Author's Accepted Manuscript

Sensitive and specific detection of microRNAs based on two-stage amplification reaction using molecular beacons as turn-on probes

Wancun Zhang, Fang Hu, Qi Zhang, Jiaqi Zhang, Yibo Mao, Peng Wang, Gu Yueqing



www.elsevier.com/locate/talanta

PII: S0039-9140(17)31195-5

DOI: https://doi.org/10.1016/j.talanta.2017.11.068

Reference: TAL18130

To appear in: Talanta

Received date: 21 August 2017 Revised date: 16 November 2017 Accepted date: 28 November 2017

Cite this article as: Wancun Zhang, Fang Hu, Qi Zhang, Jiaqi Zhang, Yibo Mao, Peng Wang and Gu Yueqing, Sensitive and specific detection of microRNAs based on two-stage amplification reaction using molecular beacons as turn-on probes, *Talanta*, https://doi.org/10.1016/j.talanta.2017.11.068

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting galley proof before it is published in its final citable form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

ACCEPTED MANUSCRIPT

Sensitive and specific detection of microRNAs based on two-stage amplification reaction using molecular beacons as turn-on probes

Wancun Zhang, Fang Hu, Qi Zhang, Jiaqi Zhang, Yibo Mao, Peng Wang, Yueqing,

Gu*

State Key Laboratory of Natural Medicines, Department of Biomedical Engineering, School of Engineering, China Pharmaceutical University, 210009 Nanjing, China

ABSTRACT

In this study, a rapid, sensitive, and specific assay for detecting miRNAs was developed based on a two-stage amplification reaction (TSAR) using molecular beacons (MBs) as turn-on probes. In the TSAR, different miRNAs can be converted to the same reporter oligonucleotides (Y), which can hybridize with the same MB. Therefore, in combination with specific templates, this method can be applied to multiplex miRNA detection by simply using the same MB. The loop region of the MB was screened by computer simulation methods. In particular, to improve the specificity of the MB in real sample analysis, the maximum similarity of the MB loop region to the human genome and human transcriptome is less than 70%. Two MBs were designed in this study. MB I, with nine flanking base pairs in its stem region, was used for real-time monitoring of the production of Y during the TSAR. MB II, with five flanking base pairs in its stem region, was used to detect the production of Y at the end of the TSAR. This assay exhibited high sensitivity with a limit of detection of 2.0×10^{-16} M and 6.7×10^{-16} M using MB I and MB II as turn-on probes, respectively. In addition, this assay can clearly discriminate single base differences in

Download English Version:

https://daneshyari.com/en/article/7677465

Download Persian Version:

https://daneshyari.com/article/7677465

<u>Daneshyari.com</u>