

Predicted feature (nt)	Sequence
E2 binding sites	ACC-N4-GGT
	ACC-N5-GGT
	ACC-N6-GGT
	ACC-N7-GGT
Polyadenylation sites	AATAAA
Sp1 binding sites	GGCGGG
NF1 binding sites	CGGAA
AP1 binding site	TGANTCA

*All motif sites displayed inside ORFs

Predicted feature (aa)	Translation
ATP-dependent helicase motif	GPPNTGKS
Cyclin interaction RXL motif	KRRLF
Metal-binding motifs	CXXC-X29-CXXC
Metal-binding motifs	CXXC-X29-CXXC
Nuclear localisation signal	RKRRR
Nuclear localisation signal	KRKR
Retinoblastoma binding domain	LXCXE

*All binding sites displayed outside ORFs

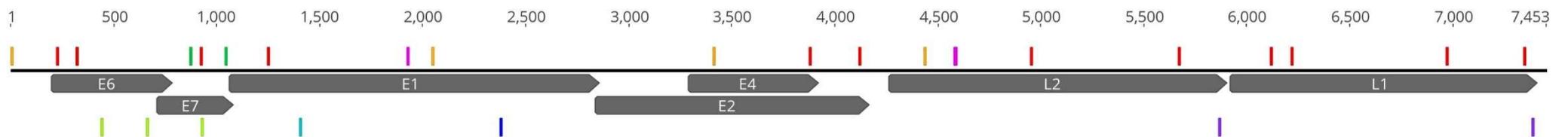


Figure S1 Felis catus Papillomavirus 6 (FcaPV6) genome configuration and nucleotide (nt) and amino acid (aa) feature location.