

Supporting Table 2. The titers of dual AAV vectors are similar to those of regular AAV vectors of normal size.

	AAV2/2			AAV2/8		
	NS ^a (GC ^b /mL)	5'-half (GC/mL)	3'-half (GC/mL)	NS (GC/mL)	5-half (GC/mL)	3'-half (GC/mL)
Average	3.8x10 ¹²	8.3x10 ¹²	4.2x10 ¹²	9.8x10 ¹²	1.1x10 ¹³	6.7x10 ¹²
s.e.m.^c	6.8x10 ¹¹	1.6x10 ¹²	9.6x10 ¹¹	2.3x10 ¹²	1.3x10 ¹²	1.4x10 ¹²
n^d preps	20	31	19	20	47	27

N.B. ^aNS: normal size; ^bGC: genome copies; ^cs.e.m.: standard error of the mean; ^dn: number of AAV preparations. The 5'-half sample includes single 5'-halves of dual AAV overlapping, trans-splicing and hybrid vectors; the 3'-half sample includes single 3'-halves of dual AAV overlapping, trans-splicing and hybrid vectors. No statistically significant differences using ANOVA were found in the titers of NS AAV2/2 or AAV2/8 compared to those of dual AAV vectors. More details on the statistical analysis including specific statistical values can be found in the Statistical analysis paragraph of the Materials and Methods section.