Gene	Oligo F	Oligo R	L	E	R2	Dr Zompo	GenBank
HSP90	CTCCATCTTGCTTCCCTCAG	TCAGTTTGGAGGAACCGAAC	146	2	0.9948	Pooc_contig353	GO349004.1
DNAJ	CACCTCCTGGAGCATCATCT	CGAAGGGCAAGCTCTACATC	243	2	0.996	Pooc_PC037B09	GO348410.1
HSP83	CGTGTGATCCAAAGCCTTCT	ACGATCCGGATGAACAACTC	248	1.9	0.999	__	KT159948
OzSP	AGATGCATCAGCCCATTTTC	GGAGGTACCACACCAACGAC	175	1.96	0.997	__	KT159949
DehSP	ATGAGAGTGGAGCTGCGTTT	AGCGACCCTTGCAGAACTTA	176	1.98	0.9973	__	KT159950
SHSP	ACCGGAGGATGTGAAGATTG	AGCTTGCTGGACAAGGTGAT	125	1.99	0.9832	__	KT159951
HSFA5	GCTCCAACAACTCCAGCTTC	CCCCTTCACAAACTCGTCAT	161	1.94	0.999	__	KT159952
LBP	ACCAGAGCTCGGTTTGAAGA	ATTCTGGTGCTTCCACCAAC	137	1.95	0.9957	__	KT159953
ABC	GCTGCTGGTTCTTCCTTGAA	AGACAATGCTGCCTCCTGTT	214	2	0.994	Pooc_PC025C02	GO346874.1
CYP	CAAAACTTGGCCTGTTGGTT	GCAGAAACCACACCCTTCAT	162	2	0.966	Pooc_PC004H08	GO346478.1
ALDH	TGATTCTGCATGCACCAGTT	ATGGGAACGAGCTACCATTG	219	2	0.99	Pooc_PC042G01	GO346135.1
CAT	CATCACATGCTGGGTTTCAC	ACCGATCCTGGACATCTGAC	178	2	0.9936	Pooc_B_c426	__
SODCP	CAATGGCTGCATATCGACTG	TGCCGGACTTTATCTTCTGG	157	2	0.9953	Pooc_PC034C12	GO347680.1
CSD1	GCTCCTGAGGATGAGATTCG	AGGCCAATAACACCACAAGC	236	1.96	0.9988	__	KT159954
FSD	TCATGACTTCTTTTGGGAATCA	CCCAACCAGATCCAAACAGT	153	2	0.9838	__	KT159955
MSD	GGCGGAGGTCATATAAACCA	ATAAGCAAGCCACACCCATC	192	1.86	0.9923	__	KT159956
GST	ACCTTCGCTAGCTTCCCTTC	AGATGGCAACAATGGGGATA	157	2	0.9963	Pooc_PC030G07	GO347013.1
GPX	TTGGAAAGGGGAAGTTGTTG	CGGGTCAAACCCTGAGATAA	232	2	0.9922	Pooc_PC006G02	GO346743.1
GSH-S	TAGGTTTGGCCAATTCTTGC	AAGGGGTGGTTCTTCCAGAT	213	2	0.992	__	KT159957
GR	AGTCCACACCAAATGGAAGC	AAGGGGAGGGAAGGGTTATT	247	2	0.9952	__	KT159958
AR	CTGCTGATTGGAGCTTAGCC	AGGGGATTTTCCTTGAAAGTG	102	1.93	0.9968	__	KT159959
APX	TCAGCTTGCTGGAGTTGTTG	CCCATGCGGTAAAAGATGTC	156	1.95	0.9997	Pooc_PC020F02	__
CAPX	GCATGATGCTGGAACGTATG	AATTTTGGGACCTCCAGCTT	228	2	0.9958	__	KT159960
Prx Q	AGGTGTCCAAGGGAGATGTG	TTGTGCGAAGCAGAATCATC	229	2	0.995	Pooc_Contig278	GO347736.1
GLP	GAAGTCGGGTTCATCACCAC	CTGCAGCTGACTGTTGAAGG	153	2	0.996	Pooc_B_c323	__
DSP5	TCTCAGGTCCGGCACTAATC	GAAAGGCTTGCTCGTATTGC	224	2	0.9992	__	KT159961
LPX	TGTAGCCACCGAAGGGATAC	GGTTGGTGGGATGAGGTAAA	172	2	0.998	Pooc_B_c10	KT159962
FtsH2	GCTCGAAGATACTCCGCAAC	GTCGGAGTGGTTCAGTTGGT	210	2	0.9973	__	KT159963
HMA	GTGATGAGAACCCCAATGCT	CGACACCAGAGCTCAAACAA	153	1.93	0.9993	__	KT159964
NRAMP1	TACTTCCTATGGGCGAGTGC	TTGCCTATGGAATCTGAACCTT	181	1.97	0.9916	__	KT159965
HMATPase	CACACCGAGCACATTGTACC	TGCTGCAGATGTAGGAATGG	181	1.98	0.9983	__	KT159966
HMATPase5	ATCGGACTCTCCTCGGTTTT	TGGCAGTGTTATGCTTCGTC	222	2	0.96	__	JF811734.1
MT3	CATGTCGACCTGTGACAACTG	TATTAATGGCCACAGGTGCAG	207	2	0.9954	__	KT159966
Fe-SP	CCATCTGATGAGGTAGTTTCAGG	CGCCAAGTCAACTTTGATAGC	188	1.9	0.9968	__	KT159967
MTP	CTCGTTTCCTGAGGTTCTGC	TTTGCTGCTGTCATGGCTAC	183	1.93	0.9989	__	KT159968