

Corrigendum

Corrigendum to: **Structural and functional similarities between HIV-1 reverse transcriptase and the *Escherichia coli* RNA polymerase β' subunit (FEBS 24219)**

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Table 1 was incorrect, it should read:

Table 1
Reaction patterns of anti-RT MAb with synthetic peptides and RT mutants

	MAbs (epitopes)				
	11G10 (193–284)	31D6 (294–318)	31G8 32E7 33D5 (294–318)	5B2 (294–318)	7D8 (334–410)
<i>A: RT peptides</i>					
Pep 1: aa 294–318 PLTEEALELAENREIL-KEPVHGVY	—	+	+	+	—
Pep 2: aa 294–305 PLTEEALELAELAE	—	+	+	—	—
Pep 3: aa 306–318 NREILKEPVHGVY	—	—	—	+	—
<i>B: RT mutants</i>					
Mut 1: 294-PLTLEAELELAELAE-305	+	—	ND	+	+
Mut 2: 294-PLTLEAQLELAELAE-305	+	—	ND	—	—
Mut 3: 294-PLTEEALEAEVLAELAE-305	+	+	ND	+	+

ELISA was used to detect MAb-binding to the peptides (A). Western blot was used to detect binding to the mutants (B). Amino acid residues are written with their single-letter abbreviation, mutations are in bold. +/— denotes binding/no binding. ND, not tested.

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