

Supporting Information

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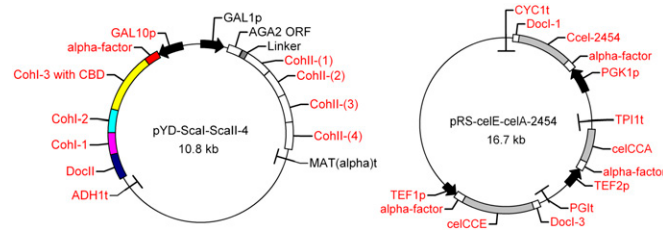


Fig. S1. Recombinant plasmids pYD-Scal-Scall-4 and pRS-celE-celA-2454 used in this study. The structures of pYD-Scal-Scall-1, pYD-Scal-Scall-2, and pYD-Scal-Scall-3 were similar to pYD-Scal-Scall-4. The only difference was that they had one, two, and three repeating CohIIs, respectively. Plasmid pYD-Scall-2 did not have miniscaffoldin I expression cassette, and its miniscaffoldin II had two repeating CohIIs. The structures of plasmids pRS-celE, pRS-celA, pRS-2454, pRS-celE-celA, pRS-celE-2454, and pRS-celA-2454 could be derived from pRS-celE-celA-2454 by removing proper gene expression cassettes. pET28-Scal, pET28-celA, and pET28-2454 were constructed based on pET28a. They were used for expression of miniscaffoldin I, celCCA, and Ccel_2454 in *E. coli*. pETduet-celE was constructed based on pETduet-1 and for *E. coli* expression of celCCE. All of the plasmids pET28-Scal, pET28-celA, pET28-2454, and pETduet-celE used T7 promoter and T7 terminator, and α -factor was not fused with genes. Yeast constitutive promoters (PGK1p, 750 bp; TEF1p, 412 bp; TEF2p, 560 bp) and the corresponding terminators (CYC1t, 249 bp; PGI, 401 bp; TPI1t, 401 bp) were PCR-amplified from the genomic DNA isolated from *S. cerevisiae* EB Y100. Galactose-inducible promoter (GAL10p, 491 bp) and the corresponding terminator (ADH1t, 324 bp) were amplified from pYES2 (Invitrogen). Yeast-secretion signal (α -factor, 270 bp) was amplified from pPICZ α A (Invitrogen). Cellulases [celCCE without native dockerin, 2,403 bp; celCCA with native dockerin (DocI-2), 1,350 bp; Ccel_2454, 2,133 bp] and CohI-2 (438 bp) were cloned from the genomic DNA isolated from *C. cellulolyticum* DSM 5812. DocI-3 (246 bp), CohI-3 with CBD (1,131 bp), DocII (492 bp), and CohII (522 bp) were amplified from the genomic DNA isolated from *C. thermocellum* ATCC 27405. DocI-1 (174 bp) and CohI-1 (414 bp) were amplified from the genomic DNA of *C. cellulovorans* DSM 3052. All of the primers are listed in Table S2–S7.

Table S1. Summary of the recombinant *S. cerevisiae* EB Y100 and *E. coli* BL21 (DE3) used in this study

Strain	Plasmid(s)	Phenotype
EBY (control)	pYD1 and pRS425	No surface display (negative control)
EBY (EA2-1)	pYD-Scal-Scall-1 and pRS-celE-celA-2454	Displays trifunctional minicellulosome with one CohII on scaffoldin II
EBY (EA2-2)	pYD-Scal-Scall-2 and pRS-celE-celA-2454	Displays trifunctional minicellulosome with two CohIIs on scaffoldin II
EBY (EA2-3)	pYD-Scal-Scall-3 and pRS-celE-celA-2454	Displays trifunctional minicellulosome with three CohIIs on scaffoldin II
EBY (EA2-4)	pYD-Scal-Scall-4 and pRS-celE-celA-2454	Displays trifunctional minicellulosome with four CohIIs on scaffoldin II
EBY (E-2)	pYD-Scal-Scall-2 and pRS-celE	Displays unifunctional minicellulosome with CBH activity and with two CohIIs on scaffoldin II
EBY (A-2)	pYD-Scal-Scall-2 and pRS-celA	Displays unifunctional minicellulosome with EG activity and with two CohIIs on scaffoldin II
EBY (2454-2)	pYD-Scal-Scall-2 and pRS-2454	Displays unifunctional minicellulosome with BGL activity and with two CohIIs on scaffoldin II
EBY (EA-2)	pYD-Scal-Scall-2 and pRS-celE-celA	Displays bifunctional minicellulosome with CBH and EG activities and with two CohIIs on scaffoldin II
EBY (E2-2)	pYD-Scal-Scall-2 and pRS-celE-2454	Displays bifunctional minicellulosome with CBH and BGL activities and with two CohIIs on scaffoldin II
EBY (A2-2)	pYD-Scal-Scall-2 and pRS-celA-2454	Displays bifunctional minicellulosome with EG and BGL activities and with two CohIIs on scaffoldin II
EBY (C ₄ doc-2)	pYD-Scal-Scall-2	Displays scaffoldin I and scaffoldin II with two CohIIs
EBY [EA2-2 (-)]	pYD-Scall-2 and pRS-celE-celA-2454	Displays trifunctional minicellulosome with two CohIIs on scaffoldin II and without scaffoldin I
<i>E. coli</i> (C ₄ doc)	pET28-Scal	Expression of miniscaffoldin I in <i>E. coli</i> BL21 (DE3)
<i>E. coli</i> (E)	pETduet-celE	Expression of CBH in <i>E. coli</i> BL21 (DE3)
<i>E. coli</i> (A)	pET28-celA	Expression of EG in <i>E. coli</i> BL21 (DE3)
<i>E. coli</i> (2454)	pET28-2454	Expression of BGL in <i>E. coli</i> BL21 (DE3)

Table S2. Primers used for constructing cellulase gene expression cassettes in pUC19

Name	Sequence
α -factor-F	CGGAATCCGATGAGATTTCTTCAATT
α -factor-R	CCGAGCTCGAGCTTCAGCCTCTCTTTTC
PGK1p-F	CGGAATTCACGCACAGATATTATAACATC
PGK1p-R	CGGAATTCGTGTTTATATTTGTTGTAAGTAG
TEF1p-F	CGGAATTCATAGCTTCAAAATGTTTCTACTC
TEF1p-R	CGGAATTCCTTTGTAATTAAGCTTAGATTAG
TEF2p-F	CGGAATTCGGGCGTACTTACATATAG
TEF2p-R	CGGAATTCGTTTAGTTAATTATAGTTCGTTG
CYC1t-F	ACGCGTCGACATCATGTAATTAGTTATGTCAGCG
CYC1t-R	CCCAAGCTTGCAAATTAAGCCTTCGAG
PGIt-F	ACGCGTCGACAACAAATCGCTCTTAAATATATACC
PGIt-R	CCCAAGCTTGGTATACTGGAGGCTTCATGAGTTATG
TPI1t-F	GCTCTAGAGATTAATAAATATATAAAAAATATTATC
TPI1t-R	ACGCGTCGACCTATATAACAGTTGAAATTTGG
celCCA-pUC19-F	CGGAGCTCCGCATCATCACCATCACCATTATGATGCTTCACTTATTCCGAATC
celCCA-pUC19-R	TGCTCTAGATTAGTTGCTTGGAGCTTACTTACC
celCCE-pUC19-F	TCCCCGGGGCATCATCACCATCACCATATAGGACAAGCATTGCCCCTT
celCCE-pUC19-R	ACGCGTCGACAGTTGGAGGAGTCACTGACCC
Ccel ₂₄₅₄ -pUC19-F	CGGAGCTCCGCATCATCACCATCACCATATGCAATACGATCAGATAGATAA
Ccel ₂₄₅₄ -pUC19-R	ACGCGTCGACAGGCAAGAGCTATAGCTATCGGAA
DocI-1-F	ACGCGTCGACTTAAAGGTGATGTTAACTCTGATGC
DocI-1-R	ACGCGTCGACTTAAGCAAGAAGTCTTTCTTTAA
DocI-3-F	ACGCGTCGACTtcccggatatgacataataaag
DocI-3-R	ACGCGTCGACTTAGTCTTTGTACGGCAATG

F, forward; R, reverse.

Table S3. Primers used for shifting cellulase gene expression cassettes into pRS425

Name	Sequence
M3-NotI-F	ATAAGAATGCGCCCGCTAAAACGACGGCCAGT
M3-NotI-R	ATAAGAATGCGCCCGCAGATCTCAGGAAACAGCTATGAC
M3-BglII-F	GAAGATCTGTAACACGACGGCCAGT
M3-BglII-R	GAAGATCTGCATGCCAGGAAACAGCTATGAC
M3-SphI-F	ACATGCATGCGTAAAACGACGGCCAGT
M3-SphI-R	ACATGCATGCCAGGAAACAGCTATGAC

Table S4. Primers for constructing miniscaffoldin I expression cassette in pUC19

Name	Sequence
GAL10p-F	CGGAATTCCTGATTAATTACCCAGAAATAAGGC
GAL10p-R	CGGAATTCCTATATTGAATTTTCAAAAATCTTACTT
ADH1t-F	ACGCGTCGACTGGACTTCTTCGCCAGAGGTTTGG
ADH1t-R	CCCAAGCTTGCATGCCGGTAGAGGTGTGTC
α -factor-F	See Table S1
α -factor-R	See Table S1
Cohl-1-F	GCTCTAGAACCAGTTGAAGCTGTAACAGCTA
Cohl-1-R	GCTCTAGATGATAGTTACTGTTCTGGGTTAACTGC
Cohl-2-F	CGGGATCCGGCGATTCTCTTAAAGTT
Cohl-2-R	CGGGATCCTTGAGTACCAGGATCTAT
Cohl-3-CBD-F	CGAGCTCGGTGTGGTAGTAGAAATTTGGCAAAG
Cohl-3-CBD-R	CGGGATCCCGGATCATCTGACGGCGGTATT
DocII-F	AACTGCAGaataaacctgtaatagaag
DocII-R	CCCAAGCTTTTACTGTGCGTTCGTAATCAC

F, forward; R, reverse.

