

**Table S1. Oligonucleotide primers used in PCR and RT-PCR reactions**

Gene Name or PGRPLC primer	Forward (F) or Reverse (R)	Sequence	Product size or location info
<i>PGRPS1</i>	F	GGACGAGCCGGCCAGGAATC	493 bp
	R	CAGCTTCGCGTACAGATAC	
<i>PGRPLD</i>	F	GACTCGGAGGATTGTCTGGA	281 bp
	R	CGGATCGACTCGGTGATAAA	
<i>PGRPLB</i>	F	CCATCCCGTACGTCATCATA	274 bp
	R	ATGTTCTTCGGTGGCAAATC	
<i>PGRPLA1</i>	F	CCGACATTCCAAGCAACTTT	280 bp
	R	ACCAGCCTAGCGTACAGCAT	
<i>PGRPLA2</i>	F	TGCTGATAACGCACATAGGC	124 bp
	R	TTGCTCGGTATGTCTTGCAG	
<i>PGRPS2&amp;3</i>	F	ACAACCTCCTGGTCGGTGAG	
<i>PGRPS2</i>	R	TCACCTGTCACAATGGTCGT	533 bp
<i>PGRPS3</i>	R	CCCCACATTAAGCTACGTTTC	423 bp
<i>PGRPLC1</i>	F	AAAGTTGGAGCCCACACCAAA	234 bp
	R	CAAAAGCCTCAGCTGTTC	
<i>PGRPLC2</i>	F	ACCATTCCCGGTTACAATTC	273 bp
	R	CTCGTGATCGGGGCGCTCGTC	
<i>PGRPLC3</i>	F	AAAGGGTTCAACGTGGACAGC	282 bp
	R	TCGTCGGTTTGTGGTGTTCGTT	
<i>PGRPLC</i> ; qRT-PCR primers	F	CCGGTTAATAACGTCATCATTGC	
	R	TCACCGTAGTTGGTGTGGTT	119 bp
	R	CGTCGTAGTTTTTGCCATCC	118 bp
	R	TCGGGACTCGAATGAAACTC	107 bp
<i>S7</i>	F	GTGCGCGAGTTGGAGAAGA	78 bp
	R	ATCGGTTTGGGCAGAATGC	
<i>GFP</i>	F	TAATACGACTCACTATAGGGCAAGACACGTG CTGAAGTCAA	~450 bp

	R	TAATACGACTCACTATAGGGGCCTGAATTTA ACCAGGAACC	
<i>BSF340/19 &amp; BSF806/26</i>	F	TCCTACGGGAGGCAGCAGT	
	R	GGACTACCAGGGTATCTAATCCTGTT	
<i>CECI</i>	F	ACCAACCAACCACCAAACAAC	168 bp
	R	CTTCTCTGCTGCCTTGAACACT	
<i>DEF1</i>	F	GCCTTTGTGCCGCTCACT	61bp
	R	GCCTTACTGTTGCAGTAACCACC	
1	F	AGGAACCTACAGCTAGAAAATG	2 <sup>nd</sup> exon
2	F	GTTAAGGTTGCCGTGGAAGC	2 <sup>nd</sup> exon
3	F	CAACTCGTCGGACATCACGTT	2 <sup>nd</sup> exon
4	R	ACCTGCCCTTGATGTACGTT	2 <sup>nd</sup> exon
5	R	ATCCTGGTAAATGTTCTTTATCAC	2 <sup>nd</sup> exon
6	F	CACGCACCTGGCAATCTAGTCT	3 <sup>rd</sup> exon
7	F	AACCCTTGATCAGCTTTATCG	3 <sup>rd</sup> exon
8	R	TGGCACACAGGACAATCATCA	3 <sup>rd</sup> exon
9	R	GGATCGATATGACCGCTACAA	3 <sup>rd</sup> exon
10	F	CCAGATCCGAGACCGTTACGG	4 <sup>th</sup> exon
11	F	AGACCGTTACGGTTAGTGACG	4 <sup>th</sup> exon
12	F	CCGGTTAATAACGTCATCATTGC	4 <sup>th</sup> exon
13	R	GTGGCAGTGTGAGCAATGA	4 <sup>th</sup> exon
14	R	ACGGCATGCGGCCTGAG	4 <sup>th</sup> /5 <sup>th</sup> exon
15	R	TGCTTTGCATGCTTGCTGA	4 <sup>th</sup> /5 <sup>th</sup> exon
16	R	TACCTGATACATGCATTTTCGTCTGA	4 <sup>th</sup> /5 <sup>th</sup> exon
17	F	CACAAAACACTTCTGACCTCTCC	4 <sup>th</sup> intron
18	F	TCTGACCTCTCCCATGCAG	4 <sup>th</sup> intron
19	F	AAACCACACCAACTACGGTGA	5 <sup>th</sup> exon
20	R	TCACCGTAGTTGGTGTGGTT	5 <sup>th</sup> exon
21	F	ACAACTTTTGATCGGTGGTG	5 <sup>th</sup> exon
22	F	AAAGTTGGAGCCCACACCAAA	5 <sup>th</sup> exon

23	F	GCTTGCCAAGGAGTACAAGC	6 <sup>th</sup> exon
24	R	CAAAAGCCTCAGCTGTTC	6 <sup>th</sup> exon
25	R	AATGTGGCAAAGCCTCAG	6 <sup>th</sup> exon
26	R	AAATGTGGCAAAGCCTCAG	6 <sup>th</sup> exon
27	R	AAAATGTGGCAAAGCCTCA	6 <sup>th</sup> exon
28	F	GGGTGCGCTTAATTCAGGA	7 <sup>th</sup> exon
29	F	ATGGATGGCAAAAACCTACGAC	7 <sup>th</sup> exon
30	R	CGTCGTAGTTTTTGCCATCC	7 <sup>th</sup> exon
31	F	ACCATTCCCGGTTACAATTC	7 <sup>th</sup> /8 <sup>th</sup> exon
32	R	CTCCGAGGCATATATTCGGTA	8 <sup>th</sup> exon
33	R	CAGTGCTTCGATTAGCCACTT	8 <sup>th</sup> exon
34	R	CAAGCTGTGCAGTGCTTCGAT	8 <sup>th</sup> exon
35	R	AAACTGGGGCAAGCTGTGCAG	8 <sup>th</sup> exon
36	R	CTCGTGATCGGGGCGCTCGTC	8 <sup>th</sup> exon
37	R	TCGGGACTCGAATGAAACTC	9 <sup>th</sup> exon
38	F	TTTAGCGACATTGCGTATCA	9 <sup>th</sup> exon
39	F	AAAGGGTTCAACGTGGACAGC	9 <sup>th</sup> /10 <sup>th</sup> exon
40	R	CCAGTGGGGCCAAGTTTTGATG	10 <sup>th</sup> exon
41	R	TCGTCGGTTTTGTGGTGTGCGTT	10 <sup>th</sup> exon
42	R	GGAAGTCATCAAGGACACTTGG	3' UTR
43	R	GGTTCTCTTTTTCCGGTTCA	3' UTR