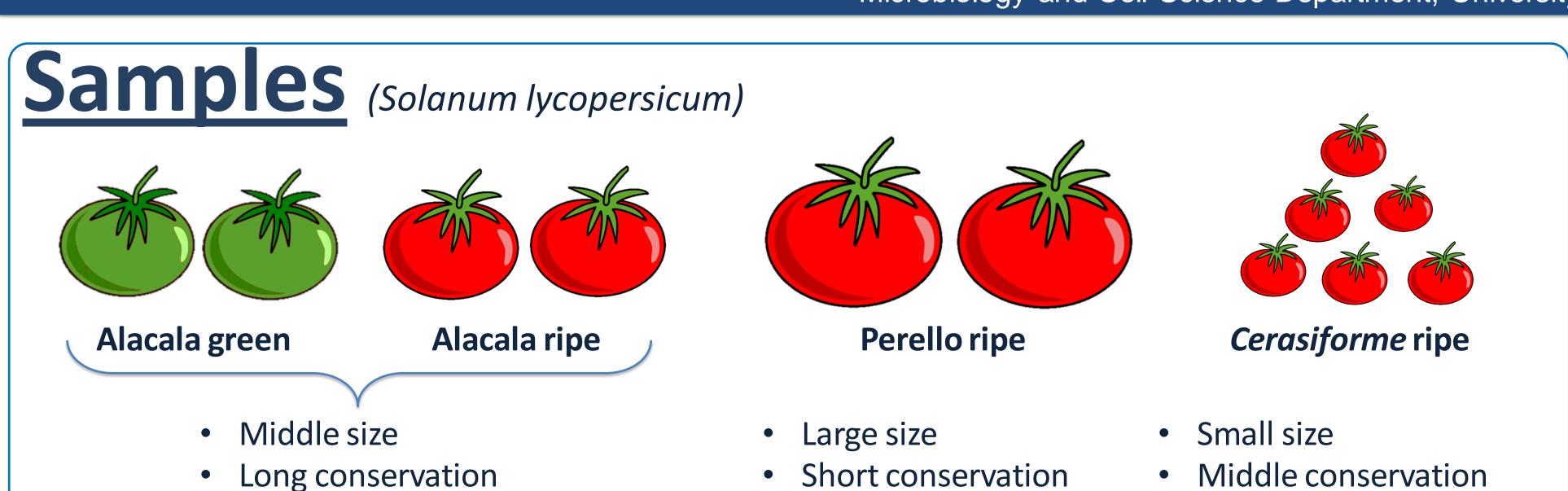


Understanding the functional role of alternative splicing in tomato ripening

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2 tomatoes



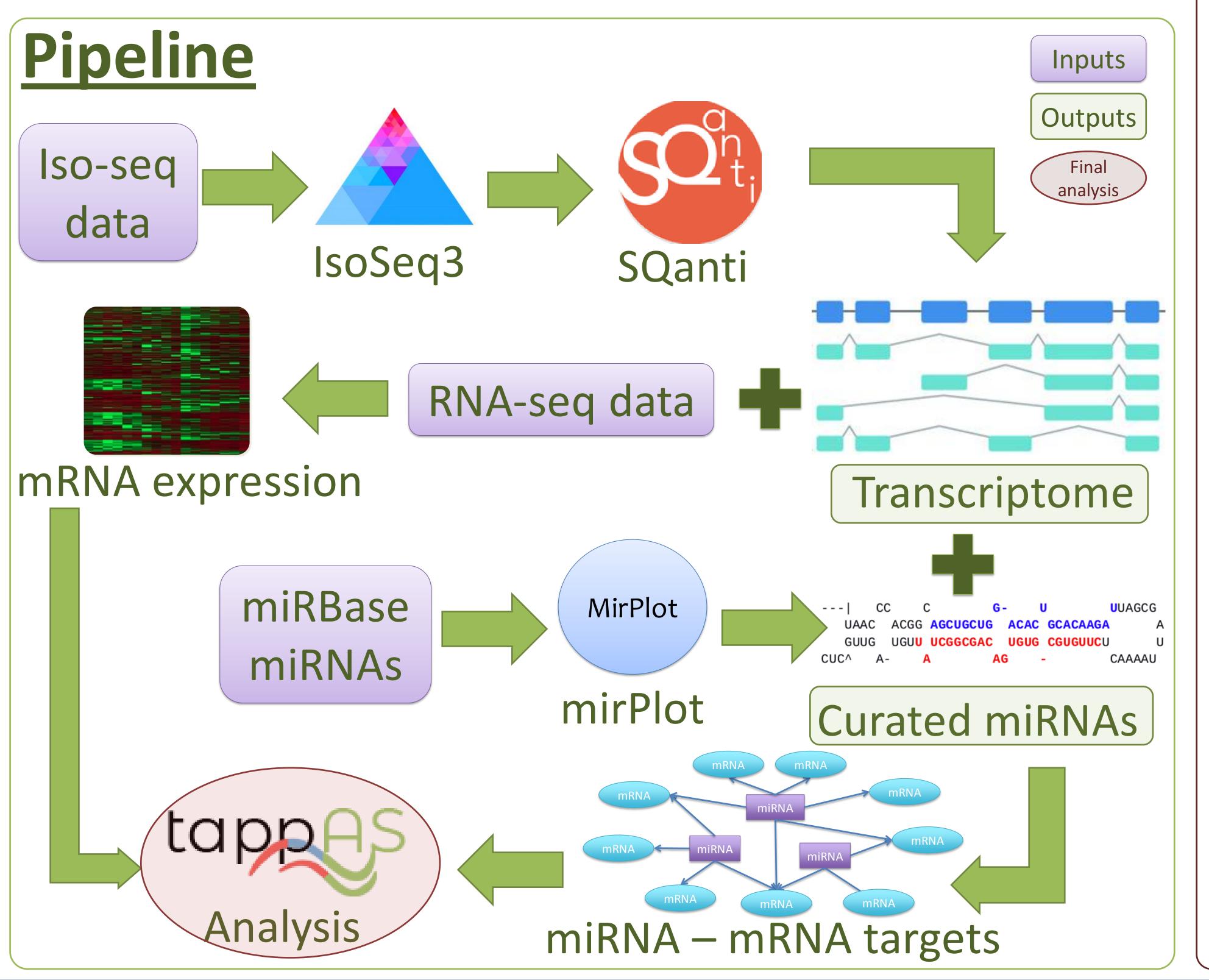
2 tomatoes for condition

- Iso-seq data.
- Whole length transcripts.
- One library for condition.

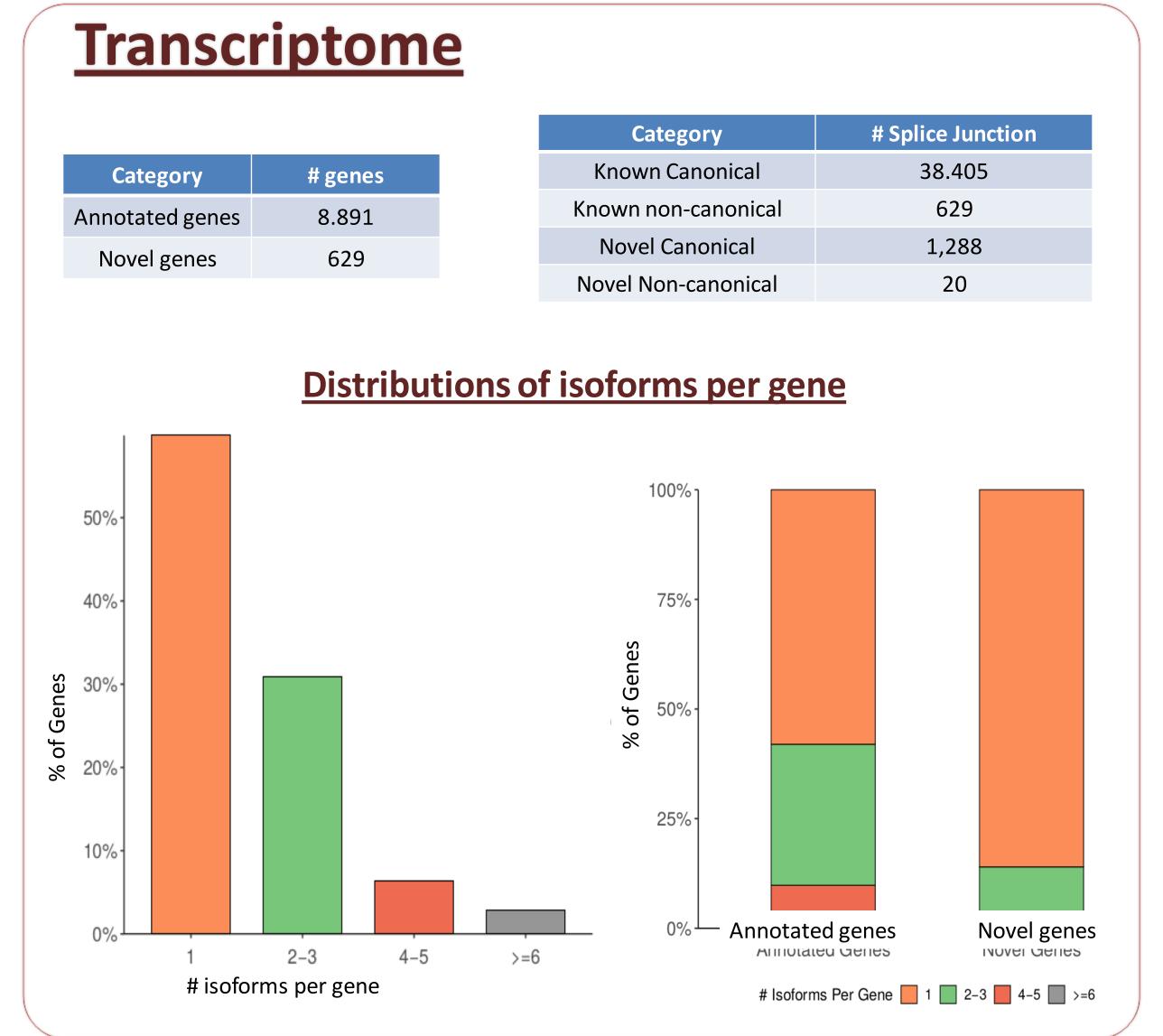
Ilumina®

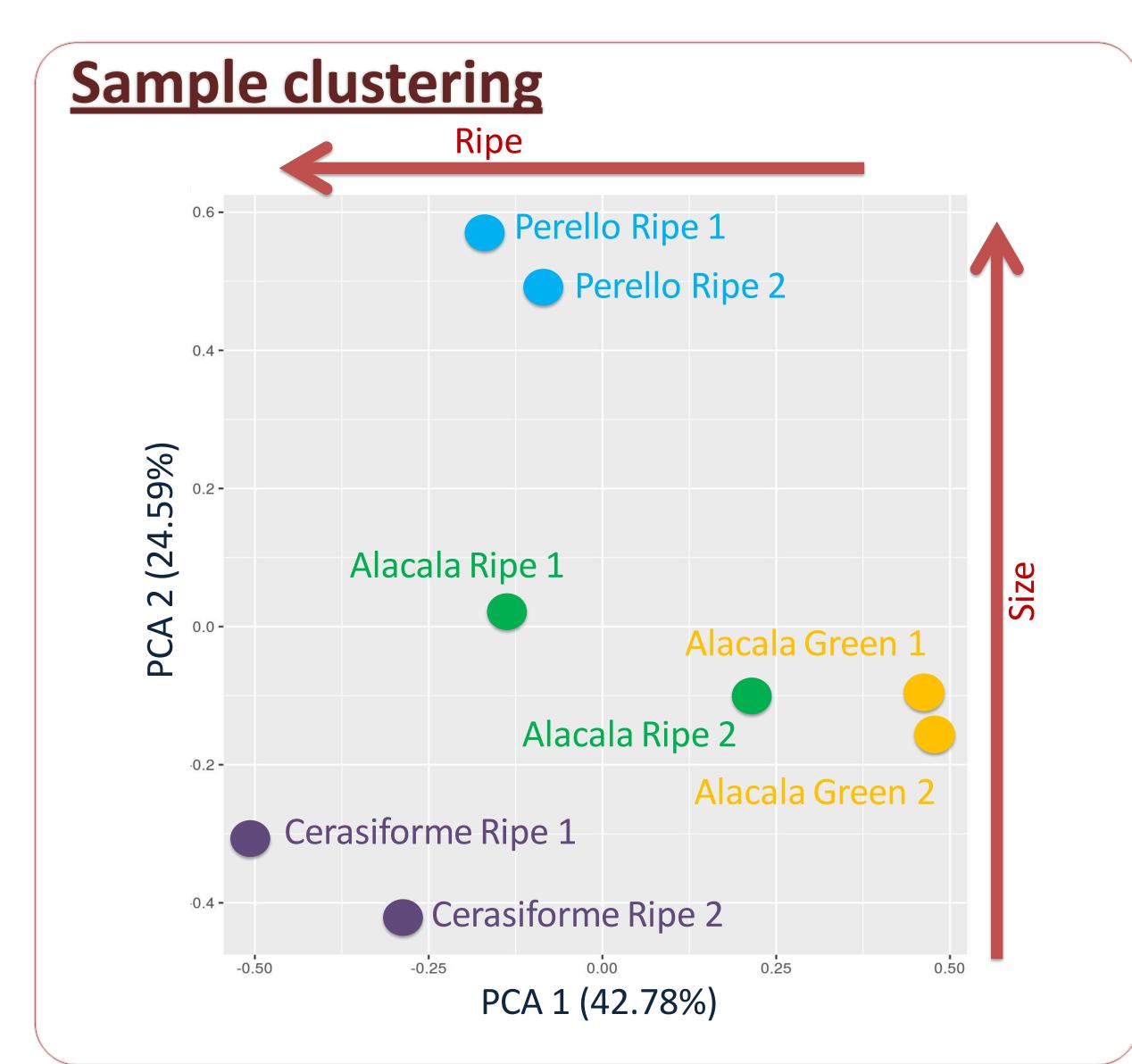
2 pools of 3 tomatoes

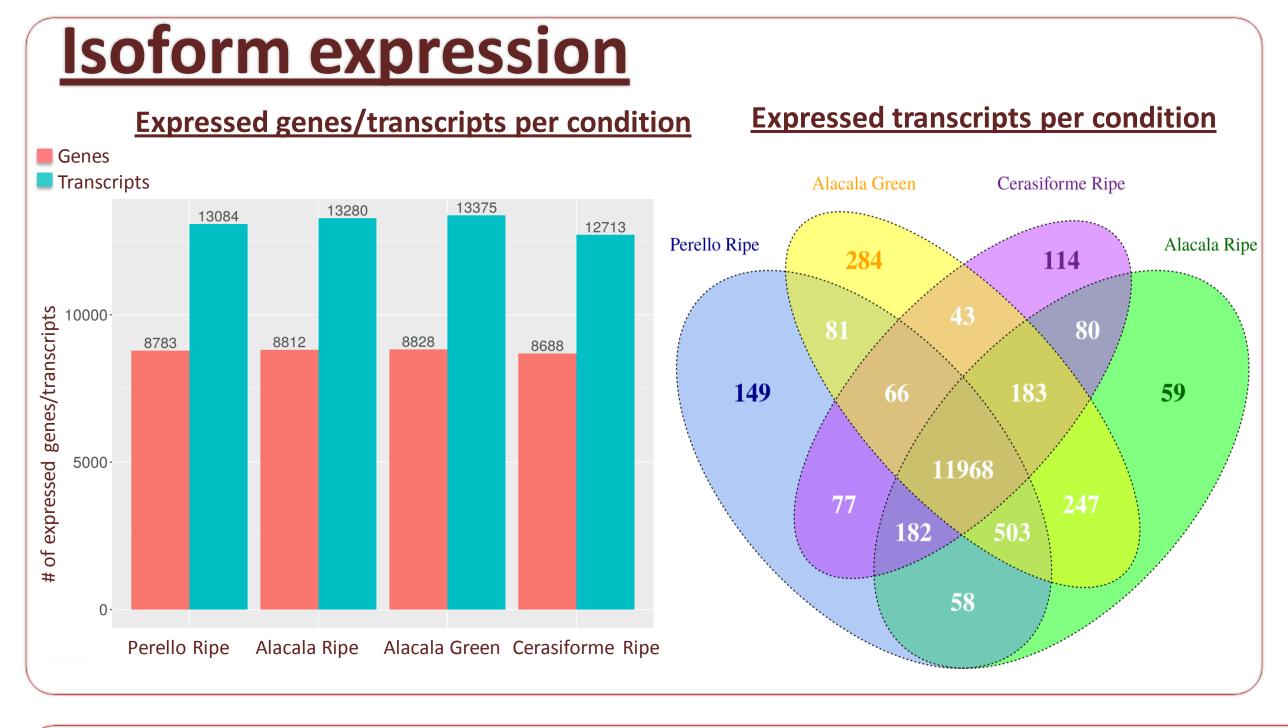
- RNA-seq data.
- Paired-end reads of 150 nts.
- One library for sample.

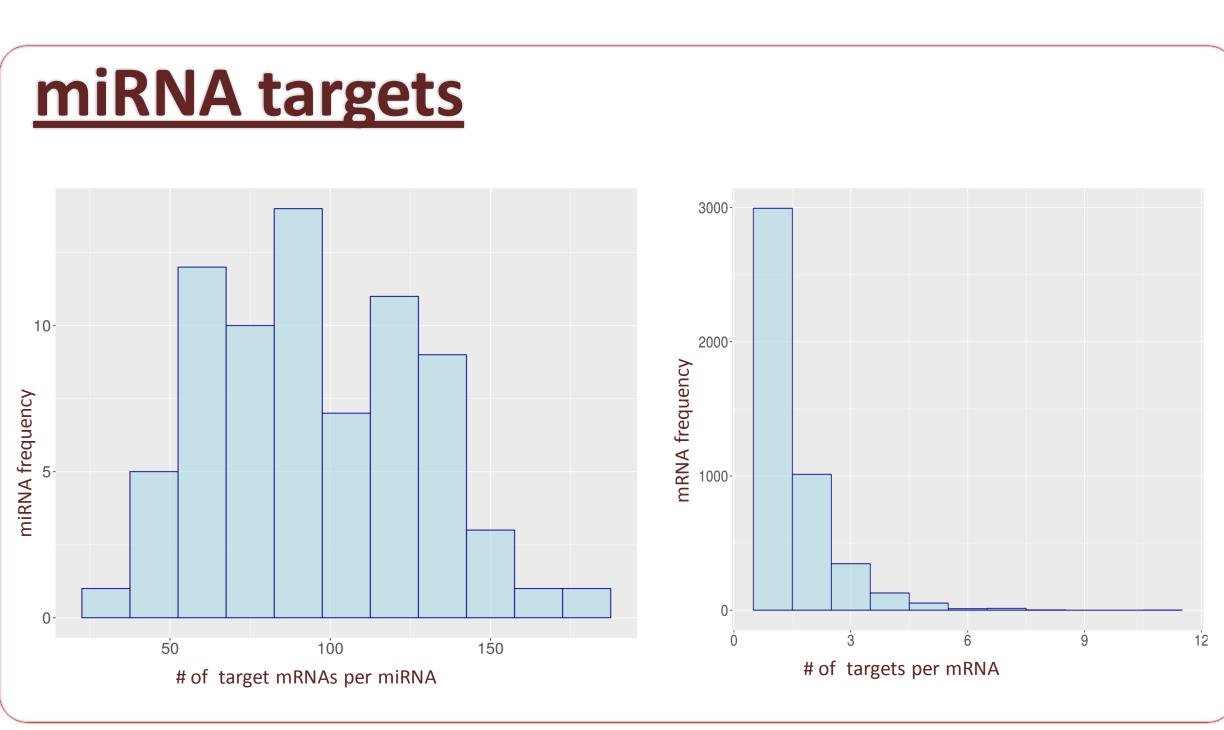


Results











Software references:

IsoSeq3 – PacBio©

