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REVIEW ARTICLE

Neutrophil Extracellular Traps and Its Correlation with Several Pathological Conditions: Prosperities and Deleterious Implications

Phey Liana, Krisna Murti, Zen Hafy, Iche Andriyani Liberty, Tungki Pratama Umar; p.1-11

Role of Genetics in Anesthesiology

Ziske Maritska, Rani Iswara, Ignatius Aldo Winardi, Yuni Dwi Marantika, Irfan Ferdinand Tambunan, Lovina, Mgs. A. Rifqi Murtadho, Sendy Aditya Nugraha, Cendy Legowo, Victor Pulpa Seda, Awang Budi Saksono; p.12-9

RESEARCH ARTICLES

Non-Synonymous Mutation Analysis of SARS-CoV-2 ORF3a in Indonesia

Hartiowidi Yuliawuri, Jeanne Elvia Christian, Nathanael Steven; p.20-7

ZEB1 is Negatively Correlated with E-Cadherin in Prostatic Anomaly Tissue

Sari Eka Pratiwi, Sri Nuryani Wahyuningrum, Rachmagreta Perdana Putri, Danarto, Didik Setyo Heriyanto, Nur Arfan, Sofia Mubarika Haryana, Indwiani Astuti; p.28-34

Photo-illuminated Glutathione Inactivates Alpha-2-macroglobulin: Spectroscopic and Thermodynamic Studies

Syed Saqib Ali, Haseeb Ahsan, Sana Ansari, KM Abdullah, Fahim Halim Khan; p.35-42

Correlation of Serum Nitric Oxide and Urine Malondialdehyde Levels in Non-Hemodialysis Chronic Kidney Disease Patients

Dinda Dwi Purwati, Arifa Mustika, Lukman Hakim, Mochammad Thaha; p.43-9

The Effect of Thelison Use in The Etiology of Lung Disorders among Homeless People

Alaa Aldin Alsafi Alalwiy, Alkhair Abd Almahmoud Idris; p.50-4

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Murray PR, Rosenthal KS, Kobayashi GS, Pfaller MA. *Medical microbiology*. 4th ed. St. Louis: Mosby; 2002.

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Molecular and Cellular Biomedical Sciences

CONTENT

REVIEW ARTICLE

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Phey Liana, Krisna Murti, Zen Hafy, Iche Andriyani Liberty, Tungki Pratama Umar
p.1-11

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p.28-34

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Syed Saqib Ali, Haseeb Ahsan, Sana Ansari, KM Abdullah, Fahim Halim Khan
p.35-42

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Dinda Dwi Purwati, Arifa Mustika, Lukman Hakim, Mochammad Thaha
p.43-9

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p.50-4

Molecular and Cellular Biomedical Sciences

Abstract

DDC 616.978

Liana P, Murti K, Hafy Z, Liberty IA, Umar TP (Department of Clinical Pathology, Faculty of Medicine, Universitas Sriwijaya/Dr Mohammad Hoesin General Hospital, Palembang, Indonesia)

Neutrophil Extracellular Traps and Its Correlation with Several Pathological Conditions: Prosperities and Deleterious Implications

Mol Cell Biomed Sci. 2022; 6(1): 1-11

Abstract (English)

Neutrophil extracellular traps (NETs) are immune components found in a variety of pathological states. It has been shown to have either beneficial or harmful implications, depending on how it is controlled and has been particularly observed in three major scenarios: infection, autoimmune disease, and cancer. In this article, we compiled some of the roles of NETs in pathological conditions, as well as the benefits of targeting them for improved patient outcomes. The role of NETs were primarily positive in infectious disease, whether caused by bacteria, virus, or fungal infection. In non-infectious inflammatory scenarios, on the other hand, it's the complete opposite, with the effects being mainly deleterious and even worse than the original disease states. Targeting NETs directly or indirectly may help to prevent complications and improve patient outcomes. A plethora of compounds, including immunomodulators, anti-thrombosis, nicotinamide adenine dinucleotide phosphate (NADPH)/reactive oxygen species (ROS) inhibitors, nuclease, and other compounds, may be used to accomplish the therapeutic goals.

Keywords: autoimmune disease, cancer, DNase, infection, neutrophil extracellular traps

DDC 617.96

Maritska Z, Iswara R, Winardi IA, Marantika YD, Tambunan IF, Lovina, Murtadho MAR, Nugraha SA, Legowo C, Seda VP, Saksono AB (Department of Biology Medicine, Faculty of Medicine, Universitas Sriwijaya, Palembang, Indonesia)

Role of Genetics in Anesthesiology

Mol Cell Biomed Sci. 2022; 6(1): 12-9

Abstract (English)

One thing that differentiates one person from another is one's genetic make-up. Genetic plays a role in every branch of medicine, including anesthesiology. An anesthesiologist must be well familiarized with hereditary (genetic) conditions, chromosomal traits, heredity-familial disorders, and even recessive variants because particular conditions might demand a different anesthetic and perioperative pharmacological management. These circumstances may lead to an opening of a rapidly expanding state of pharmacogenetics/genomics and its relevancy in anesthesia nowadays. This narrative review provides insight into the role of genetics in the field of anesthesiology.

Keywords: genetics, anesthesiology, pharmacogenetics

DDC 576.549

Yulawuri H, Christian JE, Steven N (Biomedical Science Program Study, Calvin Institute of Technology, Jakarta, Indonesia)

Non-Synonymous Mutation Analysis of SARS-CoV-2 ORF3a in Indonesia

Mol Cell Biomed Sci. 2022; 6(1): 20-7

Abstract (English)

Background: The report of mutation sites ORF3a SARS CoV-2 in Indonesia is still limited. Some research showed that mutations in ORF3a protein might alter SARS-CoV-2 pathogenesis. Observation of new variants should be conducted as a risk monitoring framework.

Materials and method: We assessed the impact of mutations in ORF3a protein by analyzing 3,751 SARS-CoV-2 DNA sequences from the GISAID database from March 2020 until July 2021. The whole-genome sequences were aligned using Clustal Omega Multiple Sequence Alignment from EMBL-EBI and analyzed using BioEdit version 7.2.5 software. The reference whole genome sequence was taken from the Genbank database with accession number NC045512. We excluded the samples containing N letters due to inaccurate reading. Effect of point mutations on protein structure was analyzed using PredictProtein (<https://predictprotein.org>) and Protein Variation Effect Analyzer (PROVEAN) v1.1.3. online software.

Results: We identified five most frequent non-synonymous mutations in ORF3a protein of SARS-CoV-2 which were Q57H (58.04%), S26L (27.25%), S220I (10.37%), D155H (8.98%), and P104S (5.47%).

Conclusion: These mutation data showed the phenomenon of amino acid changes in ORF3a SARS-CoV-2 in Indonesia until July 2021. The implication of this mutation needs to be determined in further studies.

Keywords: Indonesia, mutations, non-synonymous, SARS-CoV-2, whole genome

Molecular and Cellular Biomedical Sciences

DDC 616.65

Pratiwi SE, Wahyuningrum SN, Putri RP, Danarto, Heriyanto DS, Arfian N, Haryana SM, Astuti I (Department of Biology and Pathobiology, Faculty of Medicine, Universitas Tanjungpura, Pontianak, Indonesia)

ZEB1 is Negatively Correlated with E-Cadherin in Prostatic Anomaly Tissue

Mol Cell Biomed Sci. 2022; 6(1): 28-34

Abstract (English)

Background: Prostatic anomalies are common in tumor or infection condition. The enlargement of prostate gland affects the epithelial cell polarity that involves epithelial-mesenchymal transition (EMT). Transition into mesenchymal is mediated by transcription factor ZEB1 and E-cadherin protein. Upregulation of ZEB1 and loss of E-Cadherin expression were associated to proliferation and metastasis of malignancy cells. This study aims to describe the correlation of ZEB1 and E-cadherin expression in prostatic anomaly.

Materials and method: Samples were Formalin Fixed Paraffin Embedded (FFPE) block consist of 8 block Benign Prostatic Hyperplasia (BPH), 6 blocks High Grade Prostatic Intraepithelial Neoplasia (HGPIN) and 6 blocks Prostate Carcinoma (PCA). The blocks then sliced into 5 sections to be prepared for RNA extraction procedures. ZEB1 and E-Cadherin expression was analyzed by semi-quantitative procedures using PCR and electrophoresis. Correlation between ZEB1 and E-Cadherin expression was analyzed using Spearman's rank correlation.

Results: Relative expression of ZEB1 and E-cadherin mRNA in each group of prostatic anomaly were not significantly different ($p > 0.05$). ZEB1 and E-Cadherin mRNA expression showed a significant and moderate level of negative correlation ($p < 0.05$; $0.40 < r < 0.59$). Increasing of ZEB1 mRNA expression will be followed by decreasing of E-Cadherin mRNA expression.

Conclusion: ZEB1 negatively correlates with E-cadherin due to EMT process in prostatic anomaly. High expression of ZEB1 induced down-regulation of E-cadherin and vice versa. Various studies can be developed, especially the development of targeted therapy against ZEB1 to suppress the EMT process by increasing the expression of E-cadherin.

Keywords: epithelial-mesenchymal transition (EMT), ZEB1, E-Cadherin, BPH, HGPIN, PCA

DDC 543.5

Ali SS, Ahsan H, Ansari S, Abdullah KM, Khan FH (Department of Biochemistry, Faculty of Life Sciences, Aligarh Muslim University, Aligarh, India)

Photo-illuminated Glutathione Inactivates Alpha-2-macroglobulin: Spectroscopic and Thermodynamic Studies

Mol Cell Biomed Sci. 2022; 6(1): 35-42

Abstract (English)

Background: Glutathione (GSH) is a principle thiol-containing tripeptide (cysteine, glutamic acid and glycine) antioxidant against free radicals and other harmful oxidants in cellular defence. The alpha-2-macroglobulin ($\alpha 2M$) is large tetrameric zinc-binding glycoprotein which inhibits proteinases regardless of their specificity and catalytic mechanism.

Materials and Methods: The interaction of GSH was analyzed with $\alpha 2M$ including the structural and functional alterations in $\alpha 2M$ using various biochemical and biophysical methods. UV-visible and fluorescence spectroscopy were used to study the binding of $\alpha 2M$ with GSH and Fourier transform infrared (FT-IR) spectroscopy was explored to study the structural change induced in $\alpha 2M$.

Results: The results suggest that exposure of $\alpha 2M$ to GSH decreases the antiproteolytic potential as suggested by the amidase assay. The UV-spectroscopic study showed the formation of $\alpha 2M$ -GSH complex and fluorescence analysis showed significant quenching in fluorescence intensity of $\alpha 2M$ suggesting GSH binding and structural alteration in the protein. FT-IR spectroscopy was explored to study the structural change induced in $\alpha 2M$ which suggest that the secondary structure of $\alpha 2M$ changes upon complex formation.

Conclusion: Our studies show that interaction of $\alpha 2M$ with photoilluminated GSH results in functional and conformational changes of the protein.

Keywords: glutathione, GSH, alpha-2-macroglobulin, photo-illumination, ITC, FTIR

DDC 616.614

Purwati DD, Mustika A, Hakim L, Thaha M (Medical Program, Faculty of Medicine, Universitas Airlangga, Surabaya, Indonesia)

Correlation of Serum Nitric Oxide and Urine Malondialdehyde Levels in Non-Hemodialysis Chronic Kidney Disease Patients

Mol Cell Biomed Sci. 2022; 6(1): 43-9

Abstract (English)

Background: In 2017, about 1.2 million people died because of Chronic Kidney Disease (CKD). Patients with CKD are known to have increased levels of oxidative stress which leads to decrease in NO production. NO is a highly reactive signaling molecule and a major determinant of vascular homeostasis. Thus, the decreased NO can be a risk factor for the development of atherosclerosis and increased cardiovascular risk. Meanwhile, Malondialdehyde (MDA) is known as excellent biomarker for oxidative stress. This study aims to determine the correlation of serum total nitric oxide (NO) and urine MDA levels in non-hemodialysis CKD patients.

Molecular and Cellular Biomedical Sciences

Materials and Methods: This study was an observational clinical study with a cross sectional design. Forty-nine CKD subjects were selected by consecutive sampling. The samples for laboratory tests were collected from urine. MDA concentration was measured using the High-Performance Liquid Chromatography (HPLC) kit. NO concentration was measured with Griess reaction method and Total Nitric Oxide Parameter kit. The data were analyzed using the Statistic Package for Social Science (SPSS) software version 16.

Results: The data showed significant negative correlations between MDA with NO ($r=-0.294$; $p=0.041$).

Conclusion: There was a correlation between serum total NO and urine MDA levels in non-hemodialysis CKD patients.

Keywords: chronic kidney disease, malondialdehyde, nitric oxide, non-hemodialysis

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The Effect of Thelison Use in The Etiology of Lung Disorders among Homeless People

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Abstract (English)

Background: Thelison, the adhesive synthetic material that bind surface together, is widely used in industry and domestic purpose along with epoxy, glue and putty. The aim of this study was to detect the effect of thelison use in the etiology of lung disorders among homeless people in Khartoum State, Sudan.

Materials and method: This was a descriptive cross sectional study conducted in homeless people in Khartoum State. Sputum smears samples from 80 alcohol fixed homeless thelison user were collected. After the collection of the sample, Argyrophilic Nucleolar Organizer Region (AgNOR) technique and papanicolaou (PAP) stains were applied to each sample. A questionnaire to obtain essential data about respondents was also provided for each participant.

Results: Participant's ages range was 10-37, while the mean age was 17. The range of participants duration of use per years was 1-10, while the mean was 3 years. Number of thelesion dose use per day was ranged between 1-10, while the mean was 5 years. The majority of participants (80%) showed no cellular change and 20 % showed chronic inflammation. Results showed that 31 of the study population were males (77.5%), while the female population of the study was 9 (22.5%). The mean of AgNoR score was ranged between 1-4, while the mean AgNOR score was 1. AgNoR showed insignificant association with gender, duration and number of thelesion dose used per day ($p>0.05$), but showed significant association with cytomorphological and age ($p<0.05$).

Conclusion: AgNoR score showed insignificant association with gender, duration and number of thelesion dose used per day ($p>0.05$), but showed significant association with cytomorphological and age ($p<0.05$).

Keywords: AgNoRs score, cytological changes, sputum, homeless thelison users

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