

Alternative splicing occurs in the efflux pump transcripts in response to antifungal drugs

Supplementary Materials

Table S1. Primers used in RT-qPCR assays.

ID	Primers	Amplicon	Concentration (nM)	Reference
TERG_04309 (ABC transporter) Total expression	F: GGGTATCGTGTTGAAGCTAAGA R: CATCCAGAAAGACAGGCCATAA	114 bp	100 nM	This work
TERG_04309 (ABC transporter) (Intron-3)	F: ACACGGATTAATCGGAACACC R: AGCACAGGTATACTCACGTTTG	101 bp	100 nM	This work
TERG_04309 (ABC transporter) (Intron-4)	F: ACTCGATTAAAGAGCGAAGGA R: CCGCATTCTTGATAGGAGAGA	106 bp	300 nM	This work
TERG_04224 (ABC transporter) Total expression	F: CAACTGCTCTCCGTTCTT R: CTTCCGTCGATAGAGAATCG	125 bp	100 nM	This work
TERG_03468 (Ste14p O-methyltransferase) Total expression	F: CAACTGCTCTCCGTTCTT R: CTTCCGTCGATAGAGAATCG	100 bp	400 nM	This work
TERG_00588 (Ste2 – Pheromone Receptor) Total expression	F: TCGAGCTATCCCTACTCATC R: AATTGGAACCGCTGCTAC	107 bp	100 nM	This work
TERG_04402 (<i>gapdh</i>)	F: GCGTGACCCAGCGATGTAGT R: CGGTGGACTCGACGATGTAGT	62 bp	200 nM	[24]
TERG_05742 (<i>rpb2</i>)	F: TGCAGGAGGTTTGATGAAGA R: GCTGGGAGGTACTGTTTGATCAA	59 bp	300 nM	[24]

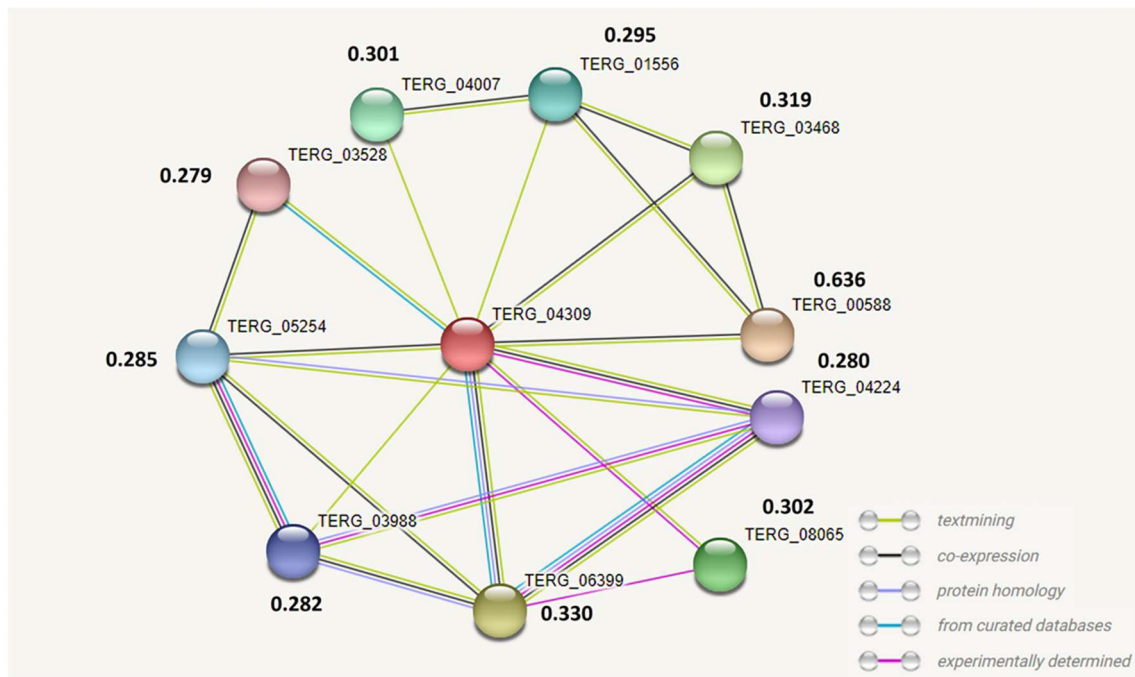


Figure S1. Interaction network of the *T. rubrum* TERG_04309 gene (Transporter ABC) with other genes. Next to each gene, the interaction score obtained is indicated (in bold). The colors of the interaction lines refer to the putative forms of that interaction (STRING software).