Result:

- 1. Proved that empty vector transformed E. *coli* osmotic shock supernatant is not able to degrade xylan at 65°C
- 2. Proved that osmotic shock supernatant of E.coli transformed by either pET22b(+)_x2 or pET22b(+)_ArfB can degrade xylan from beech wood.
- 3. Proved that osmotic shock supernatant of E.coli transformed by pET22b(+)_ManA1 can degrade mannan from *S. cerevisiae*.
- 4. Proved that XynB is active at a wide range of temperature up to 94.7°C

Procedure:

1. Prepare DNS reagent

DNS reagent recipe is from a published paper (Effects of Different DNS Reagents in Determination of Xylose Content, Wang et al.), and is shown as below.

Weight (g/L)	Component
10.00	3,5-Dinitrosalicylic acid
16.00	NaOH
5.00	Phenol
5.00	Sodium sulfite
300.00	Potassium sodium tartrate

Store at room temperature in brown bottle for a week before use.

- 2. To testify enzyme activity, we employed DNS method to measure reductive saccharide released from substrate, and indicate enzyme activity. We found that spectra of DNS reagent is somehow miscellaneous, but optic density improves at 540nm when heated with reductive saccharide.
- 3. We testified that our substrate, xylan from beech wood (product of MERYER), does not contain reductive saccharide (Figure 1A), and the osmotic shock supernatants do not contain enough reductive saccharide for DNS reagent to detect, compared to untreated samples, osmotic shock supernatant treated samples do not show prominent increase at OD 540 (Figure 1B, C).

Culture BL21(DE3) transformed with empty pET22b+ vector and apply osmotic shock procedure, test enzyme activity for both supernatant and precipitate(Figure 1B, C)

+10 µ l	Osmotic shock supernatant(S) Osmotic shock precipitate(P)				
+10 μ Ι	Xylan(0.5 g/mL)				
For each tube, extract 5 μ I , add 5 μ I DNS, incubate at 99°C for 10min as sample before enzyme					
treatment(SO and PO), the rest 15 µ I Incubate at 65°C for 6h					
+15 µ I DNS reagent					
incubate 99°C for 10min, sample named as S6 and P6					
assay by Nanodrop					

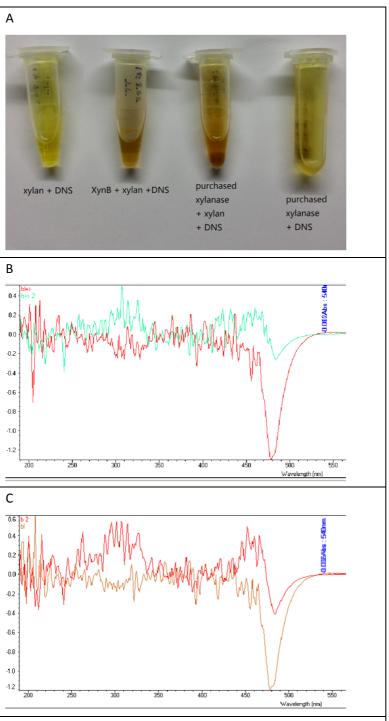


Figure 1, (A) During examination of purchased xylanase, we testified that our substrate (0.05 g/mL) does not contain enough reductive saccharide for DNS method to detect. (B) Green line is the spectra of P6, red line is the spectra of P0. (C) Brown line is the spectra of P0, red line is the spectra of P6.

Test XynB working temperature, find that XynB shows stable performance under a wide range of temperature conditions, even up to 94.7° C

Recipe(μ L):

Temperature(°C)		74.	74.	76.	78.	80.	83.	85.	90.	94.	94.
		5	9	2	1	4	0	8	8	2	7
XynB from C.st	0.1mg/mL xylan	10	10	10	10	10	10	10	10	10	10
(Group A) XynB from C.st		10	10	10	10	10	10	10	10	10	10
Xylase purchased	0.1mg/mL xylan	10	10	10	10	10	10	10	10	10	10
(Group B) Xylase		10	10	10	10	10	10	10	10	10	10
	purchased(over dose)										

Incubate in metal bath for 4.5h

Add $10\mu l$ DNS reagent to each tube, incubate at $99^{\circ}C$ for 10min

Use 99°C incubated reaction system containing following components as blank, measure OD 540 of each sample

 $10\mu L$ 0.1mg/mL xylan, $10\mu L$ H₂O, $10\mu L$ DNS reagent

Nanodrop data, wavelength = 540 nm

Temperature (°C)	OD 540		
	Group A	Group B	
74.5	0.119	0.007	
74.9	0.135	-0.002	
76. 2	0.128	-0.065	
78.1	0.127	- 0 . 106	
80.4	0.156	-0.006	
83.0	0.119	-0.002	
85.8	0.112	-0.096	
90.8	0.145	0.004	
94. 2	0.151	-0.079	
94. 7	0.172	0.01	

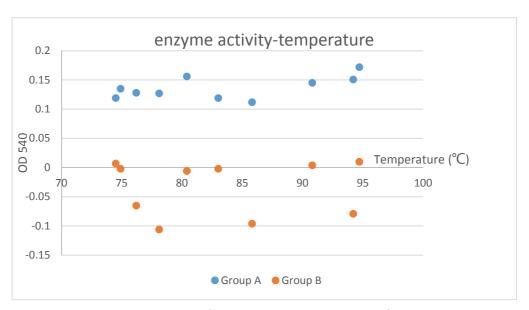


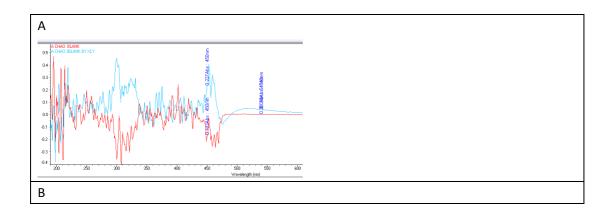
Figure 2, Enzyme shows stable performance under a wide range of temperature conditions

Verify enzyme activity of ArfB and ManA1, compared to untreated samples, OD 540 show increase after protein extraction product treatment.(Figure 3A, B)

ArfB and ManA1 are easy to form inclusion body during expression, so we tested different protein extraction methods, and used ultrasonic sound cell lysis (200W, 1h) for ArfB (product named as A CHAO)and mechanic pressure cell lysis for ManA1 (product named as M LS)to obtain best yield(Data not shown).

Recipe:

+10 μ Ι	A CHAO	H2O	M LS	H2O		
+10 μ Ι	l Xylan		Mannan			
For each	For each tube, extract 5 μ I , add 5 μ I DNS, incubate at 99°C for 10min as sample before enzyme					
treatme	treatment(A CHAO BLANK and M LS BLANK), the rest 15 µ I Incubate at 65°C for 6h					
+15 μ	μ DNS					
L	L					
incubate 99'C for 10min						
assay by	assay by Nanodrop, use $ \textcircled{\scriptsize 1} $ as blank for A CHAO and $ \textcircled{\scriptsize 2} $ as blank for M LS					



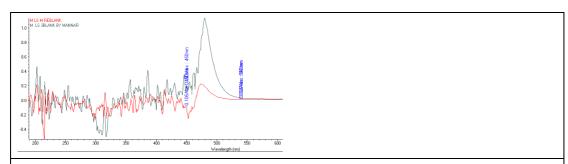


Figure 3, (A) Cyan line is the spectra of A CHAO, red line is the spectra of A CHAO BLANK.(B) Grey line is the spectra of M LS, red line is the spectra of M LS BLANK.