

SUPPLEMENTAL TABLE 1  
Primers used to sequence the genome of Entebbe bat virus

Primer	5' to 3' sequence	Primer	5' to 3' sequence
15F	TAATCGCTTGGCGTTAGTCC	5979F	CCAGACCGACACACTGACAC
859R	CCCATCTTTCCACTTTTCC	6819R	ATCTCTGCATTCTGCGTCT
767F	CAAGCCACATAACGGAAGGT	6610F	TCCCAGAAG CTATGCTCACC
1418R	TCAGAGGTGTGCAGTTGGAG	6976R	TCTGCTTGTGACCGAAGATG
1154F	GGAAGGTGTGCTACGATGCT	6726F	GATGGGGATGTTTCAGTCGT
1751R	TCACAGCTCCTTCCTGATCC	7567R	AGAAGCCCAGCCACACCTAC
1399F	CTCCAAC TGCACACCTCTGA	7420F	TCTCACTCGCCAATGTGTTC
2192R	CTTTCAATGCCCTCATGT	8257R	ACGAGTCCACCACCAAAAAC
1908F	CGGACACCAAACAGTGCTAA	7979F	GAGGACAACCTATGGCTGGA
2650R	TTCCACATTCGATGCTCAAG	8796R	CCAGTTGCTGGAAGATCCAC
2491F	GGTGATGCGTGTCATCTTC	8678F	CACACCTTTGGACAGCAGA
3299R	TTGGTCCCTGGACAGACTTC	9523R	TCCTCTGACTCAGCCATCCT
3142F	GGACAAGCACAGTTCAAGA	9327F	ATCCTCGCTGAAGCAGTCAT
3973R	AGCCAGTAATCCGAGCATTG	10125R	TCTCCTGCATGTGAGGATTG
3839F	TCCATATCCCAAGTGGTGCT	9882F	TGGATGGTAAGGGAAACTGG
4675R	CCCTTCTGGTTTATGGCGTA	10500R	GGTCTTTTCCAGGTGTCAA
4415F	GGACCTCTGAACGCTTGGAC	RACE514R	TCCAGGGTTCCACTAAACA
4867R	GAGTTTCCAGTTGCCACCAT	RACE388R	ATTGGCGATGTTCTCACTC
4580F	CAGCAACATCATGTGGGAAT	RACE10328F	TACGCGGACTACATGCAAAC
5406R	CTGGTTCAAGCAACCGATTT	-	-
5285F	TGAAGCACTCAAGGACATGC	-	-
6091R	CCAGGGATTCCATGTTGTC	-	-