

Table S2. Average Ct value, standard deviation (SD) and coefficient of variation (CV) of candidate internal reference genes in all lingonberry samples.

Gene	Mean Ct	Standard deviation	Coefficient of variation
<i>18S rRNA CL5051.Contig1</i>	29.98	1.37	4.48
<i>Actin CL1167.Contig3</i>	28.68	1.44	5.04
<i>Actin CL2126.Contig2</i>	29.04	1.41	4.90
<i>Actin CL2172.Contig2</i>	23.29	0.97	4.22
<i>Actin CL2172.Contig3</i>	25.19	1.40	5.46
<i>Actin CL3559.Contig7</i>	30.64	1.10	3.66
<i>Actin CL494.Contig13</i>	27.67	1.26	4.59
<i>Actin CL5740.Contig1</i>	28.49	1.49	5.32
<i>Actin CL5740.Contig2</i>	30.34	1.37	4.58
<i>Actin CL5740.Contig5</i>	28.46	1.62	5.75
<i>Actin CL7856.Contig2</i>	32.03	1.33	4.26
<i>Actin Unigene12465</i>	26.96	0.89	3.28
<i>Actin Unigene20323</i>	31.73	1.08	3.43
<i>Actin Unigene23839</i>	30.79	1.17	3.87
<i>Actin Unigene6171</i>	31.54	1.22	3.96
<i>Chy Unigene26262</i>	31.76	1.42	4.56
<i>Tub CL1466.Contig3</i>	29.95	1.42	4.77
<i>Tub CL1466.Contig7</i>	32.29	1.06	3.31
<i>Tub CL3192.Contig5</i>	29.57	1.68	5.76
<i>Tub CL7489.Contig2</i>	29.55	2.12	7.36
<i>Tub Unigene3128</i>	29.77	1.28	4.32