

Supporting Information

Engineering Bacteria to Control Electron Transport Altering the Synthesis of Biopolymer

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Table S1. Bacteria and Plasmid Strains.

Strain/Plasmid	Description	Source/Reference
<i>Escherichia coli</i> wild type	Plasmid Storage Strain (K12 Top 10)	Invitrogen
pMTL_83153	Modular plasmid containing pCB102, catP, ColE1 + tra, P _{fdx} + MCS	http://www.clostron.com/pMTL80000.php

Table S2. Oligonucleotide Primers used for PCR of DNA regions. All Primers used for PCR with Q5 ® Polymerase. *Exceptions used for Colony PCR with Green DreamTaq.

Primers	Sequence (5'-3')	Tm (°C)	Function
NapC_fwd_hifi	GAGCGAAATCATGGGAA ATTCTGACCGTAAG	61.8	To amplify <i>napC</i> in cloning with P_{BAD} promoter
NapC_rev_hifi	ATCTCCATGGACGCGTG ACGTTAAAAACCTGGCTC GAC	59.3	To amplify <i>napC</i> in cloning with P_{BAD} promoter
P _{BAD} _araC-fwd	CAGGAAACAGCTATGAC CGCTTATGACAACTTGAC GGC	59.3	Amplify P_{BAD} promoter
P _{BAD} _araC_rev	AATTTCCCATTTTCTCCT CTTTAATCTAGAGAATTC	58.9	Amplify P_{BAD} promoter
ColE + <i>tra</i> _F2*	CCATCAAGAAGA GCGAC	56.7	Colony PCR
pCB102_R1*	GATAGTCAAAGGCATAA CAG	55.4	Colony PCR

Table S3. Arabinose induction concentrations for *E. coli* containing Inducible Promoter Vector.

Sample	Stock arabinose (w/v)	Final arabinose Concentration
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<i>E. coli</i> (IP_0%)	0	0
<i>E. coli</i> (IP_0.000018%)	0.0018%	0.000018%
<i>E. coli</i> (IP_0.0018%)	0.18%	0.0018%
<i>E. coli</i> (IP_0.18%)	18%	0.18%

Table S4. Protein standardisation calculations using BCA assay. *Calculated using BSA equation ($y=0.0016x$). **Note samples were diluted by 2; therefore, the concentrations were doubled to obtain original solution concentrations.

Vector	Abs _{562nm} (av)	Sample conc ($\mu\text{g/mL}$)*	Original conc ($\mu\text{g/mL}$ **	mg/ml
Empty Plasmid	1.57	982.7	1965	1.97
<i>E. coli</i> (IP_0%)	1.63	1019	2038	2.04
<i>E. coli</i> (IP_0.000018%)	1.69	1053	2106	2.11
<i>E. coli</i> (IP_0.0018%)	1.65	1029	2058	2.06
<i>E. coli</i> (IP_0.18%)	1.40	873	1746	1.75

Table S5. Reagent ratios for *E. coli*_{IP} initiated Fe ATRP.

Reagent	Ratio	Mmol	Mass (mg)	Vol (μL)
PEGMA	100	0.052	17	16.2
FeCl ₃ .6H ₂ O	4.65	0.0024	0.65	-
Me ₆ TREN	13.95	0.0072	1.7	1.9
HEBIB	2	0.0010	0.22	0.15

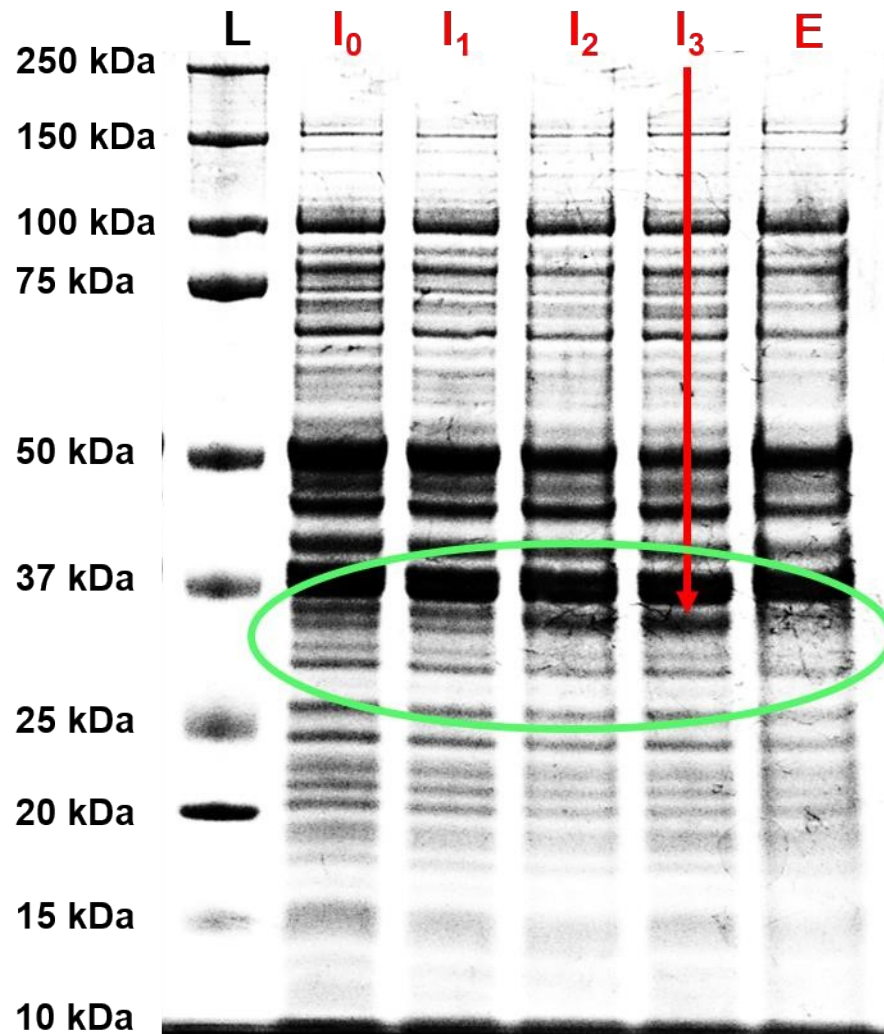


Figure S1. Protein expression analysis. SDS PAGE Gel for lysates of bacteria containing empty plasmid (E), Inducible promoter vector with 0% (I₀), 0.000018% (I₁), 0.0018% (I₂) and 0.18% (I₃) total arabinose concentration induction. Protein Gel against Precision Plus Protein™ Kaleidoscope ladder (L).

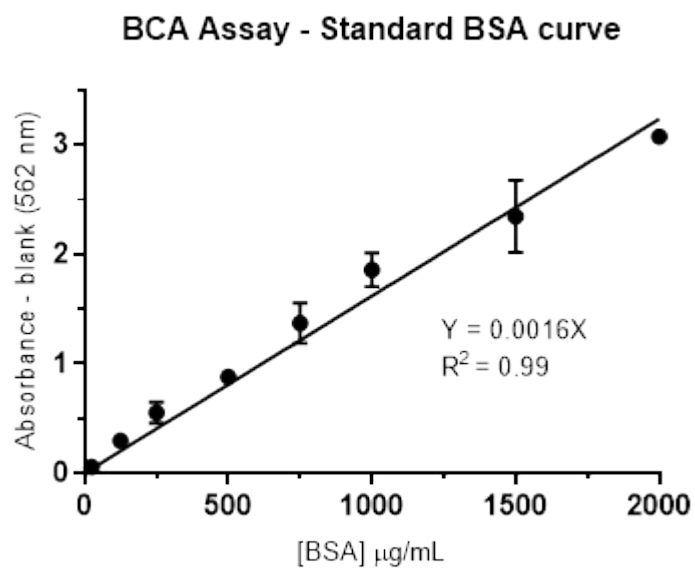


Figure S2. BCA Standard curve for BSA (Bovine Serum albumin).

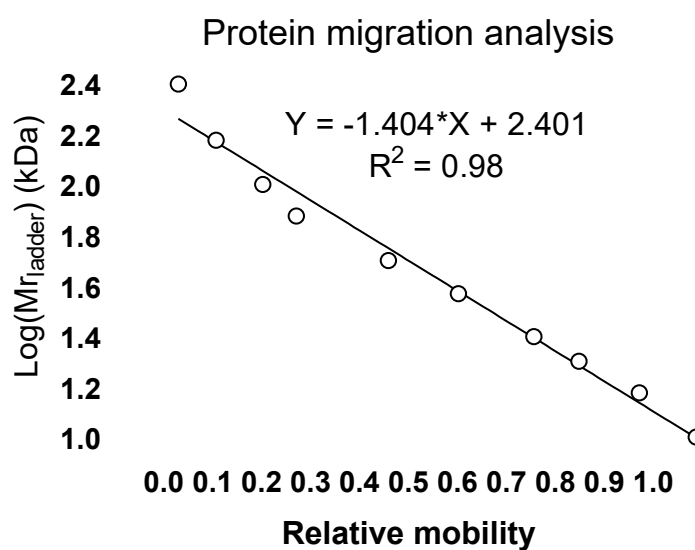


Figure S3. Protein migration analysis of protein ladder Log(M_r) against relative mobility on SDS PAGE Gel.

Calculation:

Migration (NapC) = 10.7 cm Dye

front = 17.2 cm

$$\text{Relative mobility (X)} = \frac{\text{migration}}{\text{dye front}} = \frac{10.7}{17.2} = 0.622$$

$$\text{Log}(Mr_{NapC}) = [-(1.404 * X) + 2.401]$$

$$\text{Log}(Mr_{NapC}) = (-1.404 * 0.622) + 2.401$$

$$Mr_{NapC} = 10^{((-1.404*0.622)+2.401)} = 33.7 \text{ kDa}$$

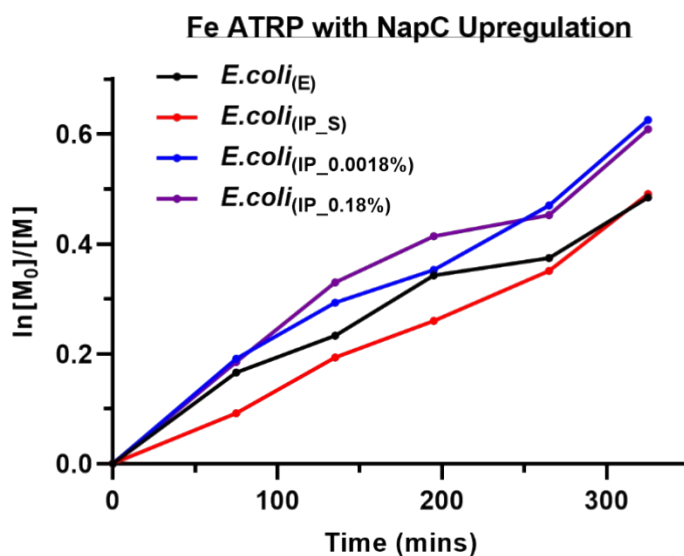


Figure S4. ^1H NMR kinetics of Fe ATRP activated by *E. coli* harbouring empty plasmids, $E. coli_{(E)}$, or inducible promoter plasmids, $E. coli_{(IP)}$ either i) suppressed by addition of glucose $E. coli_{(IP_S)}$, ii) activated by 0.0018% total arabinose concentration $E. coli_{(IP_0.0018\%)}$ or ii) activated by 0.18% total arabinose concentration $E. coli_{(IP_0.18\%)}$.

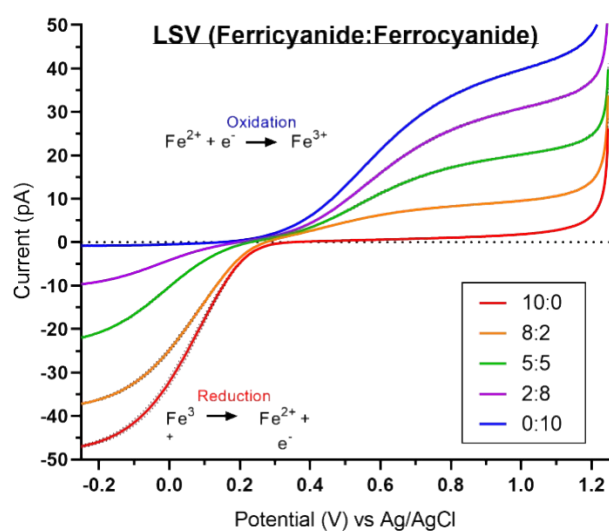


Figure S5. Linear sweep voltammogram of Current Vs Potential carried out using 3 electrode system with carbon fibre micro-disk electrode (33 μm), Ag/AgCl reference electrode and Pt counter electrode in 1X PBS electrolyte. Scans were carried out at 100 mV/s from 1.25 V to -0.25 V. 1 mM potassium ferricyanide and ferrocyanide were made in PBS (1X) and mixed in ratios (10:0, 8:2, 5:5, 2:8, 0:10) and voltammograms observed (n=3, error = SD) for each sample, where electrode was polished between each scan.

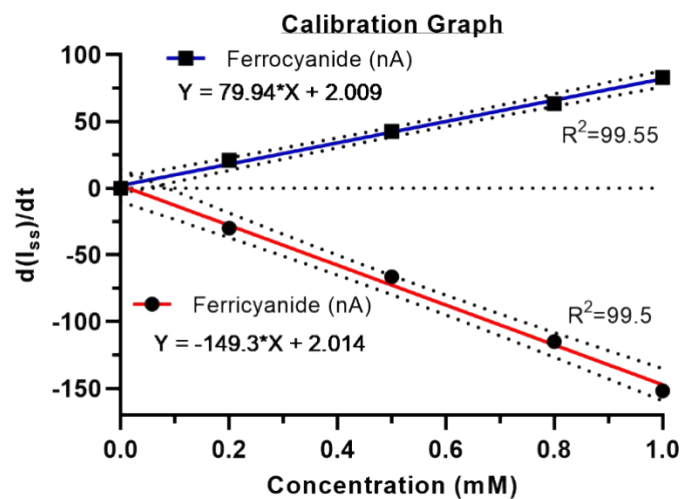


Figure S6. Calibration curve to determine Fe concentration. First derivative peaks from graphs in figures S5 and 3a against Ferricyanide or ferrocyanide concentration with line of best fit and 95% confidence limit.

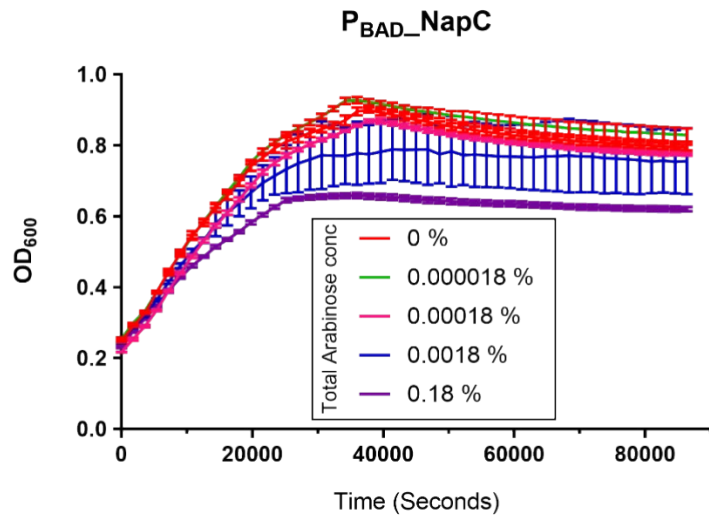
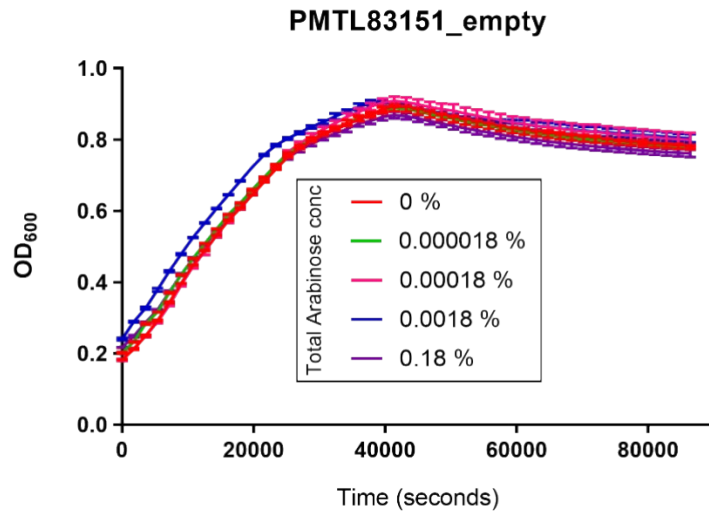


Figure S7. Arabinose toxicity study for, Top: *E. coli* containing empty plasmid and Bottom: *E. coli* containing Inducible vector Plasmid at different arabinose concentrations.