**Table S1** RT-PCRprimers for cellulase- and chitinase-like genes. GAPDH served as the internal control. Gene IDs of predicted transcripts of target genes of *Acropora digitifera* (v1.0) are enclosed within parentheses.

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| Target gene | Forward Primer (5'-3') | Reverse primer (5'-3') | Product size |
| Cellulase-like-1 (adi\_v1.03986) | CCCTGTCGCTGTTCTGAATC | GGTCGAGCCATCGTCATATT | 831 bp |
| Chitinase-like-2 (adi\_v1.09189) | CTTGTATCTTCTTGTCAGCAAGC | CGATATATCTTGATCCCGTGATAG | 1400 bp |
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