



Figure S3. The distribution of small RNAs on the sno-miR775 and (b) sno-miR779. Small RNA sequences were adopted from following samples: GSM575246 (WT), GSM154361 (*dcl1-7*), GSM1533542 (*dcl3*) GSM154364 (*dcl4-2*), GSM893124 (*rdr2*) and GSM575247 (*rdr6*). The visualization of reads mapping was carried out using Golden Helix GenomeBrowse® visualization tool v2.1.0. The light dash lines above the solid peaks in the plots indicate multi-mapped alignments.