

Tableau 1. Gènes cibles et amorces suggérées pour l'outil de typage génomique NG-STAR

Cible	Amorce	Séquence de nucléotides (de 5' à 3')	Taille (pb) de l'amplicon	Taille des allèles (pb) dans NG-STAR	Référence
<i>penA</i>	PenA-A1 (forward) PenA-B1 (reverse)	CGGGCAATACCTTTATGGTGGAAAC AACCTTCCTGACCTTTGCCGTC	669	1746-1752	Ito et al, 2005
	PenA-A2 (forward) PenA-B2 (reverse)	AAAACGCCATTACCCGATGGG TAATGCCGCGCACATCCAAAG	581		
	PenA-A3 (forward) PenA-B3 (reverse)	GCCGTAACCGATATGATCGA CGTTGATACTCGGATTAAGACG	863		
	PenA-A4 (forward)	AATTGAGCCTGCTGCAATTGGC			
<i>mtrR</i>	MTR1 (forward) MTR2 (reverse)	AACAGGCATTCTTATTTTCAG TTAGAAGAATGCTTTGTGTC	916	698 - 708	Mavroidi et al, 2001
<i>porB1b</i>	por-NGMAST-F (forward) por-NGMAST-R (reverse)	CAAGAAGACCTCGGCAA CCGACAACCACCTTGGT	737	30	Martin et al, 2004
<i>ponA</i>	ponA1-f (forward) ponA1-r (reverse)	CGCGGTGCGGAAAACCTATATCGAT AGCCCGGATCGGTTACCATACGTT	1240	75	Ropp et al, 2011
<i>gyrA</i>	GYRA-1 (forward) GYRA-2 (reverse)	AACCCCTGCCCGTCAGCCTTGA GGACGAGCCGTTGACGAGCAG	270	264	Lindback et al, 2002
<i>parC</i>	parC F (forward) parC R (reverse)	GTTTCAGACGGCCAAAAGCC GGCATAAAATCCACCGTCCCC	332	332	Ng et al, 2002
23S rRNA	gonrRNAF (forward) gonrRNAR2 (reverse)	ACGAATGGCGTAACGATGGCCACA TTCGTCCACTCCGGTCTCTCGTA	712	567	Ng et al, 2002

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