

Development of an equine IFN γ release assay (IGRA) for glanders

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すべてはキミの未来のために。



SCIENCE is here, the future is here.

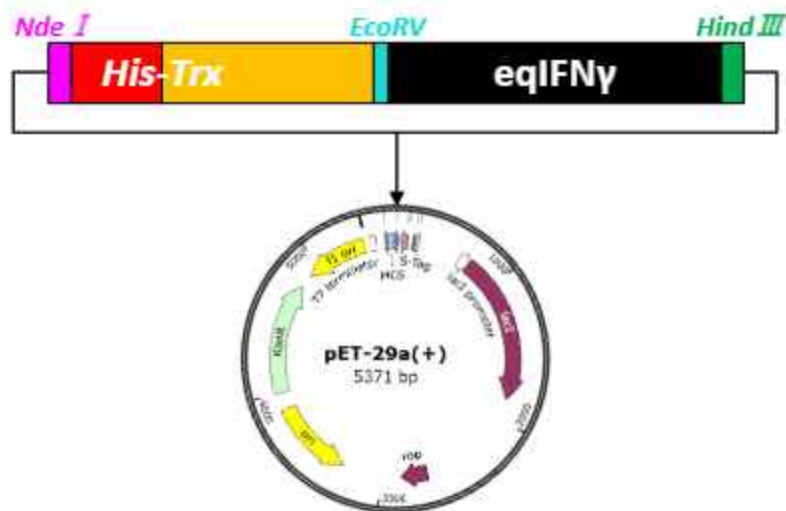
岡山理科大学

OKAYAMA UNIVERSITY OF SCIENCE

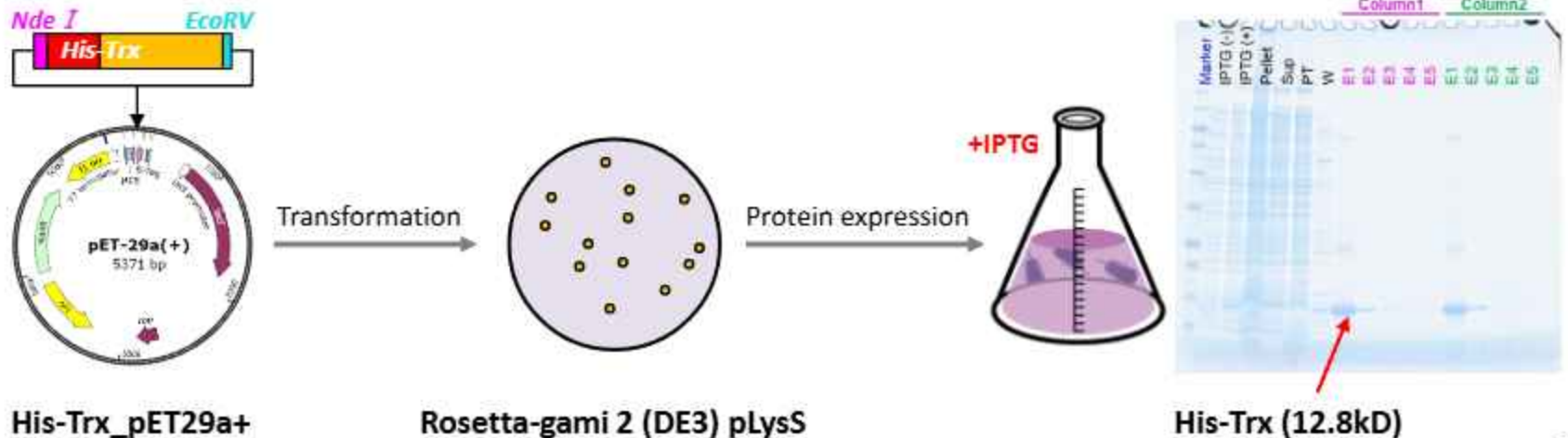
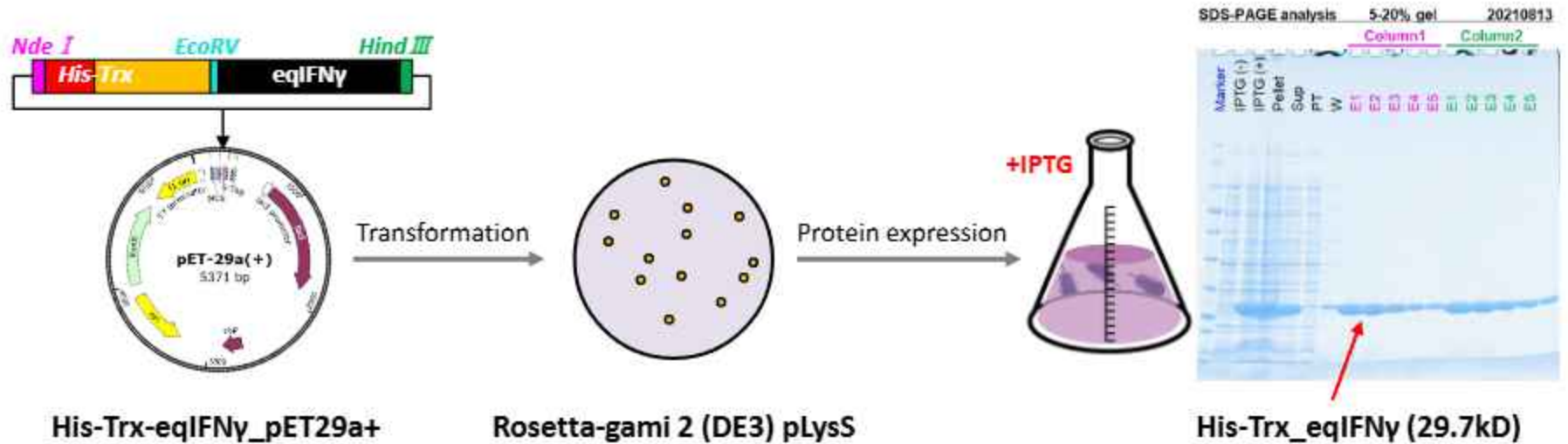
1. The eqIFN γ DNA was synthesized by referring to the nucleotide sequence from NCBI

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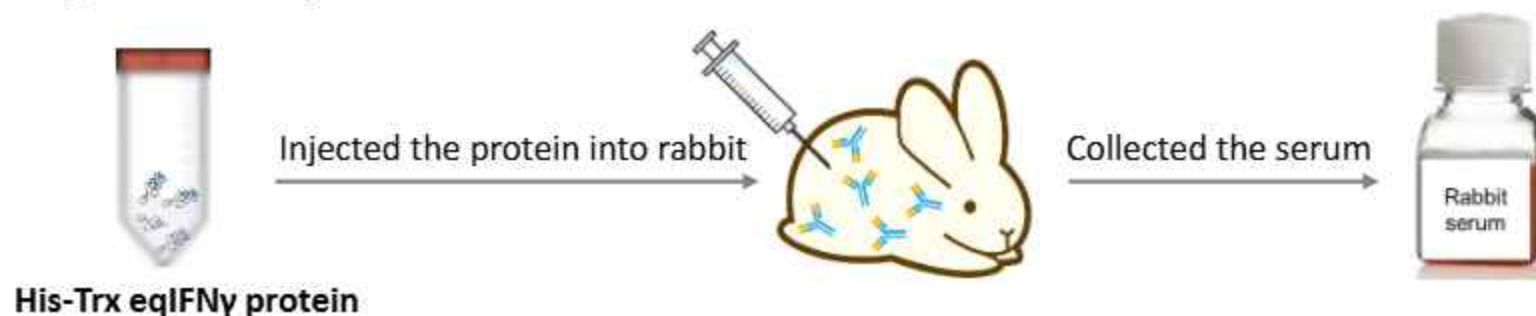
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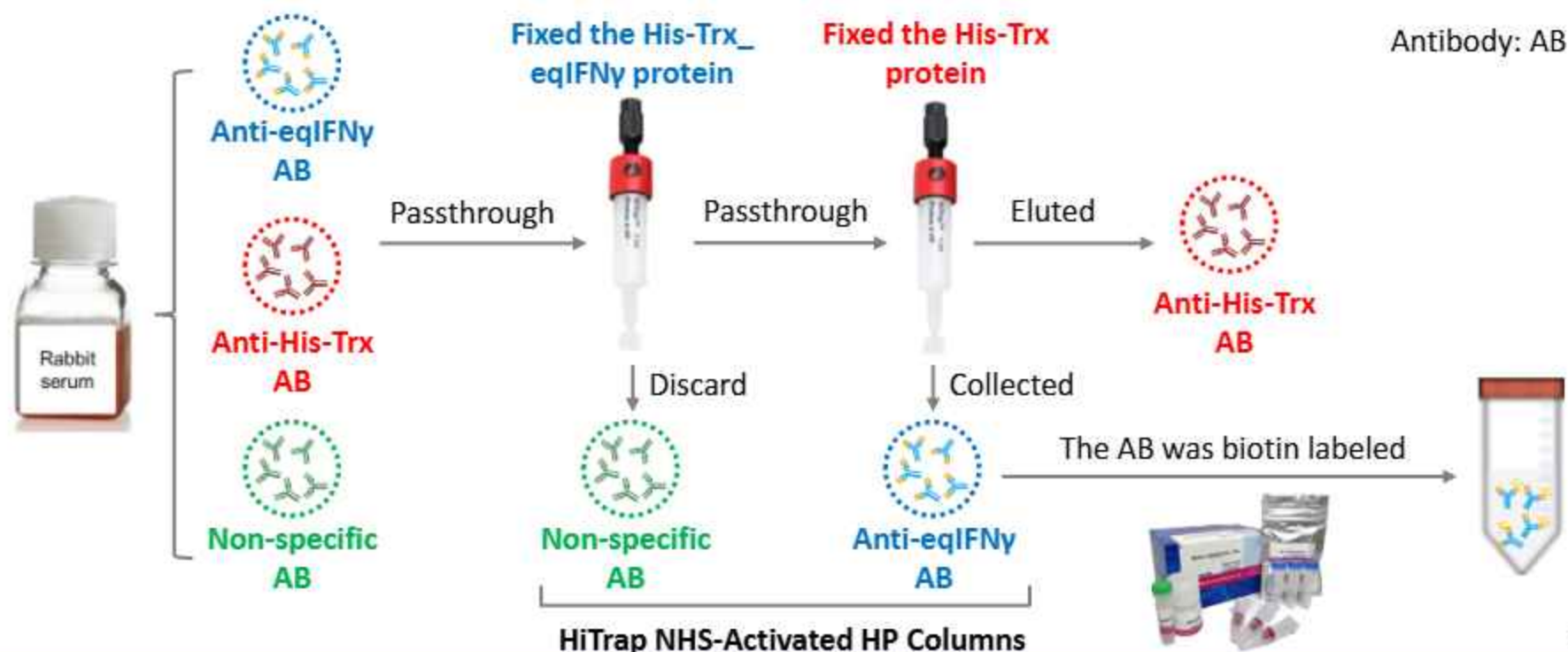
2. His-Trx_eqIFN γ and His-Trx were Inserted into the pET29a+ plasmid and the proteins were expressed, respectively



3. Injected the protein into rabbit and collected the serum



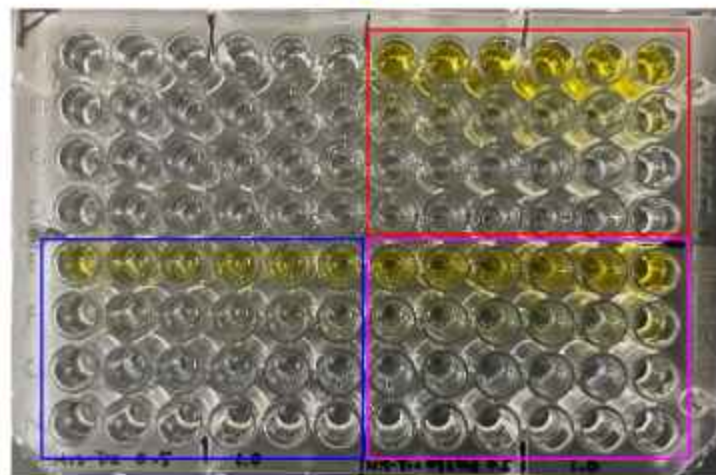
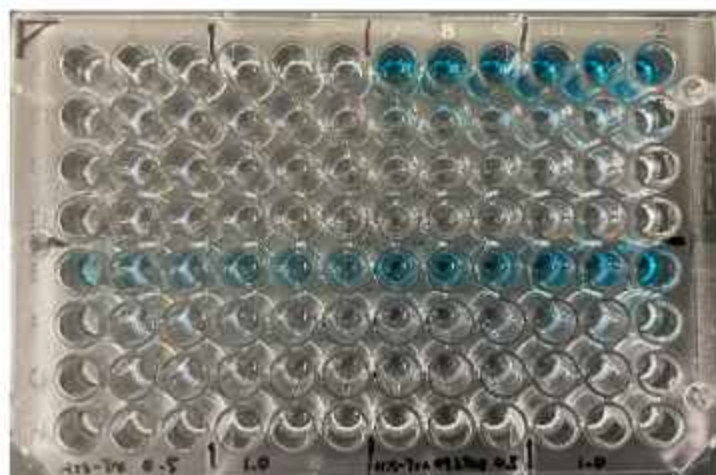
4. Specific AB were purified from serum, and the anti-eqIFN γ AB was biotin labeled



































































































3/20

Biotin Labeling Kit-NH $_2$

5. The eqIFN γ protein specific antibody was confirmed by ELISA method



The binding reaction between His-Trx protein and His-Trx-specific antibody

	1	2	3	4	5	6	7	8	9	10	11	12		
A													100 ng/mL	Anti-eIFN γ
B													10 ng/mL	
C													1 ng/mL	
D													Blank	
E													100 ng/mL	Anti-eIFN γ Anti-H-Trx
F													10 ng/mL	
G													1 ng/mL	
H													Blank	
	0.5 μ g/mL			1.0 μ g/mL			0.5 μ g/mL			1.0 μ g/mL				
	His-Trx						His-Trx-eIFN γ							
	Ag Coating (proteins)													Purified antibody conc.

The binding reaction between His-Trx-eIFN γ protein and eIFN γ -specific antibody

The binding reaction between His-Trx-eIFN γ protein and His-Trx-specific antibody

6. ELISA Development of IGRA for glanders



Overnight at 4°C



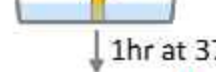
1hr at 37°C



1hr at 37°C



1hr at 37°C



0.5hr at 37°C



0.5hr at 37°C



0.5hr at 37°C



0.5hr at 37°C



0.5hr at 37°C



- Capture Ab: **Anti-eqIFNy Ab**

- Blocking: Block Ace

- Wash, add standard
His-Trx eqIFNy protein

- Wash, add detection Ab
Biotin labeled anti-eqIFNy Ab

- Wash, add Ultra-Sensitive
ABC Peroxidase

- Wash, add TMB
Substrate Solution

- Stopped the reaction
Read the OD value (450nm)

Table1. Coating the Biotin non-labeled anti-eqIFNy AB into the 96 well plate

	1	2	3	4	5	6	7	8	9	10	11	12
A	Coating anti-eqIFNy AB 4ug/ml			Coating anti-eqIFNy AB 2ug/ml			Coating anti-eqIFNy AB 4ug/ml			Coating anti-eqIFNy AB 2ug/ml		
B												
C												
D												
E												
F												
G												
H												

Table2. Dilution of standard (His-Trx eqIFNy protein)

Standard	St1	St2	St3	St4	St5	St6	St7	Blank
Conc. (ng/ml)	30	10	3	1	0.3	0.1	0.03	0



Reaction was slower than in general ELISA



1. To improve sensitivity and specificity, eqIFN γ was expressed as three fragments

eqIFN γ 1

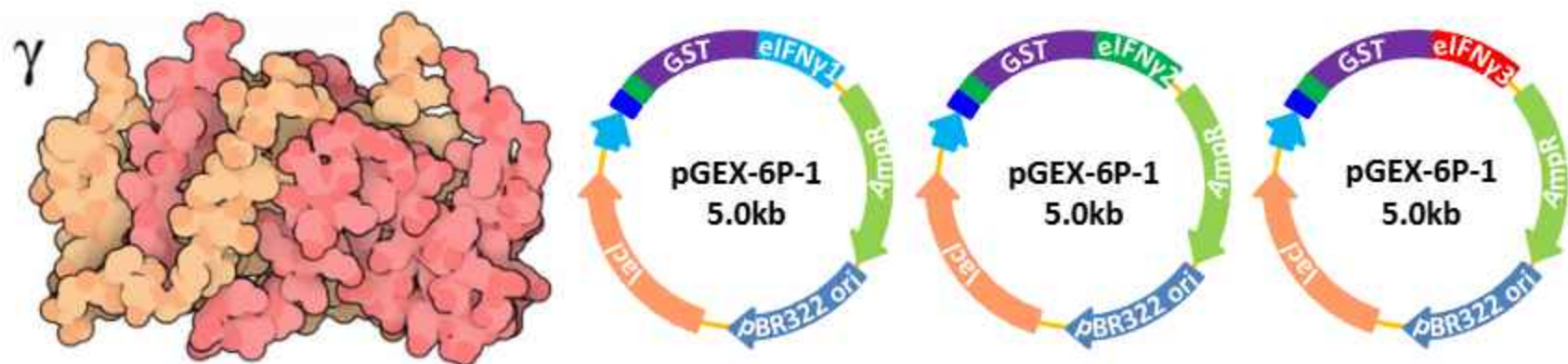
QAAFFKE IENLKEYFNA SNPDVGDGGP LFLDILKNWK EDSDKKI IQS

eqIFN γ 2

QIVSFYFKLF ENLKDNQVIQ KSMDTIKEDL FVKFFNSSTS KLEDFQKLIQ

eqIFN γ 3

IPVNDLKVQR KAISELIKVM NDLSPKANLR KRKRSQNPFR GRRALQ



- 1) Fragmented the eqIFN γ cDNA into three parts: eqIFN γ 1, eqIFN γ 2, and eqIFN γ 3.
 - 2) Inserted them into pGEX-6p-1 plasmid for individual protein expression.
- ⇒ To separately purify specific antibodies against epitopes on each fragment protein and develop a high-sensitivity, high-specificity ELISA assay.

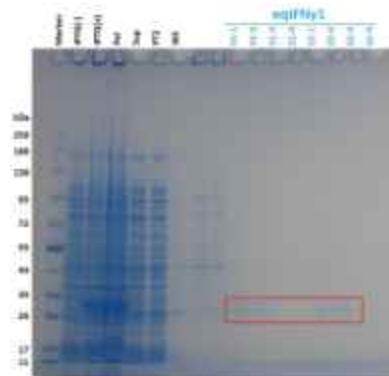
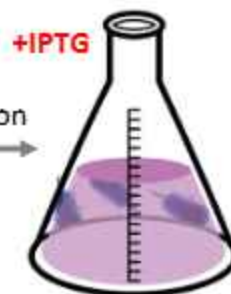
1-1. Purification of eqIFN γ 2 and 3 was successful, but eqIFN γ 1 was not purified



Transformation

Competent Cells BL21

Protein expression



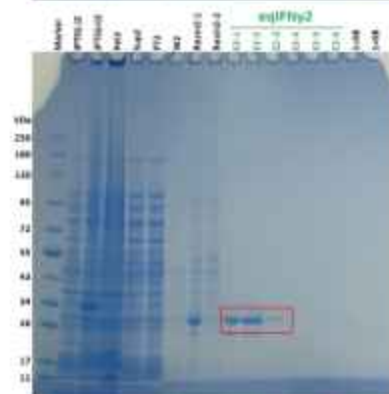
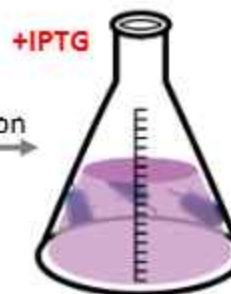
The target protein band



Transformation

Competent Cells BL21

Protein expression



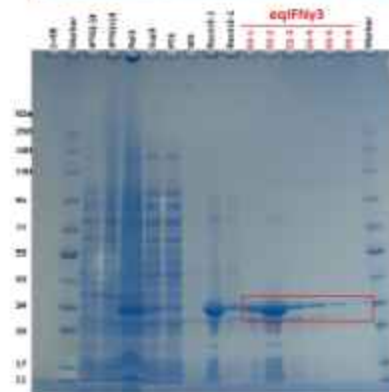
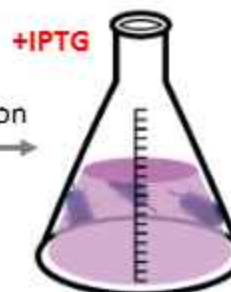
The target protein band



Transformation

Competent Cells BL21

Protein expression

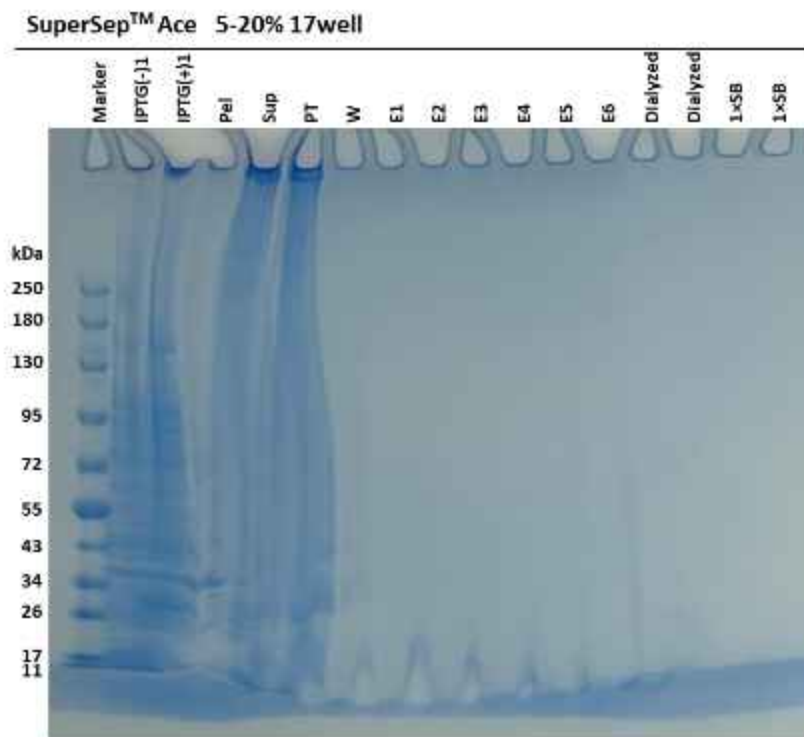
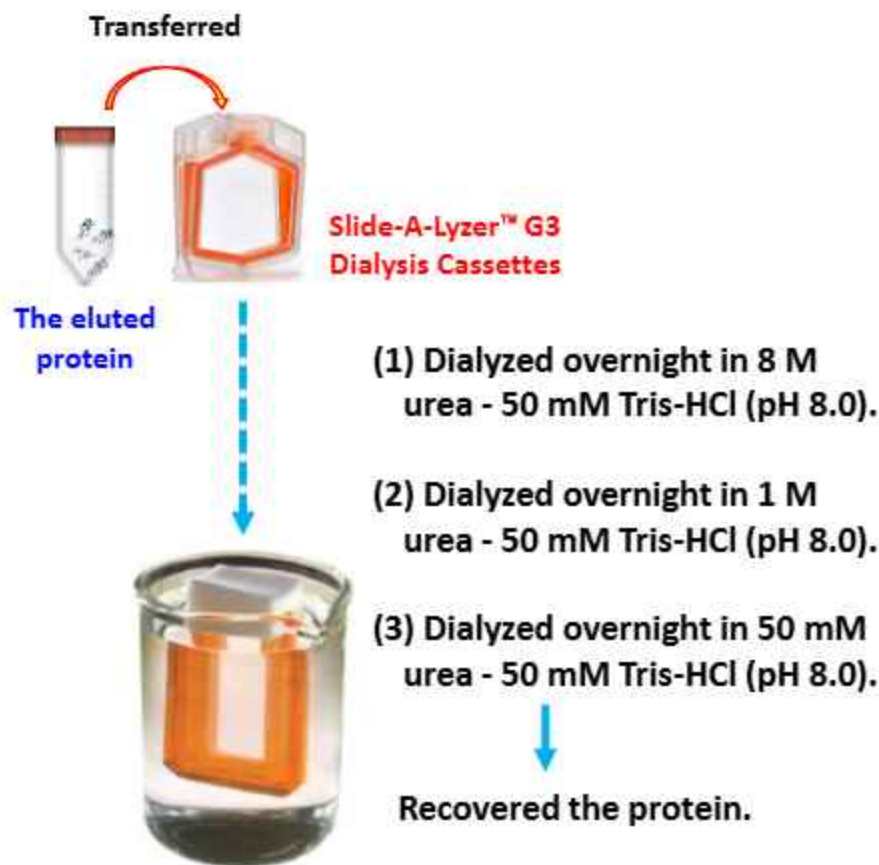


The target protein band



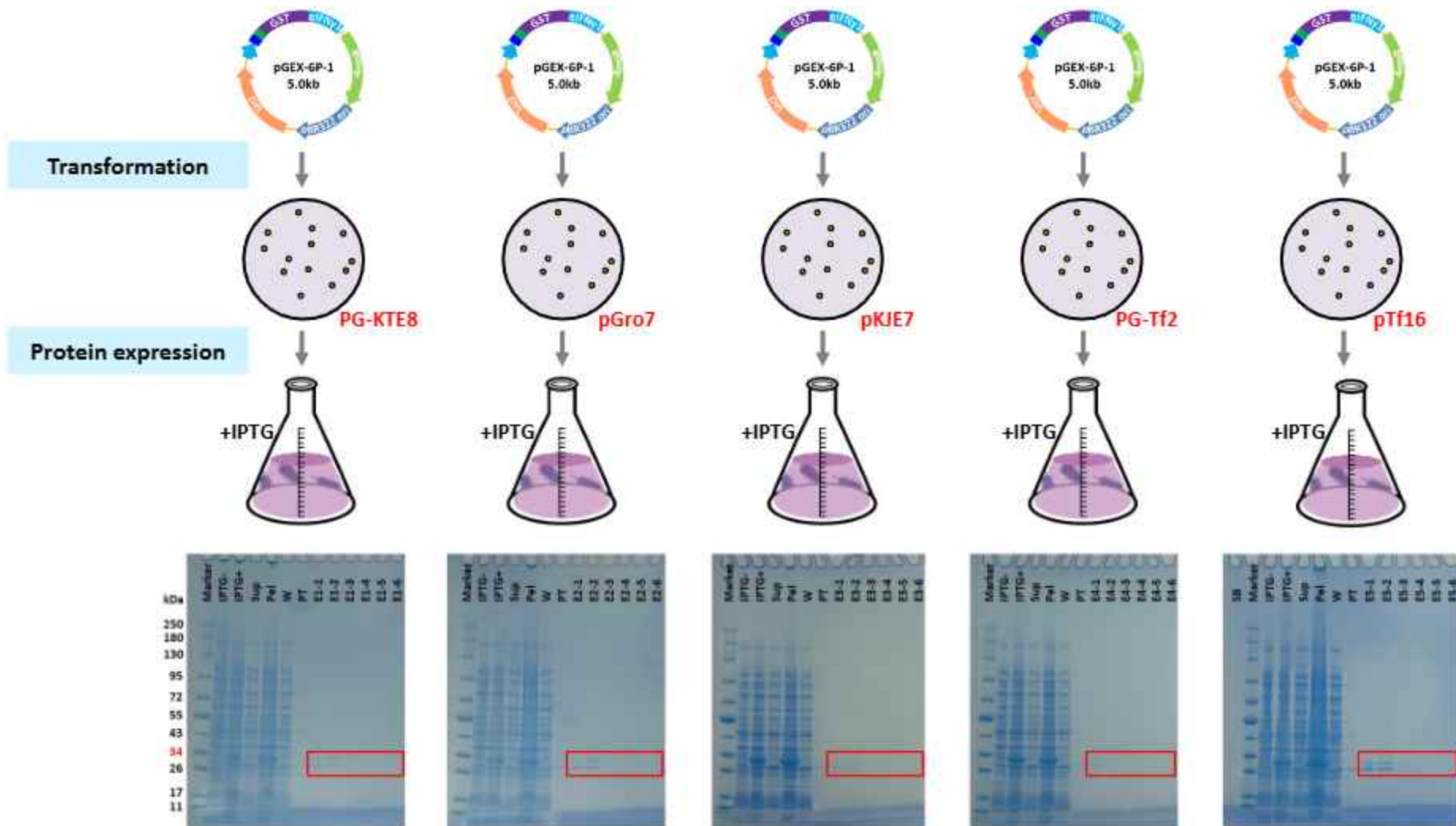
1-2. Purified eqIFN γ 1 under denaturing conditions and then performed dialysis

- 1) The cells were lysed in 6 M Guanidine hydrochloride, 50 mM Tris-HCl (pH 8.2), and the protein purified.
- 2) Supernatant after sonication was dialyzed in three steps.



The target band was not detected in the SDS-PAGE analysis.

1-3. eqIFN γ 1 was expressed & purified using Chaperone Competent Cells BL21 Set



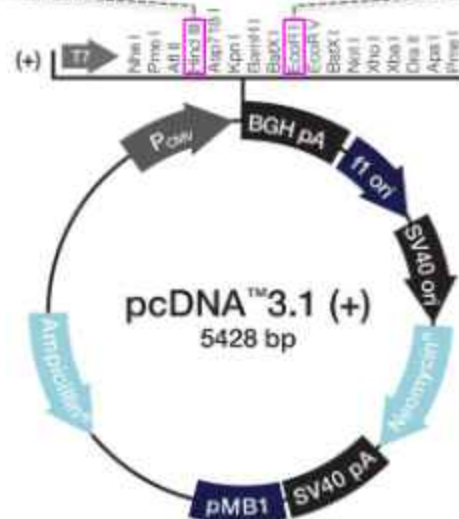
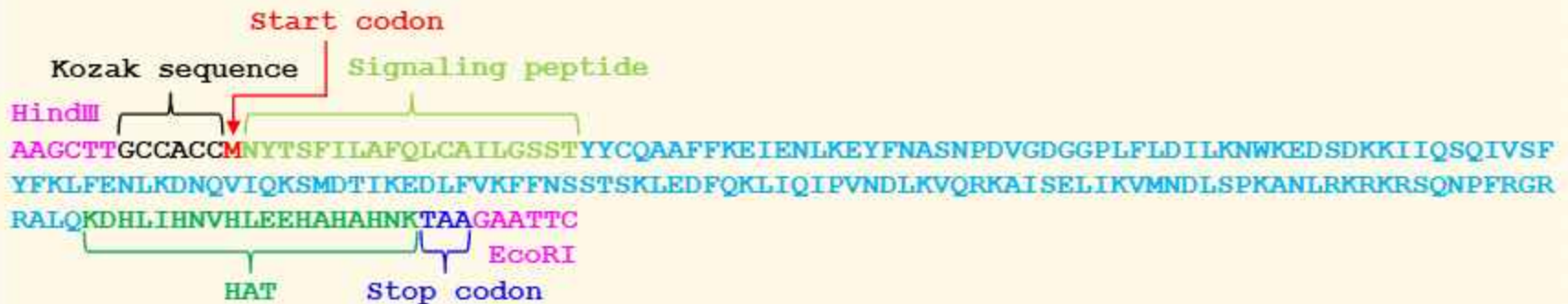
Co-expression with five chaperone plasmids was performed to promote solubilization, but it was unsuccessful.



2. eqIFN γ was expressed in mammalian cells

HindIII-eqIFN γ -HAT-EcoRI

HAT (Lys-Asp-His-Leu-Ile-His-Asn-Val-His-Lys-Glu-Glu-His-Ala-His-Ala-His-Asn-Lys)



To form the correct structure and improve antibody specificity, eqIFN γ cDNA was inserted into the pcDNA3.1 plasmid and expressed in mammalian cells.

10/20



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2-1. eqIFN γ -HAT protein expression in mammalian cells (FreeStyle CHO-S cells)

Day0



- (1) 24 h before transfection, dilute FreeStyle CHO-S cells to $5-6 \times 10^6$ cells/ml and seed into a shake flask.

At 8% CO₂, 37°C, and shaking at 135 rpm for 24 h

Day1



- (3) Just before transfection, dilute the cells to 1×10^6 cells/ml.

- (2) Dilute plasmid DNA and FreeStyle MAX reagent in OptiPRO™ SFM medium separately, then combine and incubate at room temp. for 15 mins.



- (4) Slowly add 1.2 ml of the DNA-lipid mixture while gently mixing the cells.

At 8% CO₂, 37°C, and shaking at 135 rpm for 24 h

Day2

Day3

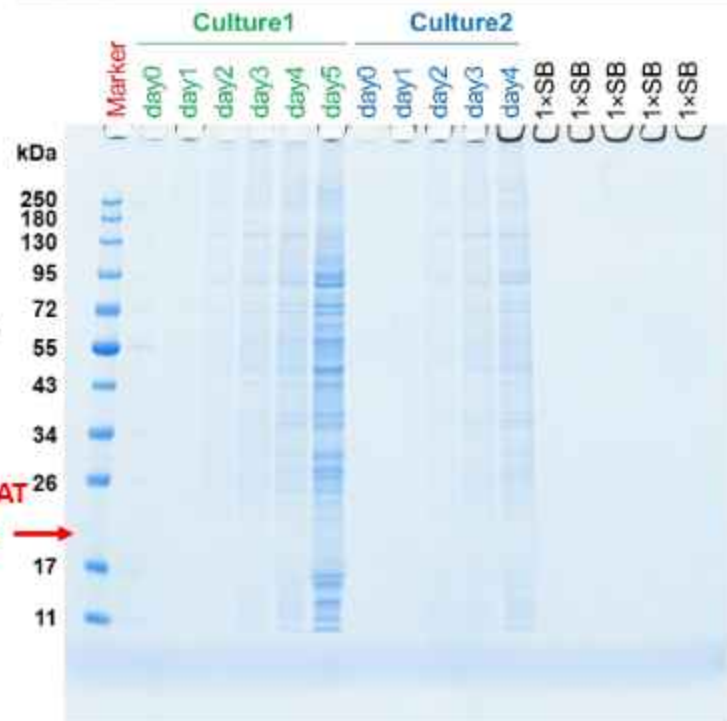
Day4

Day5



- (5) Perform cell counting and sampling every 24 hours until day 5.

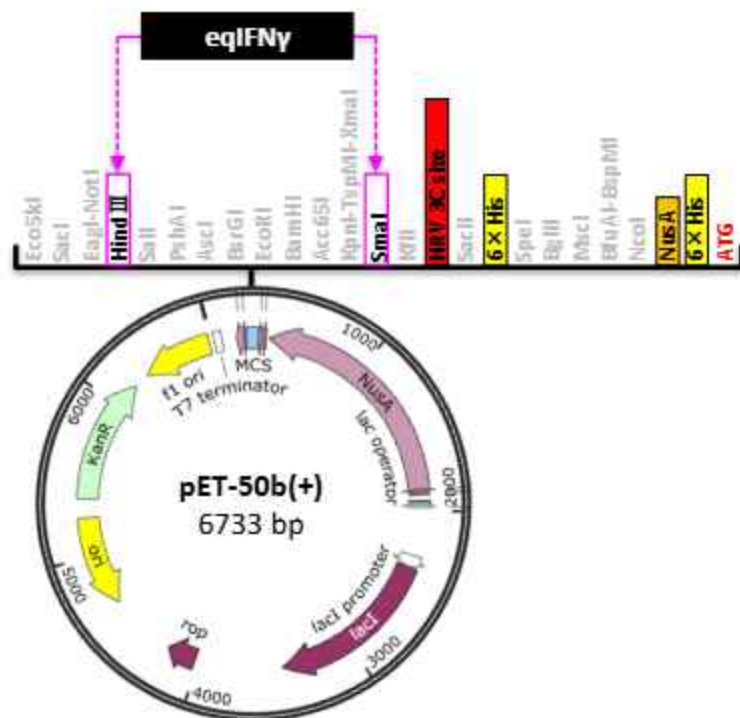
SuperSep™ Ace 10-20% 17well



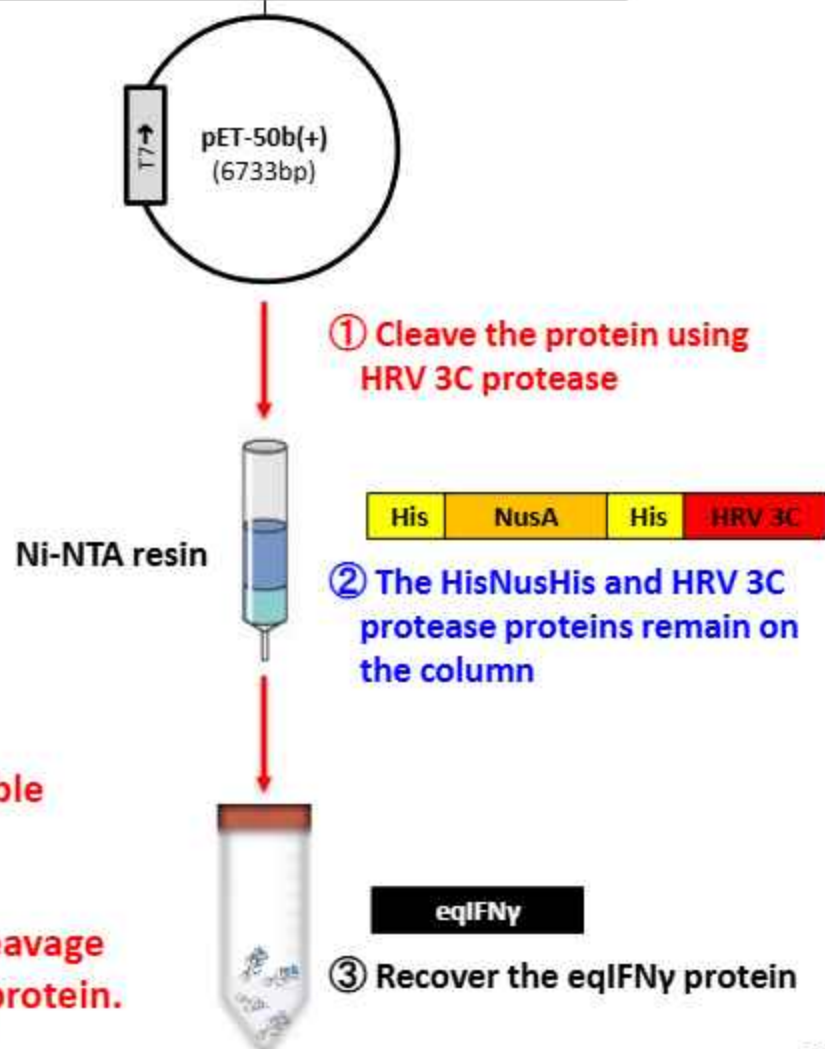
The target band was not detected in the SDS-PAGE analysis.



3. eqIFN γ was expressed by inserting it into the pET-50b(+) plasmid



- ✓ This plasmid contains an NusA-tag, allowing for stable protein expression.
- ✓ Additionally, it contains an HRV 3C site, allowing cleavage by the HRV 3C protease to recover only the eqIFN γ protein.

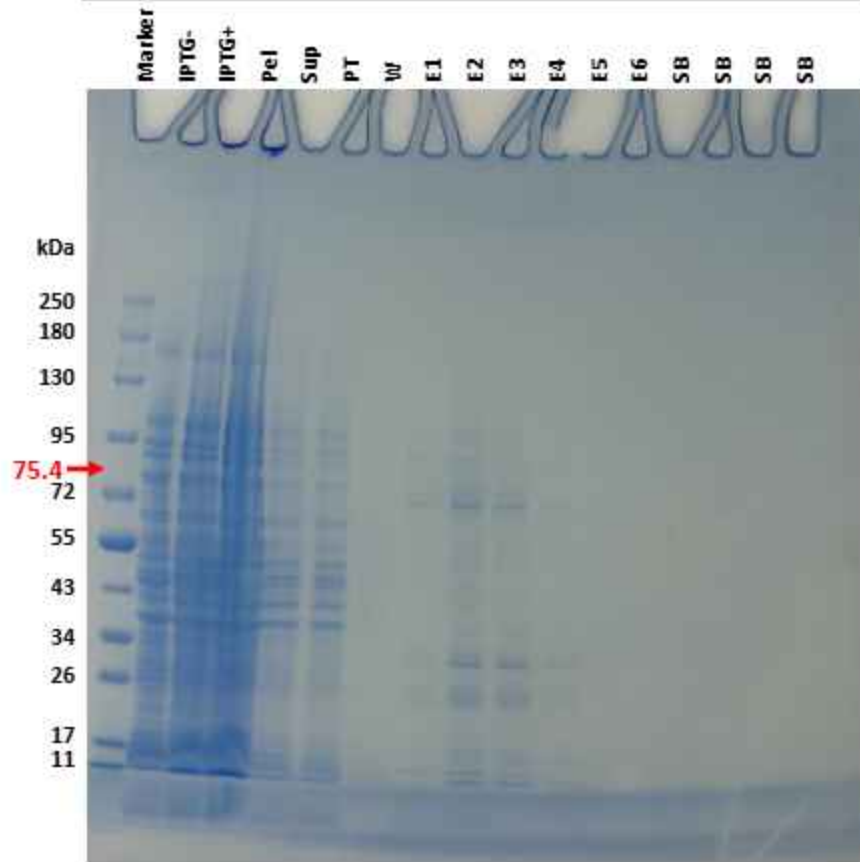


3-1. The plasmid was transformed into two types of competent cells, **BL21** and **Rosetta-gami™ 2(DE3)**, for protein expression

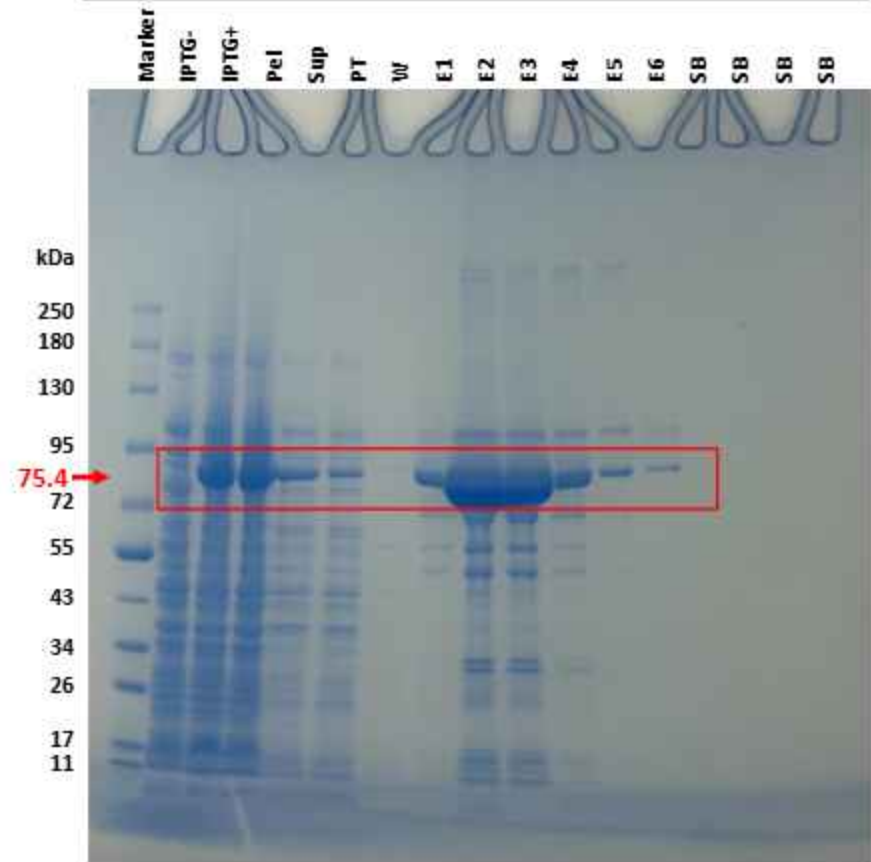
SuperSep™ Ace 5-20% 17well

eqIFN γ NUS-His \div 75.4kDa

eqIFN γ NUS-His_pET-50b(+)_BL21

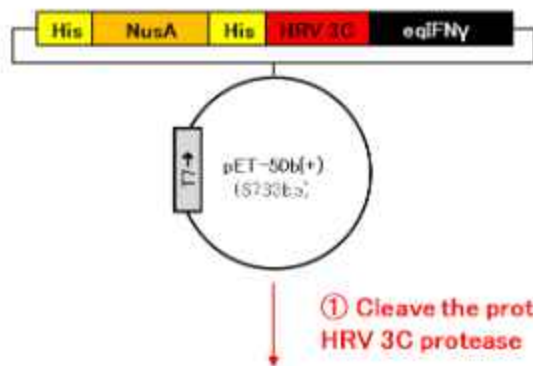


eqIFN γ NUS-His_pET-50b(+)_Rosetta-gami™ 2(DE3)



The results showed target protein expression in **Rosetta-gami™ 2(DE3)**, while no expression was observed in **BL21**.

3-2. Verification of protein cleavage conditions



Protein cleavage reaction (Buffer: 50mM Tris-HCl (pH8.0), 200mM NaCl)

Protease: Whole protein	1 : 2	1 : 5	1 : 10	1 : 20	1 : 50
Whole protein (ul)	3.6	8.9	17.9	35.7	89.3
HRV 3C protease (ul)	1.0	1.0	1.0	1.0	1.0
Buffer (ul)	45.4	40.1	31.1	13.3	-
Total (ul)	50	50	50	50	90.3

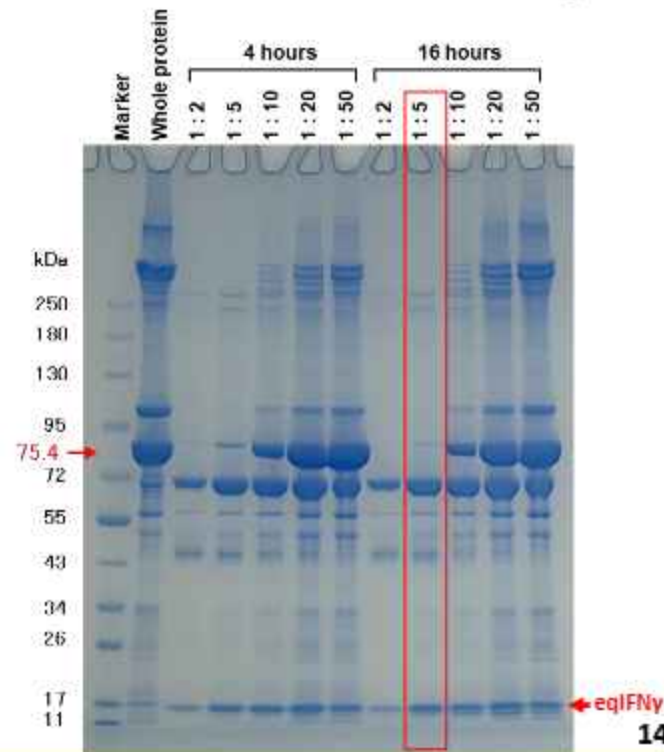
At 4°C, and rotating at 2 rpm for 16 h

※ The samples were collected at 4 hours and 16 hours, respectively, to check the cleavage status

Whole protein conc.: 1.4 mg/ml
Protease conc.: 2.5 mg/ml

The best result was obtained by cleavage at 4°C, with a 1:5 ratio, for 16 hours

Whole protein (eqIFNgNUS-His_pET-50b(+)) \approx 75.4 kDa;
eqIFNy protein \approx 16.7 kDa;
His-NUS protein \approx 58.7 kDa;
HRV 3C protease protein \approx 50.3 kDa



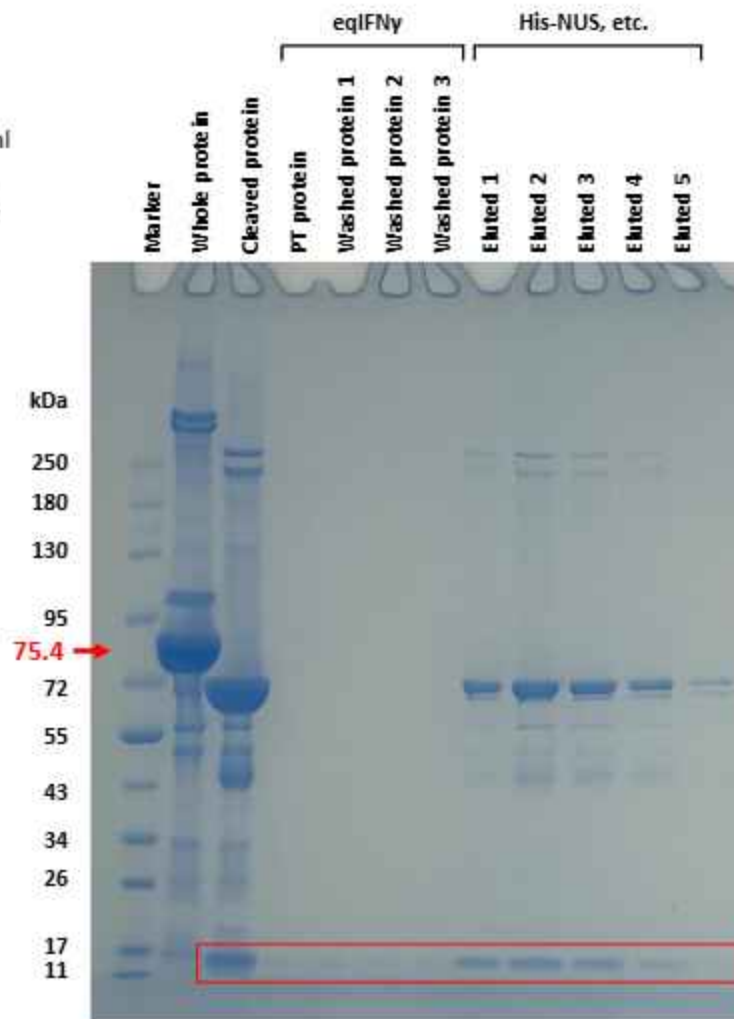
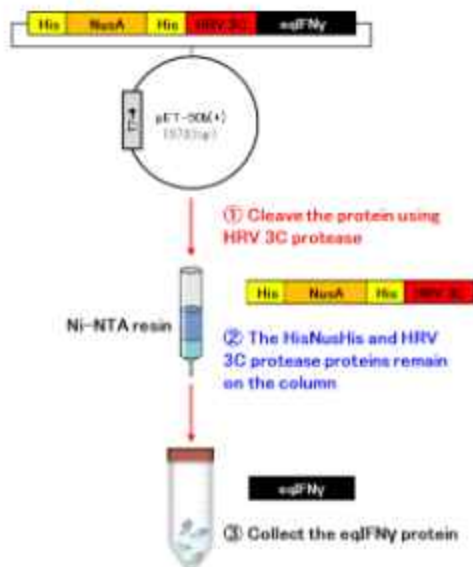
3-3. Separation and purification of cleaved protein by Ni-NTA column

Protein cleavage reaction

Protease: Whole protein	1 : 5
Whole protein (ul)	178.6
HRV 3C protease (ul)	20.0
Buffer (ul)	51.4
Total (ul)	250

Whole protein conc.: 1.4 mg/ml
Protease conc.: 2.5 mg/ml
Buffer: 50mM Tris-HCl (pH8.0),
200mM NaCl

At 4°C, and rotating at 2 rpm for 16 h



Since His-tags were attached to proteins other than eQIFNy, these proteins were expected to remain bound to the Ni-NTA column, while only eQIFNy was supposed to be recovered in the flow-through. **However, eQIFNy also remained on the Ni-NTA column and was not separated from the other proteins.**

3-4. Separation and purification of cleaved protein by Amicon® Ultra Filter

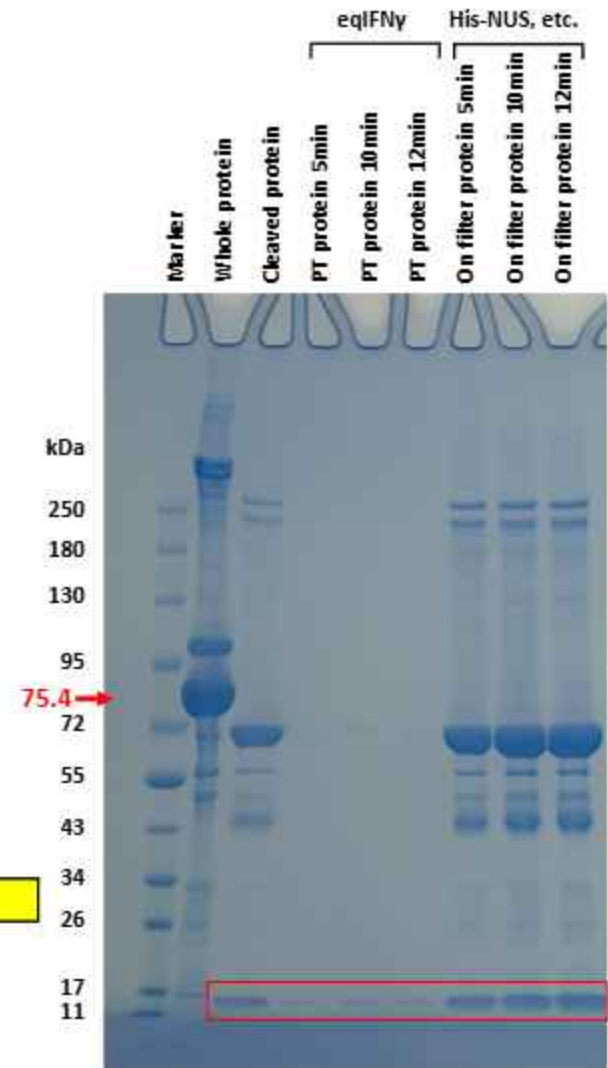


Proteins like His-Nus, being larger than 50 kDa, remain on the filter.

Being smaller than 50 kDa, eqIFN γ passes through the column into the flow-through.

Results: eqIFN γ also remained on the filter and was not separated from the other proteins.

Whole protein (eqIFNgNUS-His_pET-50b(+)) \approx 75.4 kDa;
eqIFN γ protein \approx 16.7 kDa; His-NUS protein \approx 58.7 kDa;
HRV 3C protease protein \approx 50.3 kDa



3-5. Separation and purification of cleaved protein by four kinds of conditions

Protein cleavage reaction (2 tubes)

Protease: Whole protein 1 : 5

Whole protein (ul)	178.6	Whole protein conc.: 1.4 mg/ml Protease conc.: 2.5 mg/ml Buffer: 50mM Tris-HCl (pH8.0), 200mM NaCl
HRV 3C protease (ul)	20.0	
Buffer (ul)	51.4	
Total (ul)	250	



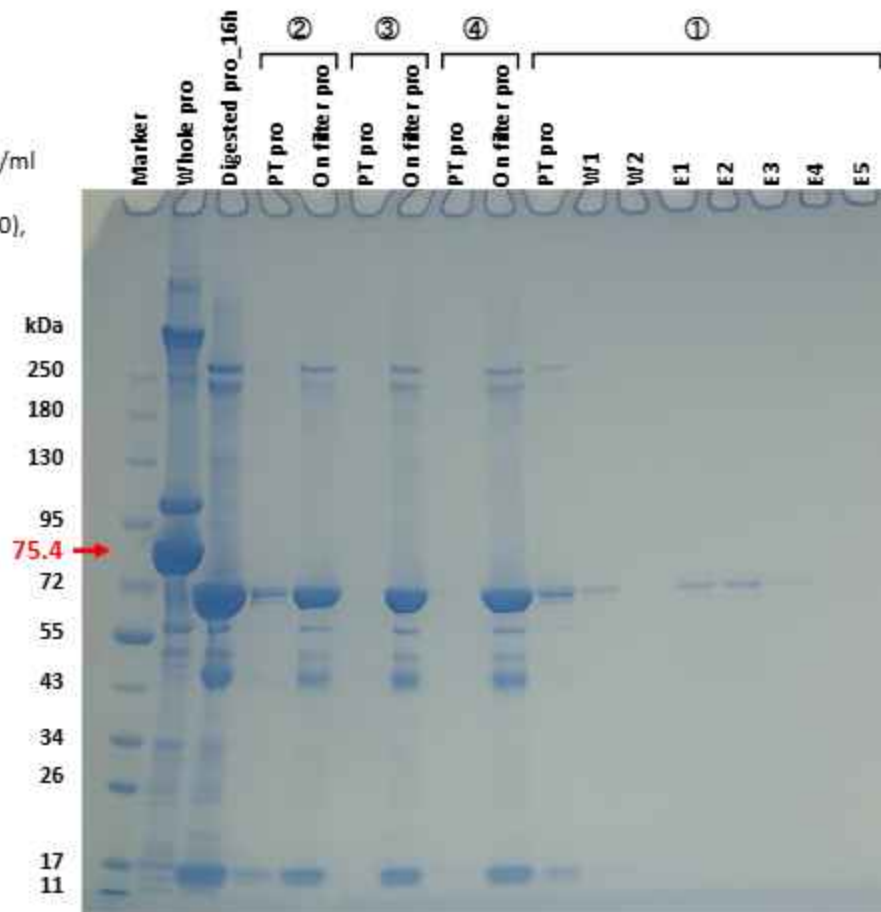
At 4°C, and rotating at 2 rpm for 16 h

1. By denaturing conditions

- ①: Protein + Denaturing buffer + Ni-NTA resin
↓ 4°C, Rotating for 1h
⇒ Purified the protein by Ni-NTA column
- ②: Protein + Denaturing buffer
↓ 4°C, Rotating for 1h
↓ Filter through an 50k Amicon tube
⇒ Recover the protein

2. high salt concentration condition

- ③: Protein + 50mM Tris-HCl (pH8.0), **500mM NaCl** buffer
↓ Filter through an 50k Amicon tube
⇒ Recover the protein
- ④: Protein + 50mM Tris-HCl (pH8.0), **1M NaCl** buffer
↓ Filter through an 50k Amicon tube
⇒ Recover the protein



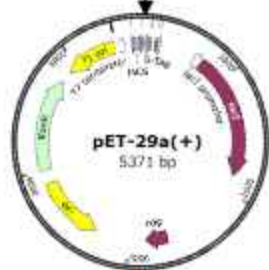
Results: eqIFN γ also remained on the filter or resin was not cleanly separated from the other proteins.

Whole protein (eqIFNgNUS-His_pET-50b(+)) \approx 75.4 kDa;
eqIFN γ protein \approx 16.7 kDa; His-NUS protein \approx 58.7 kDa;
HRV 3C protease protein \approx 50.3 kDa

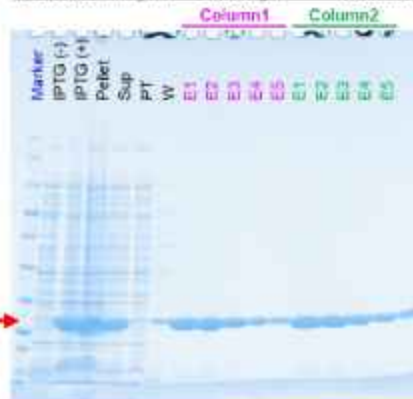


Future Research Plan 1

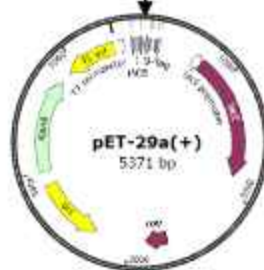
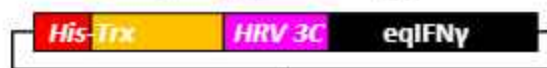
1. Identify the causes of non-specific binding between proteins and successfully isolate and purify cleaved eqIFN γ .
2. Insert the HRV 3C site between His-Trx and eqIFN γ to express the protein, and then cleave it with the protease to purify eqIFN γ exclusively.



SDS-PAGE analysis 5-20% gel 20210813



- ① Synthesize His-Trx-eqIFN γ _pET-29a(+)



- ② Express the His-Trx-eqIFN γ protein



- ③ Cleave the protein using HRV 3C protease

✗ The His-Trx and HRV 3C protease remain on the column



Ni-NTA resin

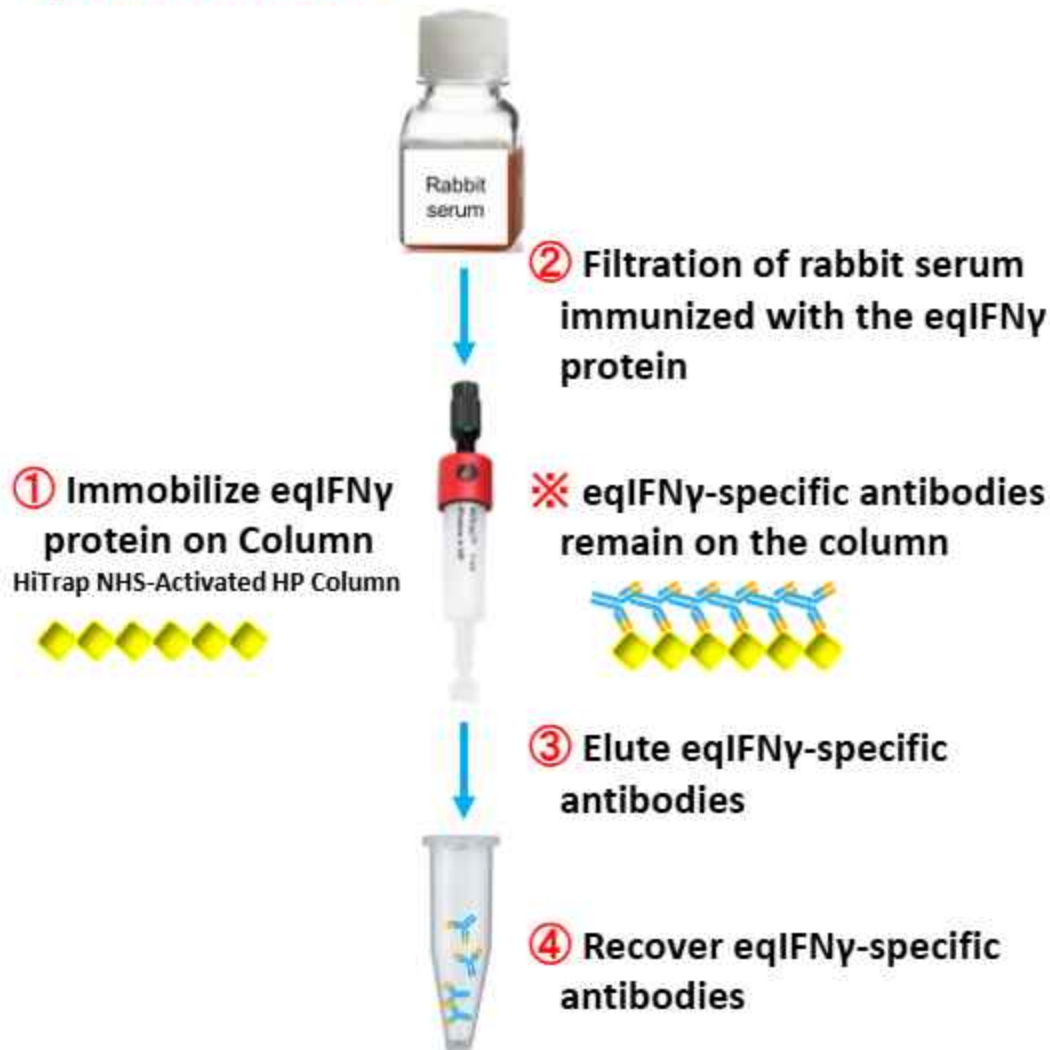


- ④ Recover the eqIFN γ protein

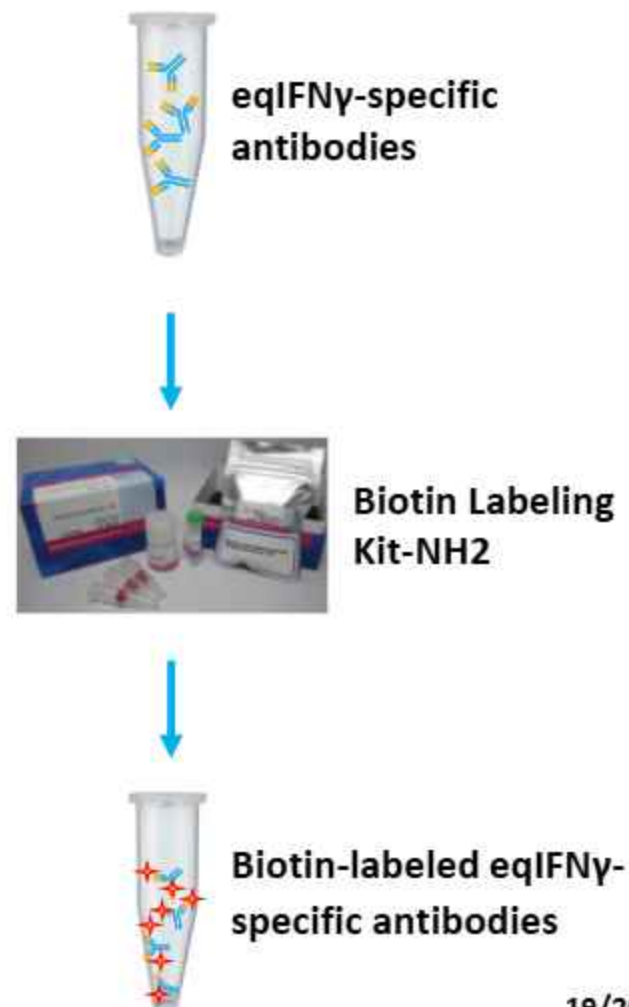


Future Research Plan 2

1. Purification of eqIFN γ -specific antibodies



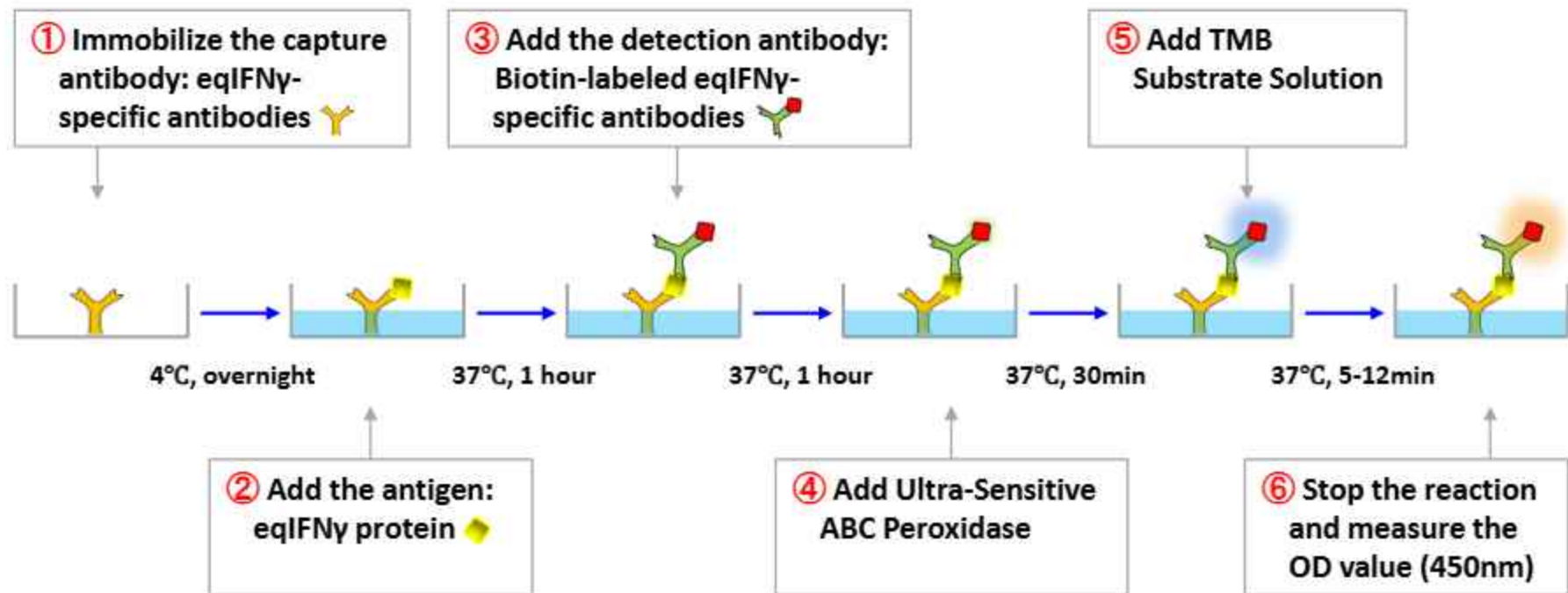
2. Label eqIFN γ -specific antibodies with biotin



Future Research Plan 2

3. Confirm eqIFN γ -specific antibodies by ELISA

Procedure of the ELISA method



4. Develop the ELISA method





Thank you for your attention

