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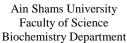
تم رفع هذه الرسالة بواسطة / هناء محمد علي

بقسم التوثيق الإلكتروني بمركز الشبكات وتكنولوجيا المعلومات دون أدنى مسئولية عن محتوى هذه الرسالة.

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Molecular study on Micro RNAs 122 and 221 as biomarkers for hepatocellular carcinoma and hepatitis C virus

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Acknowledgment

First and foremost, I feel always indebted to AUAH, the Most Kind and Most Merciful.

I'd like to express my respectful thanks and profound gratitude to **Prof. Dr. Amina M.**Medhat, Professor of Biochemistry, Faculty of Science - Ain Shams University for her keen guidance, kind supervision, valuable advice and continuous encouragement, which made possible the completion of this work.

I am also delighted to express my deepest gratitude and thanks to **Prof. Dr. Laila A. Rashed,** Professor of Medical Biochemisty and Molecular Biology, Faculty of Medicine - Cairo University, for her kind care, continuous supervision, valuable instructions, constant help and great assistance throughout this work.

I wish to introduce my deep respect and thanks to **Prof. Dr. Mahmoud M. Said,** Professor of Biochemistry, Faculty of Science - Ain Shams University, for his kindness, patience, great help, active participation and guidance, fabulous efforts with the statistical analysis and revision of the thesis.

Hanan Hanafy Abd Elhamed Zedan

Declaration

I declare that this thesis has not been submitted for a degree at this or any other university.

Hanan Hanafy Abd Elhamed Zedan

Dedication

TO...

My lovely family

For their support in every step in my life giving everything and never waiting for anything.

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Abstract

Hepatocellular carcinoma (HCC) is a common solid organ malignancy worldwide, with about 600,000 new cases diagnosed each year, and it is the fifth most frequent human cancer and a fatal disease. MicroRNAs (miRs) are small non-coding RNA segments of nearly 22 nucleotides, regulating gene expression after transcription. MiRs have an essential role in cellular proliferation, differentiation, apoptosis, as well as carcinogenesis process. The present study aimed at evaluating the expression level of miR-122, miR-221 and cyclin G1 in hepatitis C virus (HCV) and HCC patients in order to evaluate whether they could be used as sensitive biomarkers for HCC development and its different stages as surrogate biomarkers for α -fetoprotein (AFP). The study included 28 HCV patients and 36 HCC patients, further subdivided into stage I HCC patients (n=10), stage II HCC patients (n=14) and stage III HCC patients (n=12). In addition, normal healthy individuals (n=13) were recruited into this study. MiR-122, miR-221 and cyclin G1 gene expression levels were determined by quantitative real time polymerase chain reaction (qRT-PCR). The HCV patients demonstrated a significant increase in serum AFP level expression, compared to the healthy subjects. On the other hand, HCC patients manifested a sharp significant elevation in serum AFP expression level, compared to healthy subjects as well as HCV patients. In addition, HCV patients demonstrated a significant increase in serum miR-221 and cyclin G1 expression levels, compared to healthy subjects, while HCC patients manifested a significant elevation in serum miR-122, miR-221 and cyclin G1 expression levels, compared to healthy subjects. Moreover, ROC curve analysis demonstrated that serum miR-122 and cyclin G1 were able to discriminate between HCV and HCC patients. In conclusion, serum miR-122 and miR-221 may serve as non-invasive diagnostic biomarkers for HCC. MiR-122 and miR-221 were better than AFP, the gold standard biomarker for HCC, in the diagnosis of the early HCC stage with high sensitivity and specificity. Moreover, the combination between the serum biomarkers miR-122, miR-221 and cyclin G1 could be a more useful diagnostic tool for HCC better than each of them individually, and also as a substitute for AFP.

Keywords: Micro RNA 122; Micro RNA 221; Cyclin G1; Hepatocellular Carcinoma; Hepatitis C virus, α-fetoprotein.

| Abbreviation | Full term |
|--------------|--|
| AASLD | American associated for the study of liver |
| | diseases |
| ADAM10 | A disintegrin and metallopeptidase domain 10 |
| AFP | α-fetoprotein |
| AFP-L3 | Lens culinaris agglutinin-reactive AFP |
| AGO2 | Argonaute 2 |
| ALP | Alkaline phosphatase |
| ALT | Alanine aminotransferase |
| AMVRT | Avian myeloblastosis virus reverse transcriptase |
| ASSF1 | Ras association domian-contaninig protein |
| | family member 1 |
| AST | Aspartate aminotransferase |
| AUC | Area under curve |
| BCLC | Barcelona Clinic Liver Cancer |
| BMF | Bcl-2 modifying factor |
| cDNA | Covalently closed circular DNA |
| CCND1 | Cyclin D1 |
| E-CDH 1 | Epithelial cadherin |
| CDKN1C | Cyclin-dependent kinase inhibitor 1 C |
| CDKs | Cyclin-dependent kinases |
| cDNA | Complementary DNA |
| CIP | CDK interacting protein |
| CKI | CDK inhibitor |
| CLIP | Cancer of the Liver Italian Program |
| CT | Computer tomography |
| Ct | Cycle threshold |
| CTNNB1 | Catenin beta 1 |
| CUPI | Chinese University Prognostic Index |
| CUTL1 | Cut-like homeobox 1 |
| Cyc | Cyclins |
| DAA | Direct acting antiviral |
| DCP | Des-Gamma-carboxy prothrombin |
| DEAE | Diethylaminoethyl |
| DGCR8 | DiGeorge syndrome critical region gene 8 |
| DM | Diabetes mellitus |

| Abbreviation | Full term |
|--------------|--|
| E2F | E 2 factor |
| EDTA | Ethylene diamine tetraacetic acid |
| EGFR | Epithelial growth factor receptor |
| EL | Erythrocytes lysis |
| ELISA | Enzyme linked immunosorbent assay |
| G1 | Gap1 |
| G2 | Gap2 |
| GAPDH | Glyceraldehyde 3-phosphate glyceraldehyde |
| GGT | Gamma-glutamyl transferase |
| GHSS | Global health system solution |
| GPC | Glypican-3 |
| GRETCH | Group d' Etude et de Traitement du Carcinoma |
| | Hepatocellulaire. |
| HBV Ab | Hepatitis B virus anyibody |
| HBV | Hepatitis B virus |
| HCC | Hepatocellular carcinoma |
| HCV | Hepatitis C virus |
| HCV-RNA | HCV viremia |
| HDAC | Histone deacetylase |
| HEK-293 T | Human embryonic kidney 293 |
| HEPG2 | Human hepatoma G2 |
| HMOX1 | Heme oxygenase 1 |
| HNF | Hepatocyte nuclear factor |
| IFN | Interferon |
| IGFR1 | Insulin growth factor receptor 1 |
| IRES | Internal ribosome entry site |
| JIS | Japanese Integrated System |
| KIP | Kinase inhibitor protein |
| LC | Liver cirrhosis |
| LETFs | Liver-enriched transcription factors |
| LNA | Locked nucleic acid |
| LR | Likelihood ratio |
| M | Mitosis |
| MGB | Minor groove binder |
| MiRs | MicroRNAs |

| Abbreviation | Full term |
|--------------|---|
| | |
| MOPS | Morpholino propanesulfonic acid |
| mRNAs | Messenger RNAs |
| mTOR | Mammalian target of rapamycin |
| MWA | Microwave ablation |
| NAFLD | Non-alcoholic fatty liver disease |
| NCRP | National cancer registry program |
| NFQ | Non-fluorescent quencher |
| NPV | Negative predictive value |
| NTR | Non- translated RNA |
| NS | Nonstructural |
| PCR | Polymerase chain reaction |
| PEI | Precutaneous ethanol injection |
| PIVKA-II | Protein induced by vitamin K absence or |
| | antagonist II |
| PKB | Protein kinase B |
| POL II | RNA polymerase II |
| PPV | Positive predictive value |
| pRb | Retinoblastoma family of protien |
| pre-miRs | Precursor miRs |
| pri-miRs | Pri-microRNAs |
| PTEN | Phosphatase and tensin homolog |
| qRT-PCR | Quantitative real time polymerase chain |
| | reaction |
| RASSF1 | Ras association domain-containing protein |
| | family member 1 |
| RanGTP | RAS-related nuclear protein GTP-dependent |
| | transporter |
| RFA | Radio frequency ablation |
| RISC | RNA-induced silencing complex |
| RNU6B | U6B small nuclear RNA |
| ROC | Receiver operating characteristics |
| RQ | Relative quantitation |
| RT | Revers transcription |
| S | Synthesis of DNA |
| SIRT | Selective internal radiation therapy |

| Abbreviation | Full term |
|--------------|---|
| | |
| SNPs | Single nucleotide polymorphisms |
| SOCS1 | Suppressor of cytokine signaling 1 |
| SVR | Sustained virological response |
| TACE | Transarterial chemoembolization |
| TAE | Transarterial embolization |
| TGF-13 | Transforming growth factor-13 |
| TGF-β1 | Transforming growth factor beta-1 |
| TIMP3 | Tissue inhibitor of metalloproteinase 3 |
| TMB | Tetramethlbenzidine |
| TNM | Tumor-node metastasis |
| tRNA | Transfer RNA |
| UTR | Untranslated region |
| VEGFA | Vascular endothelial growth factor A |
| WHO | World health organization |
| χ2 | Pearson's chi square |
| β-МЕ | β-Mercaptoethanol |

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