

Supplementary Materials for

**Comparison of Glycoside Hydrolase family 3 β -xylosidases from basidiomycetes and ascomycetes
reveals evolutionarily distinct xylan degradation systems**

Keisuke Kojima¹, Naoki Sunagawa¹, Nils Egil Mikkelsen², Henrik Hansson², Saeid Karkehabadi²,
Masahiro Samejima^{1,3}, Mats Sandgren² and Kiyohiko Igarashi^{1,4*}

Figure legend

Figure S1. Sequence alignment of *TrXyl3A* and *PcBxl3*

The amino acid sequences of *TrXyl3A* and *PcBxl3* were aligned by Clustal W (42). The upper sequence is *TrXyl3A* and the lower is *PcBxl3*.

<i>PcBxl3</i>	1	-----A F PDCANGPLKSNLVCNASADPVSRAKALVDAL
<i>TrXyl3A</i>	1	QNNQTYANYSAQGQPDLYPETLATLTLSEFPDCEHGPLKNNLVCDSSAGYVERAQALISLF
<i>PcBxl3</i>	34	TLEELVNNTVNASPGVPRVGLPPYNNWWSEALHGVARSPGTNFSTVPGSPFSSATSFP POPI
<i>TrXyl3A</i>	61	TLEELILNTONSGPGVPRGLPNYQVWNEALHGLDRA N----FATKGGOFEWATSFPMP <i>I</i>
<i>PcBxl3</i>	94	I LGATFDDDLIHSIA T VI S TEARAFNNAGRA G LDFT T PNINPF K D P RWGR G Q E T P GEDPY
<i>TrXyl3A</i>	117	LTTAALNRTLIHOIADIISTQARAFSNSGRYGLDVYAPNVNGFRSPLWGRGQETPGEDAF
<i>PcBxl3</i>	154	HIA-QYVYQOLITGLQGGLSPDPYYKVVADCKHFAGYDLEDWHGNRMAFNAVISTODLAE
<i>TrXyl3A</i>	177	F LSSAYTYEYITGIQGGVDP-EHL K VAA T V K H F AGYD L ENWN N QSR L GF D A I IT Q QDLSE
<i>PcBxl3</i>	213	F YTPSFOSCSV R DAHVGSVMCSYNAVNGVPSCASPYLLQ D LIR D H F GL G DG-WITSDCDAV
<i>TrXyl3A</i>	236	Y YTPQFLAAARYAKSRSLMCAYNSVNGVPSCANSFLQ T LL R ESWGFP E WGYVSSDCDAV
<i>PcBxl3</i>	272	D NVF D PHNYT S TLVN A SV S LKAGTD V DCGTTYS Q TL V DAVN Q KL V TEDDVKT SM VR L YS
<i>TrXyl3A</i>	296	Y NVFNPHDYASNOSSAAASSL R AGTD I DCGOTY P WHL N ES V AGE V SR E I R SV T RLYA
<i>PcBxl3</i>	332	S LVRLGYFD S PEN Q PW R OL G W A D V NTP S AO A LA T AA E EG V V L L K N D G T L P LS R R I K H I A
<i>TrXyl3A</i>	356	N LVRLGYFD--KK N Q Y RSL G W K D V V K T D AWN I S YE A VE G I V L K N D G T L P LS K K V R S I A
<i>PcBxl3</i>	392	V VGPWANATT Q M Q NY O G IAPFLISPL Q AL Q DA G F H V S FANG T A I INST D T S G F A S A L M A
<i>TrXyl3A</i>	414	L IGPWANATT Q M Q NY Y G PAPYLISPL E AK K AG Y H V N FEL G TE I AGN S T T G FA K A I A AA
<i>PcBxl3</i>	452	K A A D A I V F A G G I D E T I E S E G H R D R D S I E W P G N Q L D L I E Q L A A L R K P L I V L Q M GG Q V D SS SS
<i>TrXyl3A</i>	474	K K S D A I I Y L G G I D N T I E Q E G A D R T D I A W P G N Q L D L I K Q L S E V G K P L V V L Q M GG Q V D SS SS
<i>PcBxl3</i>	512	L K A S K A V N A L I W G G P Q S G G T A I V N I L T G K T A P S G R L P I T Q Y P A Y V D A I P M T D A L R P
<i>TrXyl3A</i>	534	L K S N K K V N S L V W G G P Q S G G V A L F D I L S G K R A P A G R L V T Q Y P A Y V H Q F P Q N D M N L R P
<i>PcBxl3</i>	572	S S - S SPGR T Y K W Y T G T P V F D F G F G L H Y T S F K L S W A A S P P S R F D I S S L V A G A K H A G V A F T D
<i>TrXyl3A</i>	594	D G K S N P G Q T Y I W Y T G K P V Y E F G S G L F Y T T F K E T L A S H P K S L K F N T S S I L S A P H P G T Y S E
<i>PcBxl3</i>	631	L A P L F T H V A V K N S G K V T S D Y V A L F A H T T - V G P S P A P Q Q E L V A Y T R V K G I T P G R T A A
<i>TrXyl3A</i>	654	Q I P V F T E A N I K N S G K T E S P Y T A M L F V R T S N A G P A P Y P N K W L V G F D R L A I K P G H S S K L S
<i>PcBxl3</i>	690	L S V T L G S I A R V D E S G V R S L Y P G K Y S V W V D T T R E I M H T F E L T G K T T Q I L G W P Q R -----
<i>TrXyl3A</i>	714	I I P V S A L R V D S H G N R I V Y P G K Y E L A N T D E S V K L E F E L V G E E V T I E N W P L E E Q Q I K D A
<i>PcBxl3</i>	774	----- T P D A

Figure S1