

Table S7 PTERV1 distribution in chimpanzee genome

	Full-Length*	Solo LTR**
Genic	14	8
Non-genic	93	82
Total	107	90

*Full-length elements are those which extend beyond the LTR into gag and/or env portion. Because of the repetitive nature of these elements, Arachne failed to provide continuous sequence these elements.

**Solo LTR are those which consist solely of LTR sequence.