Erratum to: Human immunodeficiency virus reverse transcriptase base misincorporations can promote strand transfer

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Unfortunately in the original publication the “Full-length –Acceptor” and “Full-length +Acceptor” labels were

reversed for both the “Low dATP” and “Equal dNTPs” columns. Attached please find the corrected version of Table 1.

Table 1 Compilation of results from mutant colony frequency experiments^a

Low dATP ^b				Equal dNTPs			
# ^c	Full-length –Acceptor ^d	Full-length +Acceptor	Transfer	#	Full-length –Acceptor	Full-length +Acceptor	Transfer
1	0.140	0.094	0.182	1	0.033	0.029	0.035
2	0.130	0.118	0.141	2	0.047	0.040	0.053
3	0.146	0.130	0.133	3	0.025	0.025	0.021
4	0.103	0.100	0.091	4	0.034	0.028	0.030
Ave.	0.132	0.112	0.138	5	0.048	0.040	0.052
SD	0.020	0.017	0.037	6	0.047	0.033	0.052
				7	0.029	0.023	0.030
				8	0.034	0.028	0.036
				Ave.	0.037	0.031	0.039
				SD	0.009	0.006	0.012

^a The mutant colony frequency is defined as the number of white or faint blue colonies divided by the total number of colonies

^b Equal dNTPs refers to reactions with 100 μ M of each of the 4 dNTPs while the concentration of dATP was reduced to 1 μ M for low dATP reactions

^c Refers to the number of independent experiments performed using the stated conditions

^d Full-length refers to DNA products extended to the end of the donor template while transfer products transferred to, and were extended on the acceptor template

The online version of the original article can be found under
doi:[10.1007/s007050070113](https://doi.org/10.1007/s007050070113).

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