Reconstituted cell-free protein synthesis kit



# PURE*frex<sup>®</sup>を*用いた ジスルフィド結合含有タンパク質の合成

ジーンフロンティア株式会社



2017年12月6日 ConBio2017



# **1. Introduction of PUREfrex**<sup>®</sup>

# 2. Application of PUREfrex<sup>®</sup>

• Optimization of nt sequence at 5'-terminus of ORF

• Synthesis of proteins containing disulfide bonds

• Synthesis of antibody-related proteins



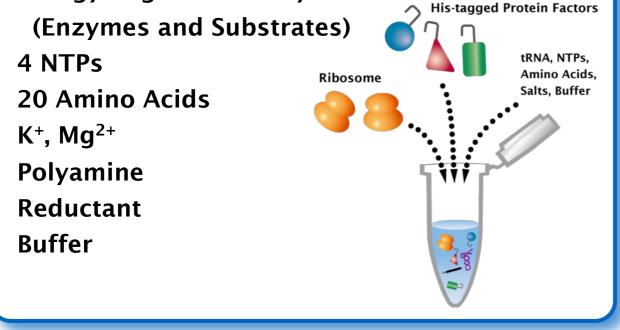
# **PURE system**

#### **Components of the original PURE system**

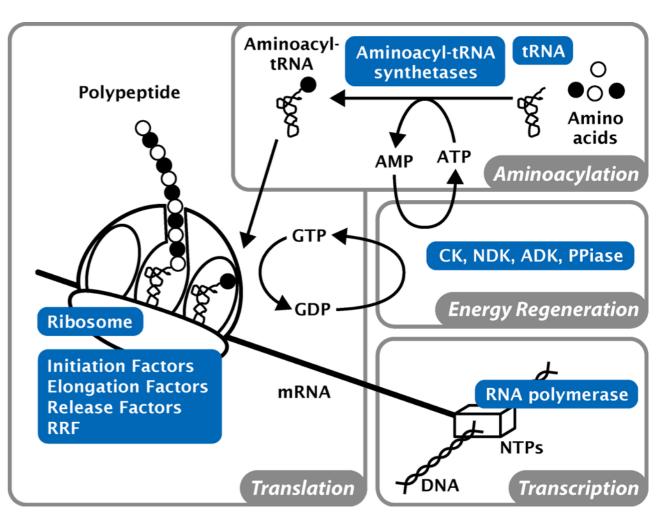
Purified Transcription/Translation Factors T7 RNA Polymerase Initiation Factors (IF1, IF2, IF3) Elongation Factors (EF-Tu, EF-Ts, EF-G) Release Factors (RF1, RF2, RF3) Ribosome Recycling Factor 20 Aminoacyl-tRNA synthetases (ARS) Methionyl-tRNA transformylase (MTF) *E. coli* Ribosome

E. coli tRNA mix

**Energy Regeneration System** 



#### Outline of protein synthesis in *E. coli*



Shimizu *et al.* (2001) *Nat. Biotechnol.*, vol.19, p.751 Shimizu *et al.* (2005) *Methods*, vol.36, p.299



#### Summary of the preparation methods of the components

|                     | Original<br>PURE system | PURE <i>frex</i> ® |
|---------------------|-------------------------|--------------------|
| Protein             |                         |                    |
| Tag                 | His-Tag                 | None               |
| Number of columns   | 1                       | > 3                |
| Wash with detergent | -                       | +                  |
| Ribosome            |                         |                    |
| Wash with detergent | -                       | +                  |
| tRNA                |                         |                    |
| Wash with detergent | -                       | +                  |

Original PURE system: Shimizu et al. (2005) Methods, vol.36, p.299



|  | Extract System | <b>Reconstituted System</b> |                    |
|--|----------------|-----------------------------|--------------------|
|  | S30 system     | Original<br>PURE system     | PURE <i>frex</i> ® |
| Typical Yield                          | 100-1000 µg/mL | 10-200 µg/mL                | 10-200 µg/mL       |
| Contamination                          |                |                             |                    |
| RNase                                  | very High      | Low                         | very Low           |
| LPS                                    | very High      | High                        | very Low           |
| Template DNA                           |                |                             |                    |
| Plasmid DNA                            | ОК             | ОК                          | ОК                 |
| PCR product                            | NG             | ОК                          | ОК                 |
| Customization of<br>Reagent            | Difficult      | Easy                        | Easy               |
| Purification of His-<br>tagged product | ОК             | NG                          | ОК                 |

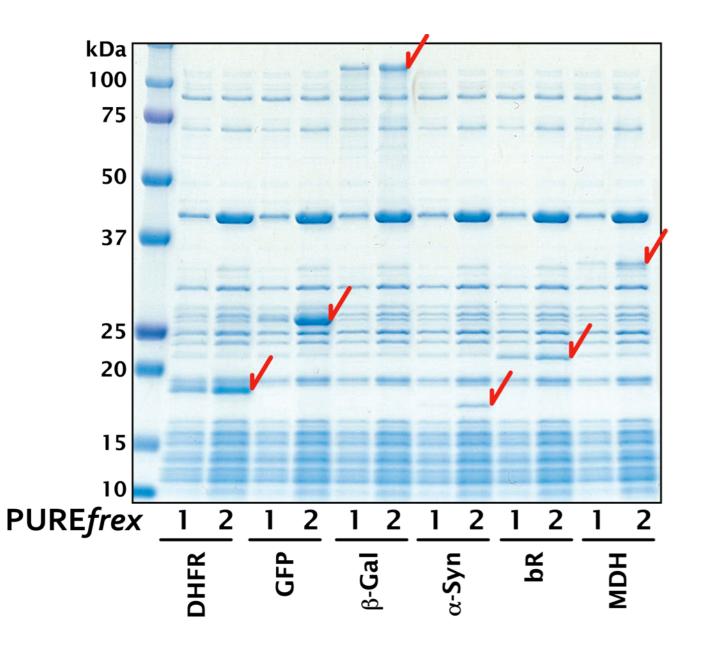
Original PURE system: Shimizu et al. (2005) Methods, vol.36, p.299

|  | Extract System | <b>Reconstituted System</b> |                        |
|--|----------------|-----------------------------|------------------------|
|  | S30 system     | Original<br>PURE system     | PURE <i>frex</i> ® 2.0 |
| Typical Yield                          | 100-1000 µg/mL | 10-200 µg/mL                | 20-1000 µg/mL          |
| Contamination                          |                |                             |                        |
| RNase                                  | very High      | Low                         | very Low               |
| LPS                                    | very High      | High                        | very Low               |
| Template DNA                           |                |                             |                        |
| Plasmid DNA                            | ОК             | ОК                          | ОК                     |
| PCR product                            | NG             | ОК                          | ОК                     |
| Customization of<br>Reagent            | Difficult      | Easy                        | Easy                   |
| Purification of His-<br>tagged product | ОК             | NG                          | ОК                     |
|  |                |                             |                        |

Original PURE system: Shimizu et al. (2005) Methods, vol.36, p.299

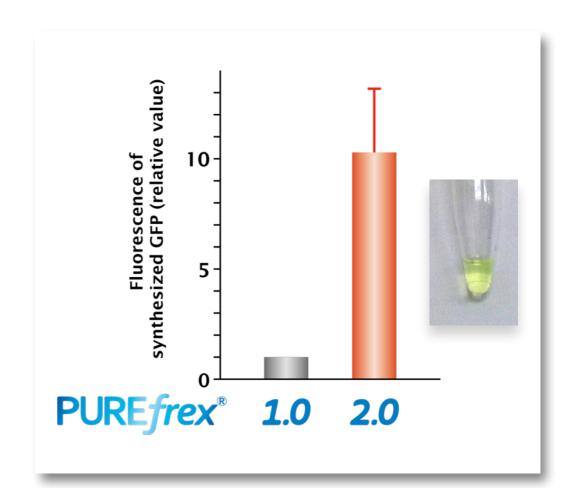
# Protein synthesis using PURE*frex*® 1.0 and 2.0

#### A) SDS-PAGE



DHFR: Dihydrofolate reductaseα-Syn: α-SynucleinGFP: Green Fluorescent ProteinbR: Bacteriorhodopsinβ-Gal: β-GalactosidaseMDH: Malate Dehydrogenase

#### **B)** Fluorescence of synthesized GFP





# **1. Introduction of PUREfrex<sup>®</sup>**

# 2. Application of PUREfrex<sup>®</sup>

# • Optimization of nt sequence at 5'-terminus of ORF

• Synthesis of proteins containing disulfide bonds

• Synthesis of antibody-related proteins

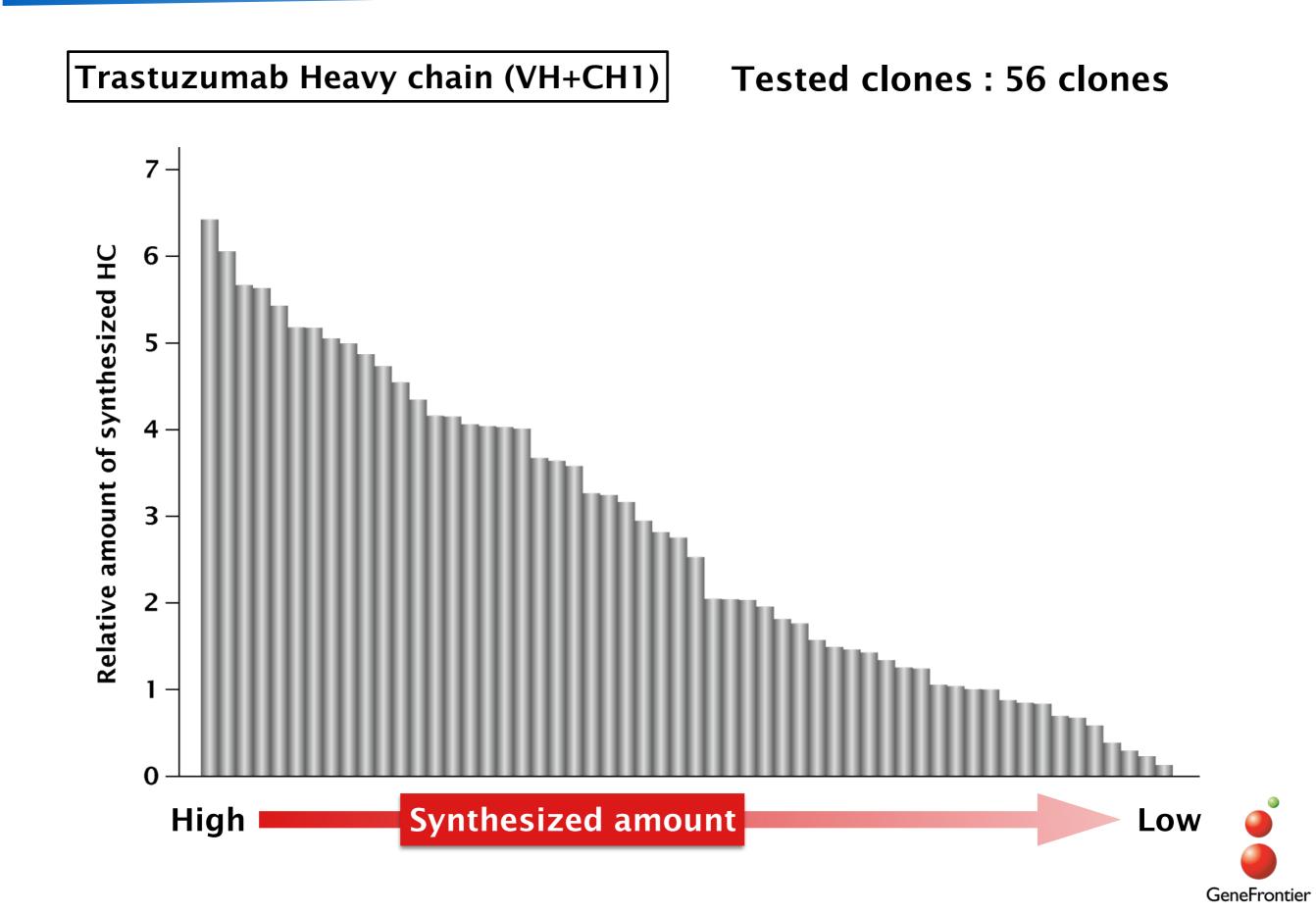


Trastuzumab Heavy chain (VH+CH1)

Met-Glu-Val-Gln-Leu-Val-Glu Val Gln Val Leu Freq. (%) Freq. (%) Freq. (%) Freq. (%) Freq. (%) codon codon codon codon codon 70 25 30 gtt 15 25 gtt gaa caa tta 30 18 70 12 18 gtc cag ttg gtc gag 17 12 17 gta ctt gta 40 10 40 gtg ctc gtg 5 cta 46 ctg Glu, Gln 65% 50% 35% Frequency Leu, Val 35% 25% 15%

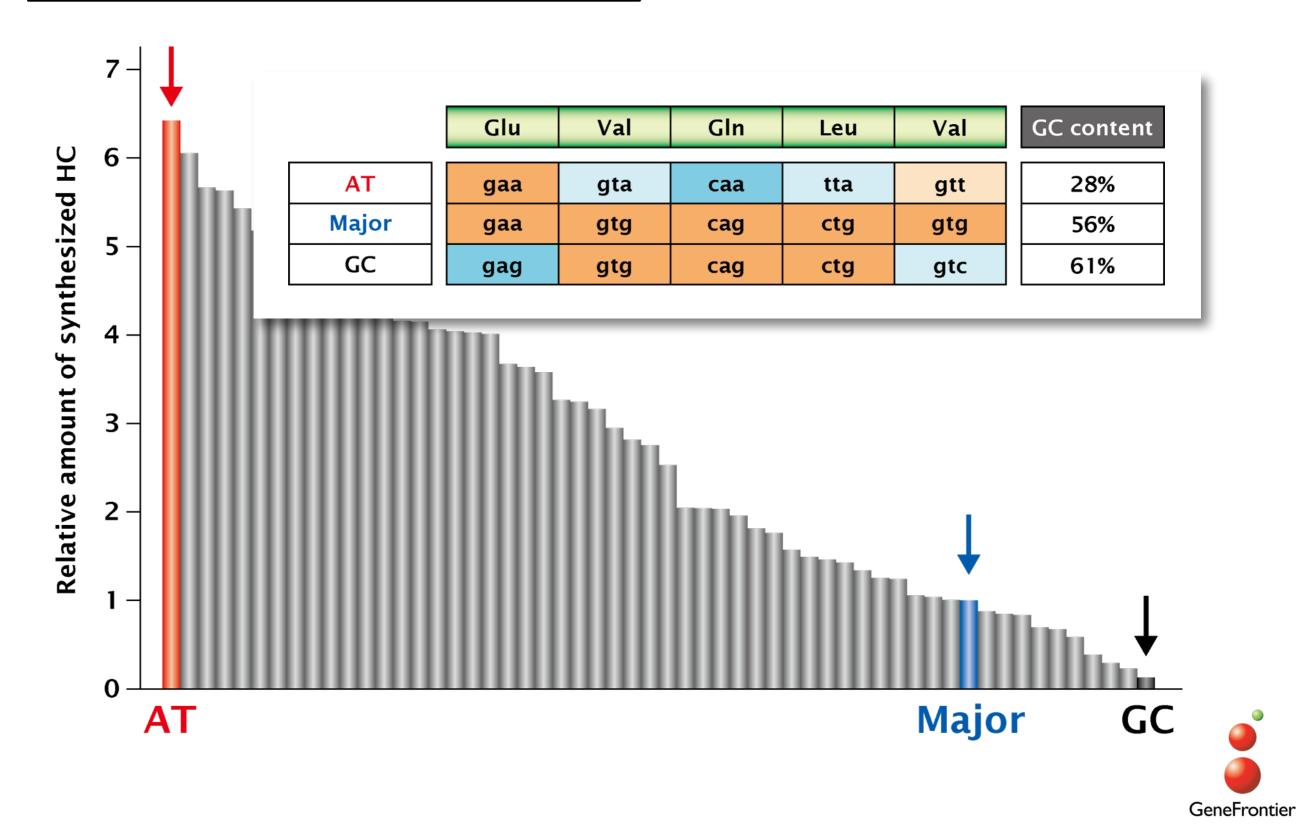
\*Frequency is calculated from Codon Usage Database in Kazusa DNA Res. Inst. (*E. coli* K-12 strain)

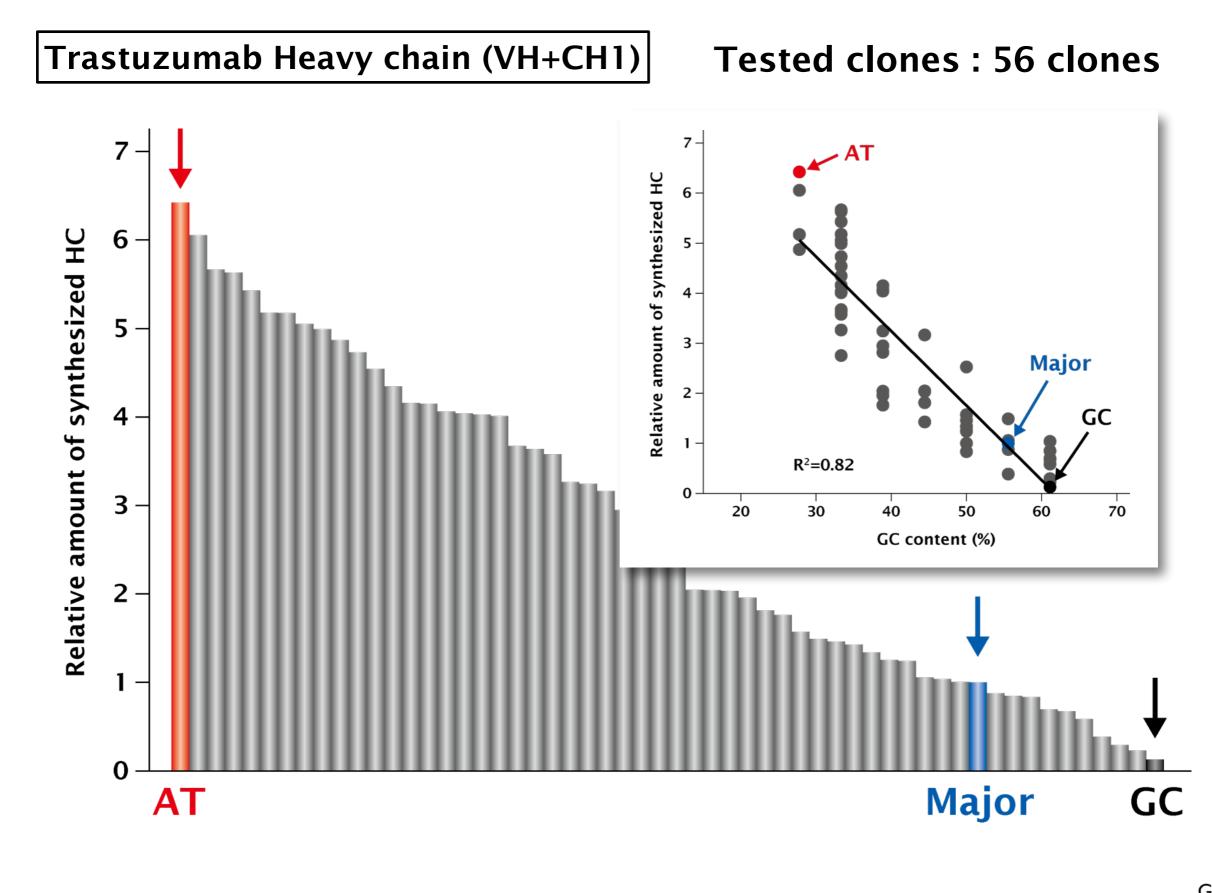




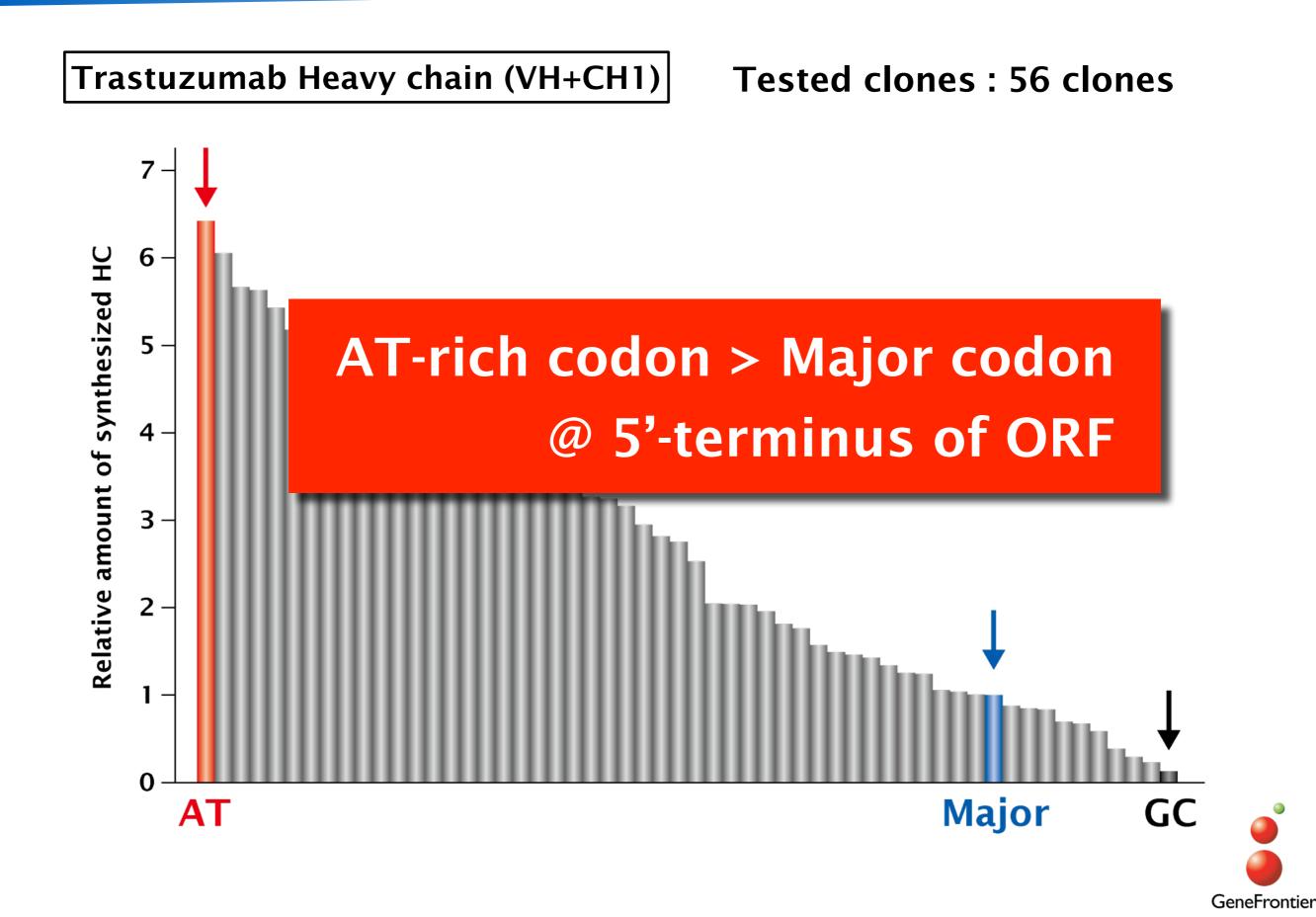
Trastuzumab Heavy chain (VH+CH1)

**Tested clones : 56 clones** 





GeneFrontier



# **1. Introduction of PUREfrex<sup>®</sup>**

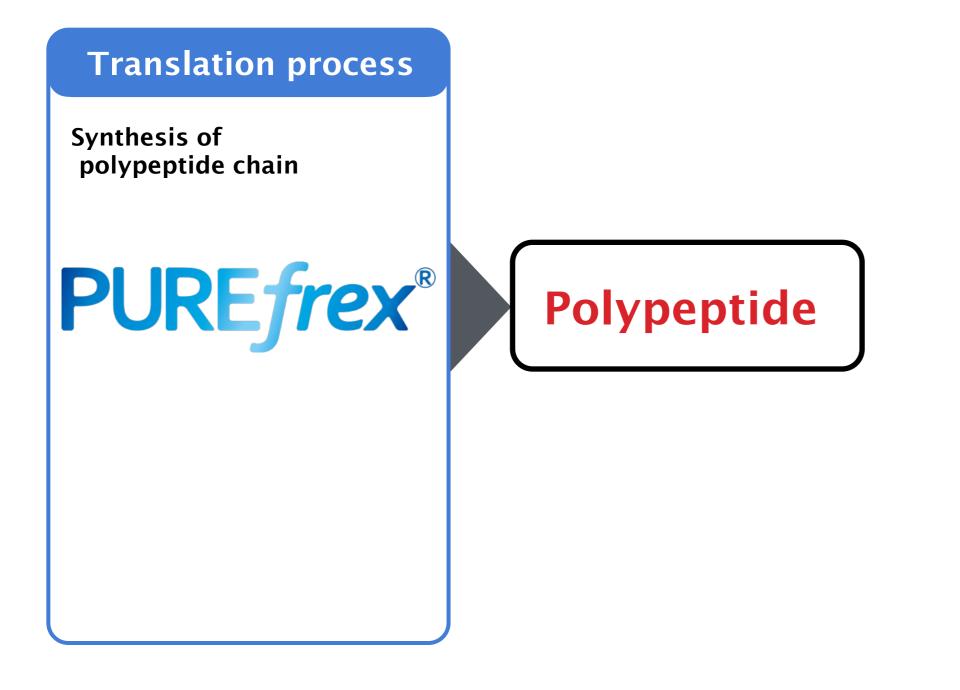
# 2. Application of PUREfrex<sup>®</sup>

• Optimization of nt sequence at 5'-terminus of ORF

• Synthesis of proteins containing disulfide bonds

• Synthesis of antibody-related proteins







### **Polypeptide** ≠ **Functional protein**



**PURE***frex*<sup>®</sup>

Synthesis of polypeptide chain

#### Common process

#### Maturation process

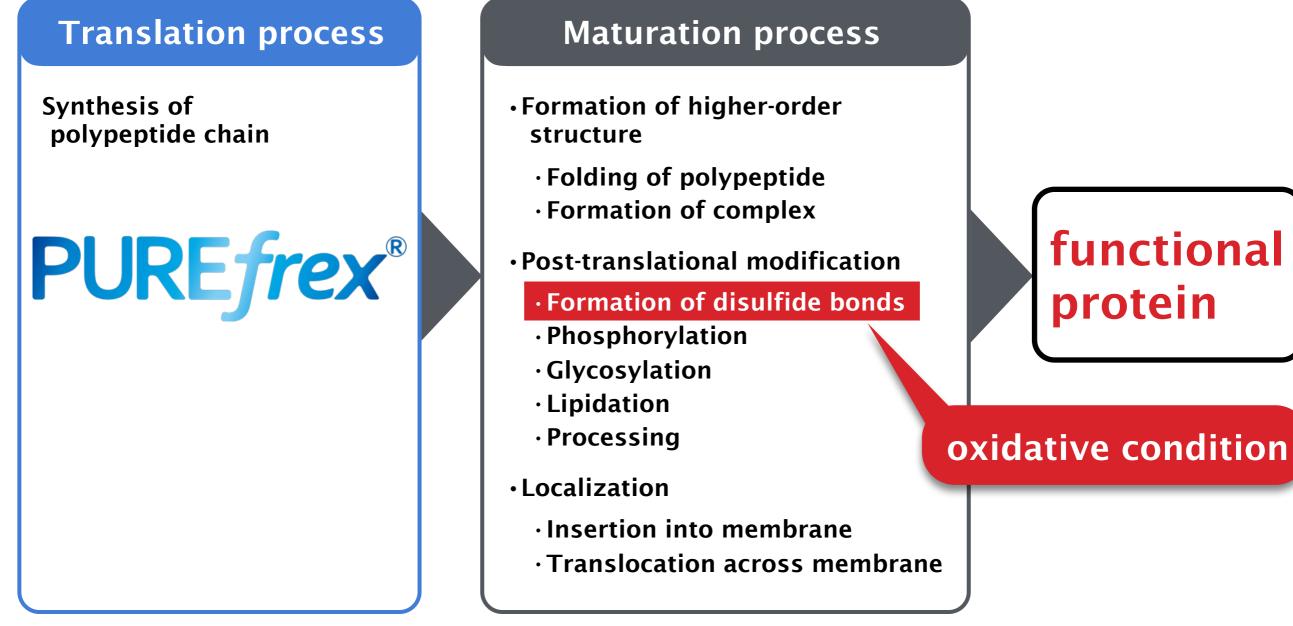
- Formation of higher-order structure
  - · Folding of polypeptide
  - Formation of complex
- Post-translational modification
  - Formation of disulfide bonds
  - Phosphorylation
  - $\cdot$  Glycosylation
  - Lipidation
  - Processing
- Localization
  - Insertion into membrane
  - $\cdot$  Translocation across membrane

#### **Diversified process**

# functional protein



### **Polypeptide** ≠ **Functional protein**

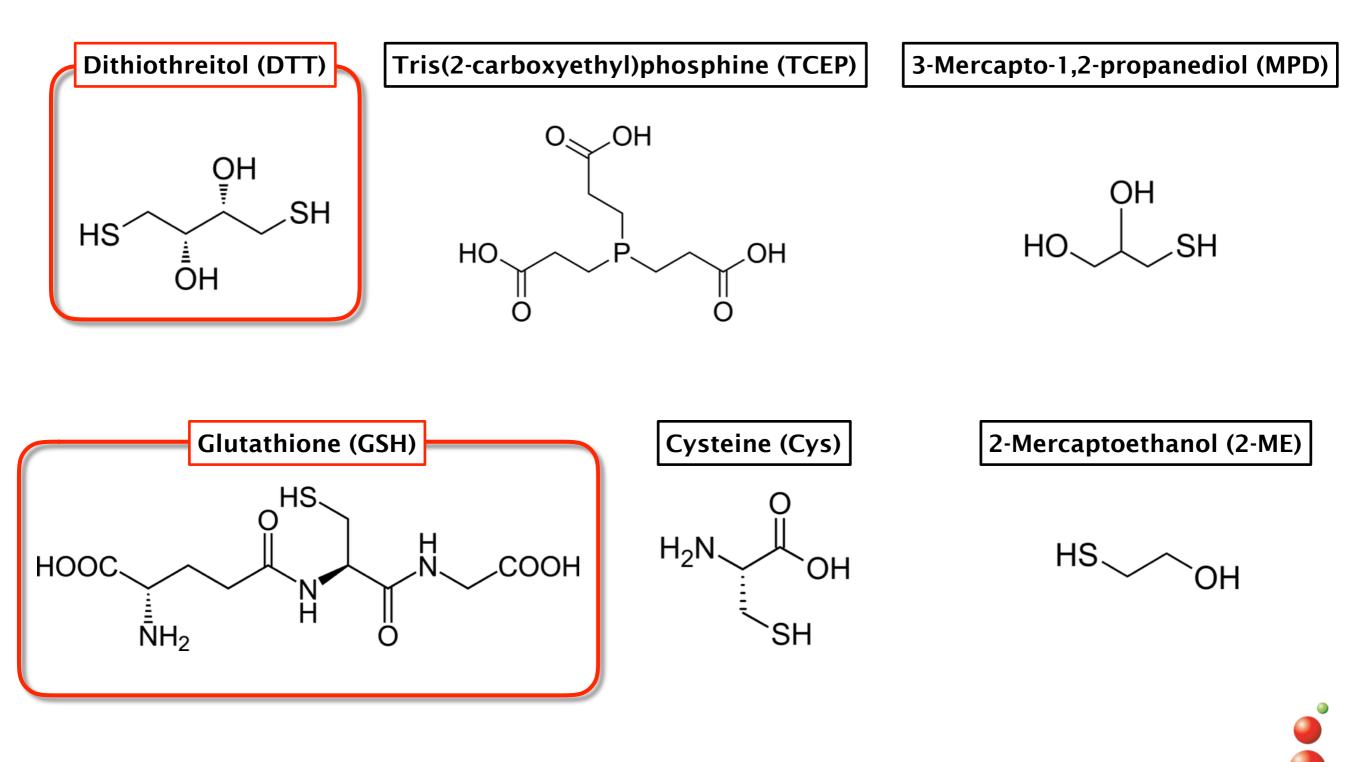


Common process

#### **Diversified process**



### <u>Reductant</u>

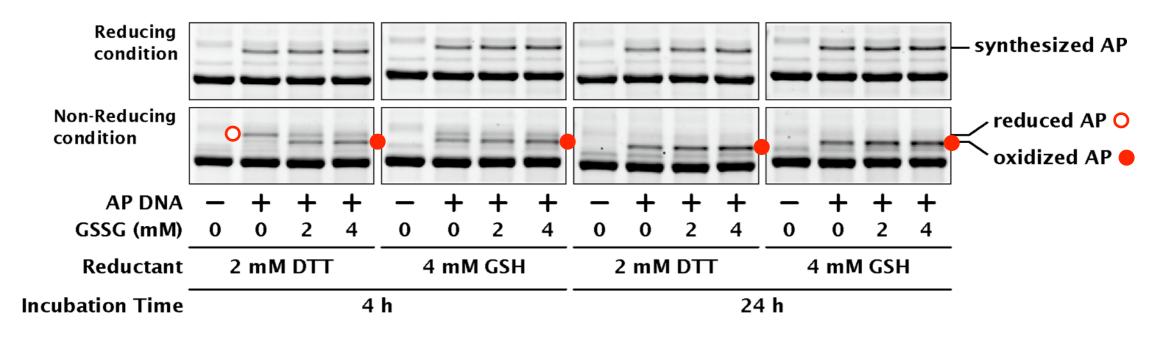


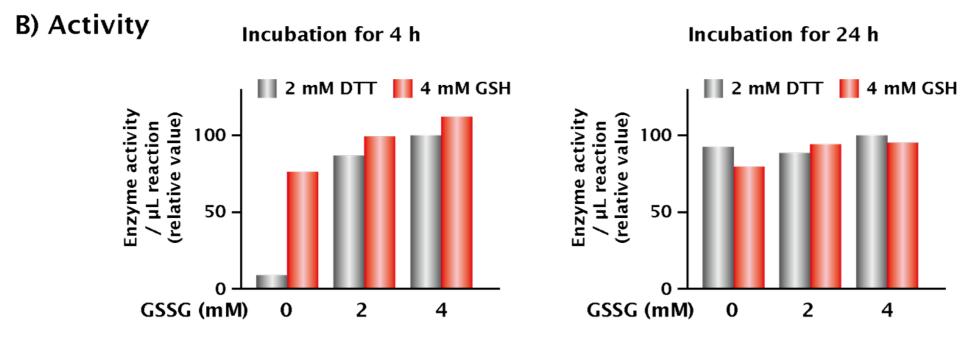
GeneFrontier

# Synthesis of functional proteins containing disulfide bonds

### *E. coli* alkaline phosphatase (AP) (2 disulfide bonds)

#### A) SDS-PAGE



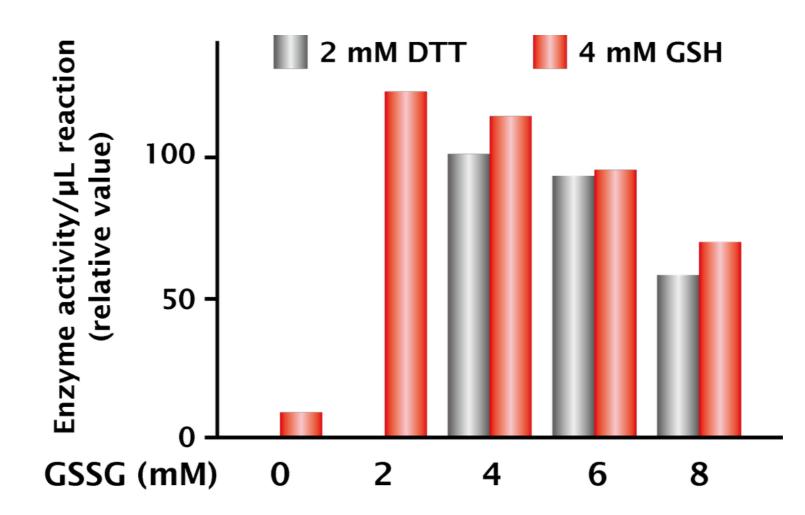




# Synthesis of functional proteins containing disulfide bonds

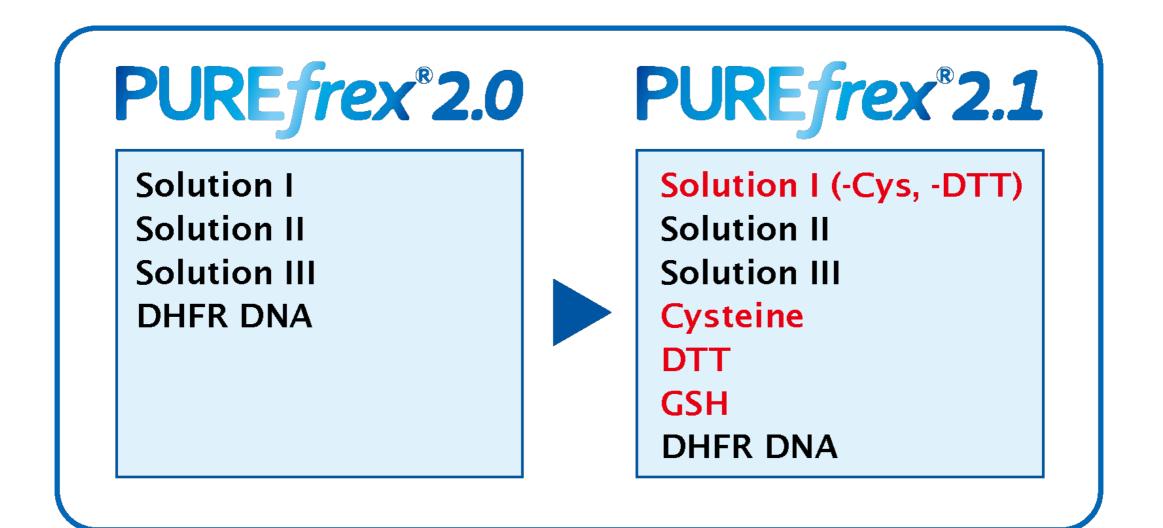
E. coli acid phosphatase (AppA) (5 disulfide bonds)

+ DsbC (*E. coli* disulfide isomerase)





### New product





# **1. Introduction of PUREfrex<sup>®</sup>**

# 2. Application of PUREfrex<sup>®</sup>

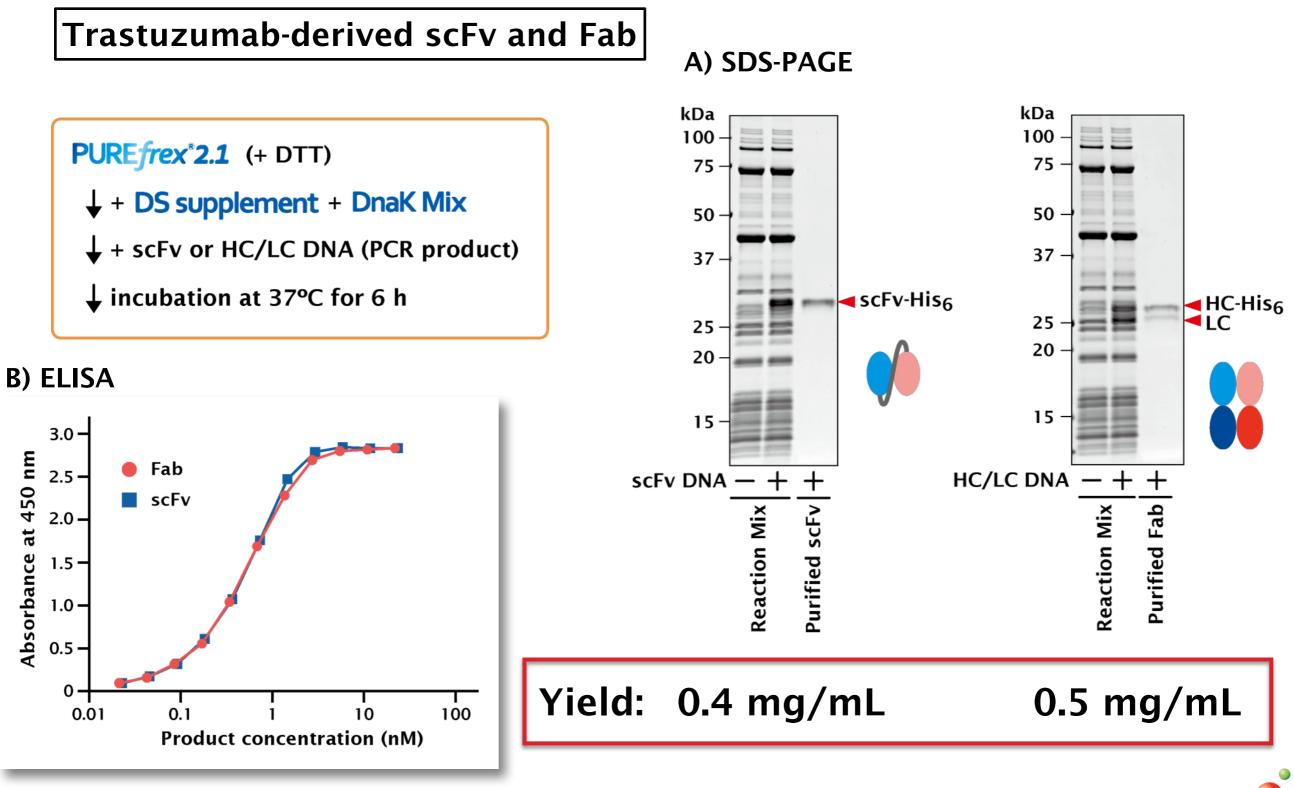
• Optimization of nt sequence at 5'-terminus of ORF

• Synthesis of proteins containing disulfide bonds

• Synthesis of antibody-related proteins

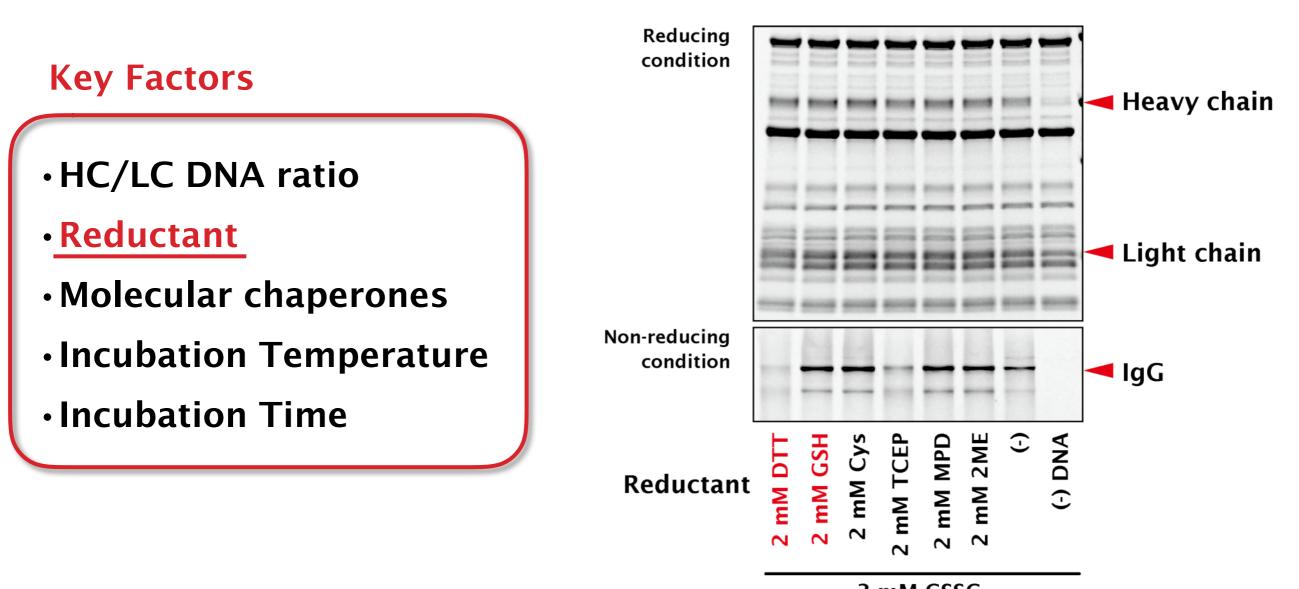


# Synthesis of scFv and Fab





### Trastuzumab-derived aglycosylated IgG



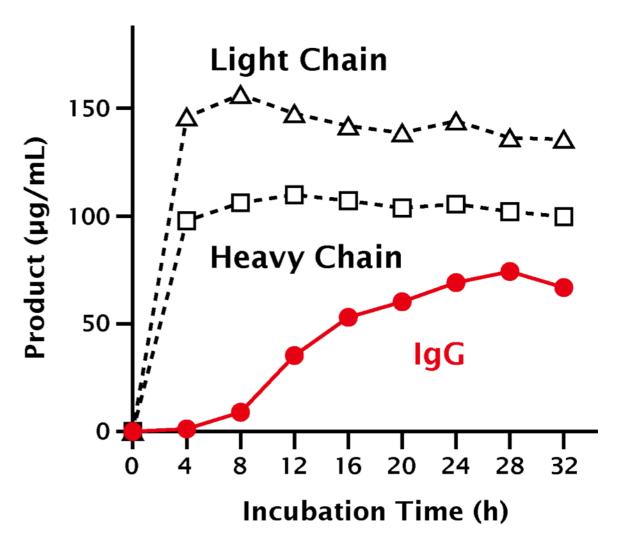
3 mM GSSG 117 µg/mL DsbC 5 µM DnaK/1 µM DnaJ/1 µM GrpE



Trastuzumab-derived aglycosylated IgG

### **Key Factors**

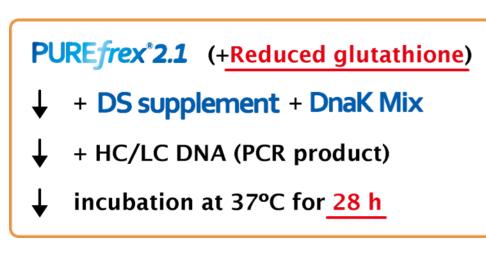
- •HC/LC DNA ratio
- Reductant
- Molecular chaperones
- Incubation Temperature
- Incubation Time



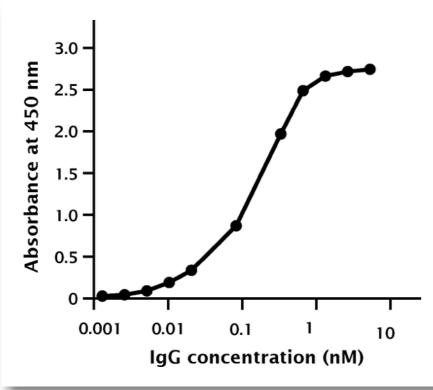


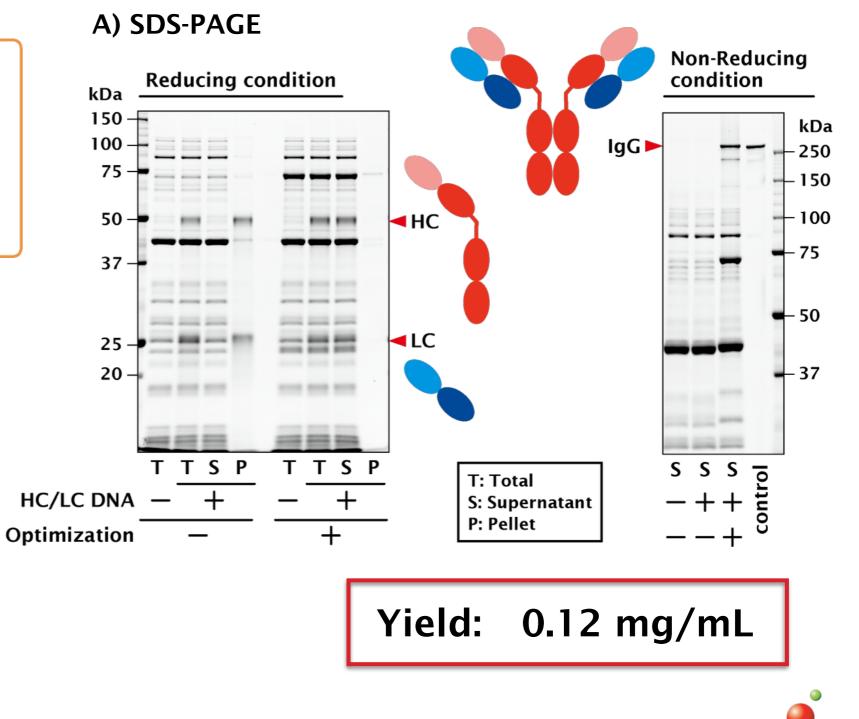
# Synthesis of aglycosylated IgG

#### Trastuzumab-derived aglycosylated IgG









GeneFrontier

# **Product List**

# タンパク質合成反応液

PUREfrex<sup>®</sup>1.0 PUREfrex<sup>®</sup>2.0

New PUREfrex<sup>®</sup>2.1

添加剤

DnaK MixDnaK/DnaJ/GrpEGroE MixGroEL/GroESDS supplementOxidized Glutathione, DsbC

