

S1 Table. Genbank ID numbers for original gene sequences and Uniprot ID numbers for amino acid sequences used in this study. The sequence coding for the extended N-terminal part of HFBVI, presumably a cell wall binding domain (amino acids 1-179), was not included in the coding sequence.

Protein	Nucleotide (NCBI*)	Peptide (Uniprot**) Reference
HFB1	Z68124.1	P52754 [1]
HFBII	Y11894.1	P79073 [2]
HFBIII	18483774	G0RVE5 [3]
HFBIV	18481287	G0RHNO [4]
HFBV	18480804	G0RBZ9 this study
HFBVI	18481131	G0RFI5 this study
HYD3	AY155498	Q6YF30 [5]
HYD4	AY155499	Q6YF29 [5]
HYD5	AY158024	Q6YD93 [5]

\* <http://www.ncbi.nlm.nih.gov/> \*\* <http://www.uniprot.org/>

- [1] Nakari-Setälä, T., Aro, N., Kalkkinen, N., Alatalo, E., et al., Genetic and biochemical characterization of the *Trichoderma reesei* hydrophobin HFB1. *Eur. J. Biochem.* 1996, 235, 248–255.
- [2] Nakari-Setala, T., Aro, N., IlmeN, M., Munoz, G., et al., Differential Expression of the Vegetative and Spore-Bound Hydrophobins of *Trichoderma Reesei* Cloning and Characterization of the Hfb2 Gene. *Eur. J. Biochem.* 1997, 248, 415–423.
- [3] Kisko, K., Szilvay, G.R., Vuorimaa, E., Lemmettyinen, H., et al., Self-assembled films of hydrophobin protein HFBIII from *Trichoderma reesei*. *J. Appl. Crystallogr.* 2007, 40, 355–360.
- [4] Espino-Rammer, L., Ribitsch, D., Przylucka, A., Marold, A., et al., Two novel class II hydrophobins from *Trichoderma* spp. stimulate enzymatic hydrolysis of poly(ethylene terephthalate) when expressed as fusion proteins. *Appl. Environ. Microbiol.* 2013, 79, 4230–8.
- [5] Fuchs, U., Czymbmek, K.J., Sweigard, J.A., Five hydrophobin genes in *Fusarium verticillioides* include two required for microconidial chain formation. *Fungal Genet. Biol.* 2004, 41, 852–64.