

## Targeting MicroRNA in Diagnosis and Therapeutics

Shaik MM\*

Human Genome Centre, School of Medical Science, Universiti Sains Malaysia, Kubang Kerian 16150, Kelantan, Malaysia

MicroRNAs (miRNAs) are part of a group of non-coding RNAs that can block messenger-RNA (mRNA) translation and affect mRNA stability. The miRNA may regulate the expressions of large non-coding RNAs as they predict to regulate at least 30% of all the human protein-coding genes by targeting their 3'-UTR sequences [1]. This indicates the role of miRNA in transcriptome networks in eukaryotic cells. The miRNA-deregulated expressions were reported in variety of complex diseases and the deregulated expressions of miRNA may be due to epigenetic changes [2].

MicroRNAs can be potential biomarkers in Neurological diseases like Alzheimer's Diseases [3] and migraine [4] where epigenetics play main role in the complexity of the disease [5-7]. Determination of circulating blood biomarkers helps to earlier diagnosis which may help in treatments earlier and prevent these devastating diseases which can have the potential to positively impact patient comfort. Additionally, the expressions of miRNAs are dysregulated in vascular diseases and they are critical modulators for vascular cell functions such as cell differentiation, contraction, migration, proliferation, and apoptosis [8]. The miRNAs play vital role in ischemic angiogenesis, vascular dysfunction, re-endothelialization and vascular neointimal lesion formation. Due to this, miRNAs may have vital role as novel therapeutic targets for vascular diseases.

Based on reasons mentioned above, there is great excitement on miRNAs research as biomarker and therapeutic target for major diseases. Regulation of mRNA levels and protein expression involves the miRNA binding step to their target and which may be affected

by polymorphisms. These polymorphisms may play vital role in pharmacodynamics and pharmacokinetics of the drugs. The multiple miRNAs may be therapeutic targets and newer drugs can inhibit miRNAs with minimal toxicity.

### References

1. Rajewsky N (2006) microRNA target predictions in animals. *Nat Genet* 38: S8-13.
2. Brueckner B, Stresemann C, Kuner R, Mund C, Musch T, et al. (2007) The human let-7a-3 locus contains an epigenetically regulated microRNA gene with oncogenic function. *Cancer Res* 67: 1419-23.
3. Maes OC, Chertkow HM, Wang E, Schipper HM (2009) MicroRNA: Implications for Alzheimer Disease and other Human CNS Disorders. *Curr Genomics* 10: 154-168.
4. Kress M, Hüttenhofer A, Landry M, Kuner R, Favereaux A, et al. (2013) microRNAs in nociceptive circuits as predictors of future clinical applications. *Front Mol Neurosci* 6: 33.
5. Shaik MM, Tan HL, Kamal MA, Gan SH (2014) Do folate, vitamins B(6) and B(1)(2) play a role in the pathogenesis of migraine? The role of pharmacoepigenomics. *CNS Neurol Disord Drug Targets* 13: 828-835.
6. Shaik MM, Gan SH, Kamal MA (2014) Epigenomic approach in understanding Alzheimer's disease and type 2 diabetes mellitus. *CNS Neurol Disord Drug Targets* 13: 283-289.
7. Shaik MM, Ahmad S, Gan SH, Abuzenadah AM, Ahmad E et al. (2014) How do periodontal infections affect the onset and progression of Alzheimer's disease *CNS Neurol Disord Drug Targets* 13: 460-466.
8. Qin S, Zhang C (2011) MicroRNAs in vascular disease. *J Cardiovasc Pharmacol* 57: 8-12.

\*Corresponding author: Shaik MM, Human Genome Centre, School of Medical Science, Universiti Sains Malaysia, Kubang Kerian 16150, Kelantan, Malaysia, Tel: 0060-09-767 6139; E-mail: [munvar.shaik@gmail.com](mailto:munvar.shaik@gmail.com)

Received October 13, 2014; Accepted October 14, 2014; Published October 22, 2014

Citation: Shaik MM (2014) Targeting MicroRNA in Diagnosis and Therapeutics. *J Biomol Res Ther* 3: e137. doi: [10.4172/2167-7956.1000e137](https://doi.org/10.4172/2167-7956.1000e137)

Copyright: © 2014 Shaik MM. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.