ERRATA

Phosphorylation of Serine Residue 89 of Human Adenovirus E1A Proteins Is Responsible for Their Characteristic Electrophoretic Mobility Shifts, and Its Mutation Affects Biological Function

CAROLYN L. SMITH, CHRISTINE DEBOUCK, MARTIN ROSENBERG, AND JEFFREY S. CULP

Department of Molecular Genetics, Smith Kline & French Laboratories, King of Prussia, Pennsylvania 19406-0939

Volume 63, no. 4, page 1574: Table 2 should appear as shown below.

TABLE 2. Transformation of REF52 cells by T24 *ras* and E1A 289R WT or mutant proteins

Protein(s) expressed	No. (%) of foci[a]
WT E1A	0
T24 *ras*	0
T24 *ras* + WT E1A	65
T24 *ras* + Ser-63 to Gly	7 (11)
T24 *ras* + Ser-69 to Gly	9 (14)
T24 *ras* + Ser-89 to Gly	40 (62)
T24 *ras* + Ser-96 to Gly	42 (64)

[a] Foci were defined as morphologically transformed G418-resistant colonies. Values are the averages from four experiments. Percentages are of foci produced by mutant E1A protein compared with those produced by WT E1A.

Nucleotide Sequence and Distinctive Characteristics of the *env* Gene of Endogenous Feline Leukemia Provirus

D. VINOD KUMAR, BRIAN T. BERRY, AND PRADIP ROY-BURMAN

Departments of Pathology and Biochemistry, University of Southern California School of Medicine, Los Angeles, California 90033

Volume 63, no. 5, p. 2382, Fig. 2, lines 3 and 4, 21st amino acid of region I:

"I" should read "I"

:

"F" "F"

Figure 2, lines 2 and 3, ninth amino acid downstream of the end of region I:

"V" should read "V"

:

"I" "I"

Figure 2, lines 1 and 2, fourth amino acid of region II:

"G" should read "G"

:

"D" "D"

Figure 2, line 1, 18 amino acids upstream of region V: "MGPNP" should read "MGPND."