

## ERRATUM

### Biological, Molecular, and Structural Analysis of a Cytopathic Variant from a Molecularly Cloned Simian Immunodeficiency Virus

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Volume 68, no. 9, p. 5512–5513: Figure 2 should appear as shown on the following pages.



**B (cont.)**

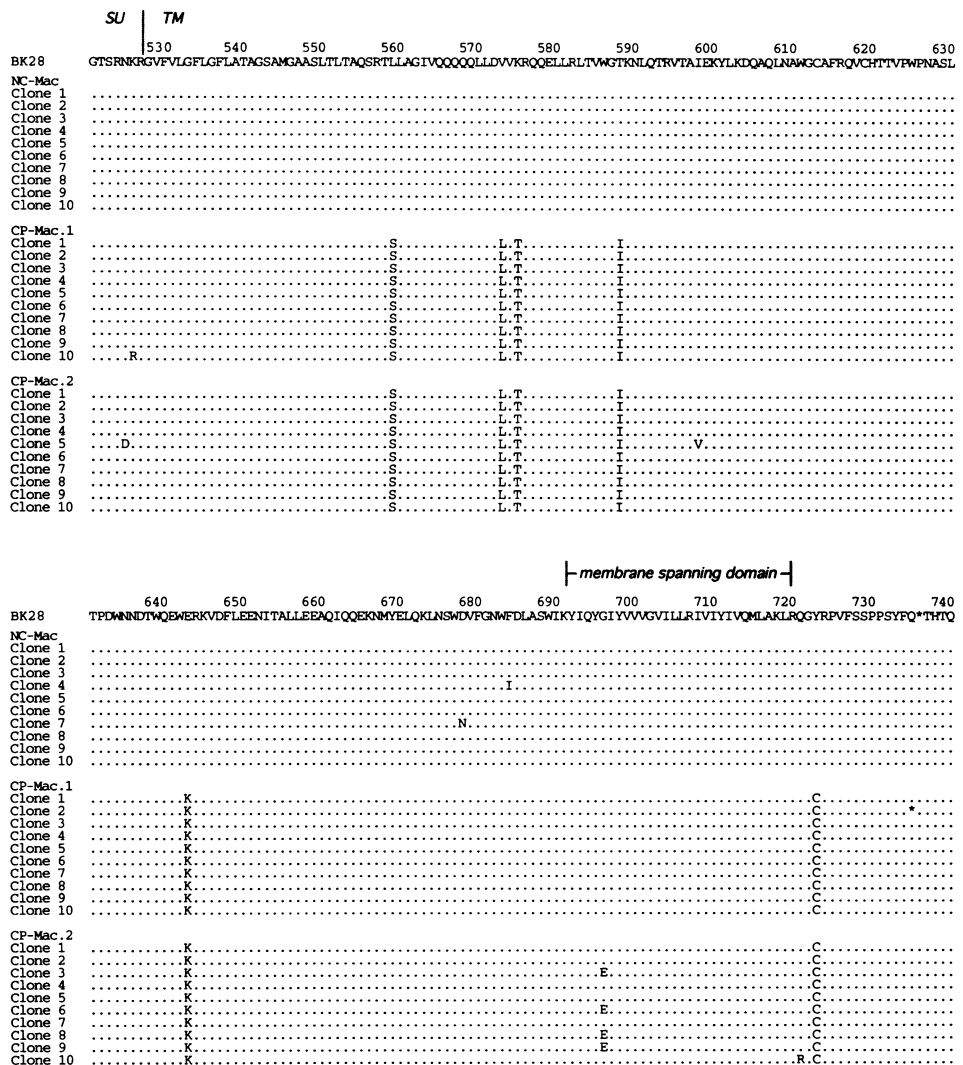


FIG. 2. Amino acid differences of the NC-MAC and CP-MAC *env* clones compared with BK28. Predicted amino acid sequences are shown for selected regions of *env* for PCR-generated clones of NC-MAC-infected cells and two cellular clones of CP-MAC-infected cells and are compared with the parental BK28 sequence. Dot, amino acid identity; asterisk, in-frame stop codon; dash, a single nucleotide deletion. Regions corresponding to variable domains in SU (39) are shown. (A) Sequences of SU from amino acids 40 to 150; (B) sequences including the carboxy-terminal half of SU (amino acids 301 to 527) and the entire TM coding region. Regions of *env* omitted from the figure contained only sporadic point mutations and are described in Results.