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Physical Mapping of β-Converting and γ-Nonconverting Corynebacteriophage Genomes

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Volume 148, no. 1, p. 135, Table 1, footnote *b*: Should read "Digestion products from γ-*tsr*-1 DNA. Since γ-*tsr*-2 does not contain HET fragments, the letter designations of its restriction fragments will, in some cases, differ from letter designations of identical fragments in γ-*tsr*-1 digests. Thus, in the *Bam*HI restriction digests, γ-*tsr*-1 fragments GHI become γ-*tsr*-2 fragments EFG."

Page 141, legend to Fig. 9, line 5: Should read "symbols are explained in Fig. 3 and 8. Note that γ-*tsr*-2 fragments EFG are equivalent to γ-*tsr*-1 fragments GHI. See footnote *b* of Table 1."

Identification of Deoxyribonucleic Acid Restriction Fragments of β-Converting Corynebacteriophages That Carry the Gene for Diphtheria Toxin

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Volume 148, no. 1, p. 160, column 1, lines 23–28, and column 2, lines 1–8: Should read "lated ATTP$_\beta$ fragment (Fig. 6). HETA, which carries the *tox* gene and includes the adjacent part of *att*P (3), hybridized to fragments A and C in the *Hae*II digests and A and B in the *Hinc*II digests of ATTP$_\beta$. HETB, which carries the portion of ATTP$_\beta$ not carried by HETA (including a part of *att*P) (3), hybridized to bands C and A in the *Hinc*II digest and band B only in the *Hae*II digest. Thus, the *Hinc*II restriction products of the ATTP$_\beta$ fragments must be ordered C-A-B as depicted in Fig. 7. An analysis of *Hae*II partial digestion products of the ATTP$_\beta$ fragment showed that *Hae*II restriction fragments were ordered B-A-C."

Page 160, column 2, lines 19 and 20: Should read ". . . gene is within *Hae*II fragments A and C and *Hinc*II fragments A and B. It was. . . ."

Page 161, Fig. 7: *Hinc*II map portion "B(0.8) A(2.4) C(0.7)" should read "C(0.7) A(2.4) B(0.8)."