

Corrigenda

Retroviral integrase domains: DNA binding and the recognition of LTR sequences

by E.Khan, J.P.G.Mack, R.A.Katz, J.Kulkosky and A.M.Skalka

Nucleic Acids Research, **19**, pp. 851–860 (1991)

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Characterization of the VirG binding site of *Agrobacterium tumefaciens*

by G.J.Pazour and A.Das

Nucleic Acids Research, **18**, pp. 6909–6913 (1990)

The authors wish to apologize for an error which occurred in the Abstract of the above paper. The last sentence in the Abstract should read as follows:

Mutational analysis indicate that the functional vir box sequence is 14 residues in length, contains a dyad symmetry and has the consensus sequence d ryTncAaTTGnAaY (r=purine, y=pyrimidine).

A complex RNA sequence determines the internal initiation of encephalomyocarditis virus RNA translation

by A.G.Evstafieva, T.Y.Ugarova, B.K.Chernov and I.N.Shatsky

Nucleic Acids Research, **19**, pp. 665–671 (1991)

This paper was inadvertently published with a typographical error in the title. The correct title is reprinted above.