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A Column: 1. We use *E. coli* MG1655 as model cells. *Cu*gA and *MSP* gene fragments are inserted in plasmid. *Cu*gA gene takes charge for *cell-free* heavy metal synthesis. In *E. coli* membrane, *MSP* can capture and absorb heavy metal ions. (Cu^{2+} , Ag^{+} , ...)

B Column: We insert heavy metal sensor in plasmid. GFP synthesizes via different heavy loss concentration. (Cu^{2+} , Ag^{+} , ...) It is used to evaluate whether the heavy metal concentration achieves government efficient standard.

C Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

D Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

E Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

F Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

G Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

H Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

I Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

J Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

K Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

L Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

M Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

N Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

O Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

P Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

Q Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

R Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

S Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

T Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

U Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

V Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

W Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

X Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

Y Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

Z Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AA Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AB Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AC Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AD Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AE Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AF Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AG Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AH Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AI Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AJ Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AK Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AL Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AM Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AN Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AO Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AP Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AQ Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AR Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AS Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AT Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AU Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AV Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AW Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AX Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AY Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

AZ Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BA Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BB Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BC Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BD Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BE Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BF Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BG Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BH Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BI Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BJ Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BK Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BL Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BM Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BN Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BO Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BP Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*)

BQ Column: Add substrate: *Bla*, *K22290* (Deverageneration *Cm*g24 and *Cys*

Part B: Mathematical modeling for E.coli survival curve analysis

Use model to fit E. coli growth curve under normal condition: [Link](#)

Part D: Improvement of Parts (MBP optimization modeling & Molecular docking)

a) Protein structure prediction

Protein structure prediction using AlphaFold 2. The figure shows the prediction of protein structure for a protein with 1000 amino acids. The color scale represents the predicted local distance difference test (pLDDT) score, ranging from 0.00 (blue) to 1.00 (red). The protein structure is shown in a ribbon representation, with the predicted structure (red) and the ground truth structure (blue) overlaid. The protein is shown in a ribbon representation, with the predicted structure (red) and the ground truth structure (blue) overlaid.

b) Protein structure prediction

Protein structure prediction using AlphaFold 2. The figure shows the prediction of protein structure for a protein with 1000 amino acids. The color scale represents the predicted local distance difference test (pLDDT) score, ranging from 0.00 (blue) to 1.00 (red). The protein structure is shown in a ribbon representation, with the predicted structure (red) and the ground truth structure (blue) overlaid. The protein is shown in a ribbon representation, with the predicted structure (red) and the ground truth structure (blue) overlaid.

c) Protein structure prediction

Protein structure prediction using AlphaFold 2. The figure shows the prediction of protein structure for a protein with 1000 amino acids. The color scale represents the predicted local distance difference test (pLDDT) score, ranging from 0.00 (blue) to 1.00 (red). The protein structure is shown in a ribbon representation, with the predicted structure (red) and the ground truth structure (blue) overlaid. The protein is shown in a ribbon representation, with the predicted structure (red) and the ground truth structure (blue) overlaid.

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