

Table S3. Percent identity matrix between the GH3 β -glucosidase. The multiple sequence alignment was performed using the Clustal Omega server (<http://www.ebi.ac.uk/Tools/msa/clustalo/>).

	KKP07226.1	KKO97705.1	KKP04308.1	KKP00605.1	KKO97043.1	KKP05725.1
KKP07226.1	100.00	30.05	28.88	25.15	23.96	25.30
KKO97705.1	30.05	100.00	46.83	30.35	31.63	31.18
KKP04308.1	28.88	46.83	100.00	26.44	28.61	27.58
KKP00605.1	25.15	30.35	26.44	100.00	41.58	40.54
KKO97043.1	23.96	31.63	28.61	41.58	100.00	50.00
KKP05725.1	25.30	31.18	27.58	40.54	50.00	100.00