

Supplemental Fig. S6 Remodeling of potential grapevine ribosome in response to ABA application in the deacclimation experiment. Presented ribosome was remodeled in Chimera based on cryo-electron microscopy structure of Arabidopsis thaliana mitochondria ribosome. Only homologous proteins shared between Arabidopsis and Vitis are shown. A-B) Potential ribosome in control samples under permissive growing condition (22 °C) for budburst; C-D) Potential ribosome in ABA-treated samples under same condition. At least one gene encoding each blue-highlighted protein exhibited significantly lower transcript abundance in response to ABA application. Abbreviations: SSU, small subunit; LSU, large subunit.