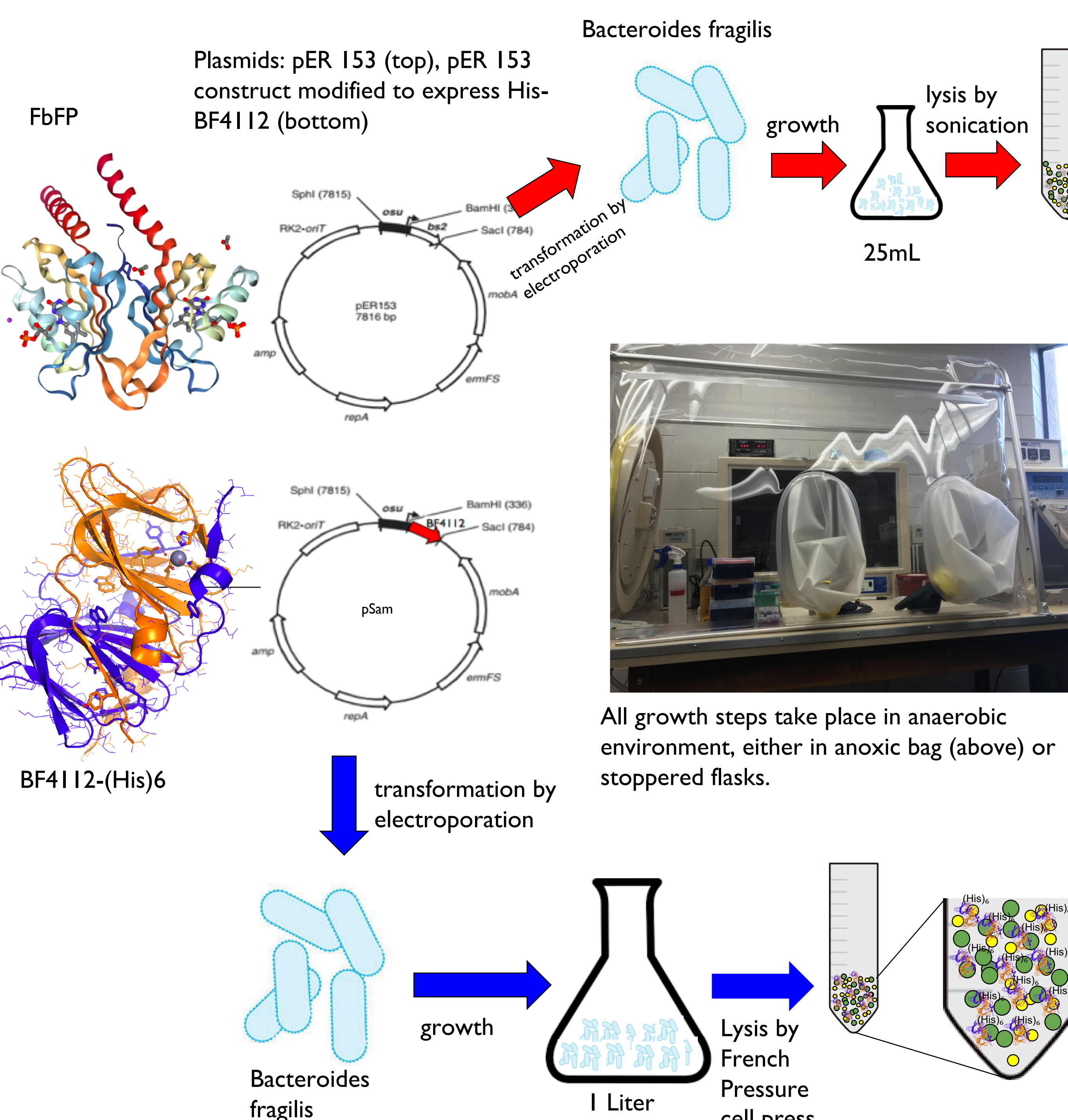


Time (h)	Fluorescence
15	1.45E+06
18	1.35E+06
21	1.05E+06
24	1.36E+06
27	1.19E+06
30	1.33E+06

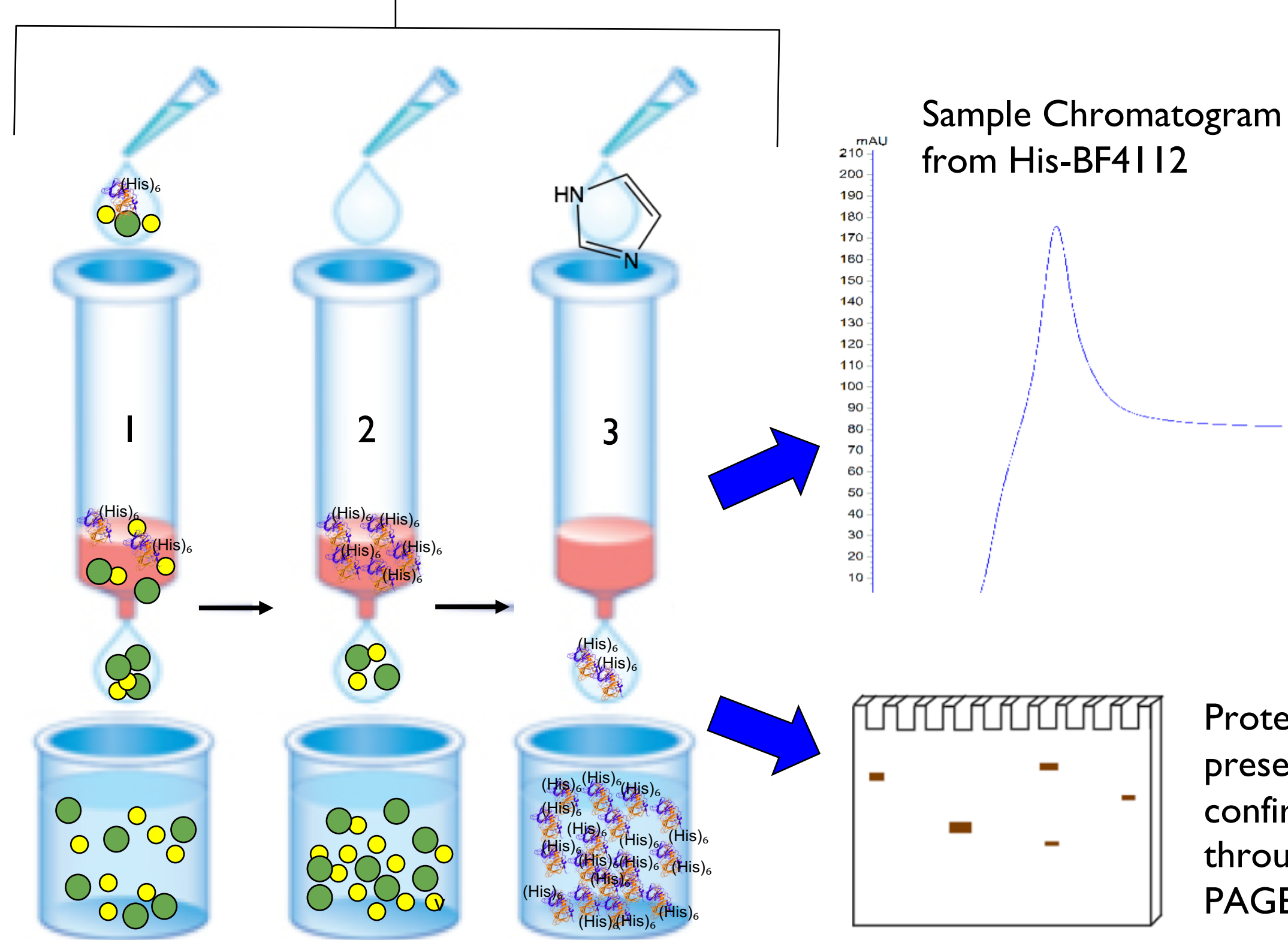
Figure 6: SDS PAGE gel showing maltose induced His-BF4112 expression, isolated from *Bacteroides fragilis*.



His-BF4112 was isolated and concentrated using NanoSep concentrator tubes. SDS PAGE confirmed protein between 10 and 15kDa and in the range of apoBF4112. Double-banding indicates the slight electrophoretic shift caused by Cys-Tyr formation. This finding confirms the expression of BF4112 in *Bacteroides fragilis* and indicates the likely formation of Cys-Tyr in native protein.



IMAC Affinity Chromatography: 1) Sample is applied to column 2) Column is washed to remove extraneous proteins, His-tagged protein stays bound 3) His-tagged protein is eluted by imidazole



- Conclusions:
- BF4112 expression can be induced in *Bacteroides fragilis* using a plasmid construct
 - Native BF4112 can be isolated and harvested
 - Cys-Tyr likely forms in native BF4112

- Future Directions:
- Confirmation of Cys-Tyr in native BF4112 by mass spectrometry
 - Further experiments into BF4112 function
 - Two hypotheses: Antioxidant or antibiotic-binding protein
 - Development of CRISPRi system to silence the gene for functional experiments
 - Confirmation of native metal ion

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