



Correction to: A candidate gene study of intermediate histopathological phenotypes in HIV-associated neurocognitive disorders

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Due to a production error data in Table 1 were not presented correctly. The corrected table is shown below.

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Table 1 Sample characteristics

Age at death	47.4 years ^a (SD = 9.2)
Length of HIV infection	12 years (SD = 6.4)
CD4+ T cell count	122 (SD = 168)
Median plasma HIV-1 RNA viral load	15,406 copies/mL
CPE	8.96 (SD = 4.08)
Global clinical rating	5.31 (SD = 1.87)
	Percent ^b of sample (<i>n</i>)
Detectable (>50 copies/mL) plasma HIV-1 RNA viral load	84% (30)
Male	81.4% (162)
Race/ethnicity	
Caucasian	50.8% (100)
African American	24.9% (49)
Hispanic	22.3% (44)
Asian/Native Alaskan	2% (4)
Major depression	
Current	23.4% (32)
Past	16.8% (23)
Substance use disorder	
Current	20.3% (31)
Past	29.4% (45)
Alcohol use disorder	
Current	9% (14)
Past	26% (40)
HIV encephalitis	9.2% (16)
HAND	74.2% (135)
ANI	14.2% (26)
MND	31% (56)
HAD	29% (53)

SD standard deviation, *CPE* central nervous system penetration or effectiveness, *ANI* asymptomatic

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