

**Table S2: List of primers and double-stranded adaptors used for AFLP**

Type of sequences	Name	Sequence
<b>Double-strand adaptors</b>	<i>Pst</i> I-adaptor	5'-CTCGTAGACTGCGTACATGCA-3'                     3'-CATCT GACGCATGT-5'
	<i>Taq</i> I-adaptor	5'-GACGATGA GT CC TGAG-3'                    3'-TACT C AGG ACTCGC-5'
<b>Pre-selective primers</b>	<i>Pst</i> I+0	GACTGCGTACATGCAG
	<i>Taq</i> I+0	GATGAGTCCTGAGCGA
<b><i>Taq</i>I-specific selective primers</b>	Taq+AA	GATGAGTCCTGAGCGAAA
	Taq+AC	GATGAGTCCTGAGCGAAC
	Taq+AG	GATGAGTCCTGAGCGAAG
<b><i>Pst</i>I-specific selective primers</b>	<i>Pst</i> I+AG	<b>6FAM</b> -GACTGCGTACATGCAGAG
	<i>Pst</i> I+CC	<b>NED</b> -GACTGCGTACATGCAGCC
	<i>Pst</i> I+GT	<b>PET</b> -GACTGCGTACATGCAGGT
	<i>Pst</i> I+TG	<b>VIC</b> -GACTGCGTACATGCAGTG

The six primer pairs used to draw the phylogenetic tree in fig 1: *Taq*+AA & *Pst*I+AG, *Taq*+AC & *Pst*I+AG, *Taq*+AA & *Pst*I+TG, *Taq*+AC & *Pst*I+TG, *Taq*+AG & *Pst*I+TG and *Taq*+AC & *Pst*I+CC