



Figure S4. Expression analysis of VviSOC1a in three independently transformed T1 tomato lines (cultivar ‘Ailsa Craig’) using real-time qPCR analysis. Bars represent the mean and standard error of three biological replicates (n = 3), where (**) and (***) indicates moderate ($P \leq 0.05$) and high ($P \leq 0.01$) significant differences compared to wildtype, respectively.