

Erratum to: Genetic structure and diversity of the *Banana bunchy top virus* population on Sumatra Island, Indonesia

Yuya Chiaki · Nasril Nasir · Henny Herwina · Jumjunidang ·
Ayaka Sonoda · Tomohiro Fukumoto · Masayuki Nakamura · Hisashi Iwai

Published online: 30 May 2015
© Koninklijke Nederlandse Planteziektenkundige Vereniging 2015

Erratum to: Eur J Plant Pathol
DOI 10.1007/s10658-015-0669-9

In the base sequence of Type A and B of Fig. 2, “T” which is the 14th from the 5’end is a mistake, and a right base is “A”. Please replace with the figure indicated here.

The online version of the original article can be found at <http://dx.doi.org/10.1007/s10658-015-0669-9>.

Y. Chiaki · A. Sonoda · T. Fukumoto · M. Nakamura ·
H. Iwai (✉)
Faculty of Agriculture, Kagoshima University, 1-21-24,
Korimoto, Kagoshima 890-0065, Japan
e-mail: topcrop@agri.kagoshima-u.ac.jp

Y. Chiaki · T. Fukumoto · M. Nakamura · H. Iwai
The United Graduate School of Agricultural Science,
Kagoshima University, 1-21-24, Korimoto,
Kagoshima 890-0065, Japan

N. Nasir · H. Herwina
Biology Department, Faculty of Mathematics and Natural
Science, Andalas University, Padang 25163 West Sumatra,
Indonesia

Jumjunidang
Indonesian Tropical Fruits Research Institute, Solok 52213
West Sumatra, Indonesia

T. Fukumoto
Fruit Tree Branch, Kagoshima Prefecture Institute for
Agricultural Development, Tarumizu, Kagoshima 891-2112,
Japan

Fig. 2 The secondary structure of the stem-loop region of DNA-U3 in BBTV Sumatra isolates. Sequence differences are indicated with boxes

