



Figure S2. Gene expression data of *VviSOC1a* over 54 different grapevine (cultivar ‘Corvina’) tissue types and developmental stages according to the grapevine expression atlas [3] and visualised on the Vitviz website. Means and standard errors represent three biological replicates (n = 3). Yellow boxes highlight tissue types displaying predominant expression.

References

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3. Fasoli M, Dal Santo S, Zenoni S, Tornielli GB, Farina L, et al. 2012. The grapevine expression atlas reveals a deep transcriptome shift driving the entire plant into a maturation program. *The Plant Cell* 24:3489-3505.