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Author Correction: Structural Features of a Bacteroidetes-Affiliated Cellulase Linked with a Polysaccharide Utilization Locus

A. E. Naas, A. K. MacKenzie, B. Dalhus, V. G. H. Eijsink & P. B. Pope

Correction to: *Scientific Reports* <https://doi.org/10.1038/srep11666>, published online 02 July 2015

This Article contains an error, where the protein name corresponding to 4IM4 (pdbid) is incorrectly given as “Lic26A-Cel5E”, and should read “CelE”.

In addition, three references were omitted from Table 2, and are given below as References 1–3.

The correct Table 2 and Figure 4 are given below as Table 1 and Figure 1 respectively.

Finally, the updated Supplementary Information file, containing the correct protein name in Figure S1, is given below.

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	Structural comparison			Enzymatic activities							Reference
	DALI Z-score	RMSD (Å)	% id	CMC (U/mg)	β -glucan (U/mg)	Filter paper (U/mg)	Avicel (U/mg)	Lichenan (U/mg)	Xylan (U/mg)	Xyloglucan (U/mg)	
<i>Bacteroidetes AC2a</i> Cellulase Cel5A	—	—	—	216.8	1471.4	0.152 ^a	0.115 ^a	839.9	nd	trace	<i>This study and</i> ⁷
<i>Paenibacillus pabuli</i> Xyloglucanase XG5 (2JEP)	49.2	1.6	34	nd	nd	—	nd	nd	nd	8700	³³
<i>Clostridium cellulovorans</i> Cellulase EngD (3NDY)	47.0	1.7	33	15	42	—	0.017	—	0.5	36	¹⁵
<i>Bacteroides ovatus</i> Xyloglucanase BoGH5A (3ZMR)	46.6	1.9	32	nd	nd	—	—	nd	—	514.2	⁴⁴
<i>Clostridium thermocellum</i> Cellulase CelE (4IM4)	46.0	1.6	32	Active	75.4	Active	—	—	Active	—	¹⁻³ , b
<i>Clostridium cellulolyticum</i> Cellulase CelCCA (1EDG)	45.9	1.9	32	101.3	104.3	—	0.028 (0.124) ^d	79.2	10.1	—	^{45,46} , c
<i>Piromyces rhizinflata</i> Cellulase CelAccl (3AYR)	43.2	1.8	30	344.9	576	0.64	1.39	542.5	106.2	—	⁴⁷

Table 1. Structural comparison of AC2aCel5A with its six closest structural homologues identified using the DALI server¹⁷, along with reported enzyme activities. RMSD; root-mean-square deviation of C-alpha atoms. The structural homologues are sorted based on the Z-score obtained in the DALI search. One Unit of enzyme activity was defined as the amount of enzyme releasing 1 μ mol of reducing sugar equivalents per minute. “nd” means not detected, whereas a hyphen, “—”, indicates “not tested”. ^a μ mol reducing sugar equivalents calculated as μ mol cellotriose + μ mol cellobiose + μ mol glucose, quantified by HPAEC-PAD⁷. ^bGH5 domain of CelE. ^cU/mg calculated from U/ μ mol reported in⁴⁵. ^dFor this enzyme data for both the full length protein (in parentheses), and the catalytic domain only were published.

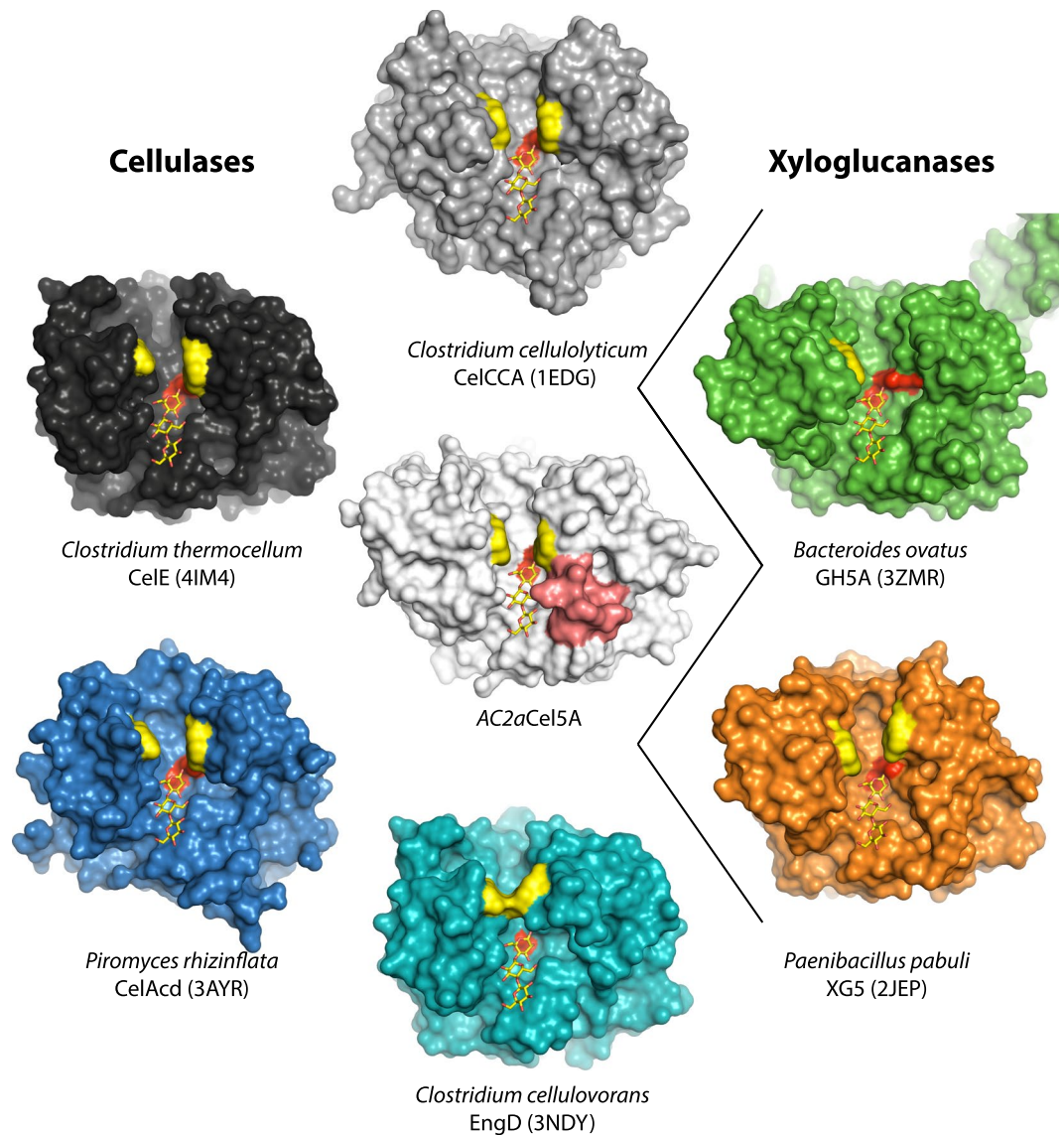


Figure 1.

References

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Additional information

Supplementary information is available for this paper at <https://doi.org/10.1038/s41598-020-62786-2>.



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