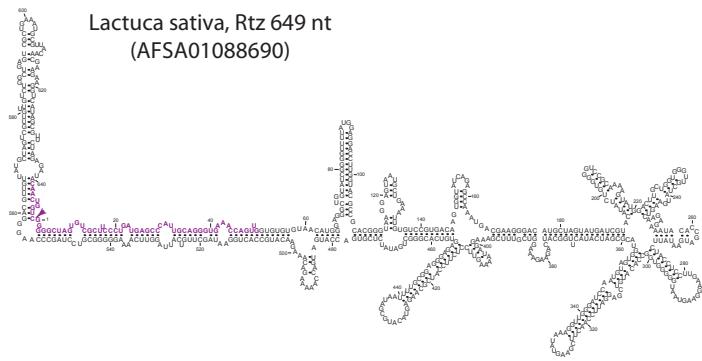
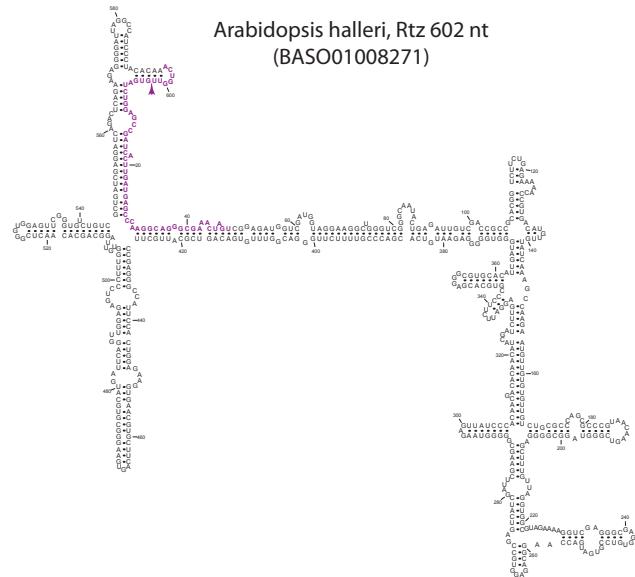


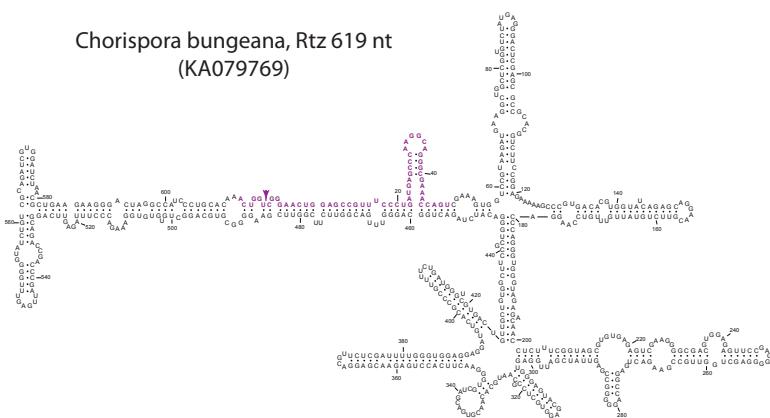
Lactuca sativa, Rtz 649 nt
(AFSA01088690)



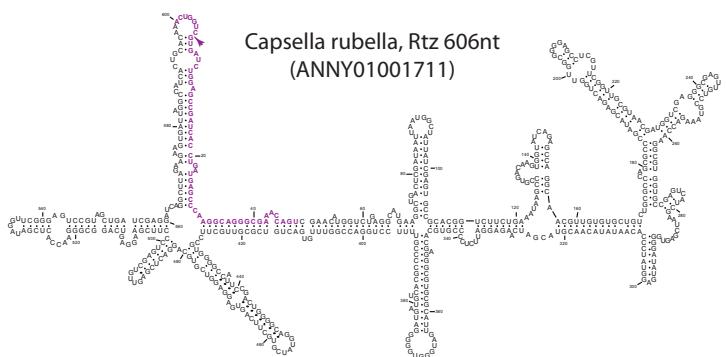
Arabidopsis halleri, Rtz 602 nt
(BASO01008271)



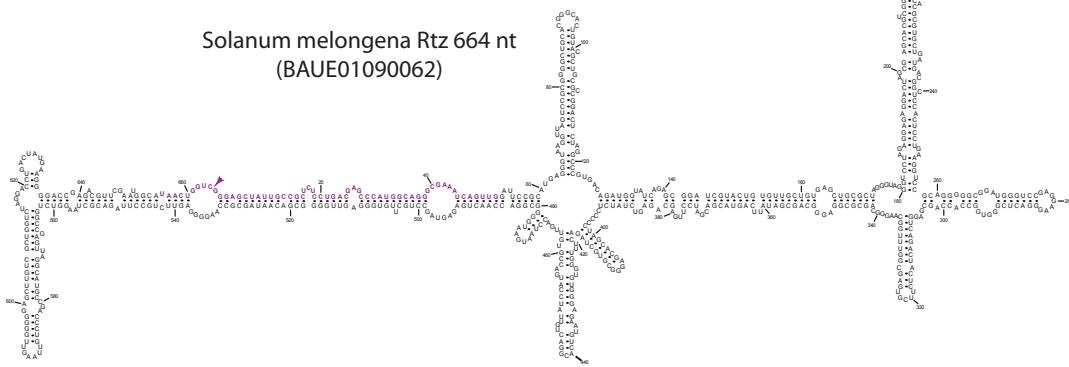
Chorispora bungeana, Rtz 619 nt
(KA079769)



Capsella rubella, Rtz 606nt
(ANNY01001711)



Solanum melongena Rtz 664 nt
(BAUE01090062)



Additional file 10

Additional file 10.

Minimum free energy secondary structure prediction of five plant circRNA retrozymes deduced from the corresponding genomic retrozyme sequences (GenBank entries are indicated below the species names). HHR sequences are shown in purple letters and the self-cleavage site is indicated with an arrow. Numbering starts at the self-cleavage site of the HHR.