

TABLE S1. Primers used for amplification and sequencing of the *SQLE* and *TERG* in this study

Forward primer	Forward primer sequence	Position	Reverse primer	Reverse primer sequence	Position
SQEL397S	5-GTTGACTGGTGGCGGTATG	1002–1020 [#]	SQEL397R	5'-GCTACGGAGTAAAAATGCCG	1315–1334 [#]
TrTERG-1	5'-ATGGTTGTAGAGGCTCCTCC	1–20*	TrTERG-1-2R	5'-GAGCTTCAGAGGTGAGGCCG	456–475*
TrTERG-2S	5'-GAACGGCAAGCTCAAGGATG	327–346*	TrTERG-2R	5'-CGGTCGCGCTTGCGGTTGTG	727–746*
TrTERG-3S	5'-CTACTCCGATCTCGACATGG	663–6825*	TrTERG-3R	5'-GGTCGAGTTTCTGAAGAGC	1054–1072*
TrTERG-4S	5'-CCAGATATTCTCGAGGAGC	979–997*	TrTERG-4R	5'-TTCACCGATACATCTGTGTC	1385–1404*
TrTERG-5S	5-GTGAAGAAAAGACAGATTAC	1307–1326*	TrTERG-5R	5'-TCATTTCTGAGAAGCCTGCT	1565–1584*

[#]Positions reflect those of the *T. rubrum SQLE* gene (GenBank accession no. XM_003233797).

*Positions reflect those of the *Trichophyton rubrum* CBS 118892 cytochrome P450 51 (TERG_01703) mRNA gene (GenBank accession no. XM_003236932.1).