

Corrigendum

Corrigendum to: Action pattern and subsite mapping of *Bacillus licheniformis* α -amylase (BLA) with modified maltooligosaccharide substrates (FEBS 26038)

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An unfortunate mistake occurred in Fig. 2. The correct figure is given below.

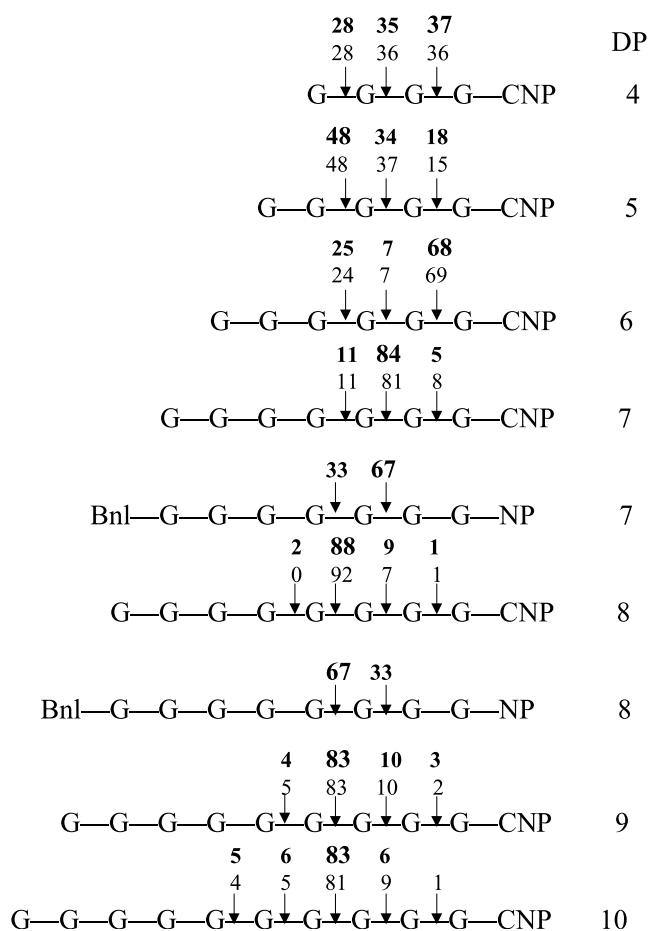


Fig. 2. Action pattern of BLA with modified maltooligosaccharides at 50°C. G, glucosyl residues; NP, 4-nitrophenyl groups; CNP, 2-chloro-4-nitrophenyl groups (NP and CNP are connected to the reducing end in β -form); Bnl groups (connected to the non-reducing end as Bnl acetals); DP, degree of polymerisation. Numbers above the arrows represent percentages of cleavage frequencies of glycosidic bonds. Bold numbers are experimental data, the bigger ones indicate the most favourable positions of cleavages. Normal numbers are related to the calculated data based on the optimised subsite map.

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