

Table S4. Boundaries and nucleotide divergences for regions of the mating-type (*mat*) chromosomes originating from three *Neurospora tetrasperma* wild-type heterokaryons.

Strain	Pseudoautosomal (PA)		Region of elevated sequence divergence		Pseudoautosomal (PA)		Introgression tract
	ID	region 1	Position	Nucleotide divergence	Position	Nucleotide divergence	
L1	1-	0.0065	1,390,001-	0.0187	6,760,001-	0.0044	1,400,001-
	1,390,000		6,760,000		7,657,888		5,490,000
L4	1-	0.0062	1,140,001-	0.0238	7,000,001-	0.0043	1,500,001-
	1,140,000		7,000,000		7,657,888		6,710,000
L9	1-	0.0068	900,001-	0.0343	6,780,001-	0.0054	1,120,001-
	900,000		6,780,000		7,657,888		6,710,000