



Supplemental Figure S2. Expression of Complex II Genes during Pollen Development. Data used to generate the digital Northern were obtained from NASC microarrays (Honys and Twell, 2004) at the Genevestigator web site (<https://www.genevestigator.ethz.ch>). Mean signal intensities were averaged from two technical replicates, except for MPG. UNM, uninucleate microspores; BCP, bicellular pollen grains; TCP, tricellular pollen grains; MPG, mature pollen grains.