

Table S1. Fungal sequences obtained with the primers ITS1F and ITS2.

Sequence identification	Length	Sequence 5'→3'
F1	183	AATGTTTCTGGCTGTAGAGGATATAACCGCAGTTGTAATCTCAGTTACTTAGCCCACTCCGAAAGGGAGATGGACAGCACCGCATAGCTTGCTTTGTGTGACATAGACTTAAATCATGACTTATTTCATTATTAACCAACCAAACTTTAACATGGATCTTGGCTTCTT
F2	194	GATCATTACAAAATGTTAAAGGGGTGCAGTTGCGTGTGCGTGAGCGACTGTGATTGCACTCTGTAAGCCCACCTCCGAAAGGGAGATGGACACGTGCGCTTCGAGCGTGCATGACATAGACTTTAAATCATGACTTAACTGATCTCTGGCTCCT
F3	170	AGGATCATTACAAATCATTGCAAGCCCAGTCAGCTTATTACTAC TGGCGAGCACATTGGCACAGGGTTGCTGCTGCGACGACCT TGATGTTACGCTTGTATGTTGACAGAGTAAATTGTGACTAAAA TATAGACAACCTTTAACATGGATCTCTGGCTCTT
F4	164	ATCATTACAAATCATTGCAAGCCCAGTCAGCTTATTACTACTGG GCGAGCACATTGGCACAGGGTTGCTGCTGCGACGACCTTGATGT TTACGCTTGTGTTGACAAAGTAAATTGTGACTAAAATAGACAACCTTTAACATGGATCTCTGGCTCTT
F5	224	AGGATCATTACGAGTTTGAACAGAGTTGAGCTGGCCTTCCG AGGCATGTGACGCTGCTCATCCACTCTACCCCTGTGCACCTTA CTGTAGGTTGGCGTGGGCTCTAACGGGAGCATTCTGCCGGCCT ATGTATACTACAAACACTTAAAGTATCAGAATGTAACCGCCT TAACGCATCTATAATACAACCTTAGCAACGGATCTCTGGCTCTT C

Table S2. One-factor ANOVA on the result of HZT removal in bioslurry for three different medias (complete media, glucose media and no-nutrient media) under sterile conditions

Source of variation	d.f.	S.S.	M.S.	f ratios
Media culture	2	31.37	15.68	5.77 ^s
Error	18	48.89	2.72	-
Total	20	80.26	-	-

d.f.= degree of freedom; S.S.= sum of squares; M.S.= mean squares; s/ns= significant or not significant at p < 0.10

Table S3. One-factor ANOVA on the result of HZT removal in bioslurry for sterile and non-sterile conditions with no-nutrient media.

Source of variation	d.f.	S.S.	M.S.	f ratios
Culture conditions	1	12.32	12.32	3.18 ^{ns}
Error	12	65.82	5.49	-
Total	13	78.14	-	-

d.f.= degree of freedom; S.S.= sum of squares; M.S.= mean squares; s/ns= significant or not significant at p < 0.10

Table S4. One-factor ANOVA on the result of HZT removal in bioslurry for inoculated and non-inoculated cultures with no-nutrient media under non-sterile conditions.

Source of variation	d.f.	S.S.	M.S.	f ratios
Culture conditions	1	5.21	5.21	0.70 ^{ns}
Error	12	88.94	7.41	-
Total	13	94.15	-	-

d.f.= degree of freedom; S.S.= sum of squares; M.S.= mean squares; s/ns= significant or not significant at $p < 0.10$