

ヒトプロテインジスルフィドイソメラーゼを添加した 再構成型無細胞タンパク質合成系(PUREflex[®])によるジスルフィド結合タンパク質合成

Synthesis of proteins containing disulfide bonds using PUREflex[®] with human protein disulfide isomerase



松本 令奈、金森 崇 (ジーンフロンティア株式会社)

Rena Matsumoto and Takashi Kanamori (GeneFrontier Corporation)

<Abstract>

PURE systemは、大腸菌のタンパク質合成に関与する因子のみから再構成した無細胞タンパク質合成系である。PURE systemにプロテインジスルフィドイソメラーゼ(PDI)として大腸菌DsbCを添加することで、活性を有したジスルフィド結合タンパク質を合成できる。複雑な構造のタンパク質や高等生物のジスルフィド結合タンパク質合成に広く対応するため、本発表ではPUREflex[®](改良型PURE system)を用い、DsbCに替えてヒトPDI(hPDI)およびその関連タンパク質を利用できるか検討した。モデル系として、9つのジスルフィド結合を持つ組織プラスミノーゲン活性化因子(tPA)の合成を行った。還元剤に還元型グルタチオン(GSH)を使用したPUREflexをベースに、酸化型グルタチオン(GSSG)存在下でhPDIを添加してtPAを合成すると、合成されたtPAはhPDIの添加濃度依存的に活性が増加した。一方、GSSG非存在下でも、hPDIおよびhEro1α(PDI酸化酵素)を添加しtPAを合成した場合、合成されたtPAは同様に高い活性を示した。すなわち、hPDIを機能させるためにはGSSGあるいはhEro1αの添加が必要であることが分かった。また、GSSG存在下でhPDIとDsbCの添加効果を比較したところ、hPDIはDsbCとほぼ同等の効果を示した。以上より、hPDI添加でも、酸化還元環境の調節やhPDI酸化酵素の添加により効率よくジスルフィド結合タンパク質の合成が可能であることが確認された。さらに他の基質タンパク質の合成についても報告する。

1. PUREflex[®]; based on the PURE system technology

Advantage

- Low level of contamination
 - High throughput preparation of proteins (including Fab, scFv, protein toxin etc.)
 - PCR products usable as a template DNA
 - Protein science research
 - In vitro display (Ribosome display, mRNA display etc.)
- (Ref: Shimizu et al. (2001) Nat. Biotechnol., vol. 19, p. 751.)

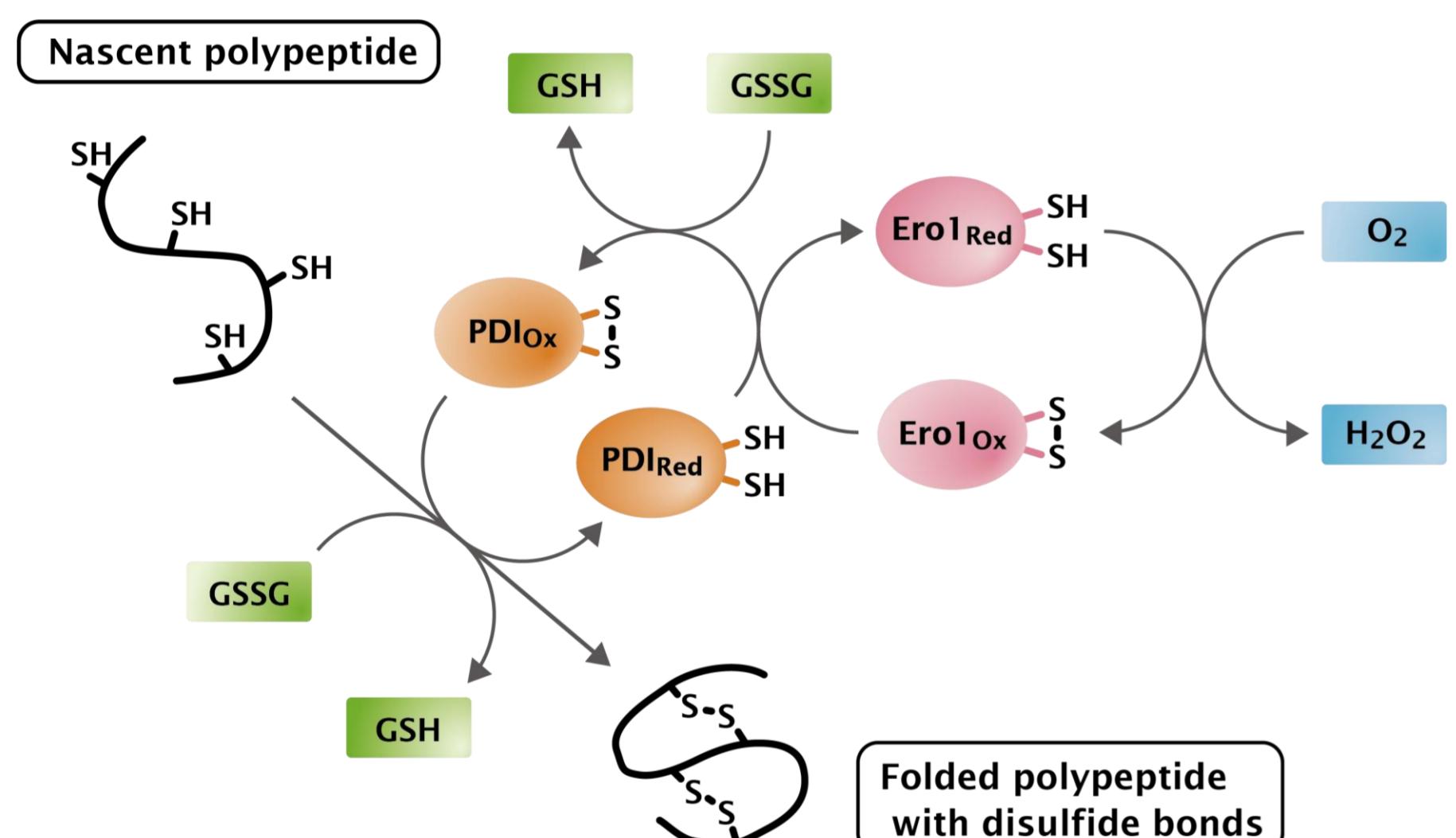
Application

PUREflex[®] 2.0 - Reductant (DTT) and Cysteine

= PUREflex[®] 2.1

It enables to select a suitable reducing agent for the protein of your interest.

2. Oxidative folding of nascent polypeptide in Endoplasmic Reticulum (ER)



2-1. PDIs and related protein used in this study

hPDI (human protein disulfide isomerase)
UniProt ID: P07237
Organism: Homo sapiens
Length: 492 a.a.
Molecular weight: 55,424 Da
(PDB ID: 4EK2)

hEro1α (human endoplasmic oxidoreductin)

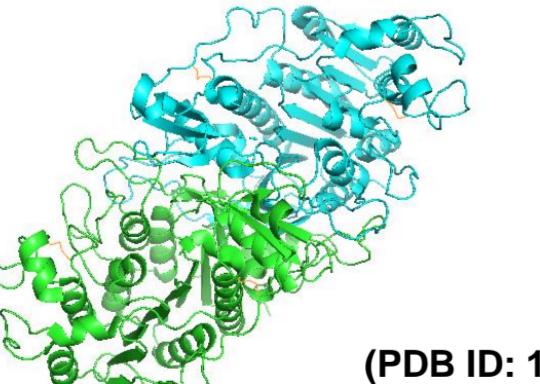
UniProt ID: Q96HE7
Organism: Homo sapiens
Length: 446 a.a.
Molecular weight: 52,122 Da
(PDB ID: 3AHQ)

DsbC (thiol:disulfide interchange protein)
UniProt ID: P0AE66
Organism: Escherichia coli
Length: 217 a.a.
Molecular weight: 23,591 Da
(PDB ID: 1EEJ)

3. Result 1: Selection of suitable reducing agent

Alkaline phosphatase (AP)

Organism: Escherichia coli
Synthesized region: 22Arg-47Lys
Length: 451 a.a.
Molecular weight: 47,330 Da
No. of disulfide bonds: 2
(PDB ID: 1AJA)



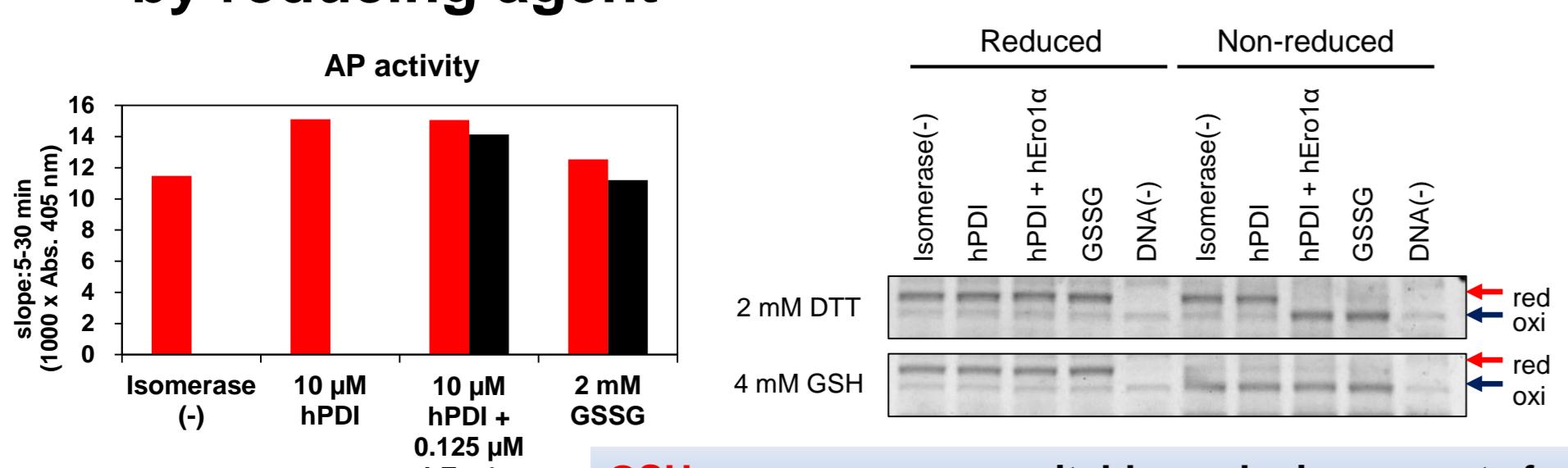
Experimental conditions of protein synthesis

No.	Factors	Basic condition	Reducant	hPDI	GSSG	hEro1α	Template DNA	Temp. & Time
1	hPDI			10 μM	(-)	(-)		
2	hPDI + hEro1α	PUREflex [®] 2.1 + 0.5 mM Cys or 2 mM DTT	4 mM GSH or 2 mM DTT	10 μM	(-)	0.2 μM	plasmid (10 ng/μL)	37°C, 4 h
3	GSSG			(-)	2 mM	(-)		

SDS-PAGE (0.5 μL of reaction mix/lane) → Gel staining (Oriole)

AP activity assay: Assay of PNPP dephosphorylation activity (0.1 μL PURE reaction mix/100 μL assay soln.)

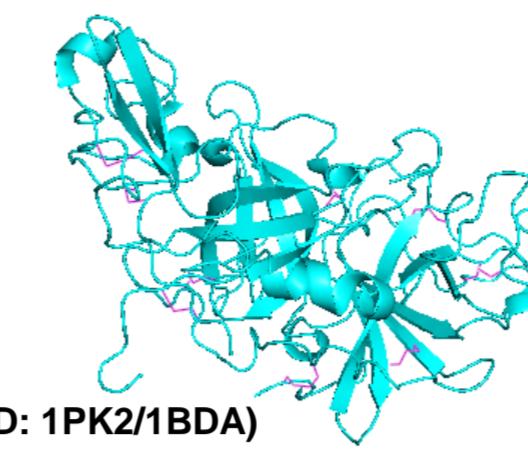
3-1. Difference in synthesized AP activity by reducing agent



4. Result 2: Disulfide bonded protein synthesis using hPDI and hEro1α with PUREflex[®]

truncated version of tissue plasminogen activator (tPA)

Organism: Homo sapiens
Synthesized region: 36Ser-40Ile/211Gly-562Pro (+FLAG)
Length: 368 a.a.
Molecular weight: 41,072 Da
No. of disulfide bonds: 9



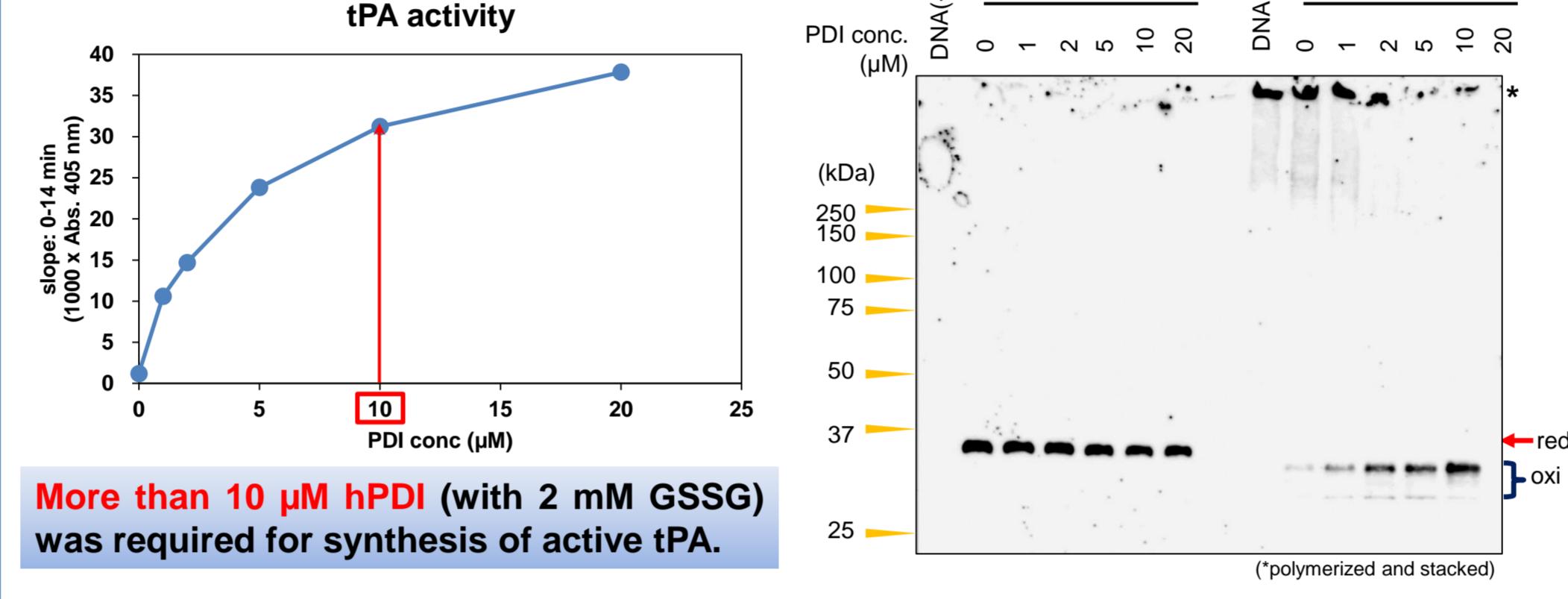
Experimental conditions of protein synthesis

No.	Factors	Basic condition	hPDI	GSSG	hEro1α	DsbC	Template DNA	Temp. & Time
1	hPDI conc.			0-20 μM	2 mM	(-)	(-)	
2	hPDI + GSSG conc.	PUREflex [®] 2.1 + 0.5 mM Cys + 4 mM GSH + DnaK mix	10 μM	0-8 mM	(-)	(-)		
3	hPDI + hEro1α conc.		10 μM	(-)	0-2 μM	(-)		
4	Incubation time		10 μM	2 mM	(-)	0.125 μM		30°C, 24 h
5	DsbC		(-)	0 mM/2 mM	(-)	20 μM		30°C, 24 h

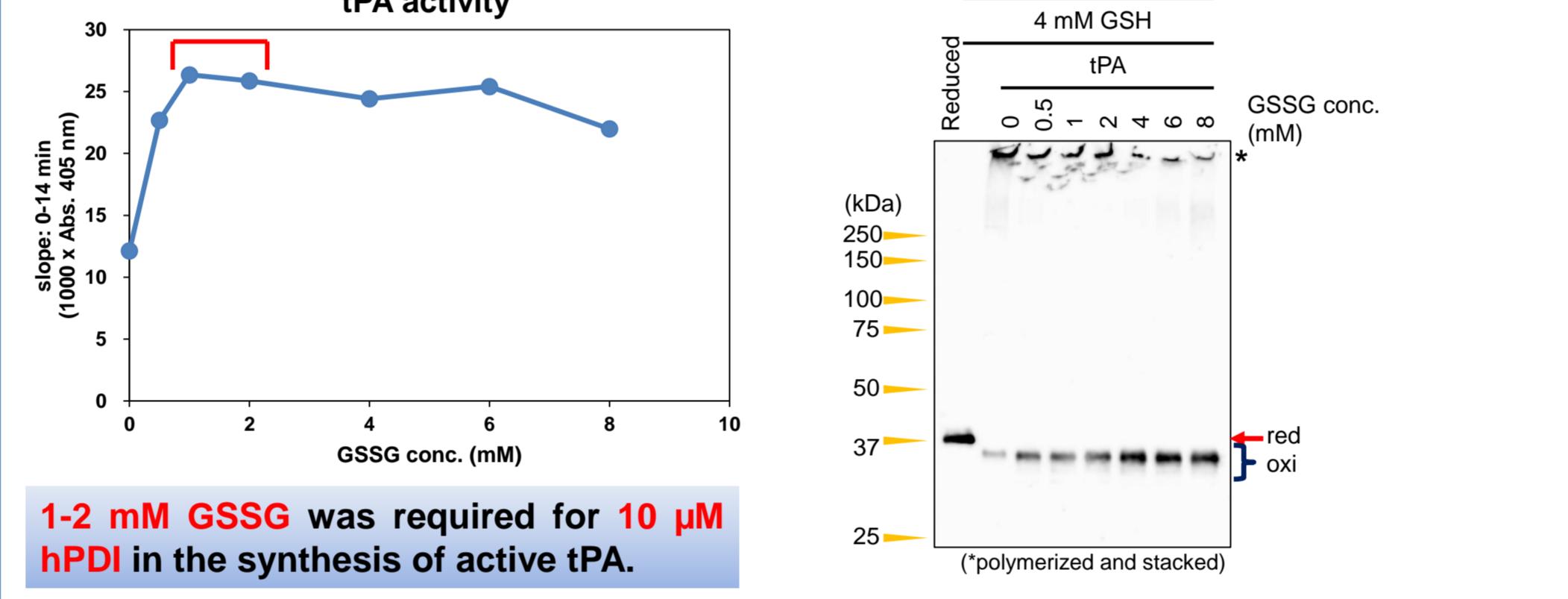
SDS-PAGE (0.25 – 0.5 μL of reaction mix/lane) → Western Blotting (anti-FLAG) or Gel staining (Oriole)

tPA activity assay: Assay of chromogenic substrate cleavage activity (2.5 μL PURE reaction mix/200 μL assay soln.)

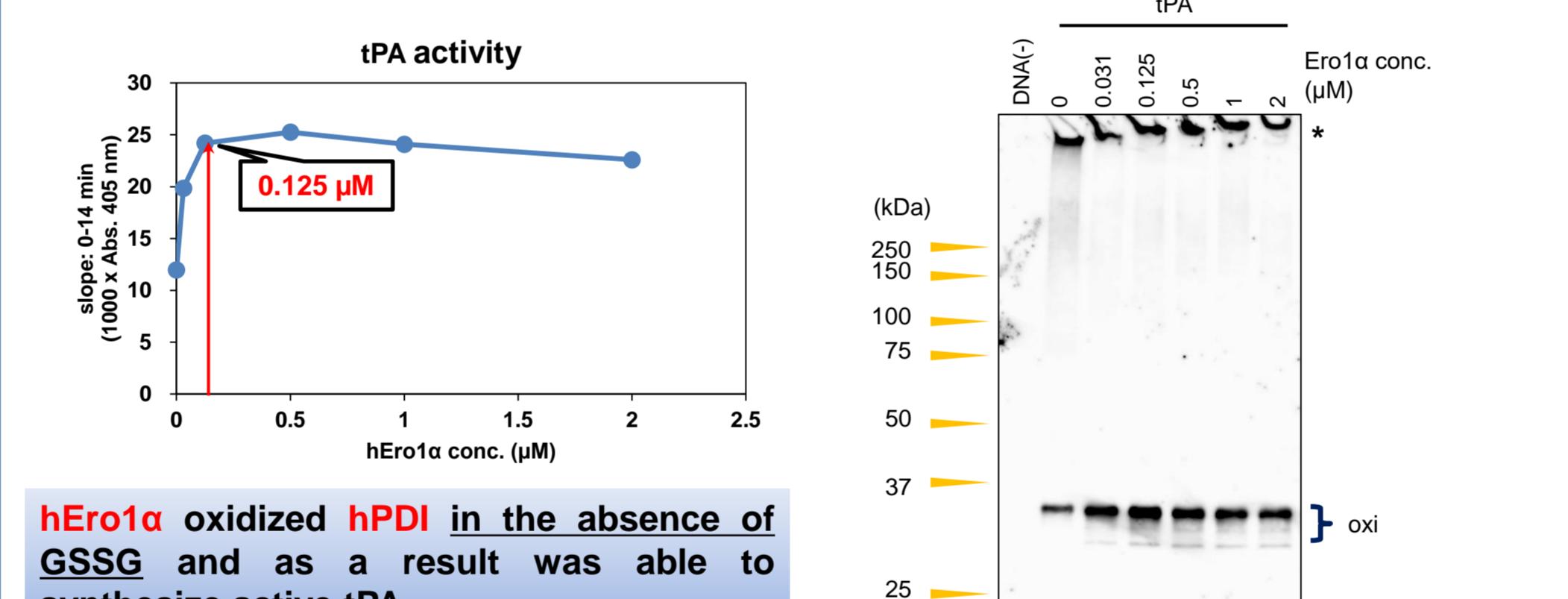
4-1. hPDI concentration



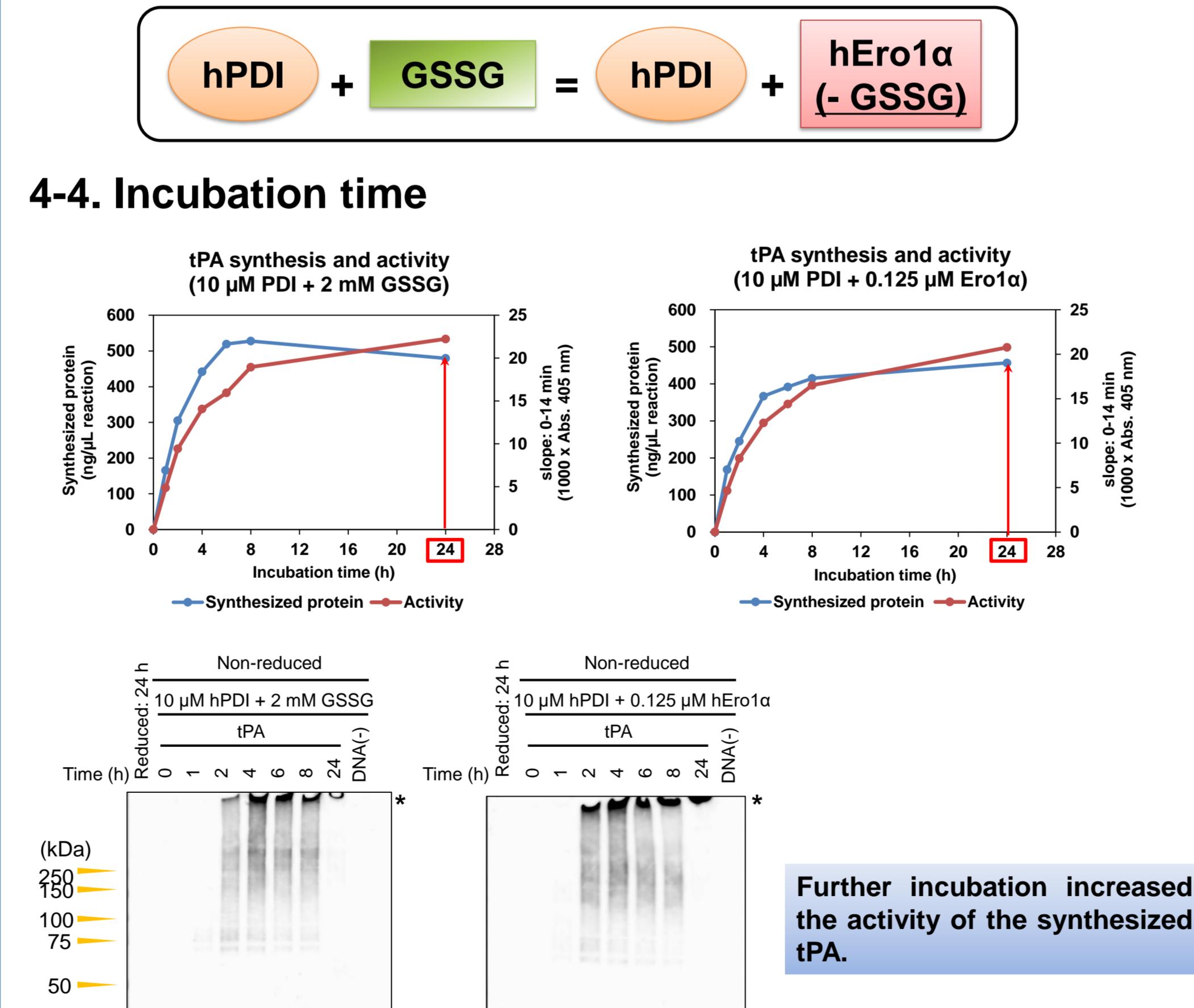
4-2. hPDI + GSSG concentration



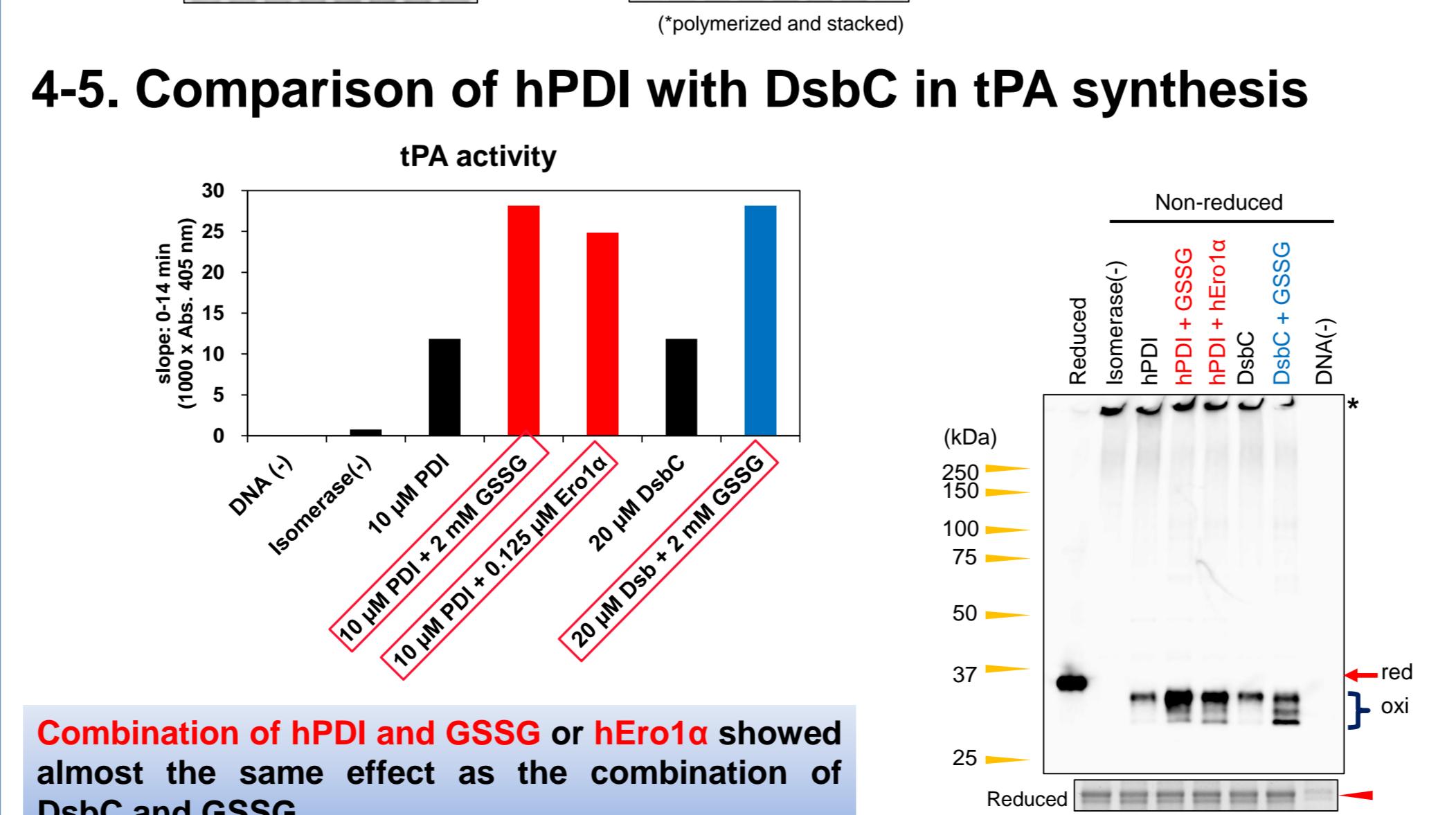
4-3. hPDI + hEro1α concentration



4-4. Incubation time



4-5. Comparison of hPDI with DsbC in tPA synthesis



5. Result 3: Examples of other disulfide bonded protein synthesis

Gaussia Luciferase (GLuc)

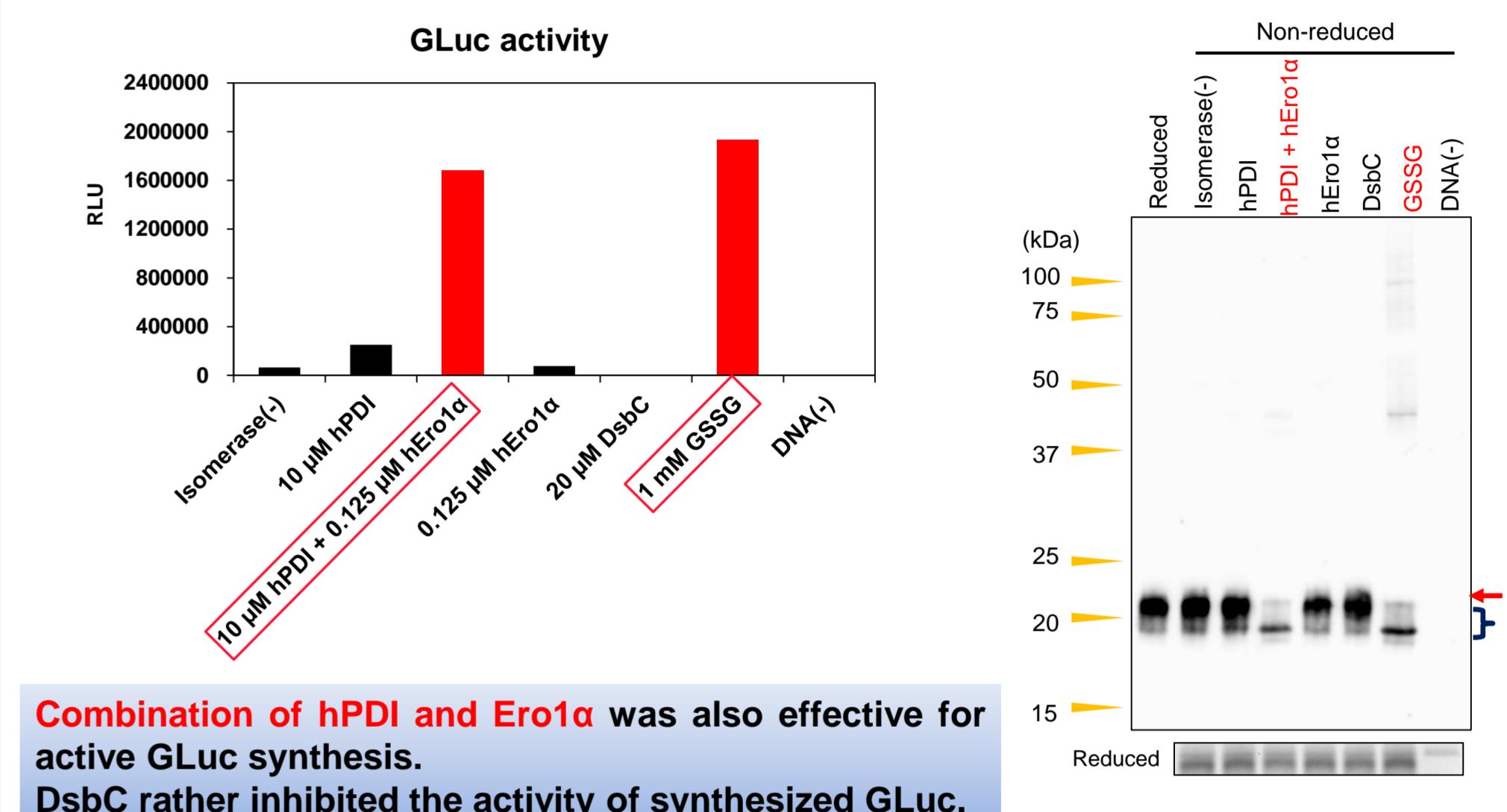
Organism: *Gaussia princeps*
Synthesized region: 18Lys-185Asp (+FLAG-His6)
Length: 187 a.a.
Molecular weight: 20,407 Da
No. of disulfide bonds: 5

Experimental condition of protein synthesis

No.	Factors	Basic condition	hPDI	GSSG	hEro1α	DsbC	Template DNA	Temp. & Time
1	hPDI		10 μM	(-)	(-)	(-)		
2	hPDI + hEro1α	PUREflex [®] 2.1 + 0.5 mM Cys + 4 mM GSH	10 μM	(-)	0.125 μM	(-)		
3	DsbC		20 μM	(-)	(-)	20 μM		
4	GSSG		(-)	1 mM	(-)	(-)	PCR product (1 ng/μL)	37°C, 4 h

SDS-PAGE (0.25 – 0.5 μL of reaction mix/lane) → Western Blotting (anti-FLAG) or Gel staining (Oriole)

GLuc activity assay: BioLux Gaussia Luciferase Assay Kit (NEB#E3300S) (2 μL PURE reaction mix/50 μL assay soln.)



Acid phosphatase (AppA)

Organism: *Escherichia coli*
Synthesized region: 23Gln-26Leu (+FLAG)
Length: 421 a.a.
Molecular weight: 45,959 Da
No. of disulfide bonds: 4

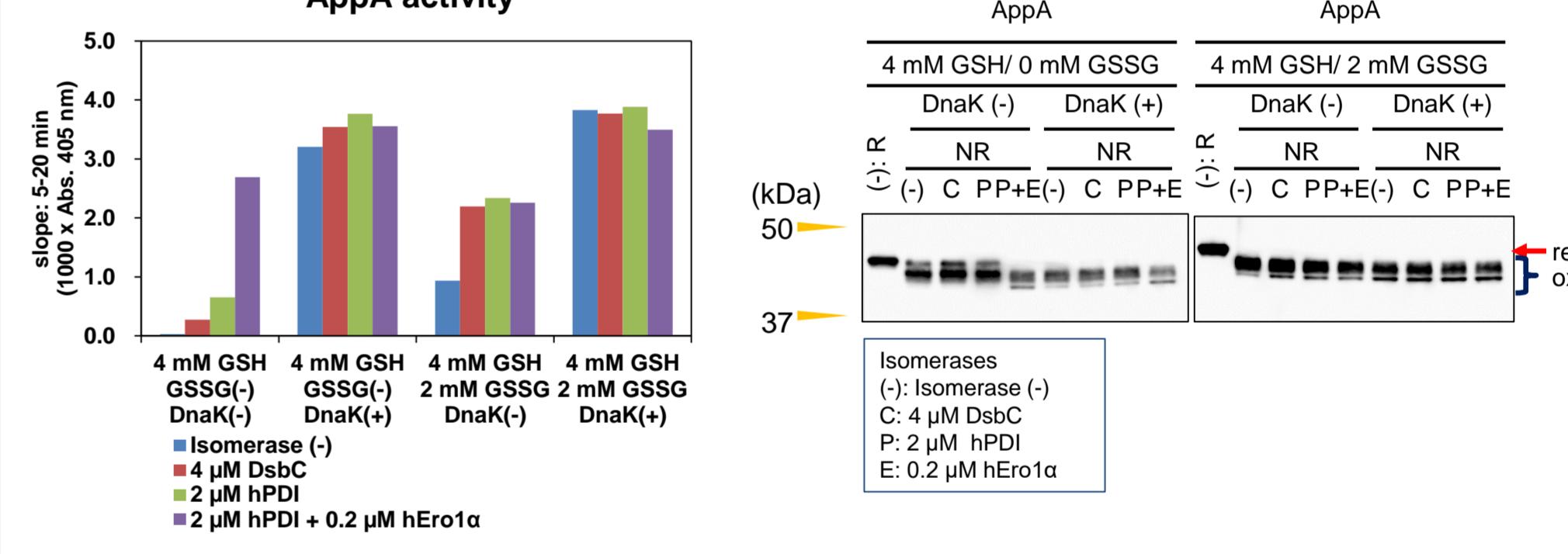
Experimental condition of protein synthesis

No.	Factors	Basic condition	hPDI	GSSG	hEro1α	DsbC	Template DNA	Temp. & Time
1	hPDI		2 μM	2 mM	(-)	(-)		
2	hPDI + GSSG		2 μM	0/2 mM	(-)	(-)		
3	hPDI + hEro1α	PUREflex [®] 2.1	2 μM	(-)	0.2 μM	(-)		
4	DsbC + GSSG	+ 0.5 mM Cys + 4 mM GSH	(-)	0/2 mM	(-)	4 μM		
5	DnaK mix	5 μM DnaK 1 μM DnaJ 1 μM GrpE	0/2 μM	0/2 mM	0/0.2 μM	0/4 μM	PCR product (1 ng/μL)	37°C, 4 h

SDS-PAGE (0.5 μL of reaction mix/lane) → Western Blotting (anti-FLAG)

AppA activity assay: Assay of PNPP dephosphorylation activity (1 μL PURE reaction mix/100 μL assay soln.)

AppA activity



6. Summary