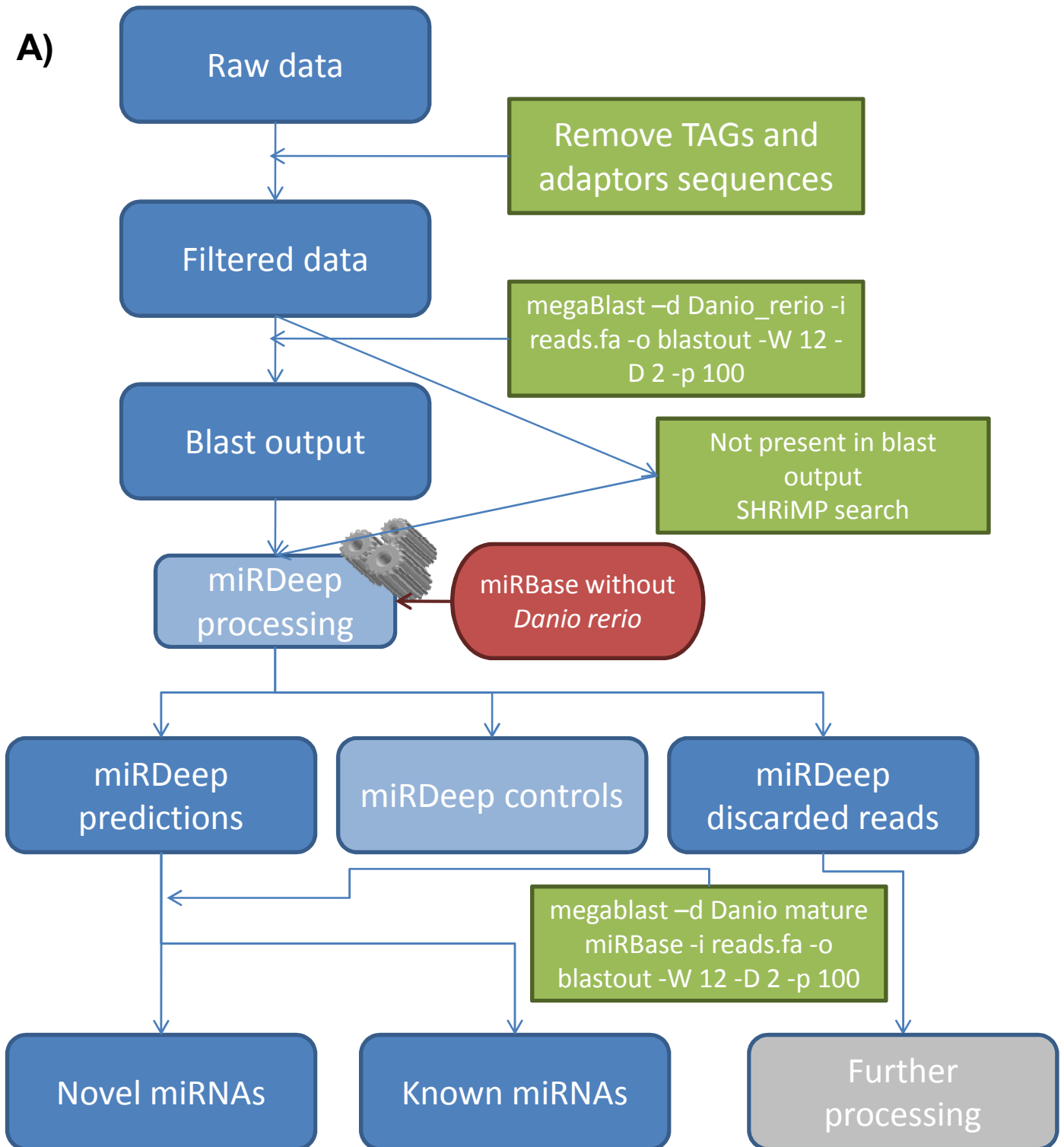


Additional File 1, Figure A1 – Data pipeline analysis.

Flowchart describing the procedure used for identify microRNAs in pyrosequencing data. **(A)** includes miRDeep analysis and **(B)** further processing of miRDeep discarded reads. The red boxes stand for data coming from databases. -i and -d stands for query file and database respectively.



B)

