

Total Extracellular Small RNA Profiles from Plasma, Saliva, and Urine of Healthy Subjects

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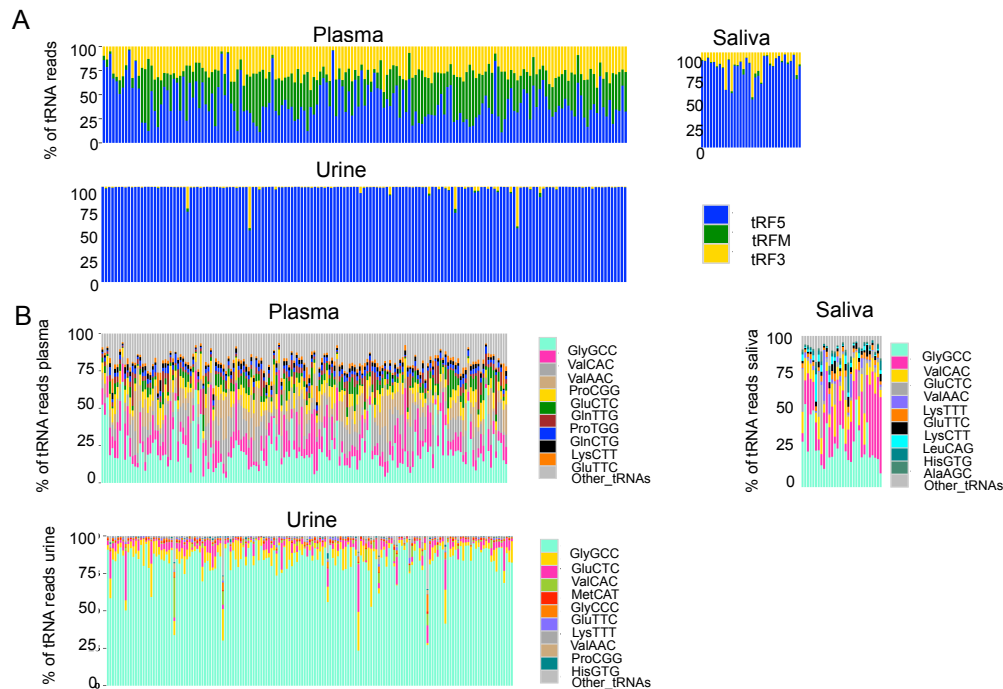
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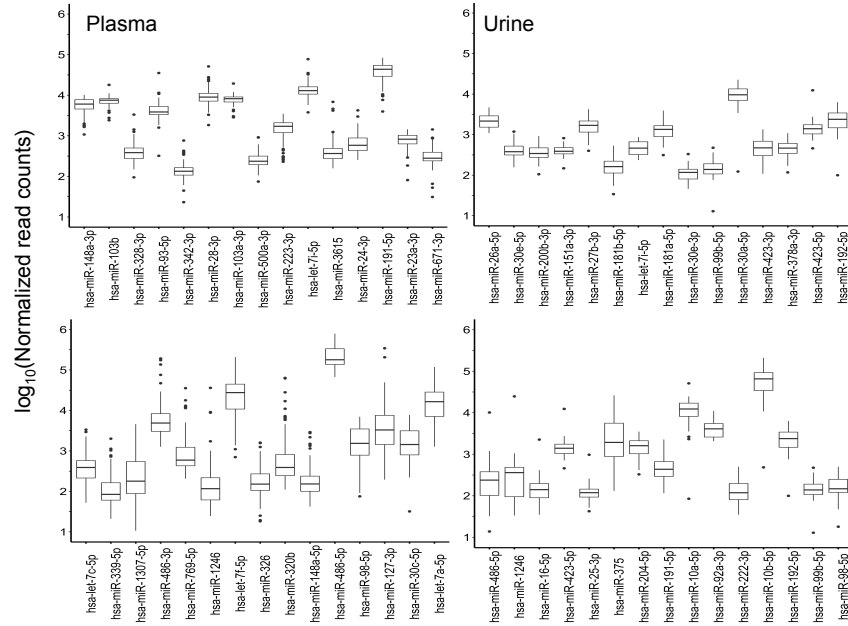
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Supplemental Figure S1

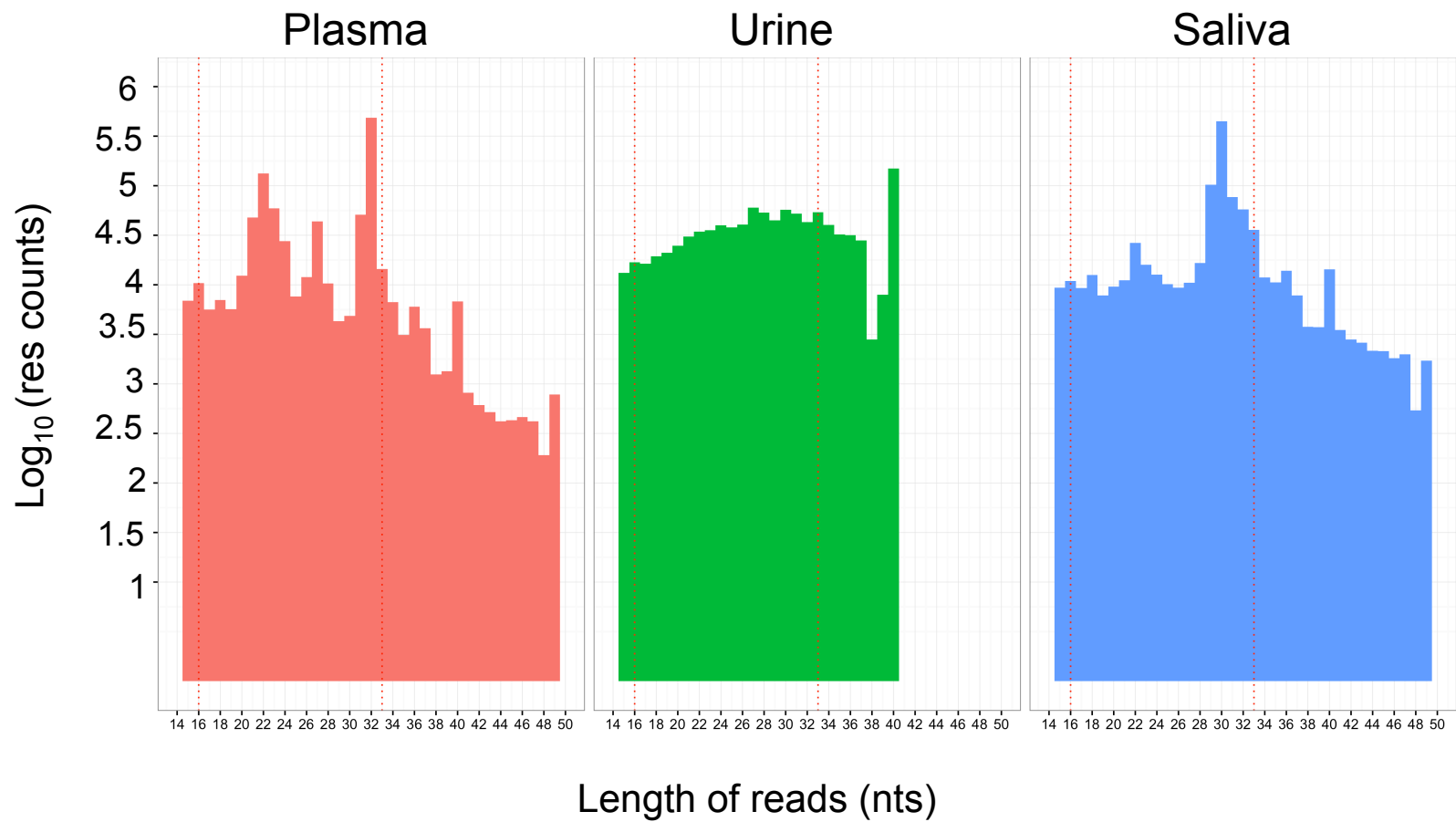


Supplementary Figure S1. Detailed analysis of the detectable tRNA fragments. Of the reads that map to tRNA fragments, (A) shows the number that aligns to the 5', 3', or middle of tRNA (tRF5, tRF3, or tRFM) for each biofluid. The fragments aligning to different detected tRNAs are shown in B.

Supplemental Figure S2



Supplementary Figure S2. CVs for miRNAs from individuals with more than 5 samples sequenced over ~70 weeks. (A) displays the 15 miRNAs with the lowest CVs, and (B) displays the 15 miRNAs with the highest CVs).



Supplementary Figure S3. Read length distribution for plasma, urine and saliva samples. The dotted red line represents the range of RNAs discussed in the manuscript. miRNAs typically have a length of 16-23 nts and the tRNA and yRNA fragments discussed in the manuscript are between 30-33 nts long.