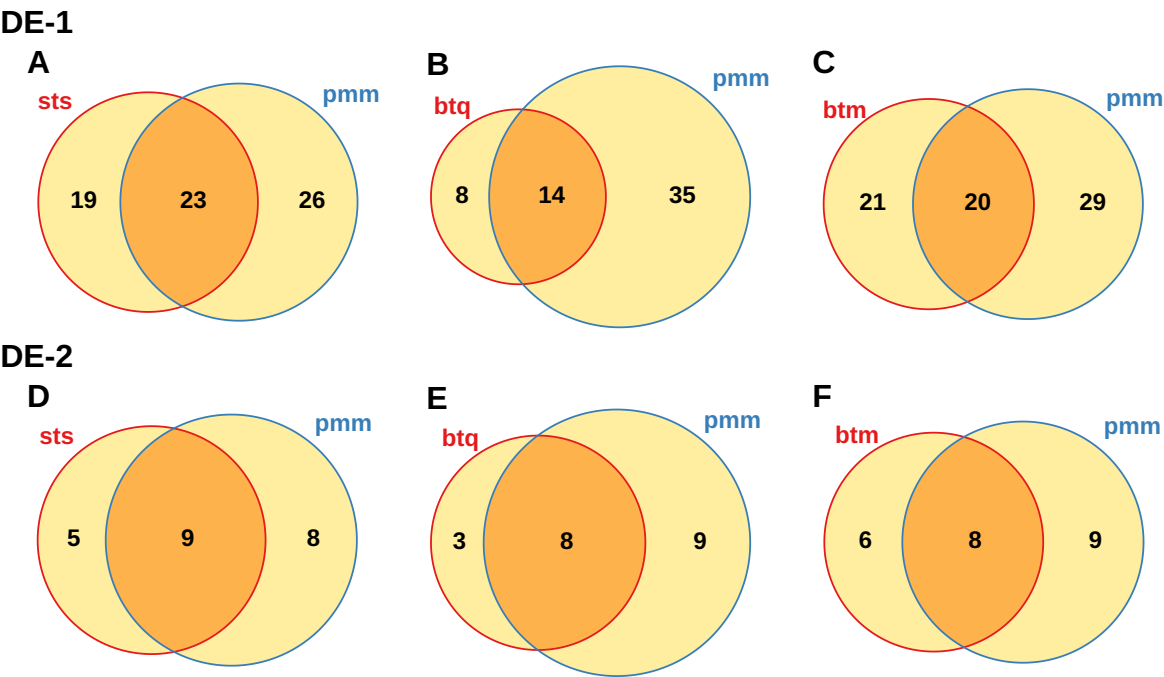


Figure S1. Venn diagrams of the differential sets in DE-1 and DE-2.



Panel 'DE-1' is showing in A to C the overlap between the three different alignment and quantification approaches and their significant microRNA candidates for the first differential analysis. The nomenclature is already described in Table 2 in the right column.

In panel 'DE-2' the same is done in D to E for the second differential analysis and again on microRNAs.

The two differential analyses differentiate by absolute count number and a more pronounced higher set number for the premature microRNAs (pmm).

Acronyms sts, btq, btm, pmm explained in supplementary Table S1.