

1 aagataatcaagaaaacaggtatagcgtttcttctacgtcagatatcaaatccgtgatcgagtgaacaatgtcaatgacaatattctacaag
2 aagaaatgtttcattcatatttcttggaatgtatcgattgatcatttgtttctcctgttttgtttctatattgttaacactacaccaatgcttctgct
3 gctttggatgacgagatagtttttttactcgacacatcatacaagacctcgtgaagtatctgtaatacacgaatcattaatacacaacagaata
4 atgcattactatagttttgagaatctcattctcatactctgtctctcttgggtcgaagaatcacacttaacataaaaaagaacaacatgctcagt
5 acacaaaggtaaagcccaccaaagagagaaaagaaaacaagacttaactataacaataaatttaatgctcgtctcaaaactcatttttcatat
6 gaatttttaacacaaacaattttcCTGGaaatctcCCGccttgatctcttataactcatctctatactccgaacatgcccctcctcgta
7 accttaacggccATG

8
9 **Supplemental Figure S2.** Location of the AW-box in *ROD1* proximal upstream region.
10 ATG, transcription start site; red, 5'-UTR; black, promoter. The AW-box motifs are highlighted
11 in blue.