

Table S2. Transposable element content of *C. amyloletus*, *C. floricola* and *C. wingfieldii* genomes. Estimates obtained from the RepeatMasker output using the combined databases: Dfam_Consensus-20170127, RepBase-20170127.

	<i>C. amyloletus</i> CBS6039			<i>C. amyloletus</i> CBS6273			<i>C. floricola</i> DSM27421			<i>C. wingfieldii</i> CBS7118		
	No. elements	length occupied (bp)	% of sequence	No. elements	length occupied (bp)	% of sequence	No. elements	length occupied (bp)	% of sequence	No. elements	length occupied (bp)	% of sequence
Retroelements:												
LINEs:												
CRE/SLACS	8	3,408	0.02	8	3,376	0.02	62	36,804	0.17	62	97,521	0.47
LTRs:												
Ty1/Copia	39	13,246	0.07	36	13,377	0.07	69	46,563	0.21	92	63,727	0.30
Gypsy/DIRS1	77	62,444	0.31	86	60,520	0.31	281	393,429	1.78	204	314,094	1.50
DNA transposons:												
Tc1-IS630-Pogo	5	829	< 0.01	9	1,693	< 0.01	1	92	< 0.01	1	62	< 0.01
Tourist/Harbinger	14	3,568	0.02	15	3,606	0.02	5	1,054	< 0.01	22	7,145	0.03
Unclassified	6	739	< 0.01	4	550	< 0.01	23	4,982	0.02	25	4,116	0.02