

B. Guo · D. A. Sleper · P. R. Arelli · J. G. Shannon
H. T. Nguyen

Identification of QTLs associated with resistance to soybean cyst nematode races 2, 3 and 5 in soybean PI 90763

Published online: 7 February 2006
© Springer-Verlag 2006

Theor Appl Genet (2005). 111:965–971

The maps of linkage groups G–O were inadvertently omitted from Fig. 2. The complete figure is reproduced here.

The online version of the original article can be found at <http://dx.doi.org/10.1007/s00122-005-0031-2>

B. Guo · D. A. Sleper (✉) · J. G. Shannon · H. T. Nguyen
Division of Plant Sciences and Center for Soybean Biotechnology,
University of Missouri-Columbia, Columbia, MO 65211, USA
E-mail: SleperD@missouri.edu
Tel.: +1-573-8827320

P. R. Arelli
USDA-ARS-MSA, 605 Airways Blvd, Jackson, TN 38305, USA

D. A. Sleper
271-F Life Sciences Center, University of Missouri-Columbia,
Columbia, TN 65211, USA

Fig. 2 Linkage map constructed from the cross Hamilton × PI 90763. QTLs are indicated by a *bar* on the right of linkage group and its 1-LOD support interval was given by the length of the bar. *Bold* SSR markers are dominant. Molecular markers Sat_036 and Satt244 produced two loci each. Faster bands (locus) are suffixed by f and slower bands by s. The two loci of Satt244 are mapped on nearly the same position. One locus of Sat_036 is mapped on the same position (D1a) as the soybean composite linkage map but the other on LG L

