MicroRNAs (miRNAs) and their isoforms/isomiRs are of increasing interest as potential biomarkers. The ability to accurately quantify miRNAs from biofluids is crucial to realizing their potential in early disease detection, diagnosis, drug development, and selection of follow-up treatment. SomaGenics’ miR-Direct® method overcomes commonly encountered problems of inconsistent RNA recovery from samples, real-time qPCR impacted by sample-born inhibitors, and by low miRNA abundance in blood.

**Technology overview**

- No solvent extraction or column purification
- Front-end processing is performed in a single tube
- miRNA concentration without total RNA purification
- miRNA capture from variable input volumes (25 to 400 µl)
- All steps until qPCR can be multiplexed for all miRNAs of interest

In the miR-Direct workflow, miRNAs are released from plasma/serum samples, captured, washed, and then assayed by quantitative RT-qPCR using SomaGenics’ circularization-based miR-ID® assays (included in miR-Direct® kits.)
miR-Direct® eliminates qPCR inhibitors

Plasma from one donor was collected in tubes containing EDTA (blue) or heparin (red). miR-Direct® allows for miRNA quantification in heparin-containing plasma, while a conventional assay using isolated total RNA from columns fails. Heparin, a widely-used anticoagulant for blood plasma collection, is a known inhibitor of qPCR enzymes. Because heparin is a highly negatively charged molecule, it is often co-purified in column-based purification kits.

miR-Direct® detection of miRNAs from plasma is quantitative and scalable

Sample-to-sample consistency (plasma input volume 50µL)

miRNAs with decreasing abundance in plasma

Proportional Ct decrease with increasing plasma volume

miR-Direct® has high sample-to-sample consistency (left graph). Measured miRNA levels increase proportionately to the input plasma volume (right graph) with an approximate 2-Ct decrease for every 4-fold increase in plasma volume.

miR-Direct® reliably detects miRNA in urine samples

miR-Direct® detects miRNAs with higher sensitivity than a competitor kit with bead-based miRNA capture followed by RT-qPCR. Urine samples from 4 healthy volunteers are analyzed. Ct values over 37 are considered background level.

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