

MK000064=C/1608

Marker: C/1608

Type: Dominant PCR

Description:

Reference: Euphytica 127: 353–365, 2002. PCR-based markers to differentiate the mitochondrial genomes of

petaloid and male fertile carrot (*Daucus carota* L.) Inga C. Bach, Annette Olesen & Philipp W. Simon,

Primers: atp1-d1 (c).....5'-CGCGTTGGGAGTACCTATTGA-3'

cmt-4 (e).....5'-TCCCGCAGGTCGTCAACAG-3'

PCR Reaction: 20 µl: [0.4 µg/ml DNA=8 ng; 0.4 µM each primer=8 pMol each; 0.025 U/µl Taq=0.5 U; 1.5 mM MgCl₂=30 nMol; 0.1 mM each dNTP=2 nMol]

PCR Program: 94°C 2:00; 35 cycles of {94°C 1:00; 55°C 1:00; 72°C 2:30}; 72°C 7:00

Screening Method: Product size by agarose gel

Product Sizes: 1608 in N cytoplasm; no product in Sp cytoplasm

Example:

Diagram of how it works:

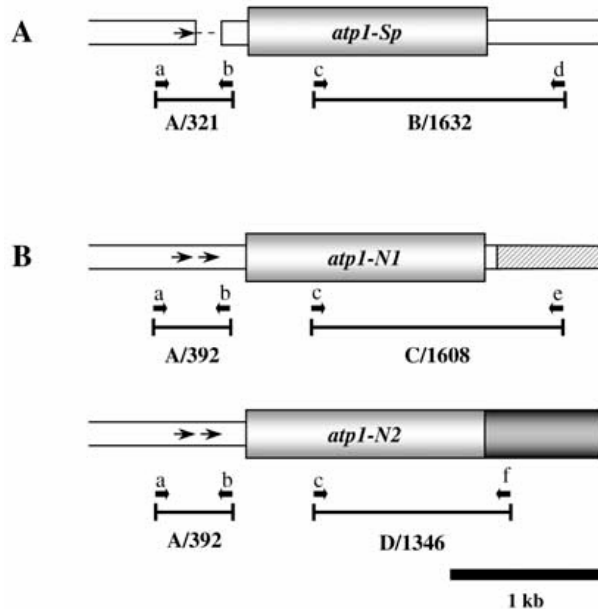


Figure 1. Structures of mtDNA *atp1* genes and flanking regions in line K826A with SpC cytoplasm (A) and line K831B with N cytoplasm (B). Open reading frames (ORFs) are indicated by large shaded boxes and intergenic regions by open boxes. Nonhomologous regions are indicated by darker shading (ORFs) or crosshatching (intergenic regions). Thin arrows indicate the sites of a 71 bp fragment that forms a direct repeat in the fertile line. The lack of this repeat upstream from *atp1-Sp* is indicated by a dotted line. Thick arrows indicate the annealing sites and orientations of the primers cmt-1 (a), cmt-2 (b), *atp1-d1* (c), cmt-3 (d), cmt-4 (e) and cmt-5 (f). PCR amplification products serving as markers are indicated by bars labeled A/321, A/392, B/1632, C/1608 and D/1346.

Genbank reference: The DNA sequences of the *atp1-Sp*, *atp1-N1* and *atp1-N2* loci have been assigned GenBank Accession Nos. [AF301602](#), [AF301604](#) and [AF301603](#), respectively.

Sequence Information:

Map Location:

Published Reference:

Other Information:

Primer Location (lab specific): Box 0 X0

PCR Program Name (lab specific):