

Table S4: Sequence identity in the ITS and D1D2 domains of the 26S rRNA gene of *Blastobotrys* species

		<b>Y6417</b>	<b>Y27120</b>	<b>YB1343</b>	<b>L1-24</b>	<b>L2-36</b>
<i>B. malaysiensis</i>	<b>Y6417</b>	1134	99%	97%	97%	97%
			0%	0%	0%	0%
<i>B. mokoensis</i>	<b>Y27120</b>	1125	1136	97%	97%	97%
		2		0%	0%	0%
<i>B. illinoisensis</i>	<b>YB1343</b>	1114	1113	1132	97%	97%
		8	6		0%	0%
<i>B. yvelinesensis</i>	<b>L1-24</b>	1113	1113	1111	1134	100%
		10	8	4		0%
<i>B. yvelinesensis</i>	<b>L2-36</b>	1113	1113	1111	1134	1134
		10	8	4	0	

Blue coloured cells represent the number and percentage of nucleotide identity

Yellow coloured cells indicated the number of gaps and associated percentage.