

PUREfres[®] : The Re-built Protein Factory

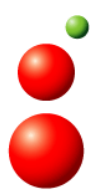


Takashi (Ebi) Ebihara, Ph.D.
COO
GeneFrontier Corporation



NextGenBiomed
24-25 of March, 2026

Corporate Summary



GeneFrontier

Founded: ***Oct 13th, 2010 (renewed)***

Shareholder: ***KANEKA Corporation (100%)***

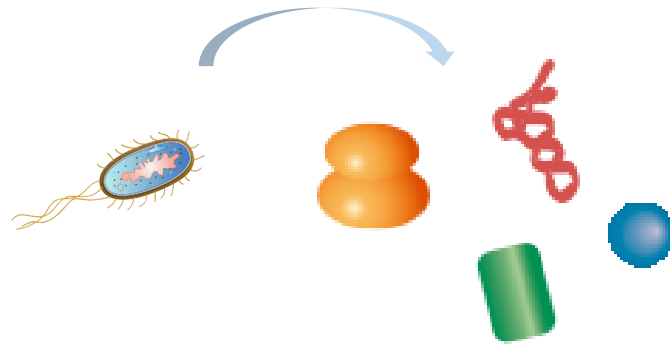
People: ***17 (Ph.D. 8, MS 1)***

Place: ***Chiba, Japan***

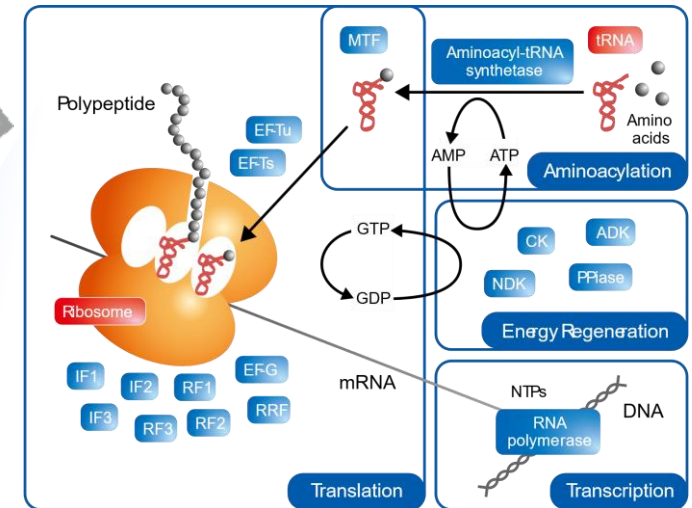
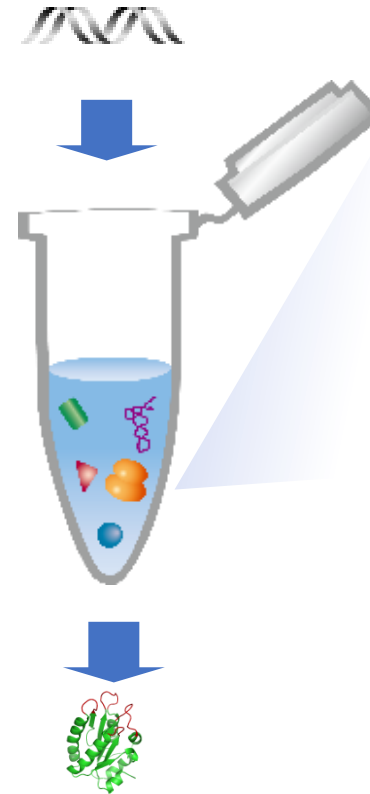
Mission: ***Rebuilding and Manipulating Biological system
for Inspiring the world!***



**Only necessary molecules
for transcription/translation**



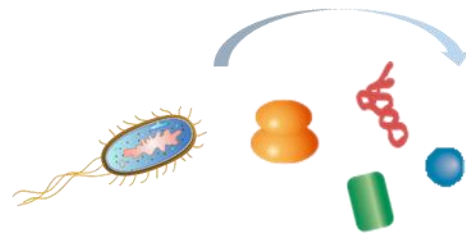
=Breaking down & Building up



PURE system

(Protein synthesis Using Recombinant Elements)

***Shimizu Y. et al. Nature Biotechnology
vol 19, p751–755 (2001)***



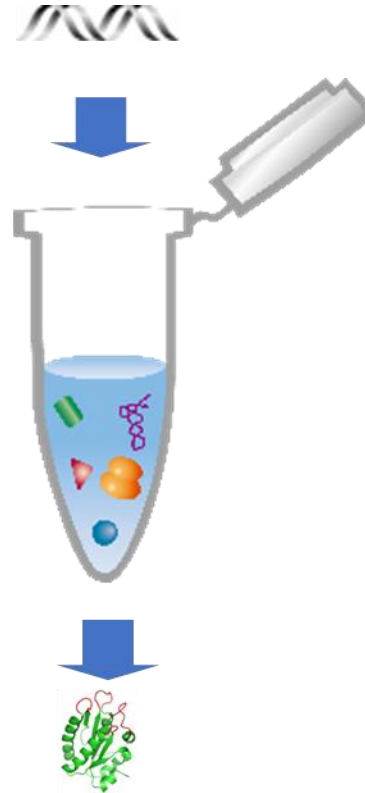
PURE system

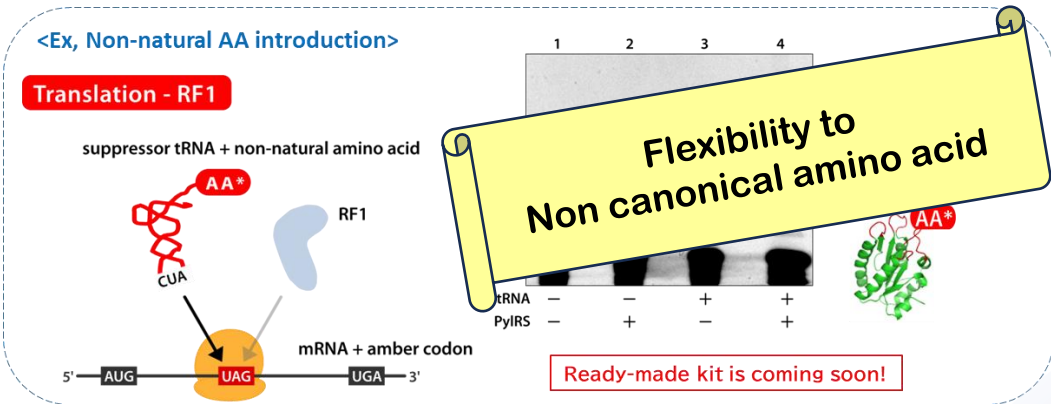


Reconstituted cell-free protein synthesis kit

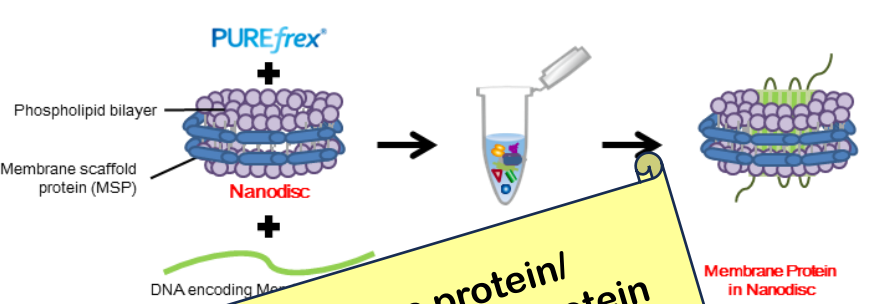
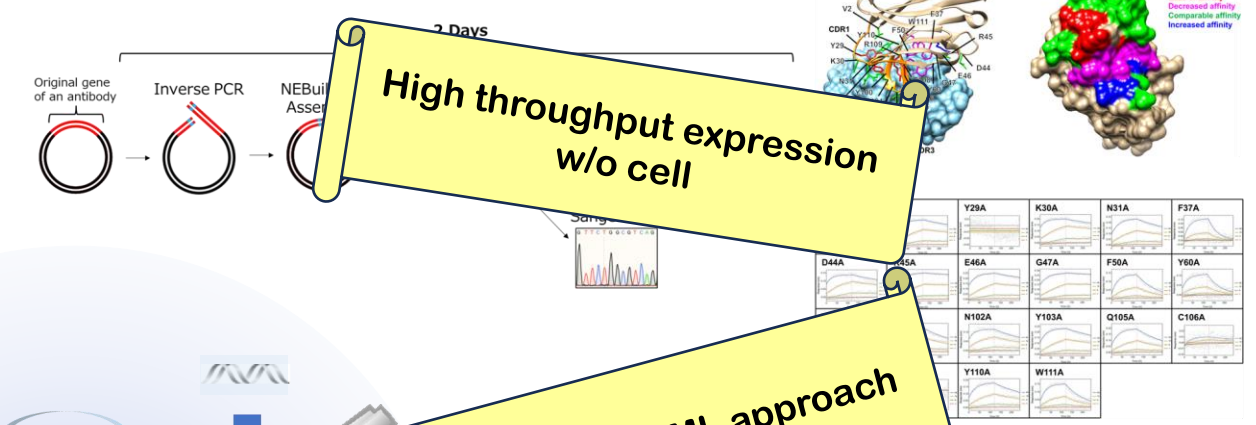
PUREfrex[®]

For Designing Central Dogma





FASTIA: Fast Interaction Analysis



**Membrane protein/
Difficult-to-express protein**

PURE system

Reconstituted cell-free protein synthesis kit

PUREfres

For Designing Central Dogma

Suitable for AI/ML approach

Speedy & Scalable & Reproducible reaction

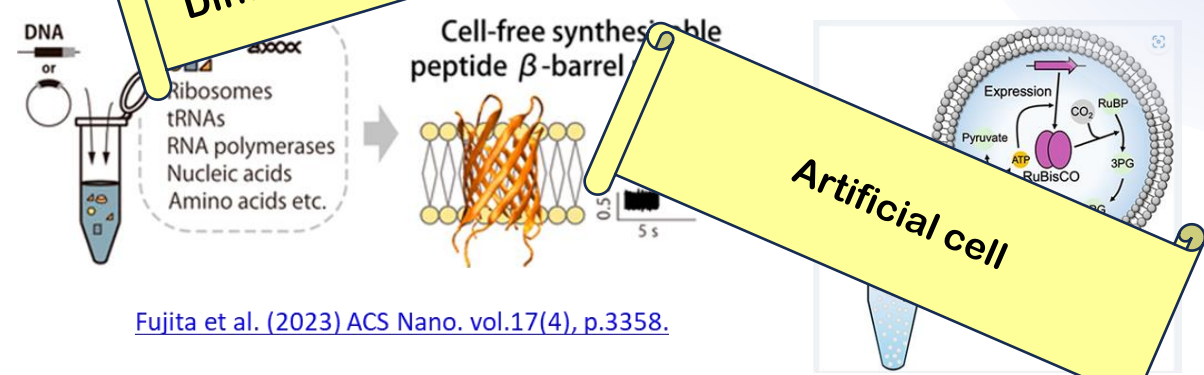
Suitable highspeed

Express

Design

Synthesized DHFR mg/mL

Ver. 1.0	Ver. 2.0/2.1
~360 mg/mL	~1000 mg/mL



In vitro display screening for Directed evolution/molecular evolution engineering

Iteration

Negative selection

FN3con-CaM-BP

- Antigen + Antigen

1. Pool synthesis

2. Protein-cDNA complex synthesis using cDNA display

900,000 sequences

PA tag

cDNA

μ L ~ L scale

Fujita et al. (2023) ACS Nano, 9(6):2846-2857.

Sugii et al. (2023) Synth. Biol. vol.8, p1

Tsuboyama et al. (2023) Nature, 620, p434.

Fujita et al. (2023) ACS Nano. vol.17(4), p.3358.

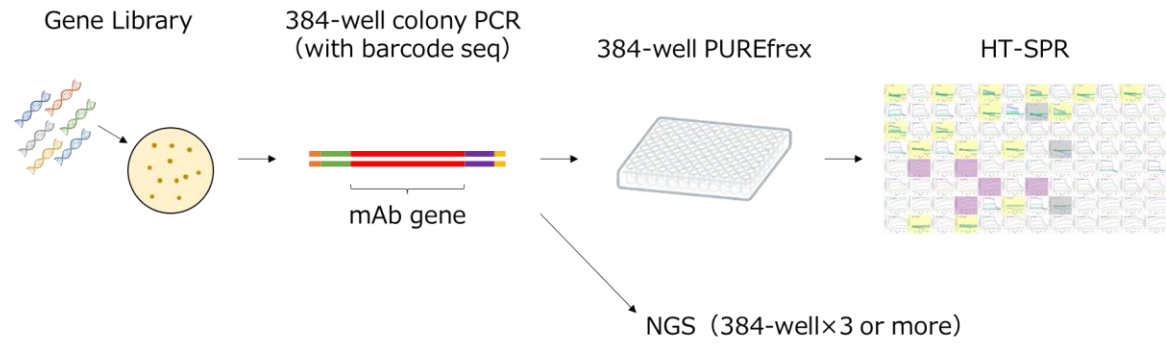
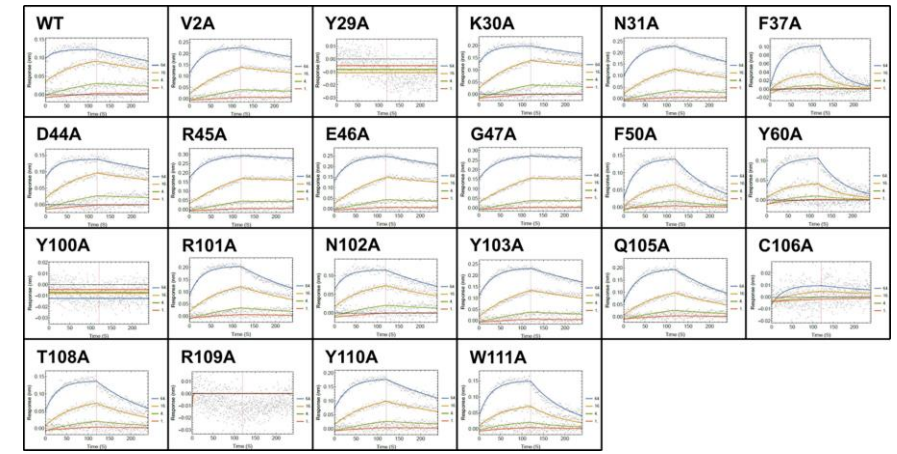
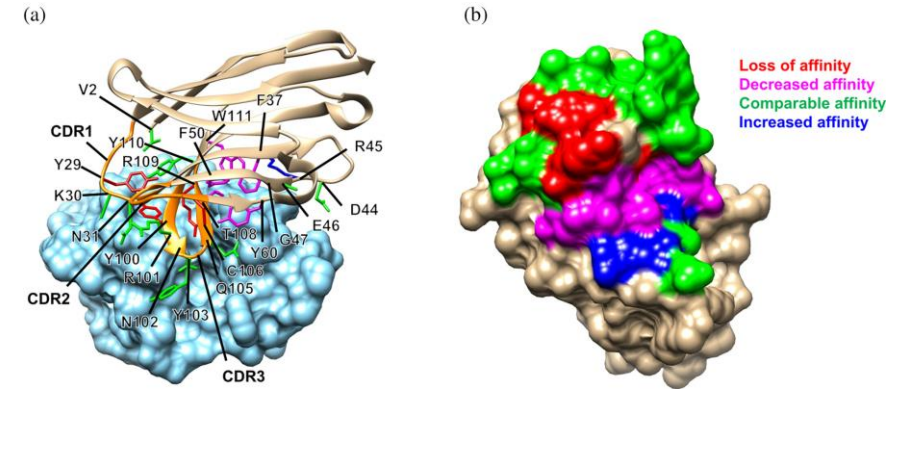
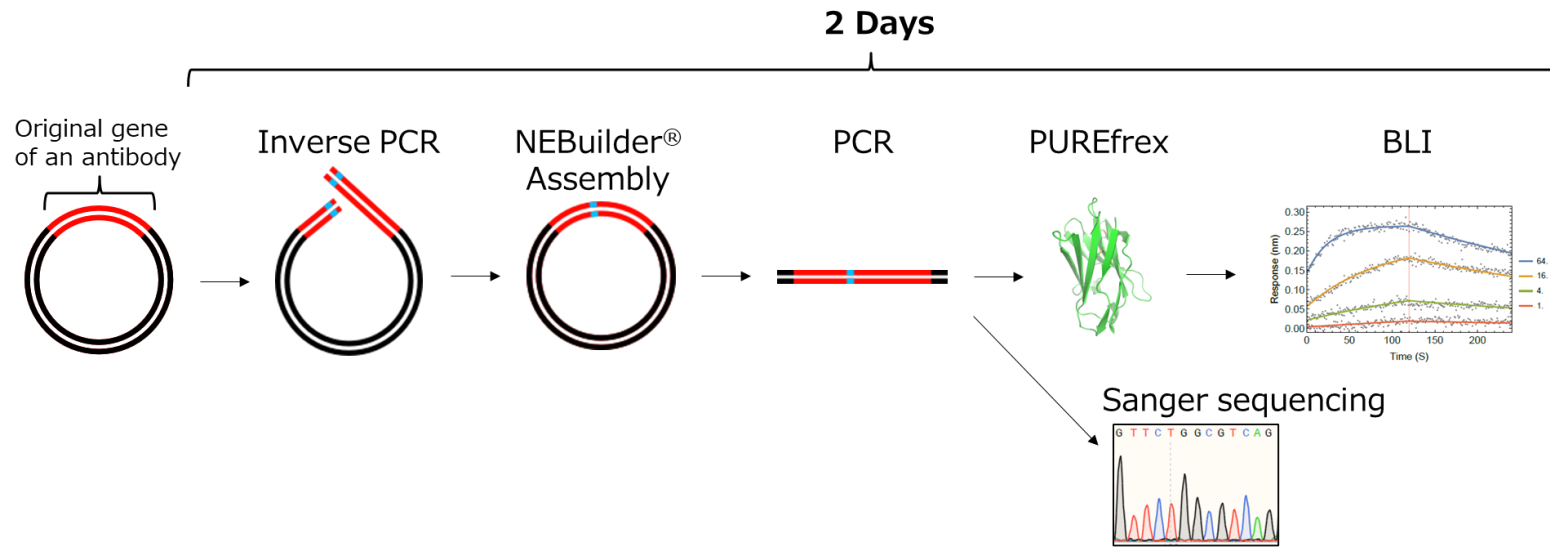
Sugii et al. (2023) Synth. Biol. vol.8, p1

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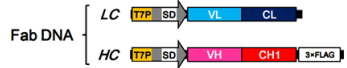
-Improve Validation from Weeks to Days-

FASTIA: Fast Interaction Analysis

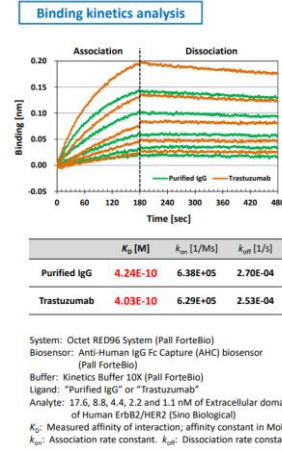
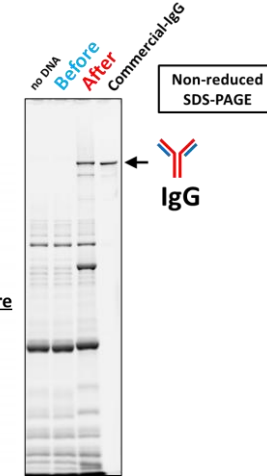
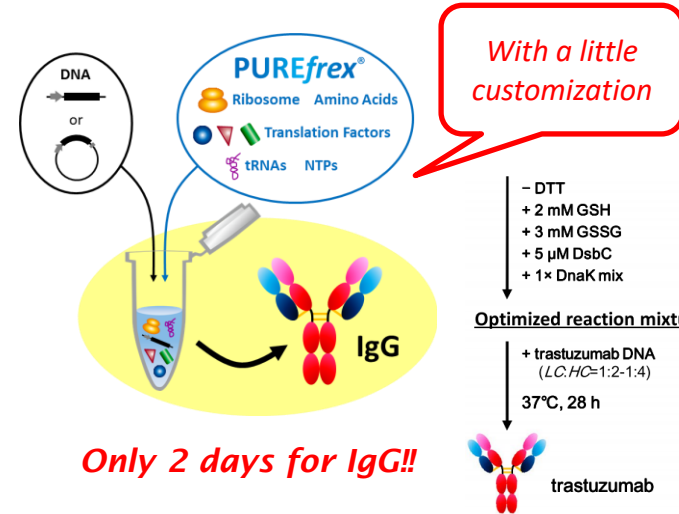
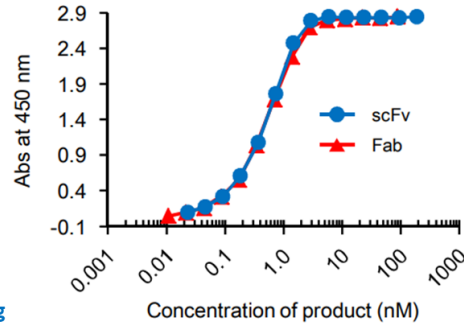


[Matsunaga et al. \(2025\) Protein Sci. Mar;34\(3\):e70065. doi: 10.1002/pro.70065.](https://doi.org/10.1002/pro.70065)

-Expression of scFv, Fab, IgG and more-

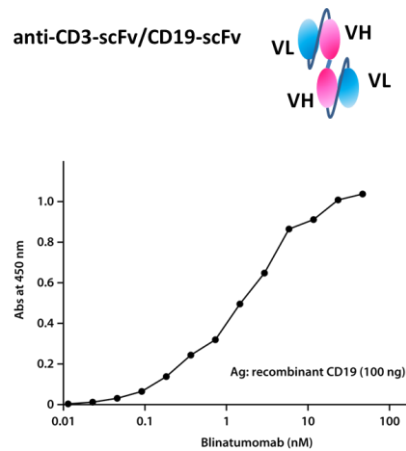


Activity

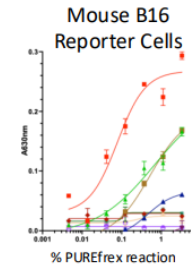
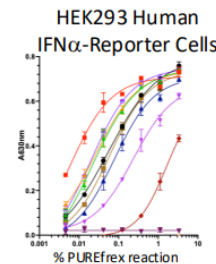


[Murakami et al. \(2019\) Sci. Rep. vol.9, p.671. \(Supplementary Information\)](#)

[Murakami et al. \(2019\) Sci. Rep. vol.9, p.671.](#)

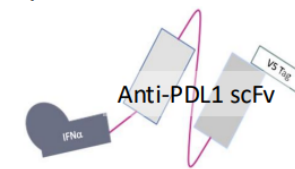


Round 1: IFN-α variants tested by in vitro transcription/translation



- Universal IFN-α
- hIFN_u_L152F
- R121K
- R121K_Q125R
- R121K_Q125R_K132T
- Y86C_R121K_Q125R
- Y86C_R121K_Q125R_K132T
- mIFN2
- IFN_u2b
- scFv control
- Human IFN_u2b (starts at 1nM)
- No DNA

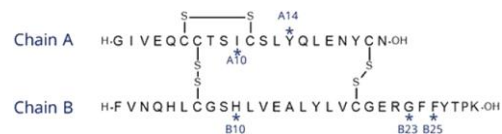
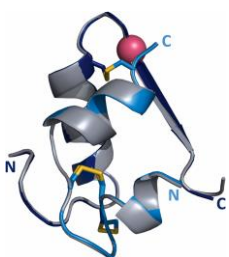
IFN-α variants were generated as IFNα-scFv fusions to identify mutations affecting activity in mouse and human cells



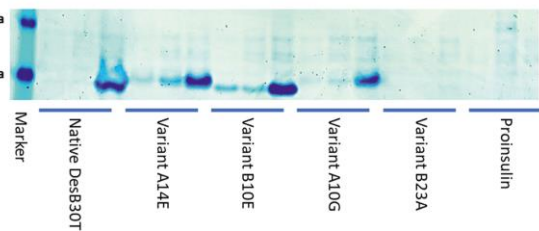
IFNα-scFv protein was generated with an anti-PDL1 antibody by in vitro transcription/translation using the PUREfres[®] system (CosmoBIO USA). Protein mixture was serially diluted and protein was captured via an anti-V5 tag coated to the wells of the plate. Proteins were assayed for PD-L1 binding to assess expression (data not shown) or the ability to stimulate an IFN-α response in receptor reporter cell lines

[Killebrew et al. \(2024\) SITC 2024 Annual Meeting \(Poster, BONUMTX.com\).](#)

-Application for complex molecule-



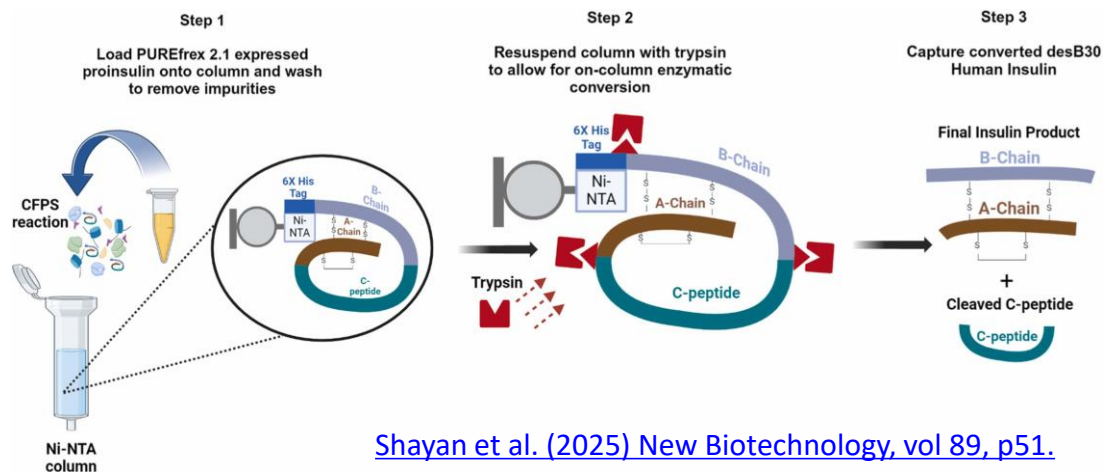
-	-	+	-	-	+	-	-	+	-	-	+	-	+	-	+	-	+	-	+	4 mM GSH/3 mM GSSG
-	+	+	-	+	+	-	+	+	-	+	+	-	+	+	-	+	+	-	+	0.375 mg/ml DsbC
+	+	-	+	+	-	+	+	-	+	+	-	+	+	-	+	+	-	+	-	2 mM DTT



[Jensen et al. \(2021\) Protein Expr. Purif., 186, 105910.](#)

	1	2	3	4	5	6	7	8	9	10
	Proinsulin Aspart	Proinsulin Lispro	Proinsulin Glargine	Regular Proinsulin	Insulin A Chain	Insulin B Chain	Insulin A Chain Heterodimer	Insulin B Chain Heterodimer	Oxytocin	Glucagon
PURE	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓
Clm24	✗	✗	✗	✗	✓	✗	✓	✓	✓	✓
BL21	✗	✗	✗	✗	✓	✗	✓	✓	✓	✗
759	✓	✓	✓	✓	✓	✓	✓	✓	✓	✗
	11	12	13	14	15	16	17	18	19	20
	Glucagon Like Peptide 1 mutant (GLP-1 mut)	Glucagon Like Peptide 1 (GLP-1)	Insulin Like Growth Factor	Growth Hormone (GH)	Leptin	Vaso-pressin	Angiotensin II	Parathyroid Hormone (PTH)	Somato-statin	Leuprolide
PURE	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓
Clm24	✓	✓	✗	✓	✓	✓	✓	✓	✓	✓
BL21	✗	✗	✗	✗	✗	✗	✓	✓	✓	✓
759	✓	✓	✗	✓	✓	✓	✓	✗	✓	✓

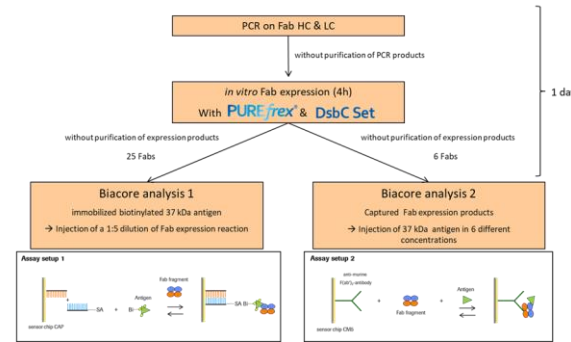
[DeWinter et al. \(2023\) ACS Synth. Biol. vol.12, 4, p1216. \(Supplementary Information\)](#)



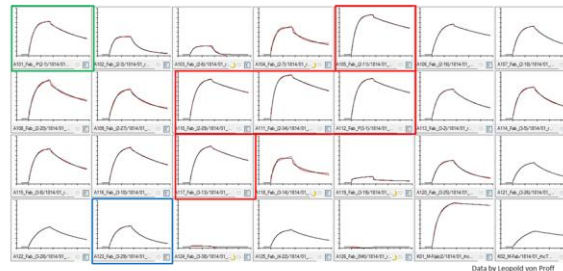
[Shayan et al. \(2025\) New Biotechnology, vol 89, p51.](#)



In vitro expression and Biacore analysis of Fab fragments



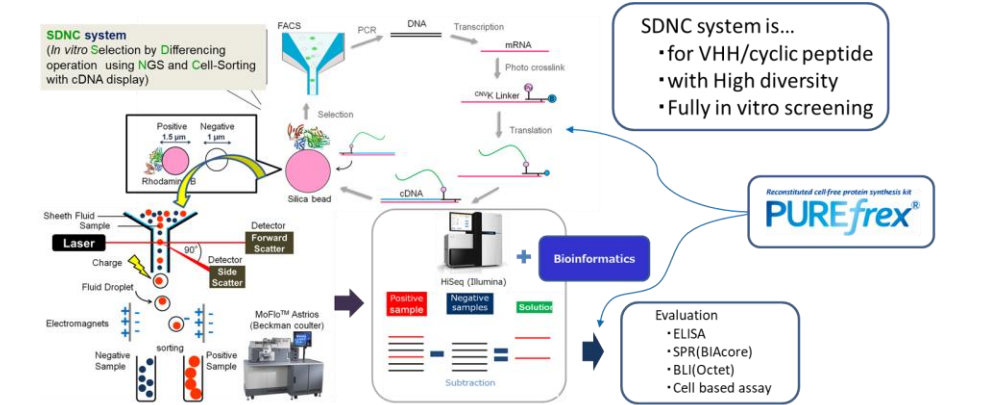
Kinetic analysis of 25 Fab binders



→ Selection of Fabs for further kinetic analysis

EME Epsilon Molecular Engineering

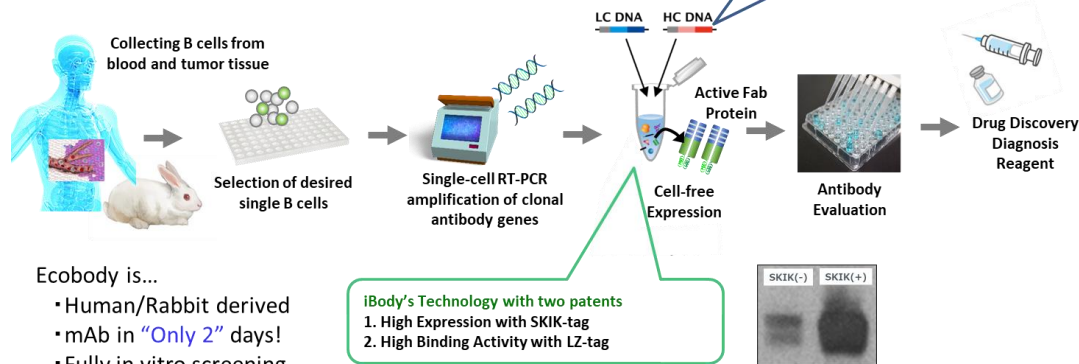
Molecular Design for Human Life



<https://www.epsilon-mol.co.jp/eng/>



iBody's Ecobody Technology



- Ecobody is...
- Human/Rabbit derived
 - mAb in "Only 2" days!
 - Fully in vitro screening
 - No culture

<https://www.ibody.co.jp/en/>

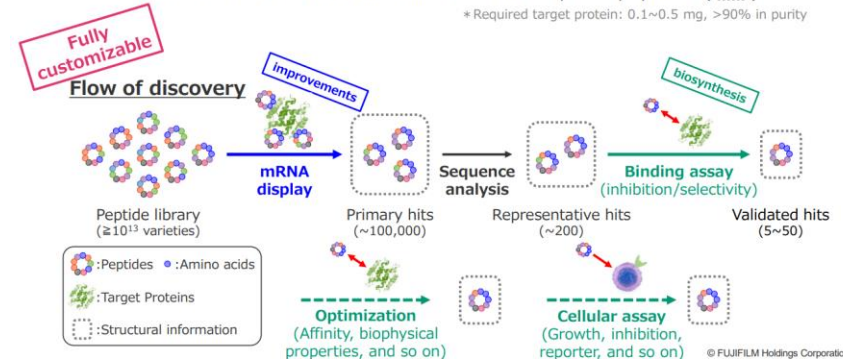
FUJIFILM peptide discovery services

collaborated with PUREfres

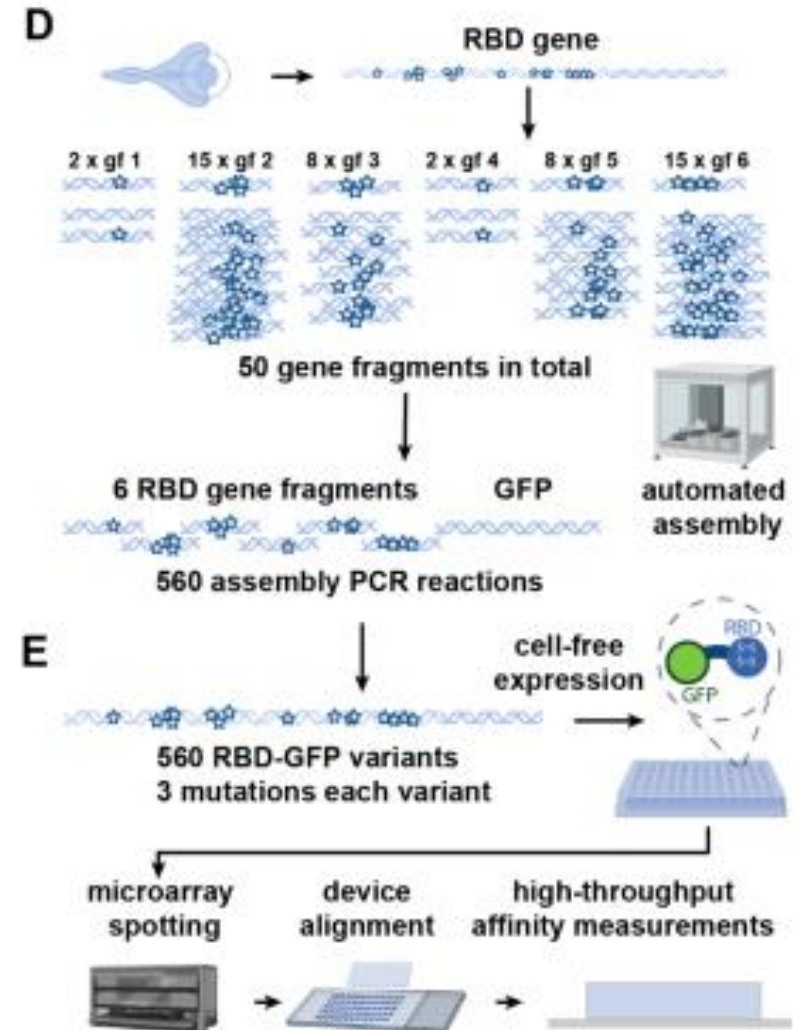
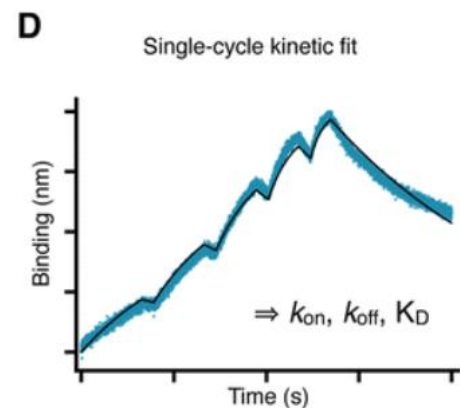
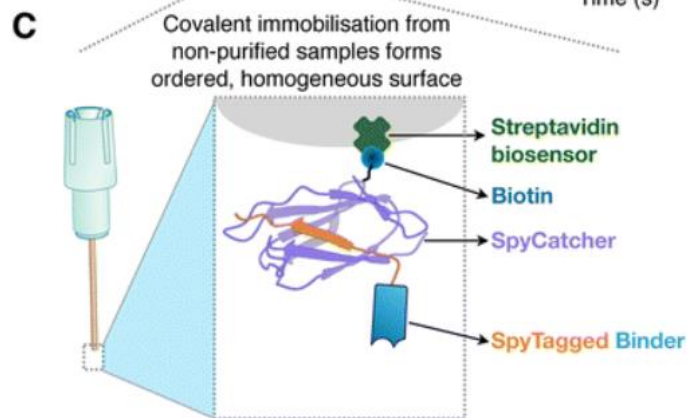
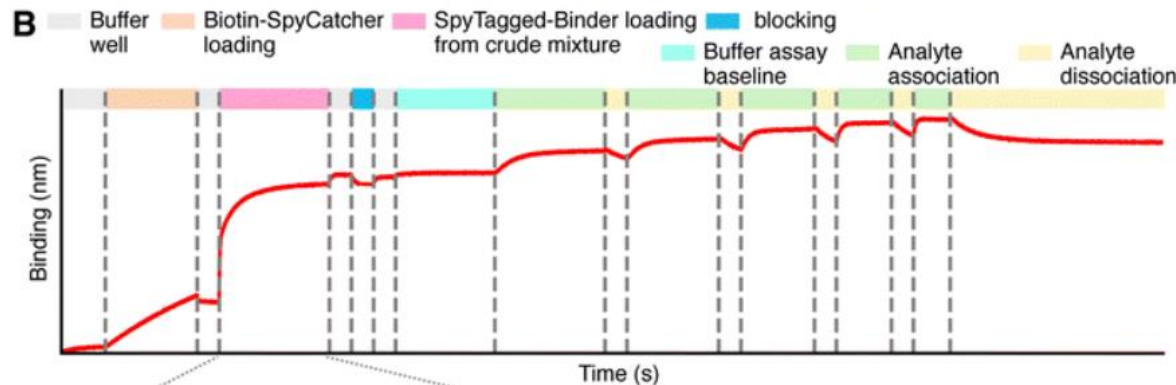
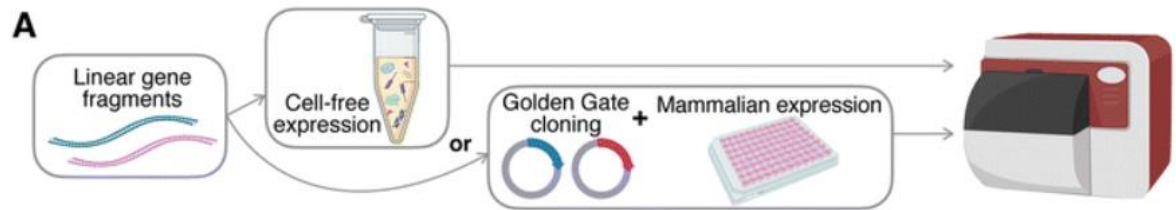
- ✓ **Innovative improvements** in mRNA display enable screening from $>10^{13}$ peptides
 - ✓ **Practical biosynthesis & assays** enable rapid selection and activity explorations.
- Peptides hits with **wide varieties** and **high-affinities** can be obtained.

We provide a CRO service, in which we receive target (🧬) from the customer* and return the structural information of the acquired peptides (📄).

* Required target protein: 0.1~0.5 mg, >90% in purity



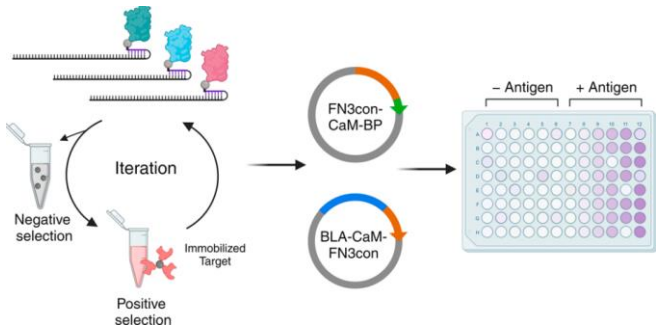
-Broad applications, yet to come!-



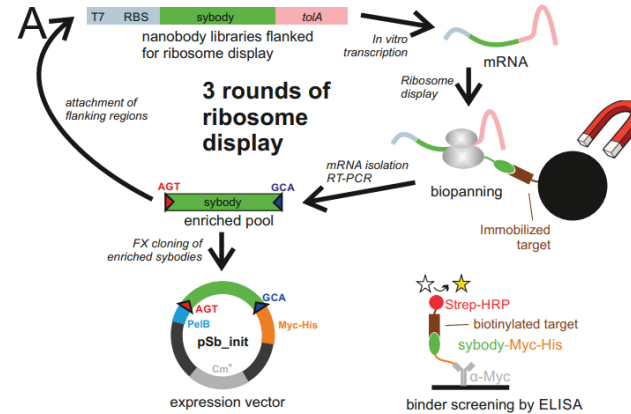
[Predenia et al. \(2025\) RSC Chem. Biol, 6\(1313\).](https://doi.org/10.1039/C5SC01313A)

[Grasemann et al. \(2025\) bioRxiv. https://doi.org/10.1101/2025.09.23.678000.](https://doi.org/10.1101/2025.09.23.678000)

-Broad applications, yet to come!-

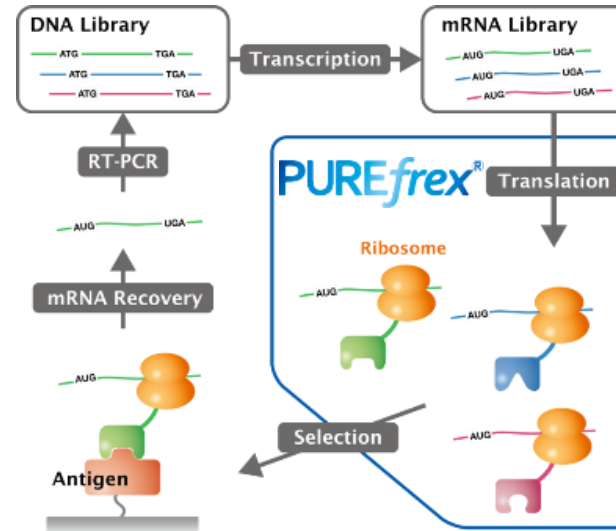


[Chui Z. et al. \(2024\) ACS Sens, 9\(6\):2846-2857.](#)

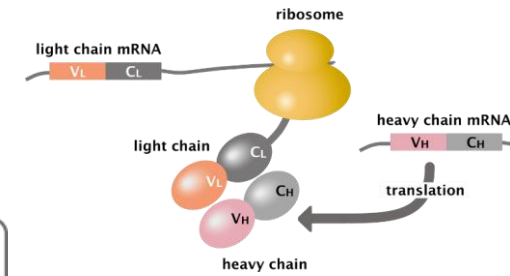


[Zimmermann I. et al. \(2018\) eLife, 7, e34317.](#)

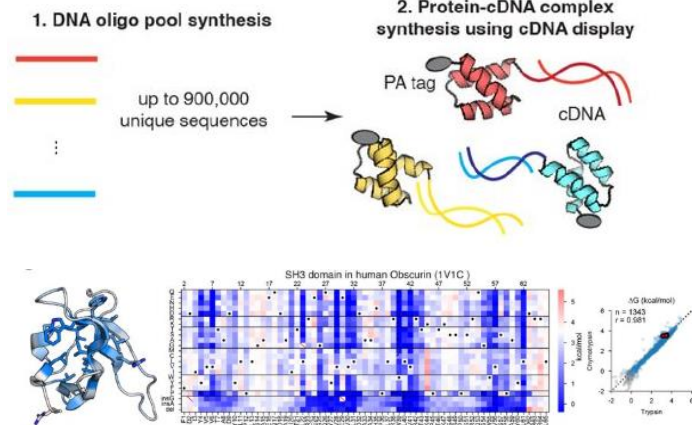
in vitro protein selection technology PUREfres[®] RD



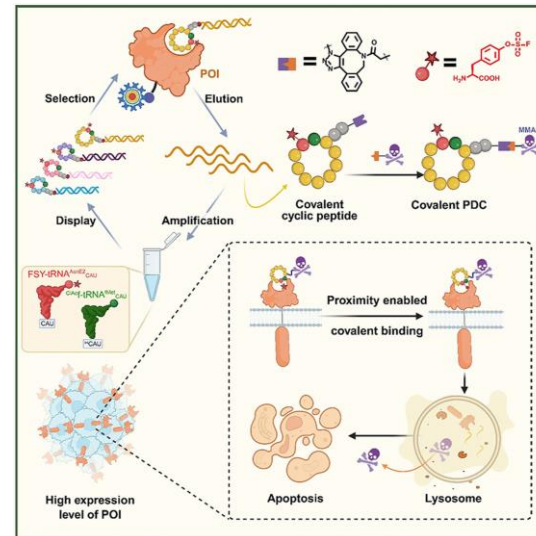
Licensed technology under JP4931135 etc.



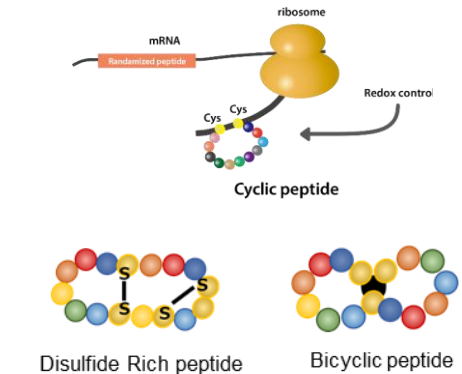
Licensed to
SUTRO
BIOPHARMA



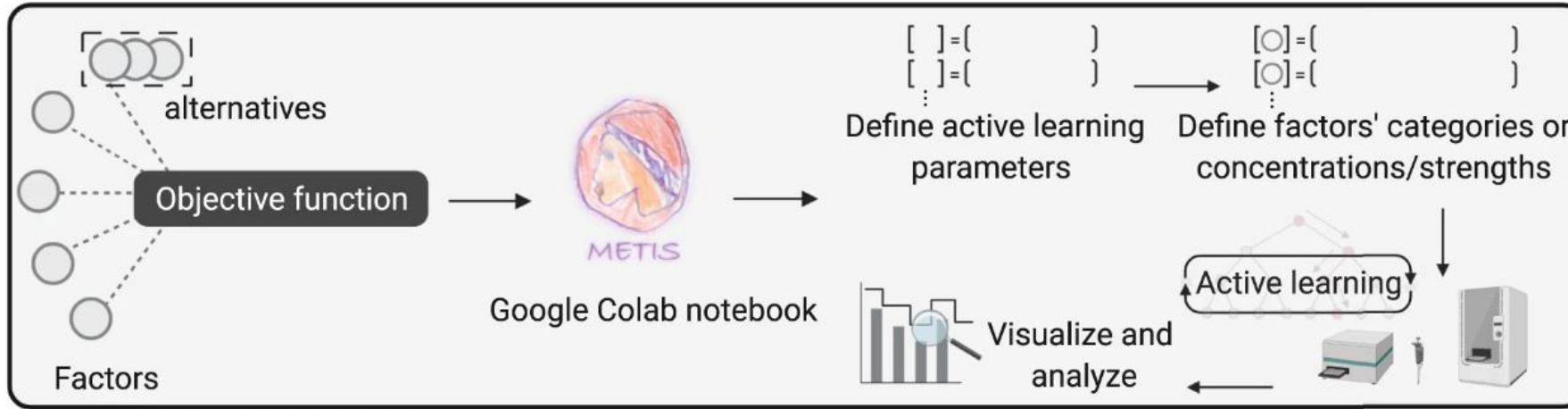
[Tsuboyama et al. \(2023\) Nature, 620, p434.](#)



[Wang et al. \(2025\) Acta Pharmaceutica Sinica B. vol.15\(5474\).](#)

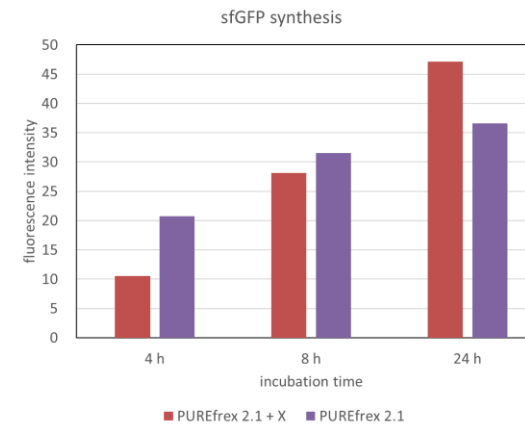
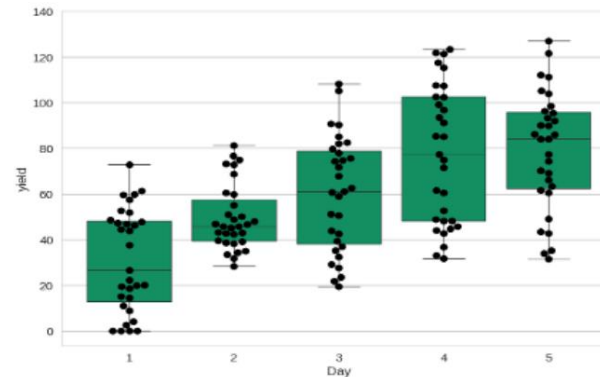


-Broad applications, yet to come!-



[Pandi A et al. \(2022\) Nature Communications, 13, 3876.](https://doi.org/10.1038/s41467-022-28388-1)

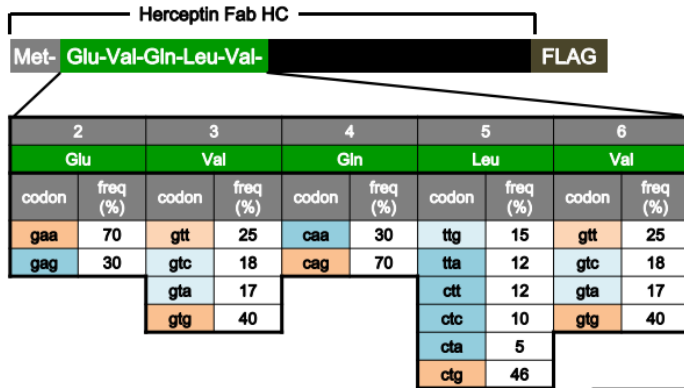
Round	3			4		5			6			
Composition No.	2	21	27	4	13	14	2	8	29	1	7	8
K Glutamate	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
Mg Acetate	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
Spermidine	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
Creatine Phosphate	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
ATP	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
GTP	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
CTP/UTP	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
tRNA	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue
IF2	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red
EF-G	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red
EF-Tu/EF-Ts	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red
T7 RNAP	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red	Red
Ribosome	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue	Blue



- ✓ Perfect fit to AI/ML approach with great controllability & reproducibility.
- ✓ Unique expression platform will give you great advantage in R&D.

-KSF; AT rich codon on N-term-

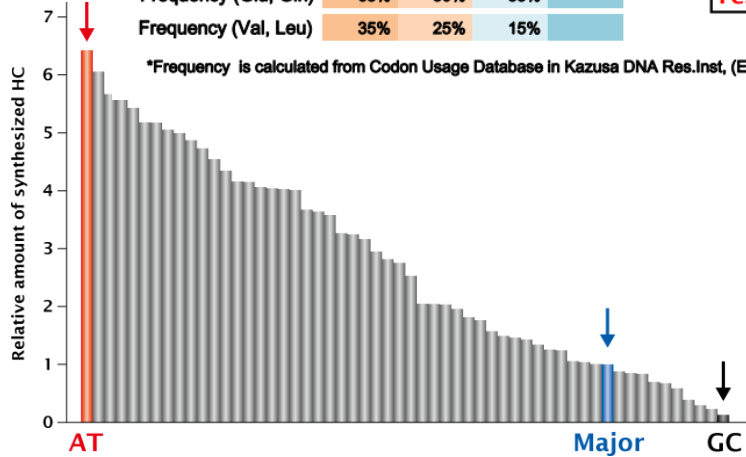
Fab Heavy Chain (Herceptin)



All clones; 384
Tested clones; 56

Frequency (Glu, Gln)	65%	50%	35%
Frequency (Val, Leu)	35%	25%	15%

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



Design of DNA template is important.

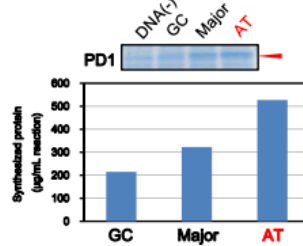
Manual is Free to download from our Web site here.



PD1

Organism: *Homo sapiens*
Synthesized region: 36Thr-150Glu(-Hisx8)
Length: 124 a.a.
Molecular weight: 14,148 Da

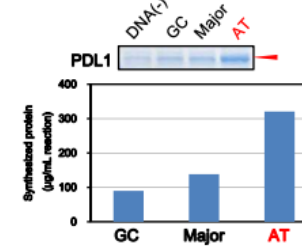
N-term type	1	2(36)	3(37)	4(38)	5(39)	6(40)	GC(%) 1-6 a.a.
GC	atg	acc	ttc	tcc	cag	gag	67%
Major	atg	acc	ttt	tct	cag	gag	56%
AT	atg	act	ttt	tca	cca	gct	39%



PDL1

Organism: *Homo sapiens*
Synthesized region: 18Ala-239Thr(-Hisx8)
Length: 231 a.a.
Molecular weight: 26,593 Da

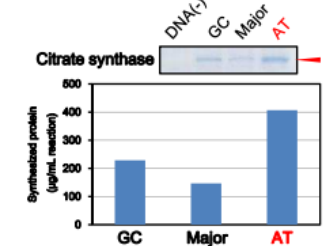
N-term type	1	2(18)	3(19)	4(20)	5(21)	6(22)	GC(%) 1-6 a.a.
GC	atg	acc	ttc	acc	gtg	acc	61%
Major	atg	gag	ttt	acc	gtg	acc	56%
AT	atg	gct	ttt	act	gta	aca	33%



Citrate Synthase

Organism: *Saccharomyces cerevisiae*
Synthesized region: 38Ser-479Asn
Length: 443 a.a.
Molecular weight: 49,346 Da

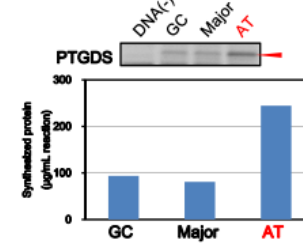
N-term type	1	2(38)	3(39)	4(40)	5(41)	6(42)	GC(%) 1-6 a.a.
GC	atg	tcc	tcc	ggc	tcc	gag	67%
Major	atg	tct	tct	ggc	tct	gaa	44%
AT	atg	tca	tca	gct	tca	gaa	39%



PTGDS

Organism: *Homo sapiens*
Synthesized region: 23Ala-190Gln
Length: 169 a.a.
Molecular weight: 18,829 Da

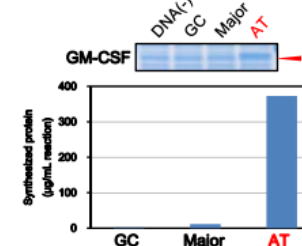
N-term type	1	2(23)	3(24)	4(25)	5(26)	6(27)	GC(%) 1-6 a.a.
GC	atg	gca	cag	gaa	gca	cag	61%
Major	atg	gag	cag	gaa	gag	cag	72%
AT	atg	gca	cct	gaa	gct	caa	50%



GM-CSF

Organism: *Homo sapiens*
Synthesized region: 18Ala-144Glu
Length: 128 a.a.
Molecular weight: 14,808 Da

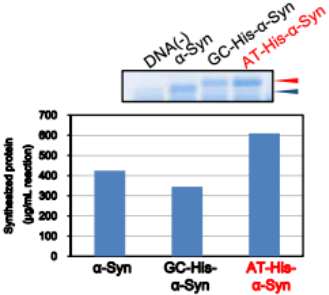
N-term type	1	2(18)	3(19)	4(20)	5(21)	6(22)	GC(%) 1-6 a.a.
GC	atg	gag	cag	gag	cgc	tcc	83%
Major	atg	gag	cag	gag	cgc	tct	78%
AT	atg	gca	cct	gct	aga	tca	50%



His-α-Synuclein

Organism: *Homo sapiens*
Synthesized region: (Hisx6)-(Gly-Ser)-2(10)Asp-140(148)Ala
Length: 148 a.a.
Molecular weight: 15,427 Da

Tag type	1	2	3	4	5	6	7	8	9	GC(%) 1-9 a.a.
GC	atg	cac	ccc	ccc	ccc	ccc	ccc	ggt	tct	59%
AT	atg	cat	cat	cat	cat	cat	cat	ggt	tct	37%



DNA design for PUREfres



Amino acid sequence entry form

Consultation is free of charge!

-KSF; Quality of DNA-

#	Construct	Size (bp)	Elegen's ENFINIA DNA	Supplier B	Supplier C
			Format	Format	Format
1	HisTEV-sfGFP(G4Y)-PPG-FLAG	978	Linear dsDNA	N/A	Linear dsDNA
2	HisTEV-PPG-sfGFP(G4Y)-FLAG	978	Linear dsDNA	N/A	Linear dsDNA
3	HisTEV-sfGFP(G4Y)-FLAG	888	Linear dsDNA	Linear dsDNA	Linear dsDNA
4	sfGFP(G4Y)-FLAG	840	Linear dsDNA	Linear dsDNA	Linear dsDNA

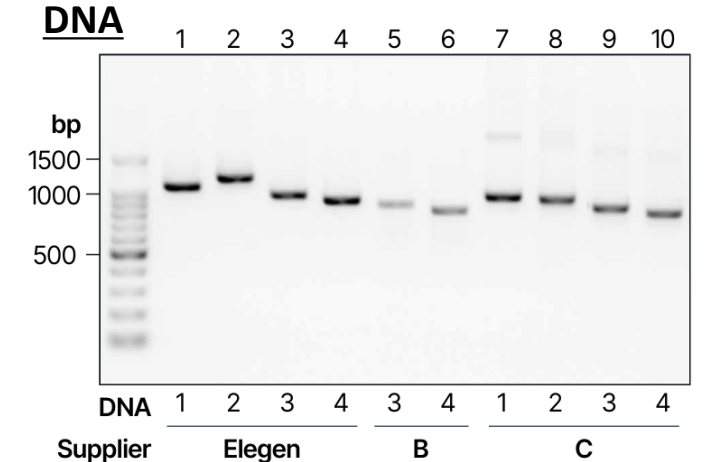
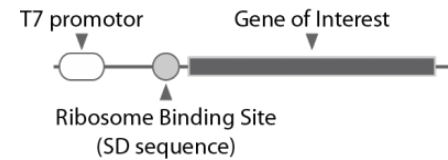
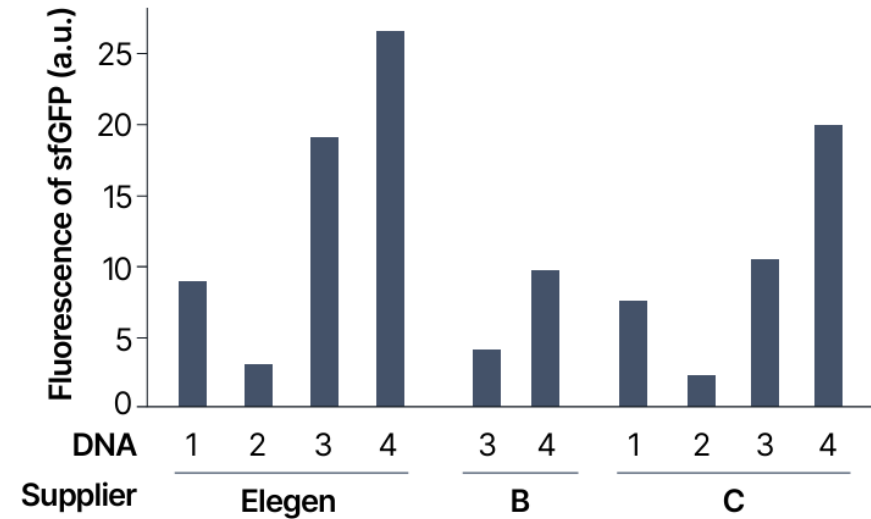
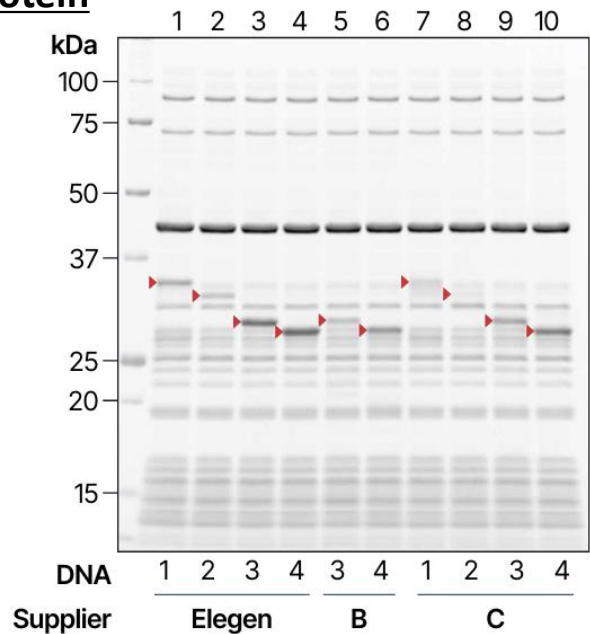


Figure 1. Analysis of DNA synthesized by three vendors. DNA synthesized by Elegen (Supplier A), Supplier B, and Supplier C was quantified using a Qubit Fluorometer (Thermo Fisher Scientific) and subjected to agarose gel electrophoresis.

Protein



To learn more about ENFINIA DNA, visit elegenbio.com or contact us at info@elegenbio.com

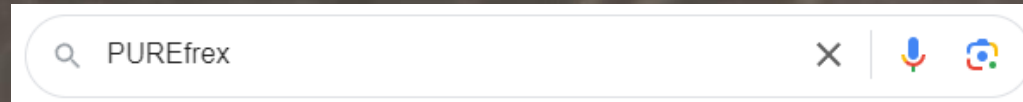
Contact information

Reconstituted cell-free protein synthesis kit

PUREfrefx[®]

*For reagent use for expression / screening of biologics
/designing central dogma*

<https://purefrefx.genefrontier.com/>



in vitro protein selection technology

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*For screening service / collaboration / technology transfer
for generation of new biologics*

Takashi Ebihara, Ph.D., COO, GeneFrontier

E-mail: ebihara@genefrontier.com





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We offer a robust VHH production platform, developed through extensive testing of diverse VHH models under a wide range of conditions:

- **mono-, bi-, and tri-specific** formats
- Isoelectric points between **pI 5 and 9**
- Diverse **hydrophobicity** properties



8 only Months

Starting from your research cell bank, we rapidly identify the optimal setup and process conditions to efficiently produce your VHH.

VHH Platform



Two expression platforms

- *P. pastoris* (*K. phaffii*)
- *E. coli*



Fermentation

- High-cell density fed-batch
- Control of critical fermentation parameters
- Animal-free media



Purification types

- Ion-Exchange
- Hydrophobic Interaction Chromatography (HIC)
- Mix mode
- Affinity



Quality control & release

- GMP QC package
- Additional custom QC development available
- QP release



Quick to clinic



Designed for Tox & PhI



GMP quality



End-to-End assistance