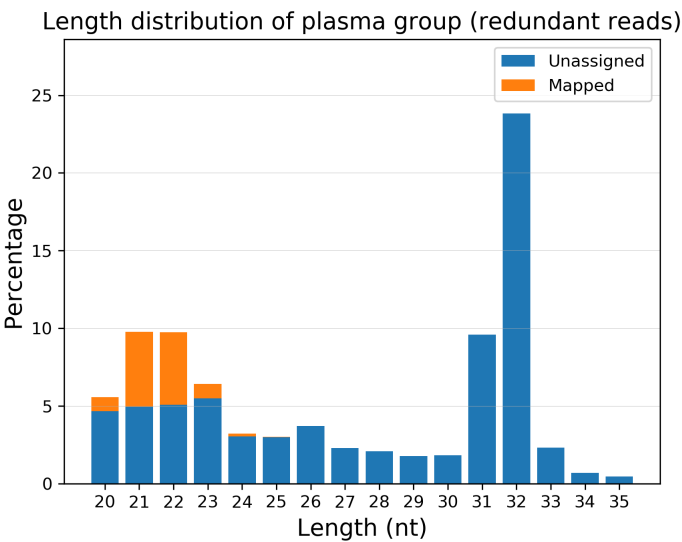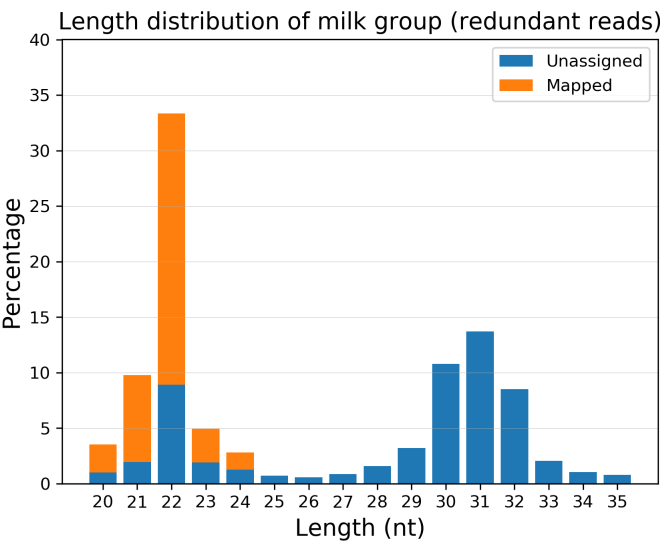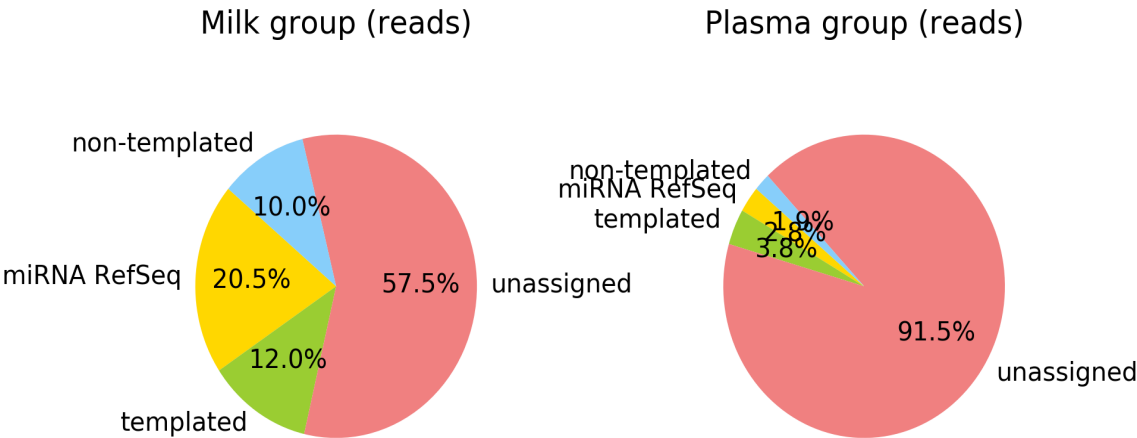# IsomiR profile report

sRNA length distribution

Mapped and unmapped reads to custom precursor arm reference DB (5p and 3p arms) in milk (left) and plasma (right) groups
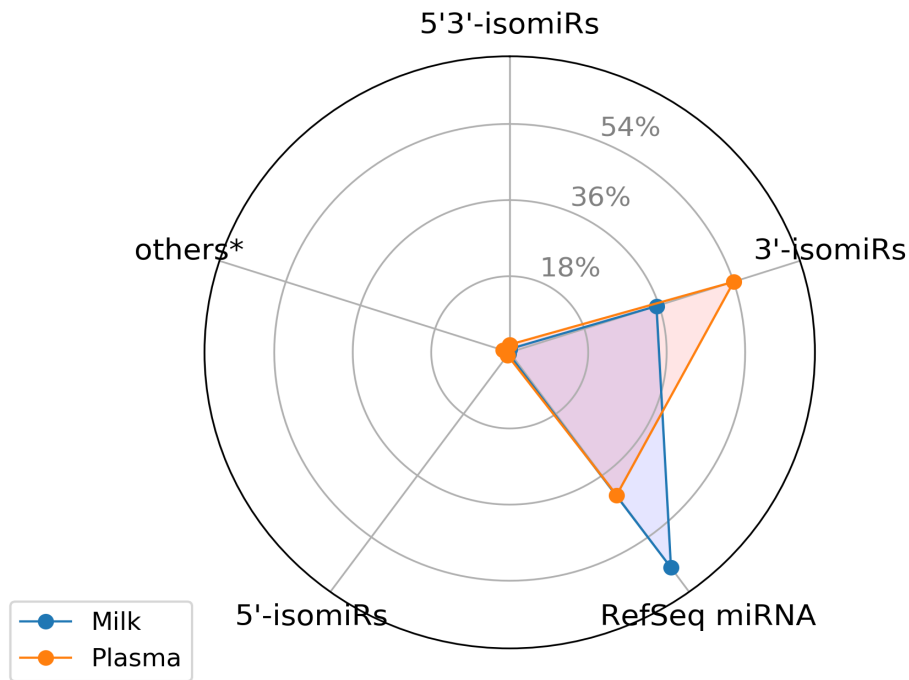
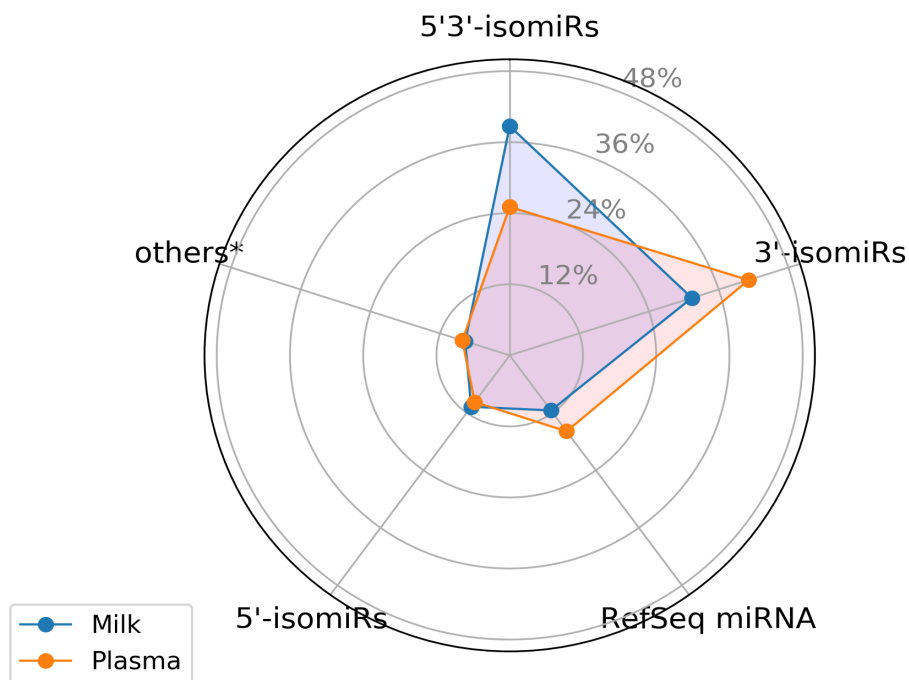## Templated and non-templated isomiRs

RefSeq miRNAs, templated isomiRs, non-templated isomiRs and unassigned sequences as percentage of total sRNA reads in milk (left) and plasma (right) groups

Templated isomiR subtypes

Templated isomiR profile (redundant reads)

Templated isomiR profile (non-redundant reads)

* IsomiRs potentially generated from multiple loci

Non-templated isomiRs

3'-end additions to RefSeq miRNAs and templated isomiRs