



U.S. ARMY COMBAT CAPABILITIES DEVELOPMENT COMMAND

Cell Free Transcription and Translation – Faster, Safer

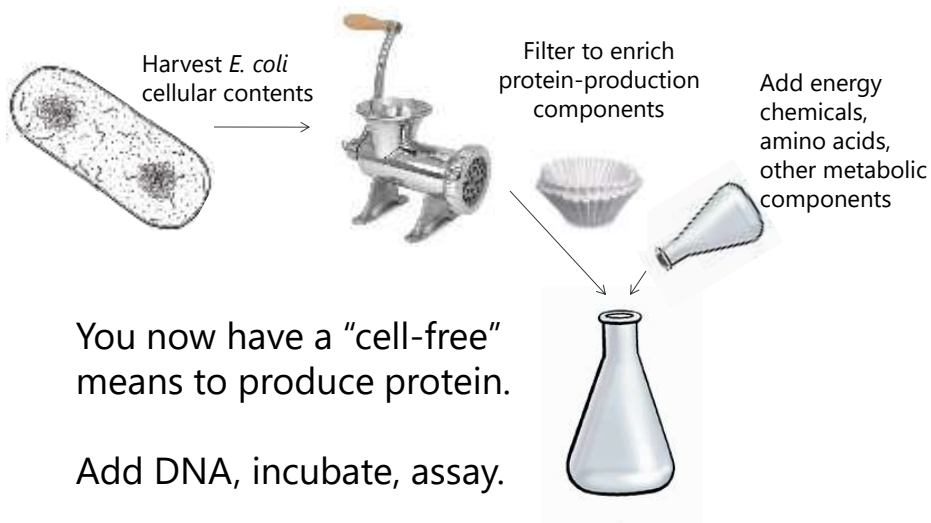
Aleksandr Miklos, Ph.D.
Research Biologist
R&T BioSciences / BioChemistry Branch

20 April 2022

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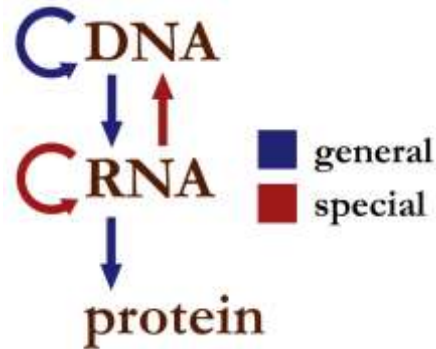
In Vitro Transcription and Translation





Cellular Lysates – Early Work

Establishing the Central Dogma of Molecular Biology



3



Cracking the Code



1961 BIOCHEMISTRY: MATTHAEI AND NIRENBERG Page: N. A. B.
CHARACTERISTICS AND STABILIZATION OF DNAASE-SENSITIVE
PROTEIN SYNTHESIS IN E. COLI EXTRACTS
By J. HUBERT MATTHAEI* and MARSHALL W. NIRENBERG
NATIONAL INSTITUTE OF HEALTH, BETHESDA, MARYLAND
Communicated by Joseph E. Smolnik, August 8, 1961

J.H. Matthaei and M.W. Nirenberg
Proc Natl Acad Sci Unit States Am, 47 (1961),
pp. 1508-1588, 10.1073/pnas.47.10.1580

1960 BIOCHEMISTRY: NIRENBERG AND MATTHAEI Page: N. A. B.
THE DEPENDENCE OF CELL-FREE PROTEIN SYNTHESIS IN E. COLI
UPON NATURALLY OCCURRING OR SYNTHETIC
POLYRIBONUCLEOTIDES
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M.W. Nirenberg and J.H. Matthaei
Proc Natl Acad Sci Unit States Am, 47 (1961),
pp. 1588-1602, 10.1073/pnas.47.10.1588

The RNA code: Nature's Rosetta Stone
"Genetics of Antibiotic and Cancer Genes" Roger College of Medicine, Boston, MA 02115
Communicated by Joseph E. Smolnik, August 8, 1961

The Nobel Prize in Physiology or Medicine 1968
The Nobel Prize in Physiology or Medicine 1968 was awarded jointly to Robert W. Holley, Har Gobind Khorana and Marshall W. Nirenberg "for their interpretation of the genetic code and its function in protein synthesis."

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Shining light in the black box; increasing productivity



Proc. Natl. Acad. Sci. USA
Vol. 71, No. 5, pp. 1803-1807, May 1974

Prolonged Transcription in a Cell-Free System Involving Nuclei and Cytoplasm

(transcription nuclei/RNA synthesis/protein synthesis)

GUANG-JER WU AND GEOFFREY ZUBAY

Department of Biological Sciences, Columbia University, New York City, N.Y. 10027

Communicated by Donald D. Brown, February 15, 1974

The Journal of Biological Chemistry
© 2000 by The American Society for Biochemistry and Molecular Biology, Inc.

Vol. 264, No. 11, Issue of April 15, pp. 6239-6244, 1999
Printed in U.S.A.

Possible Involvement of the 90-kDa Heat Shock Protein in the Regulation of Protein Synthesis*

(Received for publication, May 3, 1990)

David W. Rose[‡], William J. Welch[§], Gisela Kramer, and Boyd Hardisty[†]

From the Clayton Foundation Biochemical Institute and the Department of Chemistry, University of Texas, Austin, Texas 78712 and the Cold Spring Harbor Laboratory, Cold Spring Harbor, New York 11724

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PRODUCTIVITY BREAKTHROUGH



A Continuous Cell-Free Translation System Capable of Producing Polypeptides in High Yield

ALEXANDER S. SPIRIN, VLADIMIR I. BARANOV, LUBOV' A. RYABOVA, SERGEY YU. OVODOV, YULY B. ALAKHOV

A cell-free translation system has been constructed that uses a continuous flow of the feeding buffer [including amino acids, adenosine triphosphate (ATP), and guanosine triphosphate (GTP)] through the reaction mixture and a continuous removal of a polypeptide product. Both prokaryotic (*Escherichia coli*) and eukaryotic (wheat embryo, *Triticum sp.*) versions of the system have been tested. In both cases the system has proven active for long times, synthesizing polypeptides at a high constant rate for tens of hours. With the use of MS2 phage RNA or bromo mosaic virus RNA 4 as templates, 100 copies of viral coat proteins per RNA were synthesized for 20 hours in the prokaryotic or eukaryotic system, respectively. With synthetic calcitonin messenger RNA, 150 to 300 copies of calcitonin polypeptide were produced per messenger RNA in both types of continuous translation systems for 40 hours.

EXPRESSION OF ALIEN GENES IN LIVING cells is often subject to a number of limitations. The product polypeptide can be unstable in a given cell, and in some cases the product is toxic to the cell.

ed if translation were possible in cell-free systems. Unfortunately, however, different versions of cell-free translation systems that have been previously described have a general shortcoming, namely, a low yield of the

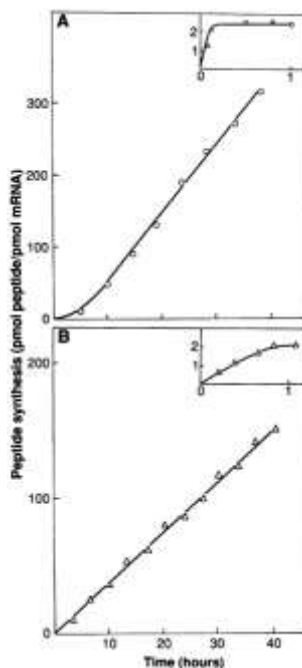


Fig. 2. Kinetics of the ValP-calcitonin synthesis in the continuous cell-free translation systems (40 hours). Inset (same axes units). Kinetics of the polypeptide synthesis in the standard cell-free system of the same composition and volume. (A) Synthesis of calcitonin in the *E. coli* system. All reaction conditions were the same as in Fig. 1A, except that 0.06 nmol of synthetic calcitonin mRNA was added (instead of viral RNA), and the PM-10 membrane was used instead of the PM-30 for removing the product from the translation cell. (B) Synthesis of calcitonin in the wheat system. All reaction conditions were the same as in Fig. 1B, except that 0.06 nmol of synthetic calcitonin mRNA was added (instead of viral RNA), 75 mM potassium acetate was found to be optimal for this synthesis, and the PM-10 membrane was used instead of the XM 50 for removing the product from the translation cell.

Spirin, Alexander S., et al. "A continuous cell-free translation system capable of producing polypeptides in high yield." Science 242.4882 (1988): 1162-1164.

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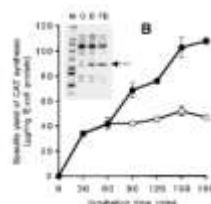
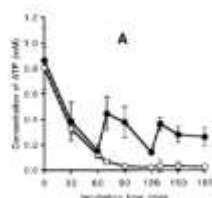


Regeneration of Adenosine Triphosphate from Glycolytic Intermediates for Cell-Free Protein Synthesis

Dong-Myung Kim, James R. Swartz

Department of Chemical Engineering, Stanford University, Stanford, California 94305-5025, USA; telephone: 650-723-5388; fax: 650-729-0568; e-mail: seawater@chemengr.stanford.edu

Received 14 September 2000; accepted 25 January 2001



Kim, Dong-Myung, and James R. Swartz. "Regeneration of adenosine triphosphate from glycolytic intermediates for cell-free protein synthesis." *Biotechnology and bioengineering* 74.4 (2001): 309-316.

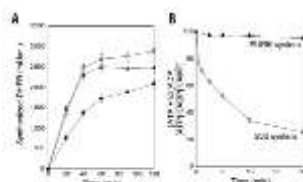


Figure 2. Probability and energy transduction of the PPR system. (A) Time course of DMT synthesis. The value of reactions using ribosome concentrations of 1.0 μM (\bullet), 2.0 μM (\blacktriangle), or 3.0 μM (\blacklozenge) reaction of ribosomes contained in a ribosome cage is plotted against time. The reaction rates decrease as PPR 2, PPR 3, and PPR 4 are released from the cage. (B) and (C) show the energy transduction in the PPR and D20 systems. The decrease in the energy charge of each autarkic unit is monitored.

Cell-free translation reconstituted with purified components

Yoshihiro Shimizu¹, Akio Inoue², Yukihito Terada¹, Tazutomo Suzuki², Takashi Yokoyama¹, Kazuo Nishikawa³, and Takuya Ueda^{2*}

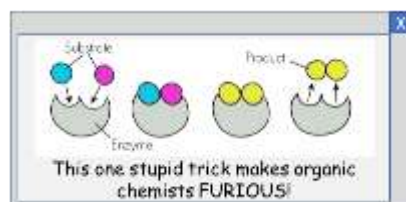
We have developed a protein-synthesizing system reconstituted from recombinant tagged protein factors part of a homogeneity. The system was able to produce proteins at a rate of about 700 $\mu\text{g}/\text{g}/\text{h}$ in a batch mode without the need for any supplementary apparatus. The protein products were easily purified within 1 h using affinity chromatography to remove the tagged protein factors. Moreover, analysis of a release factor allowed efficient incorporation of an ornithine aminic acid using suppressor transfer RNA (tRNA).

Shimizu, Y., Inoue, A., Tomari, Y. et al. Cell-free translation reconstituted with purified components. *Nat Biotechnol* 19, 751–755 (2001).
<https://doi.org/10.1038/90802>

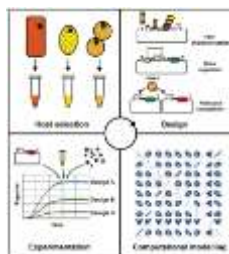
<https://doi.org/10.1038/90802>



Metabolic Engineering

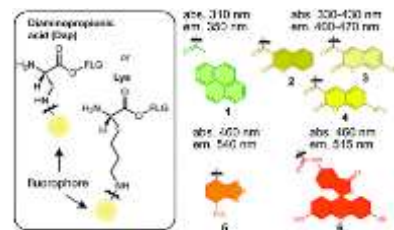


Continued Basic Research (Engineering Translation!)



"Breadboarding" / Screening

Fielding

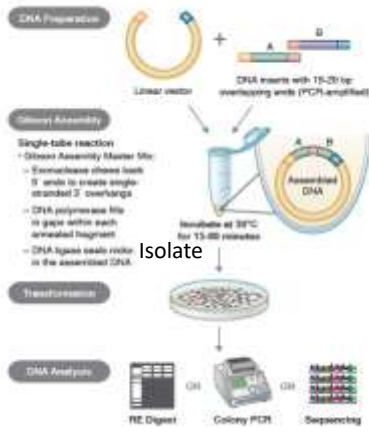


Lee, Joongoo, et al. "Ribosome-mediated incorporation of fluorescent amino acids into peptides in vitro." *Chemical Communications* 57.21 (2021): 2661-2664.

Moore, Simon J et al. "Cell-free synthetic biology for in vitro prototype engineering." *Biochemical Society transactions* vol. 45,3 (2017): 785-791. doi:10.1042/BST20170011



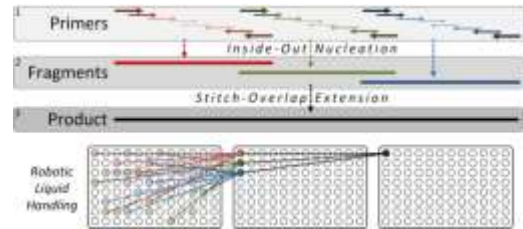
Testing an Engineered Protein



Overnight culture, grow to density, induce, lyse, purify, label, test.



Stitch together a linear ORF from primers on a robot,



Express *in vitro*

Throw that PCR product into a tube with lysate and reaction buffer, purify, label, test.

Miklos, A. E., R. A. Hughes, and A. D. Ellington. "Design and assembly of large synthetic DNA constructs." *Current Protocols in Molecular Biology* (2012): Unit3-23.

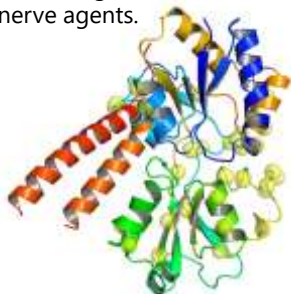
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Screening for Biosensors

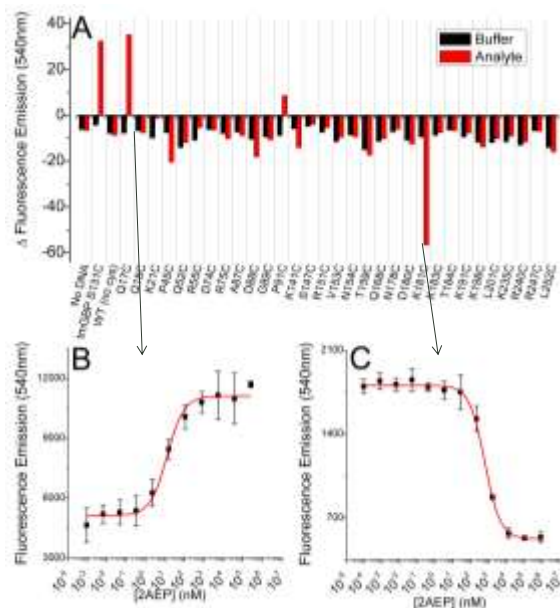


Phosphonate-binding protein
Starting point for sensing
pesticides and nerve agents.



This type of screening can take months with conventional methods, days with *in vitro* expression

Alicea, I., et al. (2011). Structure of the Escherichia coli phosphonate binding protein PhnD and rationally optimized phosphonate biosensors. *Journal of molecular biology*, 414(3), 356-369.



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Using Syn Bio in the Field - The Question:



If I had an engineered organism which could turn latrine contents into freshly-crimped 5.56 rounds, would someone let me USE it?



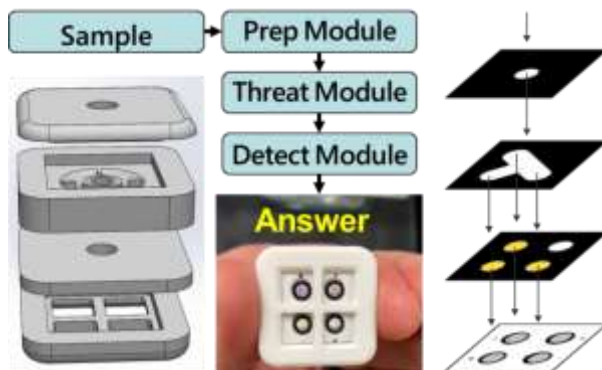
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Cell-Free Syn Bio Sensing



Lysates and energy mixes can be stable when freeze-dried, and you can even embed these reagents in paper before freeze-drying, so we can make shelf-stable devices that run various sensing schemes.



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ACKNOWLEDGEMENTS



Lysate Team

Dr. Nathan McDonald
Dr. Patricia Buckley
Katherine Rhea
Dr. Stephanie Cole



Funding:

Defense Threat Reduction Agency

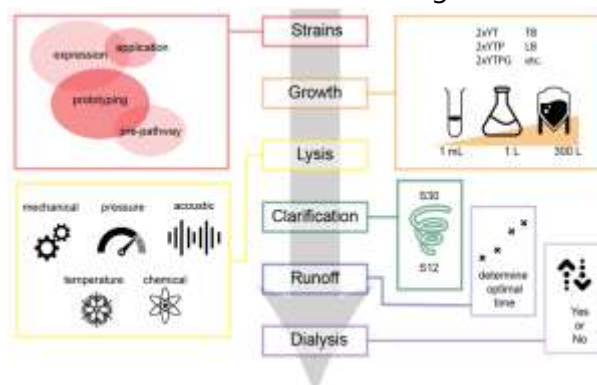
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In-house experience making cell-free systems



- CBC established protocol for generating *E.coli* cell-free lysate
- CBC experience in creating novel cell-free lysate for *V. natriegens*
- Known key considerations:
 - Tuning mRNA and protein degradation rates
 - DNA stabilization
 - Regeneration of energy (ATP)



Cole et al. 2020

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Non-traditional cell free lysate systems



Organism	Type	Proteins Made
<i>Bacillus megaterium</i>	Gram positive	GFP, mCherry
<i>Bacillus subtilis</i>	Gram positive	GFPmut3b, renilla luciferase
<i>Clostridium autoethanogenum</i>	Gram positive	Luciferase, metabolic enzymes
<i>Corynebacterium glutamicum</i>	Gram positive	eGFP
<i>Escherichia fergusonii</i>	Gram negative	eGFP
<i>Klebsiella oxytoca</i>	Gram negative	eGFP
<i>Lactococcus lactis</i>	Gram positive	None (only transcription active)
<i>Pantoea agglomerans</i>	Gram negative	eGFP
<i>Pseudomonas fluorescens</i>	Gram negative	GFP, apolipoprotein, pancreatic RNase, p37a, glucokinase, peptidases
<i>Pseudomonas putida</i>	Gram negative	sfGFP
<i>Salmonella enterica</i>	Gram negative	eGFP
<i>Streptomyces</i> species	Gram positive	eGFP, sfGFP, metabolic proteins
<i>Sulfolobus solfataricus</i>	Archaeal thermophile	ORF 104, ORF 143
<i>Sulfolobus tokodaii</i>	Archaeal thermophile	Polyphenylalanine
<i>Thermococcus kodakaraensis</i>	Archaeal thermophile	Chitinase
<i>Thermus thermophilus</i>	Gram negative	Polyphenylalanine
<i>Vibrio natriegens</i>	Gram negative	sfGFP, eGFP

Cole et al. 2020

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New Reason to go cell-free – Risk Reduction



Yersinia pestis

Gram-negative enterobacteria,
Causative agent of the plague

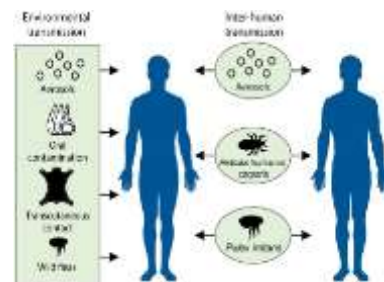
- Route of transmission proceeds typically through flea bites

Encodes a variety of virulence factors including multiple virulence plasmids

- pCD1/pMT1/pPCP1
- Pigmentation locus (pgm)
- F1 proteinaceous capsule
- Type III secretion system
- Lipopolysaccharide

Experiences dramatic environmental shifts during lifecycle
from flea hindgut through human infection

- Temperature transitions range from 21°C-37 °C



Barbieri et al. 2020

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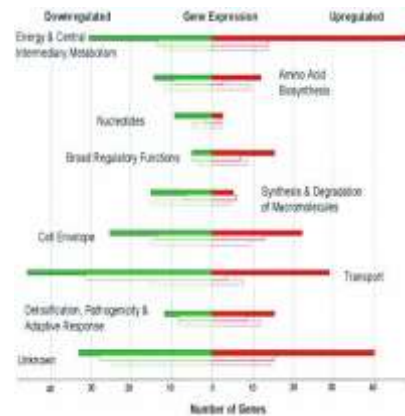
BROAD EXPRESSION CHANGES ACROSS TEMPERATURES



Y. pestis transitions from ~21°C to 26 °C to 37 °C during the infection of a human host from a carrier flea.

The shift in temperature results in dynamic gene expression changes associated with pathogenesis

- Stimulates expression of F1 capsule antigen which is essential for evading phagocytosis
- Upregulation of T3SS effector proteins
- Modifications to the lipopolysaccharide to a low immunostimulatory form

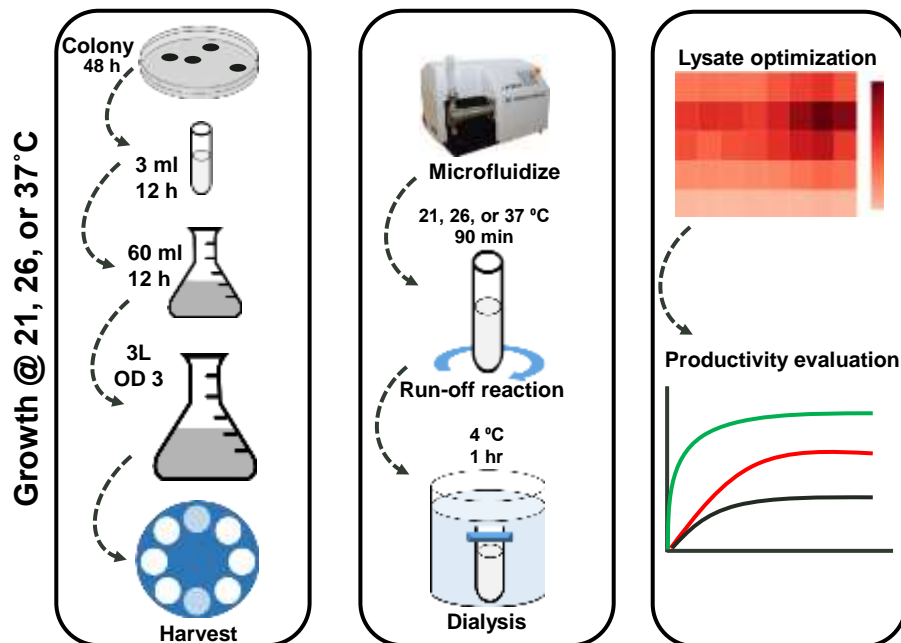


Motin et al. 2014

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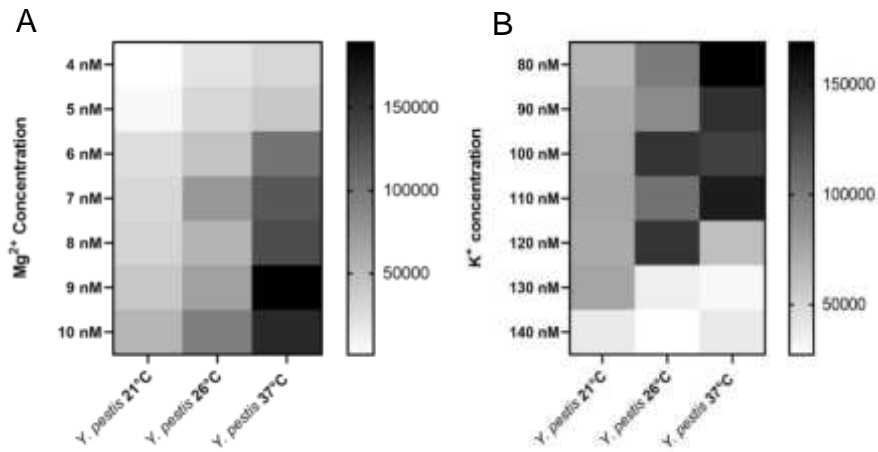
Generating a *Yersinia pestis* CO92- extract



18



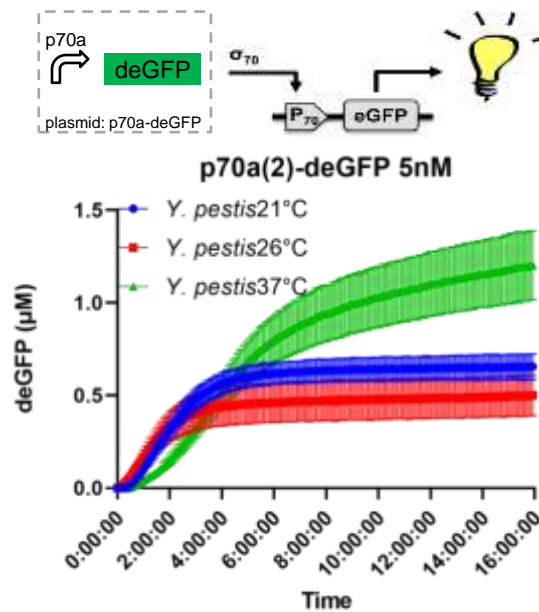
OPTIMIZATION OF EXTRACTS



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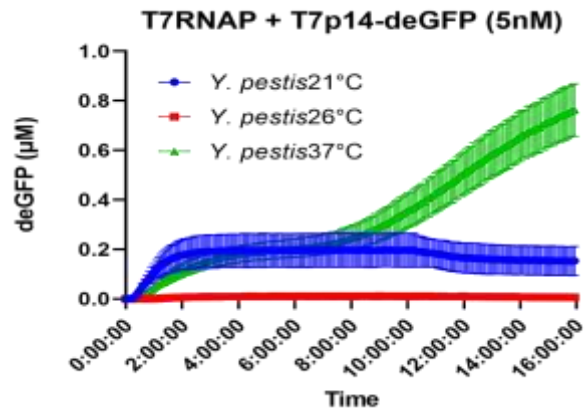
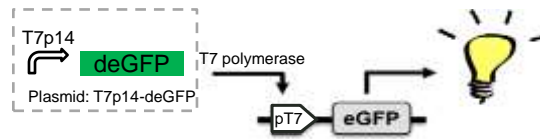
Y. Pestis CO92: 37°C lysate is most productive



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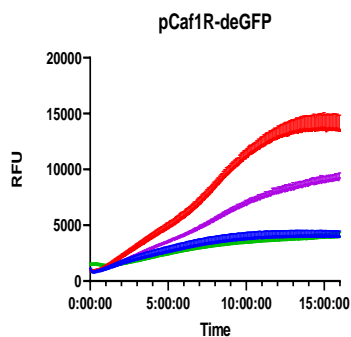
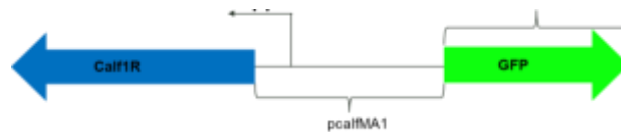
T7 promoters less active in *Y. pestis* CO92- lysates



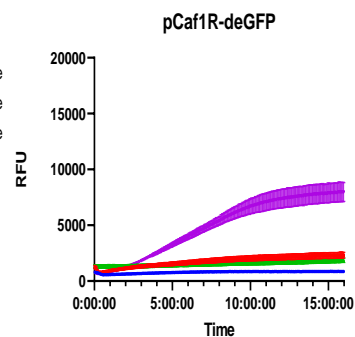
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Yp Lysates work with Native Promoters



Cell-free reaction run at 29°C



Cell-free reaction run at 37°C

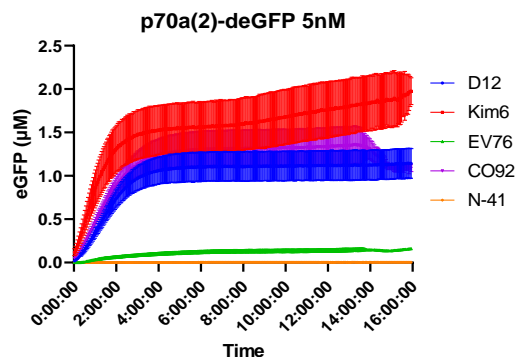
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Extract Productivity is Strain Specific



Strain	Plasmids	Attenuation
D-12	pCD1/pMT1	pCD1-
Kim6	pCD1/pMT1	Lcr-
EV76	pCD1/pMT1	Pgm-
CO92	pCD1/pMT1	Pgm-
N-41	pPCP/pCD1/pMT1	Pgm-



All extracts grown at 21°C

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<https://doi.org/10.1021/acssynbio.1c00505>

Technical Note

Cell-Free Protein Systems from *Yersinia pestis* are Functional and Growth-Temperature Dependent

Nathan D. McDonald,[†] Katherine A. Rhea,[†] Kimberly L. Berk, Julie L. Zacharko, and Aleksandr E. Miklos^{*}

Cite This: <https://doi.org/10.1021/acssynbio.1c00505>

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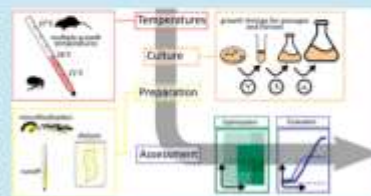
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Supporting Information

ABSTRACT: Cellular lysates capable of transcription and translation have become valuable tools for prototyping genetic circuits, screening engineered functional parts, and producing biological components. Here we report that lysates derived from *Yersinia pestis* CO92[†] are functional and can utilize both the *E. coli* σ^{70} and the bacteriophage T7 promoter systems to produce green fluorescent protein (GFP). Because of the natural lifestyle of *Y. pestis*, lysates were produced from cultures grown at 21 °C, 26 °C, and 37 °C to mimic the infection cycle. Regardless of the promoter system the GFP production from 37 °C was the most productive and the 26 °C lysate was the least. When reactions are initiated with 5 nM of DNA, the GFP output of the 37 °C lysate is comparable with the productivity of other non-*E. coli* systems. The data we present demonstrate that, without genetic modification to enhance productivity, cell-free extracts from *Y. pestis* are functional and dependent on the temperature at which the bacterium was grown.

KEYWORDS: cell free lysates, in vitro transcription and translation, alternative chassis organisms



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CONCLUSIONS AND A QUESTION



- Cell-free production systems can be generated from *Yersinia pestis* extracts
- *Yersinia pestis* extracts are able to express deGFP from native *E. coli* σ_{70} and bacteriophage T7 promoters,
- Growth temperature impacts productivity of the extracts
- *Yersinia pestis* strains have variable production in cell-free expression tests
- How much "live organism" work could be replaced with cell-free?