

Additional file 7: Table S3

Genomic PCR genotyping the four scaffolds (M, F, 33, X) in *T. reesei* wild isolates and industrial strains.

Wild isolate	Origin	Mating type	PCR primers A + B (F)	PCR primers C+ D (X)	PCR primers C + B (33)	PCR primers A + D (M)	<i>re</i>
CBS999.97	French Guiana	<i>MAT1-1</i>	+	+	-	-	+
CBS999.97	French Guiana	<i>MAT1-2</i>	-	-	+	+	-
QM6a	Solomon Islands	<i>MAT1-2</i>	-	-	+	+	-
G.J.S. 86-404	French Guiana	<i>MAT1-1</i>	-	-	+	+	-
G.J.S. 86-410	French Guiana	<i>MAT1-1</i>	-	-	+	-	-
G.J.S. 84-473	French Guiana	<i>MAT1-1</i>	-	-	-	-	-
G.J.S. 89-7	Brazil Para	<i>MAT1-2</i>	-	-	+	+	-
G.J.S. 97-178	Brazil Para	<i>MAT1-2</i>	-	-	+	+	-
G.J.S. 85-249	Indonesia Celebes	<i>MAT1-1</i>	-	-	+	+	-
G.J.S. 85-229	Indonesia Celebes	<i>MAT1-2</i>	-	-	+	+	-
G.J.S. 85-236	Indonesia Celebes	<i>MAT1-2</i>	-	-	+	+	-
G.J.S. 93-23	New Caledonia	<i>MAT1-2</i>	-	-	+	-	-
QM9414		<i>MAT1-2</i>	-	-	+	+	-
RUT-C30		<i>MAT1-2</i>	-	-	+	+	-