

Improving the efficiency and sustainability of chitin bioconversion through a combination of *Streptomyces* chitin-active-secretomes and mechanical-milling

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Fig. S1:

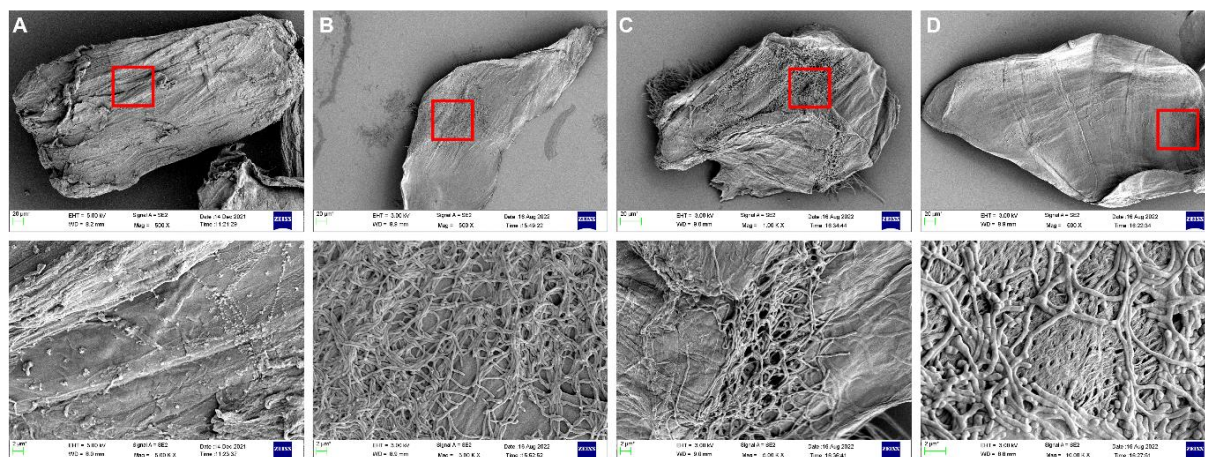


Fig. S1. FE-SEM micrographs showing the degradation of β -chitin by the *Streptomyces* spp. A – untreated β -chitin particle; B-D – β -chitin treated with *Streptomyces* sp. UH6, *S. coelicolor* and *S. griseus*, respectively. The upper panel represents the untreated and treated chitin particles, while the lower panel presents the close-up images of the target area (shown in red frame). In each panel, the scale bars of the images in the upper panel is 20 μ m and for those in the lower row, they are 2 μ m.

Fig. S2:

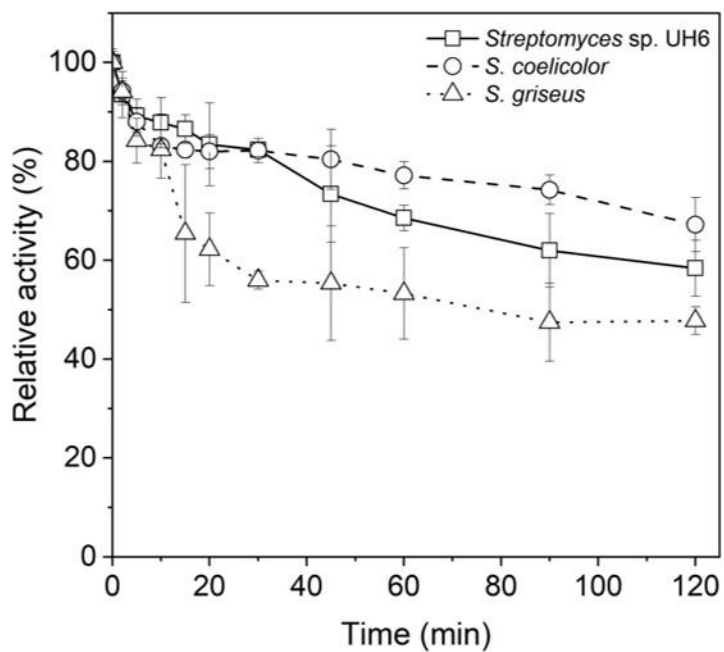


Fig. S2. Thermal stability of the *Streptomyces* spp. secretomes was assessed by pre-incubating the secretomes at 50°C for different time-intervals. The residual activity was estimated using Schales' assay. All assays were performed in triplicates, and error bars indicate standard deviation.

Fig. S3:

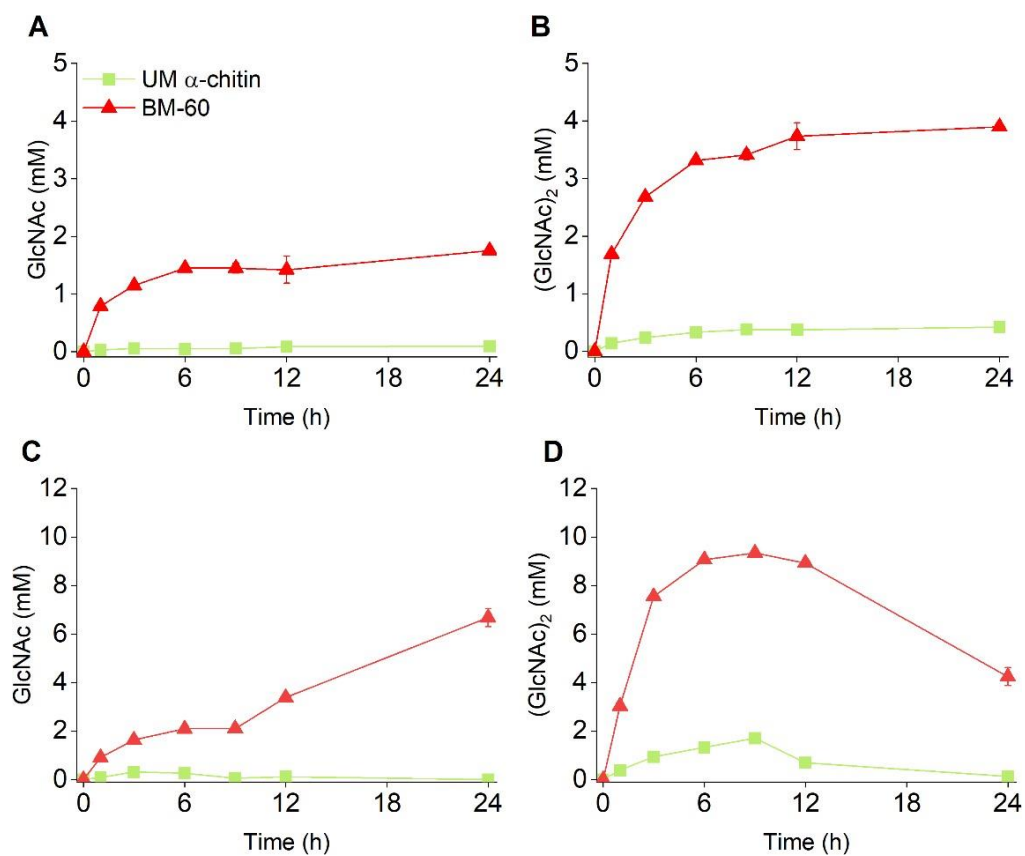


Fig. S3. A comparative analysis of GlcNAc and (GlcNAc)₂ production from un-milled α -chitin (light green) and the ball-milled α -chitin, BM-60 (red), using chitin-active-secretomes produced by *Paenibacillus* sp. LS1 (A and B) and *P. chitinolyticus* (C and D). All experiments were performed in biological triplicates and the error bars represent standard deviation.

Fig. S4:

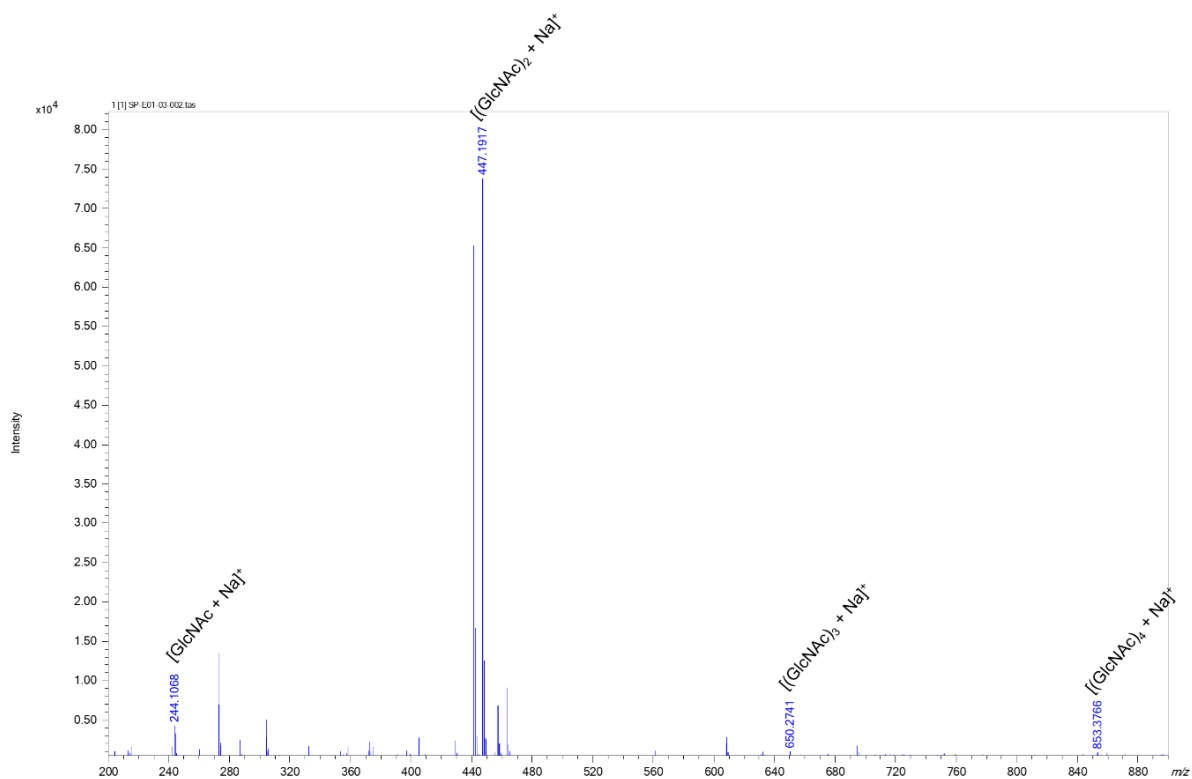


Fig. S4. MALDI-TOF-MS analysis of CHOS generated by the chitin-active-secretome of *Streptomyces* sp. UH6 from the ball-milled chitin substrate, BM-60. All identified masses were adducts of sodium [M+Na]⁺ and labelled accordingly.