

Rebuilding Expression System & Its Applications For R&D Of Biologics.

Reconstituted cell-free protein synthesis kit



Biologics US 2024 3-4 of Oct, 2024

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



Corporate Summary



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

Shareholder: KANEKA Corporation (100%)

People: **14** (Ph.D. 8, MS 1)

Place: Chiba, Japan





Mission: Rebuilding and Manipulating Biological system

for Inspiring the world!

For more information





-Customize expression toolbox for your research-

Only necessary molecules for transcription/translation

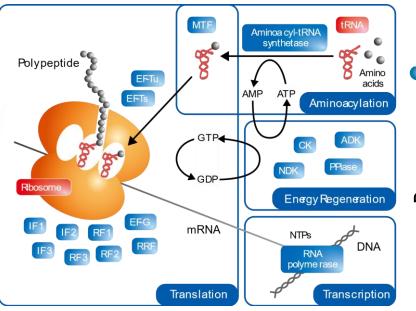


- ✓ Rebuilt cell-free system
- ✓ Tunable for your biologics
- ✓ Simple, Fast
- ✓ Suited for High throughput system
- ✓ Very low contaminants

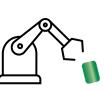


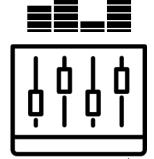


Totally constructive, molecular based system







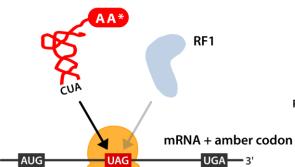


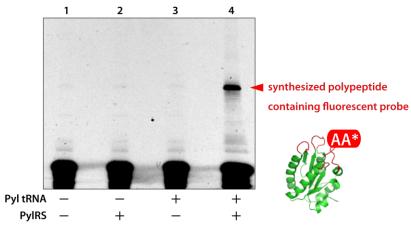
Solve of the state of the state



Translation - RF1

suppressor tRNA + non-natural amino acid





Ready-made kit is coming soon!

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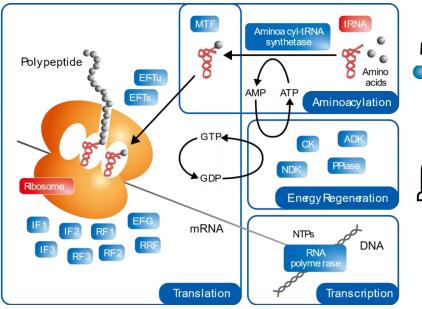


For more information



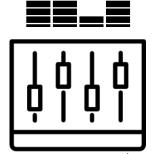


Totally constructive, molecular based system

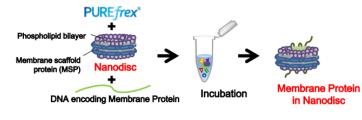








<Ex, Membrane protein with Nanodisc; artificial membrane-like structure>



10 μM Nanodisc CLDN1-AT

| (-) (+) (-) (+) | to | tal | SI | ΙÞ | ppt | | |
|-----------------|-----|-----|-----|-----|-----|-----|--|
| | (-) | (+) | (-) | (+) | (-) | (+) | |

The condition of membrane protein synthesis

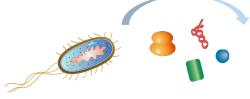
| Reaction mix | Template DNA | Incubation |
|--|--------------|------------|
| PURE <i>frex</i> ® 2.0 +Nanodisc (MSP1E3D1-His POPC*, final 10 μM) | PCR product | 37℃, 4 h |

*Ref: Denisovet al. (2007) J.Biol.Chem., vol. 282, p. 7066.

Solubilized hCLDN1 was synthesized using PURE frex® and Nanodisc.

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Only necessary molecules for transcription/translation

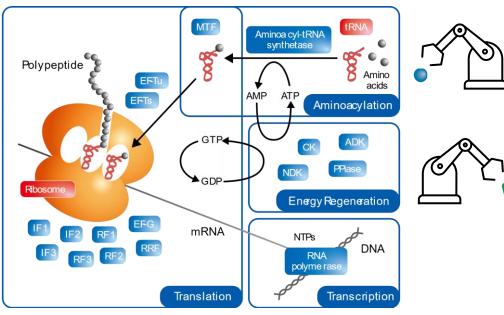


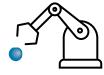
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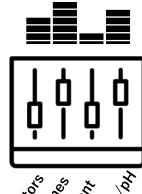
Totally constructive, molecular based system





For more information





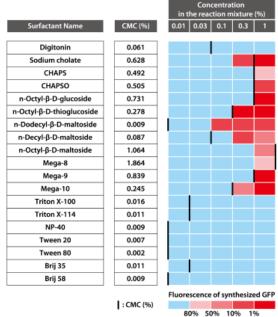
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Experimental conditions for protein synthesis

| Reaction mixture | Incubation | Template DNA |
|--|-------------|--------------------------------------|
| PUREfrex®2.1 (4 mM GSH) + Sufractants | 37°C 4 h | sfGFP PCR product (1 ng/µL) |

→ Measurement of GFP fluorescence

- · Most surfactants did not inhibit the protein synthesis reaction by PUREfrex® below the CMC.
- Some surfactants such Triton X-100 and Tween 20 could be used even above the CMC.



100%: (-) surfactant

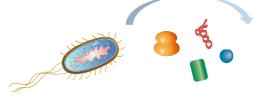
For more information





-Customize expression toolbox for your research-

Only necessary molecules for transcription/translation

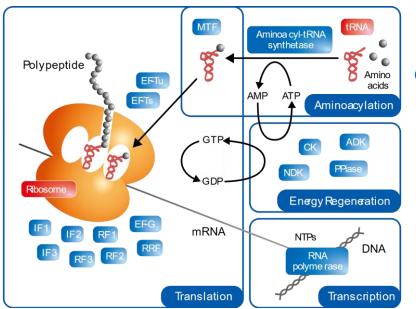


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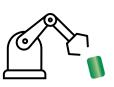


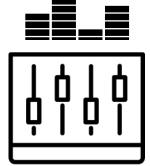


Totally constructive, molecular based system









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<u>Versatile and Robust</u> <u>Platform for protein synthesis</u>

<u>Huge potential as</u> <u>New platform in Biotech industry</u>









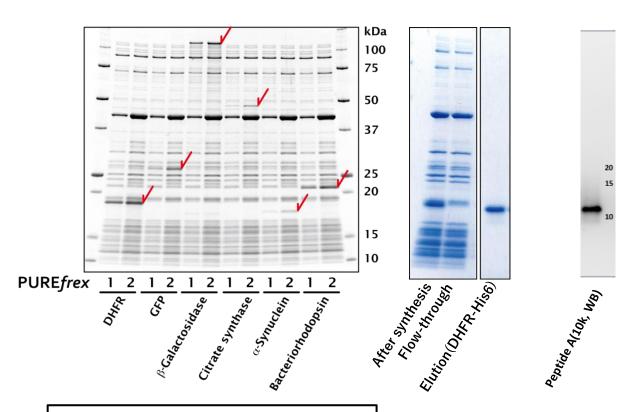


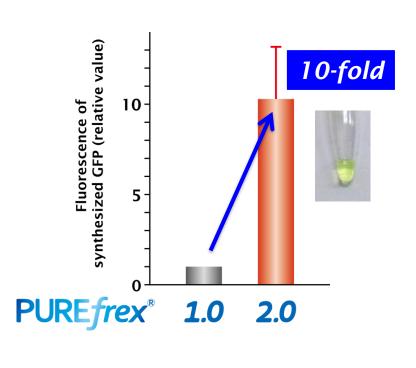




-Having good productivity-



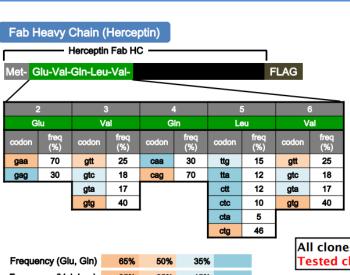




- Reaction at 37°C for 4 h
- 0.5 µL of reaction mix/lane
- stained with Oriole (Bio-Rad) and analyzed with an image analyzer (LAS)
- ✓ Good expression for many proteins, small to large.
- ✓ Good purity with simple purification.
- ✓ Good productivity, ~g/L.







All clones; 384 Tested clones; 56 Frequency (Val, Leu)

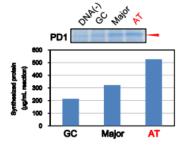
*Frequency is calculated from Codon Usage Database in Kazusa DNA Res.Inst, (E. coli K-12 strain) Major

Design of DNA template is important. Manual is Free to download from our Web site here

Organism Homo sapiens 36Thr-150Glu-(Hisx8) 124 a.a.

14,148 Da Molecular weight

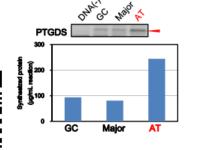
| N-term | 1 | 2(38) | 3(37) | 4(38) | 5(39) | 6(40) | GC(%) 1-6 a.a. |
|--------|-----|-------|-------|-------|-------|-------|-------------------|
| type | Met | Thr | Phe | Ser | Pro | Ala | 1-6 a.a. |
| GC | alg | 800 | ttc | toc | cog | 909 | 67% |
| Major | alg | acc | ttt | tct | ccg | gcg | 56% |
| AT | atg | act | ttt | tca | cca | gct | 39% |



PTGDS

Homo sapiens Synthesized region 23Ala-190Gln 169 a.a. 18.829 Da

| N-term type | 1 | 2(23) | 3(24) | 4(25) | 5(26) | 6(27) | GC(%) 1-6 a.a. |
|----------------|-----|-------|-------|-------|-------|-------|-------------------|
| type | Met | Ala | Pro | Glu | Ala | Gin | 1-6 a.a. |
| GC | atg | gca | ccg | gee | gca | cag | 61% |
| major | atg | gcg | ccg | gaa | gcg | cag | 72% |

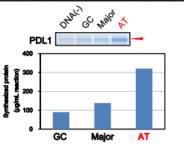


PDL1

Organism Homo sapiens Synthesized region 18Ala-239Thr-(Hisx8) 231 a.a.

26,593 Da Molecular weight

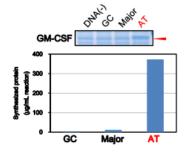
| N-term | 1 | 2(18) | 3(19) | 4(20) | 5(21) | 6(22) | GC(%) |
|--------|-----|-------|-------|-------|-------|-------|----------|
| type | Met | Ala | Phe | Thr | Val | Thr | 1-6 a.a. |
| GC | atg | gog | ttc | acc | gtg | acc | 61% |
| major | atg | gcg | ttt | acc | gtg | acc | 56% |
| AT | atg | gct | ttt | act | gta | aca | 33% |



GM-CSF

Organism Synthesized region 18Ala-144Glu 128 a.a. Molecular weight 14.608 Da

| 1 | N-term | 1 | 2(18) | 3(19) | 4(20) | 5(21) | 6(22) | GC(%) | ı |
|---|--------|-----|-------|-------|-------|-------|-------|----------|---|
| | type | Met | Ala | Pro | Ala | Arg | Ser | 1-6 a.a. | l |
| 1 | GC | atg | gog | ccg | gcg | ege | toc | 83% | ı |
| 1 | major | atg | gcg | ccg | gcg | ege | tct | 78% | l |
| 1 | AT | ato | gca | oct | act | aga | tca | 50% | ı |

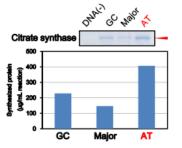


Citrate Synthase

Organism Saccharomyces cerevisiae Synthesized region 38Ser-479Asn Length 443 a.a.

49,346 Da Molecular weight

| N-term | | 2(36) | 3(39) | 4(40) | 5(41) | 0(42) | GC(%) |
|--------|-----|-------|-------|-------|-------|-------|----------|
| type | Met | Ser | Ser | Ala | Ser | Glu | 1-6 a.a. |
| GC | atg | toc | toc | gog | toc | gag | 67% |
| major | atg | tet | tct | gcg | tct | gaa | 44% |
| AT | atg | tca | tca | gct | tca | gaa | 39% |

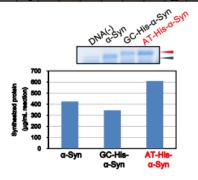


His-α-Synuclein

Synthesized region (Hisx6)-(Gly-Ser)-2(10)Asp-140(148)Ala

Length 148 a.a. Molecular weight 15,427 Da

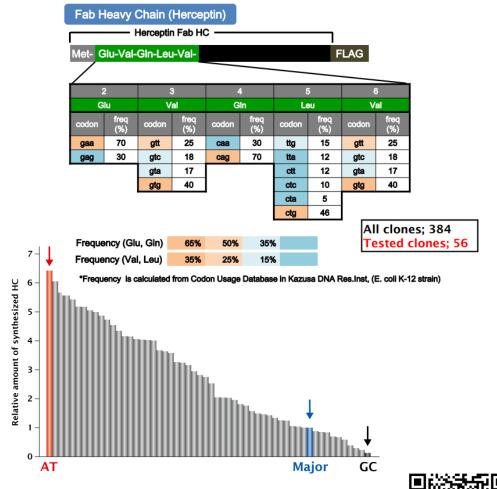
| Tag | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | GC(%) |
|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|----------|
| type | Met | His | His | His | His | His | His | Gly | Ser | 1-9 a.a. |
| GC | atg | CBC | cac | cac | CBC | cac | CAC | ggt | tct | 59% |
| AT | atg | cat | cat | cat | cat | cat | cat | ggt | tct | 37% |





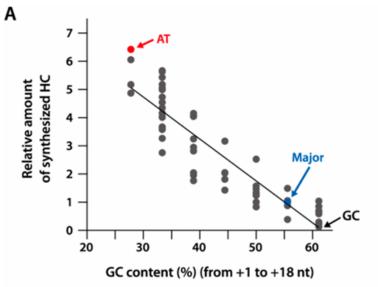
-KSF; AT rich codon on N-term-

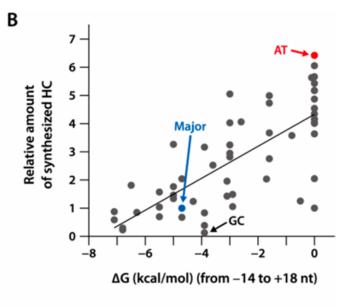




Design of DNA template is important.

Manual is Free to download from our Web site here.





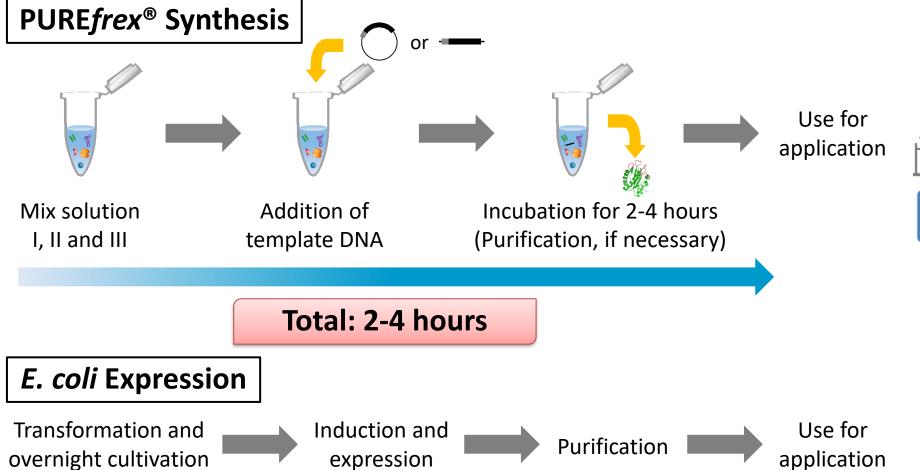
| | 1 | 2 | 3 | 4 | 5 | 6 | | | |
|-------|-----|-----|-----|-----|-----|-----|--------|---------------|--|
| Name | Met | Glu | Val | Gln | Leu | Val | GC (%) | ΔG (kcal/mol) | |
| AT | atg | gaa | gta | caa | tta | gtt | 28 | 0.0 | |
| Major | atg | gaa | gtg | cag | ctg | gtg | 56 | -4.7 | |
| GC | atg | gag | gtg | cag | ctg | gtc | 61 | -3.9 | |

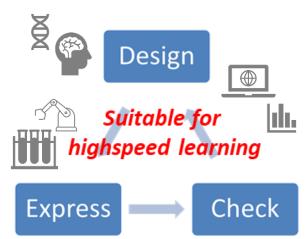
Murakami et al. (2024) Int. J. Mol. Sci. 2024, 25(10), 5264

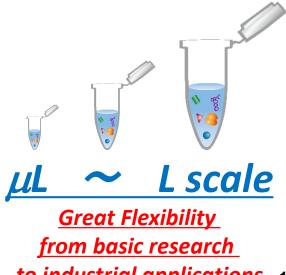


-Improve Expression from Days to Hours-









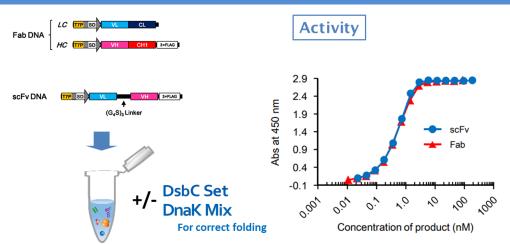
to industrial applications





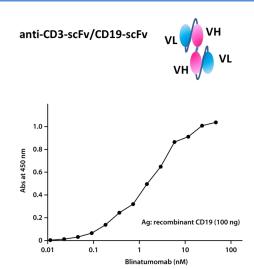
PUREfrex®

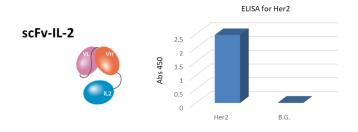
-Expression of scFv, Fab and more-

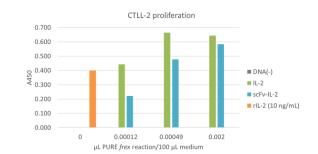


Murakami et al. (2019) Sci. Rep. vol.9, p.671. (Supplementary Information)

| | 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 | X | Х | X | X | ✓ | Х | ✓ | ✓ | ✓ | ✓ |
| BL21 | X | Х | X | Х | ✓ | X | ✓ | ✓ | ✓ | X |
| 759 | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | Х |
| | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 |
| | Glucagon Like Peptide 1 mutant (GLP-1 mut) | Peptide 1 | Insulin Like Growth Factor | Growth Hormone (GH) | Leptin | Vaso- pressin | Angiotensin II | Parathyroid Hormone (PTH) | Somato- statin | Leuprolide |
| PURE | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ |
| Clm24 | ✓ | ✓ | Х | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ |
| BL21 | X | X | X | X | X | Х | ✓ | ✓ | ✓ | ✓ |
| 759 | ✓ | ✓ | х | ✓ | ✓ | ✓ | ✓ | х | ✓ | ✓ |







Internalization analysis

Anti-HER2-IgG

BT-474 cells were surface-labeled at 4°C for

60 min with the binding medium (DMEM, 3% BSA, 20 mM HEPES (pH 7.4)) containing

10 nM of "purified IgG" or "Trastuzumab".

Cells were washed five times with the

binding medium and incubated at 37°C for

3h. Cells were then fixed and processed for

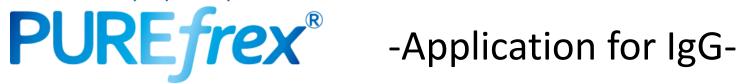
dual-label indirect immunofluorescence

microscopy. CD63 (Lamp3) is marker of late

endosome

endosome

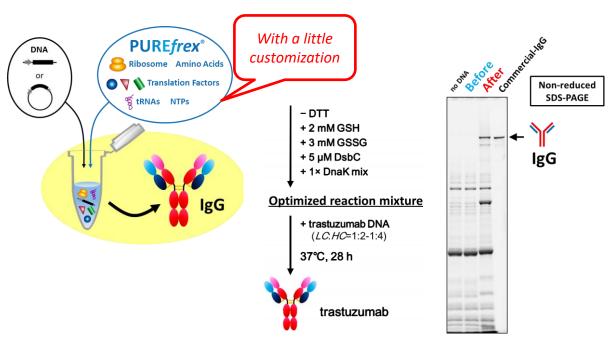
CD63



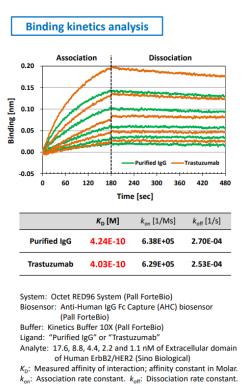


Trastuzumab

Purified IgG



Only 2 days for IgG!!



endosomes and lysosomes. White arrows indicate partial co-localization of anti-HER2-IgG with CD63. Bar indicates 20 µm. Murakami et al. (2019) Sci. Rep. vol.9, p.671.

✓ Full size IgG can be synthesized.

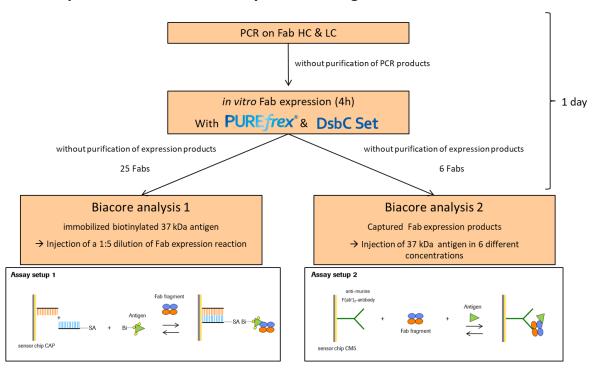
Poster PSSJ 2017



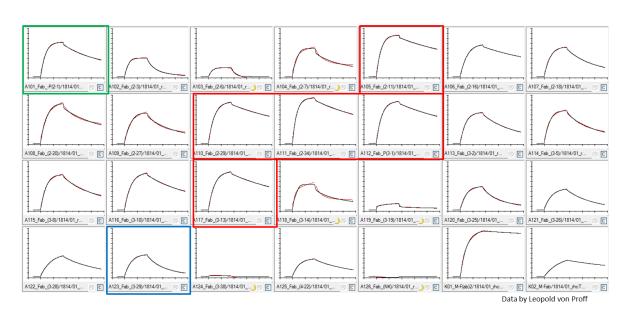




In vitro expression and Biacore analysis of Fab fragments



Kinetic analysis of 25 Fab binders



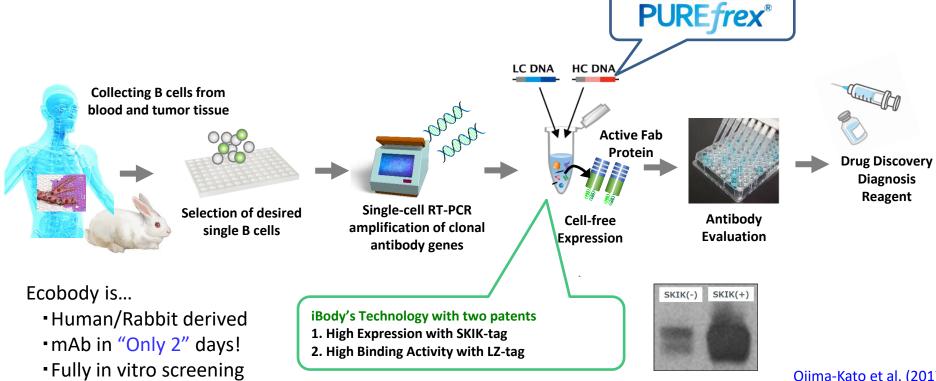
→ Selection of Fabs for further kinetic analysis











Ojima-Kato et al. (2017) Sci. Rep., 7, 13979. https://www.ibody.co.jp/en/

No culture



✓ Active Fab is expressed/screened in HT manner.

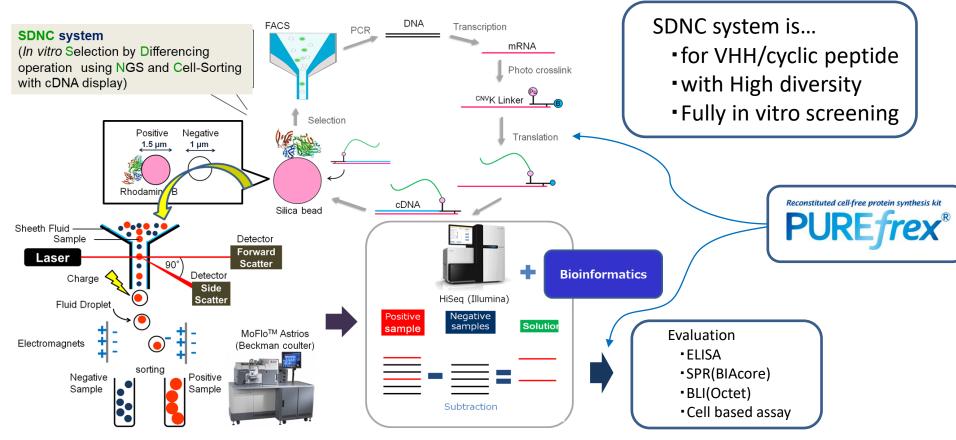




Epsilon Molecular Engineering

Molecular Design for Human Life





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



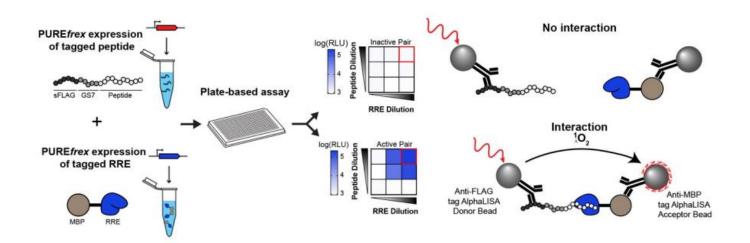
✓ PURE frex is applied for cDNA display based screening.





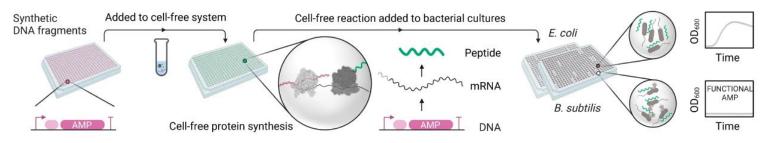


-Broad applications, yet to come!-

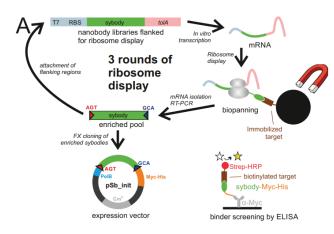


Wong et al. (2024) bioRxiv https://doi.org/10.1101/2024.03.25.586624.

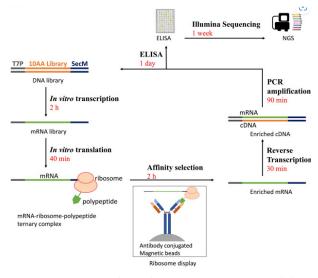
WET LAB EXPERIMENT: cell-free production and activity test of AMPs (24 hr)



Pandi et al. (2023) Nat Communications. vol.14(7197).



Zimmermann I. et al. (2018) eLife, 7, e34317.

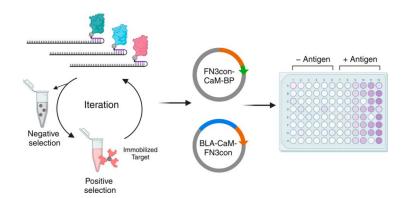


Jia B. et al. (2024) J Biosci Bioeng, 137(4):321-328.

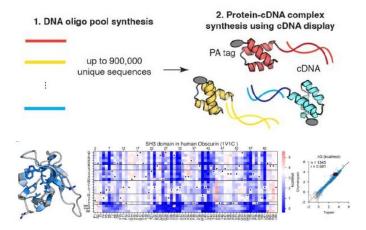


-Broad applications, yet to come!-





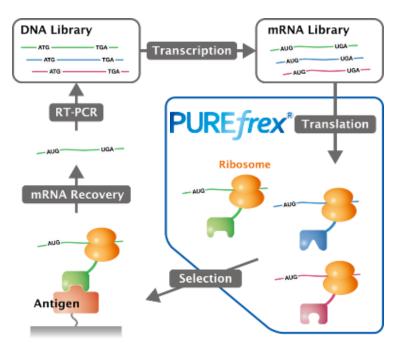
Chui Z. et al. (2024) ACS Sens, 9(6):2846-2857.



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

in vitro protein selection technology



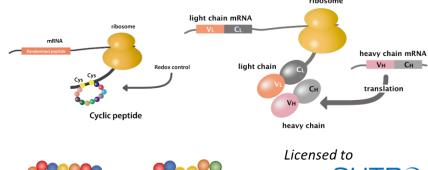


Licensed technology under JP4931135 etc.

- Advanced screening system for Biologics
- mAb (scFv / Fab)
- VHH
- Cyclic peptide

High Selection Efficiency

- Completely molecular based system
- >10¹² diversity







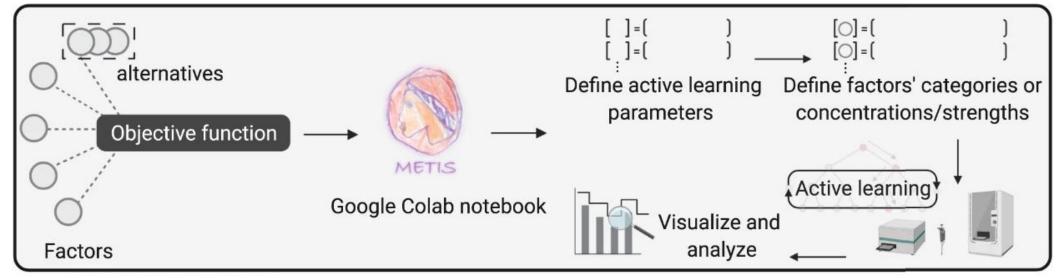
Bicyclic peptide



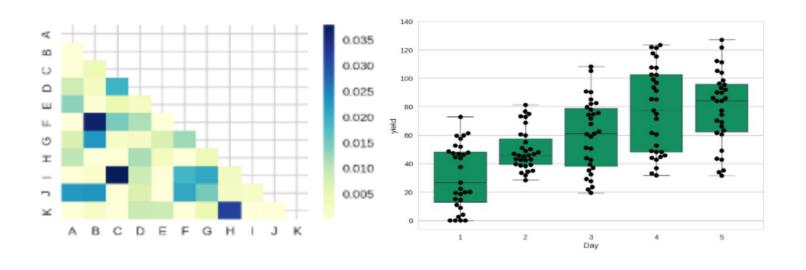


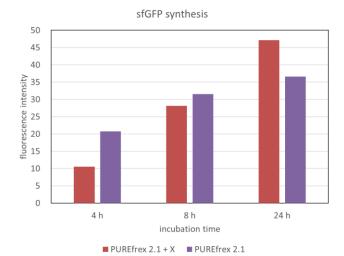
-Broad applications, yet to come!-





Pandi A et al. (2022) Nature Communications, 13, 3876.





Contact information





For reagent use for expression / screening of biologics

https://purefrex.genefrontier.com/





PURE frex RD For screening service / collaboration / technology transfer for generation of new biologics

Takashi Ebihara, Ph.D., COO, GeneFrontier

E-mail: ebihara@genefrontier.com