



Figure S3 - Semi-quantitative RT-PCR analysis of *CcGolSI* using cDNA from *C. canephora* clones D^T (clone 14) and D^S (clone 109A). The water deficit treatments were: non-stressed (NS; \pm -1.35 MPa), moderate stress (MS; \pm -2.35 MPa), severe stress (SS; \pm -4.3 MPa) and recovery (REC; 72 h following re-irrigation after severe stress); see Materials and Methods for details. The *GAPDH* gene was used as an internal control to normalize the expression level of the target gene (*CcGolSI*) between different treatments, as described by Cruz *et al.* (2009).