

Name	Oligo	Sequence 5' - 3'	Accession number	PCR efficiency	Reference
FIV gag	Forward Reverse Probe	GCAGAAGCAAGATTTGCACCA TATGGCGGCAATTTTCCT FAM-TGCCTCAAGATACCATGCTCTACACTGCA-BHQ1	M25381	1.021	[38]
RPP30	Forward Reverse Probe	GCCAAGTGTGAGGGCTGAAA GTGGACGGACTACCAAATGGA HEX-CCAGTCCTCGTCAGCGCTCCCTTC-BHQ1	XM_003994205	1.048	-
OASL	Forward Reverse	GTGAAACATCGGCCAACCA TGGCTTTCACGTAAGTGCAGGTA	ENSFCAT0000009643	0.950	-
ACHE	Forward Reverse	GTGCTCGGGTCTATGCCTACA GCCCAAAGATGAACTCGATCTC	ENSFCAG0000023546	1.001	-
TGM2	Forward Reverse	CCCGAGTGGTGACCAACTATAAC GCGGAAGTACTCGATGAGCAA	ENSFCAT0000004809	1.029	-
HMG2	Forward Reverse	AAGCTGATGCTGGCAAGGA TCTGTGCTGGTCTGTTTTGG	ENSFCAT0000032519	0.964	-
CDKN1A	Forward Reverse	GGCAGACCAGCATGACAGATT CAGATTAGGGCTTCCTCTTGA	ENSFCAT0000025177	0.987	-
ZFP36	Forward Reverse	CCACCCCAAGTACAAGACAGAAGT GATGCTCTGGCGAAGCACAT	ENSFCAT0000031857	0.939	-
BCL6	Forward Reverse	CGTGATGGCTACGGCTATGTAC CGGCTTCACTGGCTTGATA	ENSFCAT0000006812	0.931	-
CSF1	Forward Reverse	GATGGAGACTTCGTGCCAAATT AGGTAGCACACGGGATCTTTCA	ENSFCAT0000010774	0.923	-
IFI44	Forward Reverse	ACACGTACAGGACATATCCATTAAGG CTCATGCAGGCCATTGTG	ENSFCAT0000027884	0.951	-
CXCL11	Forward Reverse	ATTGTTCAAGGTTTCCCATGTT GCTTTCTCAATATCTGCCACTTTCA	ENSFCAT0000026314	1.032	-
ABL	Forward Reverse	TGTGGCGAGTGGTGATAAATACAC TCCACTCACCATTCTGGTTGTAA	ENSFCAT0000005306	0.985	[43]
B2M	Forward Reverse	CGCGTTTTGTGGTCTTGGT AAACCTGAACCTTGGAGAATGC	NM_001009876	0.953	[43]
RSP7	Forward Reverse	GTCCCAGAAGCCGCACTTT CACAAATCTCGCTCGGGAAAA	NM_001009832	0.991	[43]

Additional file 2