

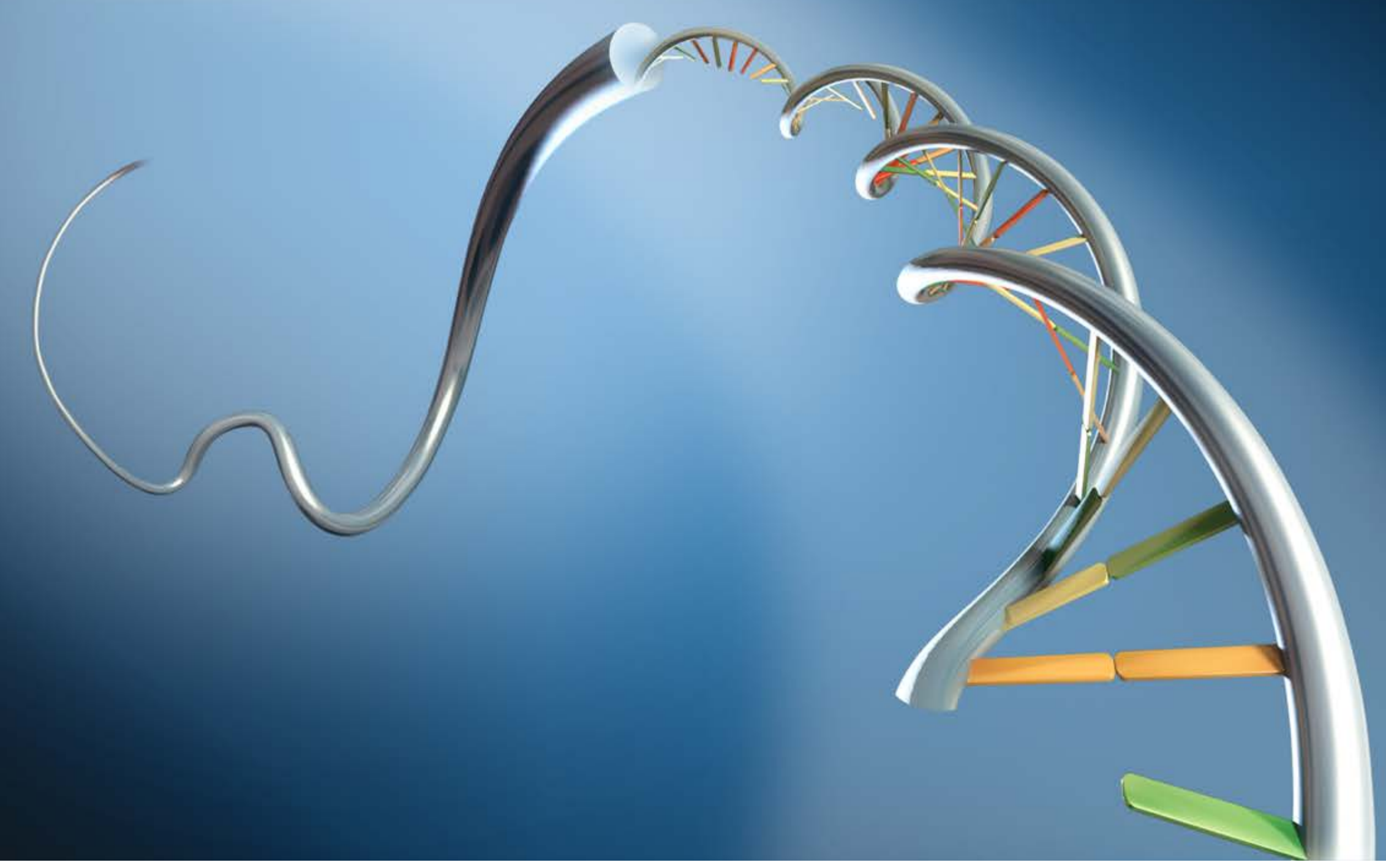
miRiam – NEW TRENDS IN microRNA QUANTIFICATION

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Introduction

MicroRNAs (miRNAs) are small non-coding RNA molecules playing an important regulatory role in gene translation through silencing or degradation of target mRNAs. They are involved in a wide range of biological processes, including differentiation and proliferation, metabolism, hemostasis, apoptosis or inflammation and also in pathophysiology of many diseases. Recently, numerous studies have suggested circulating miRNAs as promising diagnostic and prognostic biomarkers of cardiovascular diseases, neurological disorders, cancer, metabolic syndrome and many other diseases. Monitoring the level of specific miRNA together with protein-based biomarkers may represent more efficient tool for diagnosis of these diseases and prognosis estimation.

Current methods of miRNA determination are either low in specificity and sensitivity or very expensive and high-technology demanding like Next-generation sequencing (NGS) or qRT-PCR which is considered to be the gold standard for miRNA expression analysis.

Method

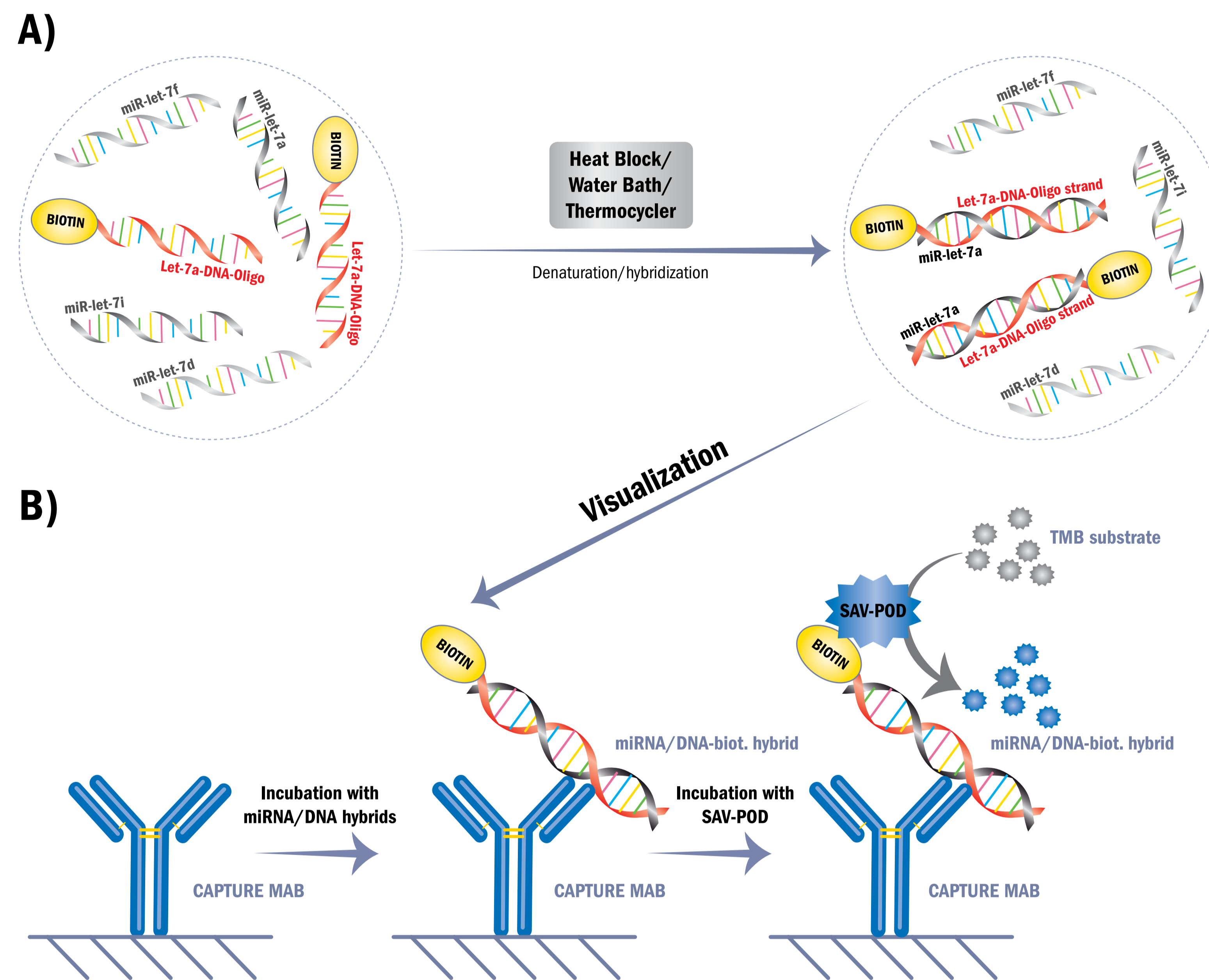
We are introducing a novel, immunoassay-based method of miRNA quantification which involves hybridization of miRNA isolated from a patient sample to complementary biotinylated DNA oligonucleotide probe. The DNA/RNA hybrids are then transferred onto a stationary solid phase coated with monoclonal antibody specific to perfectly matched DNA/miRNA hybrids. After washing, the solid phase is incubated with streptavidin-HRP conjugate and the resulting complexes are visualized (after another washing step) by a chromogenic substrate. Our immunoassay exhibits superior analytical specificity, limit of detection as low as 0.1 attomol/μl miRNA, excellent analytical characteristics and strong correlation with the qRT-PCR method (Pearson correlation coefficient >0.9).

Moreover, the assay can be run on common immunoassay analyzers, is compatible with standard clinical workflow, does not require amplification steps and results are obtained in less than three hours including miRNA profiling. Our method enables to analyze miRNA using conventional immunoassay analyzers, thus, it can promote utilization of miRNA biomarkers in clinical and laboratory practice.

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miRNA quantification



Hybridization of miRNA to complementary biotinylated DNA (A)

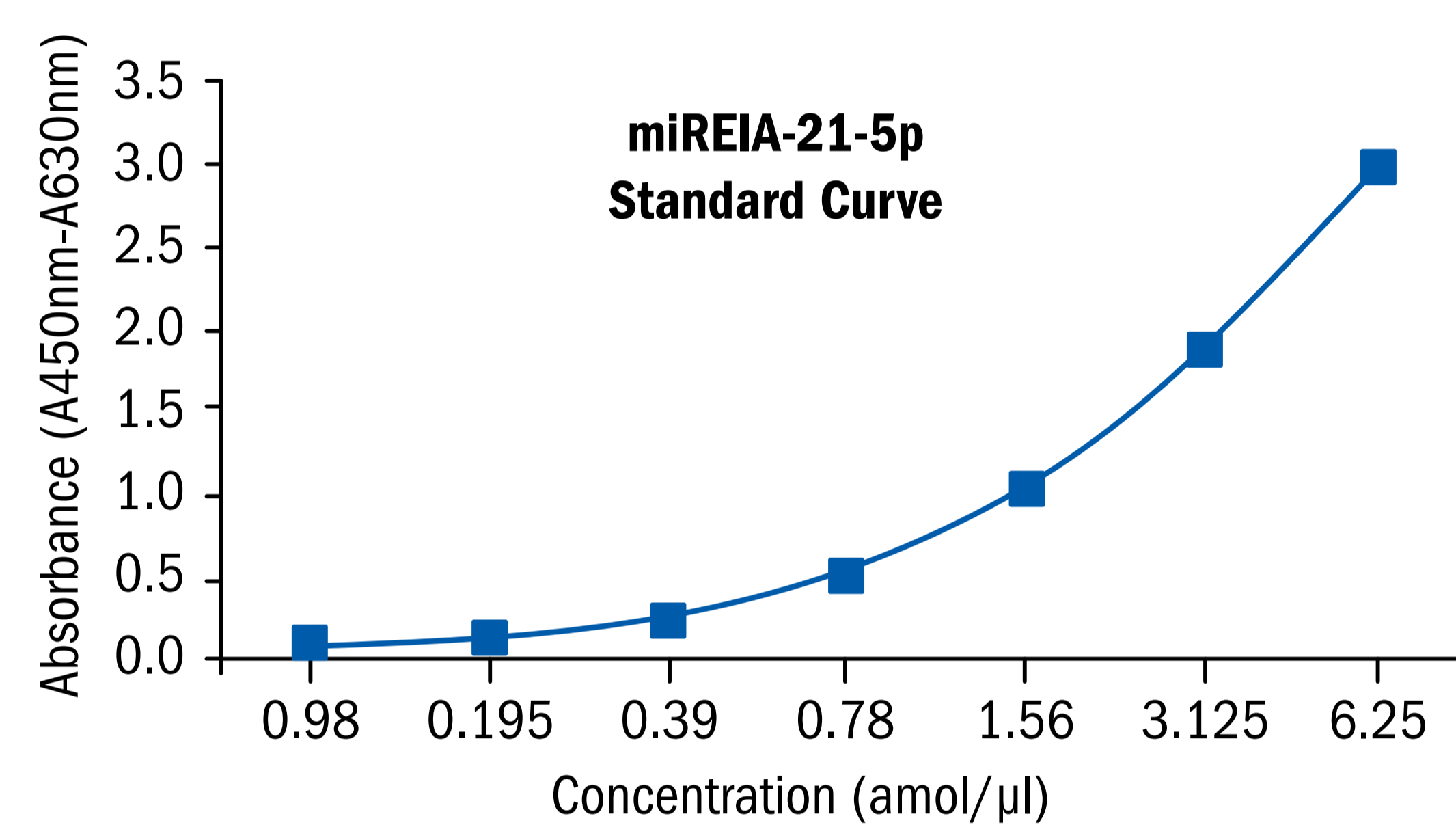
Defined amount of specific biotin-labeled DNA oligonucleotide is hybridized to miRNA isolated from blood sample.

miReia – miRNA enzyme immunoassay (B)

The hybridization mixture is transferred onto a stationary solid phase coated with monoclonal antibody specific to perfectly matched RNA/DNA-biotin hybrids. In the next step, the solid phase is washed and subsequently incubated with streptavidin-HRP conjugate. Finally, the resulting complexes are visualized by chromogenic substrate 3,3',5,5'-tetramethylbenzidine (TMB).

miREIA-21-5p

C) Calibration range



The calibration range of the miREIA-21-5p assay is 0.098-6.25 amol/ul.

D) Dilution linearity

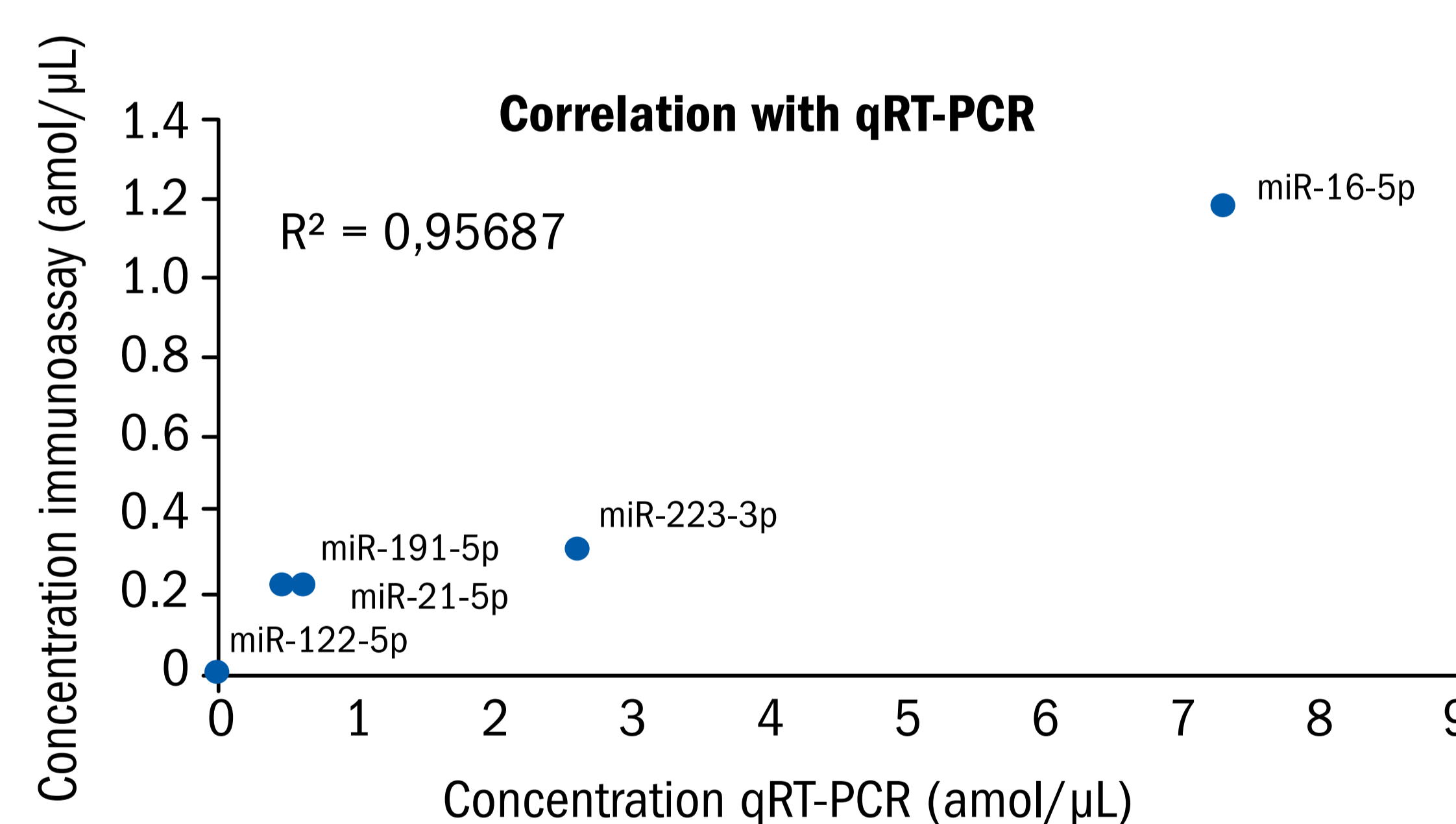
Two microRNA isolates from whole blood samples were serially diluted.

Sample	Sample dilution	Observed concentration amol/μl	Expected concentration amol/μl	Recovery O/E (%)
Sample 1	-	10.2	-	-
	2	5.0	5.1	99
	4	2.7	2.6	105
	8	1.3	1.3	100
Sample 2	-	20.4	-	-
	2	10.1	10.2	99
	4	4.8	5.1	94
	8	2.6	2.6	100

Recovery was determined to be 94 – 105%

E) Comparison with qRT-PCR

Five different miRNAs (miR-122-5p; miR-21-5p; miR191-5p; miR-223-3p and miR16-5p) isolated from whole blood were quantified by miREIA and by qRT-PCR using LNA nucleotide probes from Exiqon.



We found a strong correlation between qRT-PCR and miREIA-21-5p, the Pearson correlation coefficient being >0,9.

F) Total Assay Time

LNA qRT-PCR	miReia
Adjust each of the template RNA samples to a optimal concentration (15 min)	Hybridization (10 min)
Reverse transcription (65 min)	Preparation and pipeting of sample to the plate (15 min)
Preparation of samples after reverse transcription and PCR reagents (20 min)	Incubation of hybrids with monoclonal Antibody (60 min)
Preparation the PCR reaction plate (15 min)	Incubation with SAV-POD (30 min)
Run the PCR reaction plate (90 min)	Incubation with substrate (10 min)
Total time to result 3,5 hours	Total time to result 2 hours

We compared the total time needed for miRNA quantification by two different methods: qRT-PCR and miREIA. The total assay time of miREIA is significantly shorter when compared to qRT-PCR.

Conclusion

- ▶ We are introducing a novel method for miRNA quantification
- ▶ miREIA assays exhibits excellent analytical characteristics and strong correlation with qRT-PCR