

**Table 1** Volume usage in single- vs. double-stranded icosahedral families

	Number of Families	Average (%)	Standard deviation (%)	Spearman's rank correlation between volume usage and genome length	
				$\rho$	<i>p</i> -value
Single-stranded viruses	11	49	25	-0.3	0.37
Double-stranded viruses	13	24	10	0	1
All viruses	24	35	22	-0.17	0.42