## **Supporting Information**

## Biotransformation of geniposide into genipin by immobilized *Trichoderma reesei* and conformational study of genipin

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Figure S1: UV-VIS spectrum of genipin in MeOH.

Figure S2: FTIR spectrum of genipin.

Figure S3: EIMS spectrum of genipin.

Figure S4: <sup>1</sup>H-NMR spectrum of genipin in CDCl<sub>3</sub>.

Figure S5: HPLC chromatograms of genipin using chiral columns under different conditions.

Figure S6: <sup>1</sup>H-NMR spectrum of acetylated genipin in C<sub>5</sub>D<sub>5</sub>N.

Figure S7: <sup>1</sup>H-NMR spectrum of geniposide in CDCl<sub>3</sub>.

Figure S8: <sup>1</sup>H-NMR spectrum of genipin in DMCO-d<sub>6</sub>.

Figure S9: <sup>1</sup>H-NMR spectrum of genipin in C<sub>5</sub>D<sub>5</sub>N.

Figure S10: <sup>1</sup>H-NMR spectrum of genipin in DMSO-d<sub>6</sub>.

Table S1: Crystallographic data of genipin.



FIGURE S1: UV-VIS spectrum of genipin in MeOH.



FIGURE S2: FTIR spectrum of genipin.



FIGURE S4: <sup>1</sup>H-NMR spectrum of genipin in CDCl<sub>3</sub>.



FIGURE S5: HPLC chromatograms of genipin using chiral columns under different conditions. (a) Entry 1; (b) Entry 2; (c) Entry 3.



FIGURE S6: <sup>1</sup>H-NMR spectrum of acetylated genipin in C<sub>5</sub>D<sub>5</sub>N.



FIGURE S9: <sup>1</sup>H-NMR spectrum of genipin in C<sub>5</sub>D<sub>5</sub>N.



FIGURE S10: <sup>1</sup>H-NMR spectrum of genipin in DMSO-d<sub>6</sub>.

Empirical formula	$C_{11}H_{14}O_5$
Formula weight	226.22
Temperature	293(2) K
Wavelength	0.71073 Å
Crystal system	monoclinic
Space group (number)	P2 (1)
Unit cell dimensions	a=7.563(7) Å
	b=17.799(17) Å
	c=8.151(8) Å
	$\alpha=90^{\circ}$
	$\beta = 93.720(17)^{\circ}$
	$\gamma = 90^{\circ}$
Volume	1094.9(19) Å <sup>3</sup>
Z	4
Calculated density	$1.372 \text{ mg/m}^3$
Absorption coefficient	$0.109 \text{ mm}^{-1}$
F(000)	480
Crystal size	$0.402 \times 0.369 \times 0.211 \text{ mm}$
$\theta$ range for data collection	2.29-25.50°
Limiting indices	$-9 \le h \le -9, -21 \le k \le -20, -6 \le -1 \le -9$
Reflections collected/ unique	5288/3614 [R(int)=0.0465]
Data/restrictions/parameters	3614/1/295
Refinement method	Full-matrix least-squares on F <sup>2</sup>
Goodness-of-fit on F <sup>2</sup>	1.083
Final R indices $[I > 2\sigma(I)]$	R1 = 0.0757; wR2 = 0.1930
R indices (all data)	R1 = 0.0882 wR2 = 0.2022
Largest diff. peak and hole	0.457 and $-0.226 \text{ e} \cdot \text{\AA}^{-3}$

TABLE S1: Crystallographic data of genipin.