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## From the Publisher's Desk



#### Welcome to Biotechnology Kiosk!

The November, 2020 issue of BK is now online for our readers with the regular features.

This issue contains a cutting-edge review on regulating pro-inflammatory cytokines and health effects news and views on the current topics that include latest research breakthroughs in different areas of medicine and biotechnology that include targeted cancer drugs and cardiovascular disease. We look forward to receiving your feedback. We do hope that you will enjoy reading this issue of Biotechnology Kiosk. Please do write to us with your comments. Your suggestions are always appreciated.

# Dr. Megha Agrawal & Dr. Shyamasri Biswas.

Co Editors-in-Chief, Biotechnology Kiosk



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### Pro-Inflammatory Cytokines as Potential Early Biomarkers and Therapeutic Targets to Combat Ischemic Stroke

Dr. Megha Agrawal\* Chief Editor Biotechnology Kiosk, USA

#### Abstract

Ischemic stroke is a serious medical condition and widely considered one of the most common causes of death and disability in the world today. There have been notable research advances in stroke so far and studies have shown that stroke's complex pathophysiology process involves the oxidative stress and inflammatory reaction. However, despite the progress in stroke research, currently there are no established biochemical factors available that can be employed in the early diagnostics and intervention in stroke. Mostly, stroke diagnosis is based on neuroimaging, which is not a rapid tool to diagnose stroke. This decreases the survivability rate. Further, conventional therapeutic approaches for ischemic stroke management are based on restoring blood flow to the affected brain area and these therapies are effective only during a limited time window. Hence, this procedure results in benefiting only a very small percentage of patients. In view of these limitations, the ongoing research has focused on seeking alternative treatment methods that can reduce stroke brain damage and improve patients' outcome. To this end, research goals are targeted towards gaining insights into the inflammatory response triggered by cerebral ischemia that is supposed to play an important role in the progression of stroke, and also the subsequent study of inflammatory molecules in the acute phase of stroke. In this mini-review, we describe the inflammatory processes occurring during ischemic stroke along with the potential for pro-inflammatory cytokines to become stroke biomarkers as well as interesting neuroprotective therapeutic targets that could be blocked or stimulated to modulate inflammation after stroke. Finally, we present a perspective briefly discussing some viewpoints on future studies in the ongoing field of stroke research.

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#### Introduction

Ischemic stroke represents a serious medical condition that is known to cause disability in elderly people (over 65 years of age) and considered the third most common cause of death in the world. According to an estimate by the World Health Organization, one in six people globally can be vulnerable to suffer from stroke in their lifetime. Ischemic stroke is known to be the most frequent kind of stroke that affects approximately 85-90% of patients. It can be caused by any of the medical conditions that include cardiogenic embolism, cerebral microcirculatory impairment or cerebral microangiopathy, atherosclerosis of extra- and intracranial arteries, and also blood clotting disorders that have been suggested as the cause of ischemic stroke [1-4].

The pathophysiology of stroke has been a subject of immense research interest. Studies conducted stroke on pathophysiology have suggested it to be a complex process and the key mechanisms involved in neuronal damage are suggested to be constituted by oxidative stress and inflammatory reaction [5]. It is understood that brain and immune cells produce reactive oxygen species (ROS) during ischemia and asphyxia. Such reactive species stimulate endothelial cells and cause oxidative stress. Consequently, it has been observed that the activity of ROS during ischemic stroke can cause primary vascular damage. It can also trigger the development of the inflammatory response that is connected with the acute immune response. In this process, glial cells (microglia, astrocytes) that are activated in ischemia together with blood cells

(leukocytes) and endothelial cells have been shown to synthesize a number of biochemical mediators and markers of inflammation including cytokines, chemokines and proinflammatory enzymes. Further, some genetic factors have been found to play significant roles in the development of inflammation in stroke that results in inflammatory reactions. In this regard, many genes have been demonstrated that can contribute to both the etiology of stroke and the size of the ischemic area, which affects the patient's late outcome. All these suggest that chronic activation of inflammation can be a factor that can significantly influence the advancement and impact of stroke risk factors [5-7].

While, there are notable advances made in a broad range of molecular diagnostic methods, biochemical factors that exhibit proven benefits for the rapid and early diagnostics, intervention and treatment of stroke have not yet been clearly established. To bridge the gap, studies related to the diagnostic and prognostic value of proinflammatory factors of neuronal, glial and neuroendocrine origin along with inflammatory markers in ischemic stroke are of huge current research interests [6, 7]. A special research focus is on cytokines, as regulatory proteins that are produced by different types of cells. These cytokines are recognized and bound by specific receptors that are located on cell membranes. A number of pro-inflammatory cytokines are known that include chemokines, interleukins, interferons, as well as cell differentiation and growth factors. Studies have suggested that

they can transduce signal to the cell nucleus by binding to the receptor in which expression of genes encoding acute phase proteins (APPs) occurs. In addition, they can be triggered by the transcription factor (NFκB). Further, it has been found that the inflammatory response in the brain changes over time. These observations imply that the knowledge concerning if and when the level of cytokine increases or decreases in the course of the acute phase of cerebral stroke could be immensely helpful that could pave the way to determine a possible role of proinflammatory cytokines as early biomarkers and also the target of a potential therapeutic intervention in the acute phase of cerebral stroke [8-11].

Here, we present an overview of the role of pro-inflammatory cytokines as early inflammatory factors for possible stroke biomarkers and novel therapeutic targets. Cytokines that could potentially act as different phases of inflammatory markers and influence the final prognosis could very well provide an alternative treatment method for stroke.

#### Stroke Biomarkers and Therapeutic Targets: Technology Challenges and New Innovations in Stroke Management

Biomarkers have huge pharmacological and clinical significance as they are important indicators of normal biological processes, pathogenic processes, or pharmacologic responses to therapeutic interventions. They can be objectively measured and evaluated as characteristics of important biomedical processes. Clinical biomarkers are employed for practical applications of diagnostics. These biomarkers can be detectable molecules from different biological fluids such as blood, urine, or saliva. To be clinical-grade biomarkers, it is extremely important that they have high specificity and sensitivity for the indicated process together with accuracy in their measurements and reproducibility, and easily interpretable by clinicians [10, 12, 13].

With respect to stroke biomarkers, they are representative of the brain injury processes triggered by stroke. For clinical applications, stroke biomarkers are considered in different phases that include both the diagnosis and the prognosis of stroke. Further, in the case of ischemia, the first few days after the onset of stroke require most important therapeutic decisions that need rapid biomarkers [13]. However, in practical applications, there is a possibility that stroke biomarkers may change dramatically in a short window of time especially during the hyper acute and acute phases compared with other phases. This poses a challenge for identifying optimal treatment windows for clinical biomarkers in each phase of stroke and stroke subtype (Figure 1) [13].

While, there has been some research progress in this direction, no biomarkers for stroke are currently available to use in clinical practice [12]. This is a big technology gap that currently exists. As regards to new ideas and looking ahead, researchers have shown that blood brain barriers 'BBB' disruption favors the release of brain antigens into the peripheral circulation. This event makes these molecules promising stroke-associated biomarkers. These biomarkers could reflect in the circulatory system, where pathological

processes take place in brain after the ischemic event. Further, another thought is that both local and peripheral inflammatory

response could produce systemic indicators that might serve as stroke biomarkers [10].



Figure 1: Schematic illustrations showing potential pathways for optimizing stroke biomarker clinical trials, where biomarkers can be identified in each phase of stroke (pre-stroke, stroke, post-stroke) and stroke subtype (ischemic, subarachnoid hemorrhage, intracerebral hemorrhage). RCT: randomized controlled trial [**Source**: Transl. Stroke Res. (2020)].

Recent research innovations in highthroughput technologies has paved the way to advance our understanding of the pathophysiology of this complex disease. It has allowed generating a significant amount of data and information at different molecular levels. It is believed that the integration of these multi-omics data can be leveraged to study thousands of proteins (proteomics), genes (genomics), RNAs (transcriptomics) and metabolites (metabolomics). Concurrent studies of these multi-omics data are expected to lead to revealing interaction networks between the molecular levels. It is believed that the strategy based on integrated analysis of multi-omics data using proteomics, metabolomics, transcriptomics and genomics would pave the way to useful insight into stroke pathogenesis,

identification of therapeutic targets and biomarker discovery [12].

While different mechanisms are thought to be involved in the pathophysiology of stroke due to its complexity, there is an increasing evidence that suggests an important role of inflammation in the progression of stroke. This leads to a rapid growth in the research in therapeutic targeting of post ischemic inflammation in acute stroke as a potential neuroprotective strategy [11]. Especially, integrated analysis that combines proteomics. aenomics. transcriptomics and metabolomics is considered to extend to study proinflammatory cytokines that have been found altered after ischemic stroke and have shown a potential role in the detection and

modulation of inflammation. This could make them attractive candidates for both potential biomarkers and novel therapeutic targets for stroke management [10, 11].

#### The Occurrence and Role of Inflammatory Condition in Ischemic Stroke

The biochemical processes of pathophysiology of ischemic stroke of the brain correspond to a complex cycle that include interconnected molecular and cellular mechanisms (Figure 2) [11]. Previous studies have suggested that there is a vital role of an inflammatory condition at all stages of the ischemic cascade. This role involves right from the starting from the cessation of flow to the late regenerative processes that are associated with the repair of ischemic tissues [10, 11]. The main mechanism of cell death in cerebral infarction is supposed to be the energy deficit caused by the loss of neurons ability to synthesize ATP (adenosine triphosphate). Further, it has been observed that while there is an increased extracellular concentration of K<sup>+</sup> ions and the uncontrolled influx of Na<sup>+</sup>, Ca<sup>2+</sup>, Cl<sup>-</sup> ions, the activity of Na<sup>+</sup>/K<sup>+</sup> - ATPase decreases in this process. Subsequently, loss of membrane potential along with gradual depolarization of the cell membrane occurs. This results in an increased flow of sodium that allows for transport of water to cells by an osmotic process, which leads to the development of cytotoxic edema. Consequently, loss of membrane integrity and cell death occurs as a result of the accumulation of Na<sup>+</sup> and Ca<sup>2+</sup> ions leading to organelle degeneration. Catabolic enzymes are activated by the influx of Ca<sup>2+</sup> ions. This happens by the production of arachidonic acid and increasing the

production of ROS mainly in neurons (Figure 2) [11, 14, 15].

An increased production of ROS triggers a cascade of events. First, it enables the passage through the membrane of ions and low-molecular-weight substances that cause necrosis. Secondly, depending on the degree of neurons damage, it leads to a programmed death of the nerve cell. Further, studies have shown that the excitotoxicity and an increased ROS results in activating microglia and astrocytes, which secrete cytokines, chemokine and matrix metalloproteinases (MMP) (Figure 2) [10, 11]. Consequently, these inflammatory mediators have been shown to induce the expression of cell adhesion molecules on the endothelial surface (P-selectin, E-selectin, endothelial-leukocyte adhesion molecule (ELAM-1) and intercellular cell adhesion molecules-1 (ICAM-1)), which have been attributed as enabler of the neutrophils to infiltrate ischemic areas of the brain [10].

Furthermore, this cascade leads to the release of hazard signals by necrotic neurons that in turn activates the immune system and secrete molecular patterns (DAMP). DAMP then induces the Toll-like receptors (TLRs) on microglia that stimulates the NF-kB to synthesize most of the pro-inflammatory cytokines (Figure 2) [11]. Additionally, NLRP3 inflammasome activates caspase-1 that subsequently results in maturation and secretion of IL-1ß and IL-18 cytokine. Proinflammatory cytokines (IL-1ß, interleukin 6 (IL-6), IL-18 and tumor necrosis factor (TNFα)) and chemokines are produced by M1 microglia, while M2 microglia produce antiinflammatory cytokines (interleukin 10 (IL-

10), interleukin 4 (IL-4) and transforming growth factor-beta (TGF- $\beta$ )). These antiinflammatory cytokines are released a few days after acute brain damage that lead to inhibition of inflammation [16, 17].



*Figure 2:* A schematic depiction of the mechanisms of pathophysiology and the role of proinflammatory cytokines involving ischemic brain damage [**Source:** Clin Interv Aging. (2020)].

As Figure 2 schematically illustrates, MMP which is secreted by microglia, astrocytes

and neurons mediates in the destruction of the base plate. This results in increasing BBB

permeability that facilitate the entrance of additional peripheral immune cells into the area affected by stroke. brain This phenomenon is attributed to two factors. It is suggested that acute ischemia of the brain not only causes a local inflammatory reaction, but it also triggers a systemic immune response [11]. Studies have shown that both congenital and adaptive immunity are involved during the stroke. However, they do not affect the acute phase of damage. But, according to latest theories, it is considered that modulation of adaptive immunity can have a significant protective effect on the ischemic brain. This creates bright prospects for new stroke therapies. However, it is also speculated that such immunomodulation could produce some side effects as well. Therefore, it is thought that a better understanding of the interaction between the immune system and the ischemic brain is essential that can help exploit the full therapeutic potential of stroke immunology [10, 14-17].

#### Immune System in the Production of Pro-Inflammatory Cytokines

A significant amount of research is currently clinical underway that involves and experimental studies to gain insights into the complex role of the immune system in pathophysiological changes that occur after acute ischemic stroke of the brain. These studies are targeted to better understand molecular signals that are generated by cerebral ischemia. Consequently, the activated components of the congenital immune system lead to amplification of inflammatory cascades and tissue damage, while these simultaneously processes

stimulate a potentially harmful and cytoprotective immune adaptive response [11, 18].

There are a number of cells that are involved in the production of cytokines and are representatives of the congenital and adaptive immune response. These include microglia that are a type of resident marrow cells of central nervous system (CNS). Microglia's main roles correspond to removing debris, repairing damaged cells and also acting as CNS homeostasis regulator. They produce pro-inflammatory mediators such as TNF- $\alpha$ , IL-1 $\beta$ , IL-6 and ROS, NO after a stroke (Figure 2). Notably, studies have suggested their important contribution to the inhibition of inflammation and tissue repair [17, 19, 20]. Also, it has been studied that perivascular macrophages occur between the vascular primarv membrane and the brain surface, where M1 macrophages are known to produce proinflammatory cytokines (IL-1β, interleukin 12 (IL-12), interleukin 23 (IL-23) and TNF- $\alpha$ ), chemokine, ROS and NO (Figure 2), which promote the immune response of helper T cells type 1 (Th1) [17]. Further, mastocytes or mast cells (MC) that are found in meninges and cerebral vessels have been found to store vasoactive substances (histamine), anticoagulants (heparin), protease (MMP2, MMP9) and cytokines (TNF- $\alpha$ ). It has been reported that they are capable of phagocytosis, antigen presentation that can modulate the adaptive immune response [21]. Two different populations of tissue macrophages classical are and proinflammatory monocytes that participate in the production of TNF-α [17]. It has been observed that after ischemia of the brain,

quick recruitment of inflammatory monocytes happens to the site of damage where they activate macrophages [21].

Researchers have shown а combination of congenital and adaptive immunity dendritic cells (DC) produce antigen-presenting cells (APC) that generate antigen, containing a MHC class II antigen complex. The presentation of T-cells antigen CD4+ enables a process of clonal expansion in lymphatic organs promoted by autocrine interleukin 2 (IL-2) production [17]. Further, it has been shown that T cells cause ischemic damage after entering the brain parenchyma. This happens when T cells release proinflammatory agents and cytokines such as ROS, interferon  $\gamma$  (IFN- $\gamma$ ), TNF- $\alpha$ , IL-1 $\beta$ , interleukin 17 (IL-17) and interleukin 21 (IL-21). Auxiliary T cells (Th) are known to be responsible for coordinating and modulating immune responses and depending on molecular signals, different subpopulations of Th-cell effectors may develop that can produce cytokines. Type 1 (Th1) cells TNF produce IFN-y and and their development is supported by IL-12. These cells have been observed to produce cytotoxic cytokine IL-17 that contribute to acute ischemic brain damage [17, 22-24].

# Which Inflammatory Cytokines can be targeted to Develop Stroke Biomarkers and Therapeutics for Clinical Trials?

A large body of research evidence suggests that ischemic stroke condition could be attributed to the induction of cytokines and chemokines that may contribute to the pathogenesis of adverse remodeling, and systolic and diastolic dysfunction. It has been discussed in detail in many research reports

that inflammatory cytokines modulate phenotype and function of all myocardial cells, suppressing contractile function in cardiomyocytes, inducing inflammatory activation in macrophages, stimulating microvascular inflammation and dysfunction, and promoting a matrix-degrading phenotype in fibroblasts [25]. Therefore, the research focus has been on the important proinflammatory cytokines, interleukin-1 (IL-1), which is produced in the CNS by microglia, astrocytes, endothelium and neurons. This cytokine has two different forms that include one intracellular (IL-1α) and one secreted (IL- $1\beta$ ). It is believed that both forms act through the IL-1 type I receptor (IL-1RI), which is expressed by immune and endothelial cells [10]. Further, the pro-inflammatory cytokines tumor necrosis factor (TNF)-α, interleukin (IL)-1, and IL-6 have been extensively implicated in the pathogenesis involving ischemic stroke and it is thought that they could potentially be very effective stroke biomarkers as well as novel therapeutic targets [10, 25].

Recent research has suggested that TNF- $\alpha$ -mediated adverse remodeling and heart failure progression may involve effects on cardiomyocytes, macrophages, and the extracellular matrix [25]. The inhibition of TNF- $\alpha$  seems to be a promising tool for stroke treatment. In a clinical trial with targeting the IL-1 system in human heart failure patients, treatment of patients with prior myocardial infarction and evidence of active inflammation with the anti-IL1 $\beta$ monoclonal antibody demonstrated favorable effects that showed reduced risk of the composite endpoint (non-fatal myocardial infarction, non-fatal stroke or death) by 15% in comparison with standard treatment. The study supported the role of IL-1 $\beta$  in the pathogenesis of atherothrombotic disease and also generated observations that may be important in understanding and treating heart failure [26-28]. These data suggest high

promise of IL-1 and TNF- $\alpha$  and to corroborate the findings, it could be worth to perform a clinical trial in ischemic stroke patients to evaluate the biomarkers and therapeutic potential of inhibiting IL-1 and TNF- $\alpha$  [10, 25].



Figure 3: A schematic depiction of IL-6 signaling and cellular actions in the failing heart. IL-6 has been known to be with both pro- and anti-inflammatory actions that can stimulate fibroblast proliferation and ECM synthesis, and promotes cardiomyocyte hypertrophy. [**Source:** Cardiovasc Drugs Ther (2020)].

Another inflammatory cytokine, IL-6 has been shown to exhibit important modulatory effects on several different cell types in heart failure. Both pro- and anti-inflammatory effects of IL-6 are observed. Trans-signaling has been suggested to exert pro-inflammatory actions, whereas canonical IL-6 signaling may be anti-inflammatory [25]. Recent studies have shown that IL-6 exerts negative inotropic effects in cardiomyocytes and promotes a hypertrophic response through the gp130/STAT3 pathway. However, it has been observed that it may also exert protective actions, mediated through preservation of mitochondrial function (Figure 3) [25]. For example, IL-6 promotes proliferation and stimulates ECM synthesis in fibroblasts. It is believed that the combined effects of IL-6 on cardiomyocytes and fibroblasts could play an important role in the pathogenesis of HFpEF. Additionally, it has been reported that IL-6 has potent modulatory effects on macrophages and lymphocytes. The in-vivo role of IL-6 in inflammation has been found to be dependent on the relative contribution of macrophage and lymphocyte subsets [29-34].

Studies related to targeting IL-6 in human failure have heart suggested extensive evidence of a crucial role of IL-6 in the post-inflammatory hepatic acute phase response, which is implicated in autoimmunity. Based on this data, IL-6 has been considered by the researchers as an attractive therapeutic target in many conditions associated with inflammation [25]. In a clinical trial, the IL-6 receptor neutralizing antibody tocilizumab was approved as effective therapy for patients with moderate to severe rheumatoid arthritis and temporal arteritis, and for the treatment of the cytokine release syndrome associated with CAR-T therapies. Therefore, experimental cell evidence demonstrating involvement of IL-6 signaling in the pathogenesis of heart failure suggested has that treatment with tocilizumab may also be effective in heart failure patients [25, 34]. Thus, all data suggest that IL-6 could potentially be a good biomarker for stroke diagnosis and prognosis. However, further studies need to be done to better evaluate its therapeutic effect [10].

With respect to other antiinflammatory cytokines, such as IL-4 and IL-10, all experimental studies seem to indicate that the administration of IL-10 has a neuroprotective effect, but its application as a new stroke therapy needs to be further studied before its translation to the clinical setting [10].

#### Conclusion and Future Perspective

Recent research advances in stroke have clearly implicated inflammation in the pathophysiology of stroke. Research evidence points out to the dual role of inflammation during ischemic stroke, which displays both beneficial and also adverse effects according to the phase of the stroke. Anti-inflammatory strategies have been suggested as promising tools for regulating the inflammatory processes that take place during the acute phase of stroke. To this end, several therapeutic strategies have been shown to be effective in both experimental models of ischemia and clinical trials. However, the realization of translation from bench-to-bedside is yet to happen in stroke research. In this context, the current focus is to gain a better understanding of the inflammatory mechanisms involved during ischemic stroke that would make the inflammatory cytokines as not only possible biomarkers but potential therapeutic targets. The knowledge of modulation of the possible therapeutic targets could be leveraged in the fight against the disease and its complications.

As regards the future studies, one area could be developing sensitive and rapid blood tests to diagnose and aid stroke prognostication, which is currently lacking and a major cause of concern world-wide. A number of advantages could result from the elaboration of robust biomarkers for stroke. To this end, future research might focus on their discovery. Effective stroke biomarkers would not only provide potential methods for stroke diagnosis and prognosis, but they would also contribute to the understanding of the pathophysiology of this devastating disease.

As the previous studies have shown, the inflammatory response to stroke is extremely complex, with multiphasic proinflammatory responses. So, future studies may focus on the determination of cytokine levels in the acute phase of stroke that can contribute to a thorough understanding of their role in the immune response observed during brain ischemia. Such an advancement can also be leveraged for the identification of interactions between the immune system and brain cells during post-stroke recovery and regeneration. Therefore, they would serve as prognostic factor in acute cerebral ischemia. With respect to inflammatory cytokines to be developed as biomarkers, IL-1 and IL-6 potentially appear as good prognostic factor

in stroke. However, further research is necessarv to assess the therapeutic implications of these cytokines. Further, research could focus on facilitating experimental models based on inhibition of TNF- $\alpha$ , CCL-2, CCL-3 and CCL-5 that could be a promising tool in the treatment of acute stroke. Moreover, as the current research data indicate. administration of the recombinant human IL-1 receptor antagonist may have important implications in reducing the inflammatory response. However, more clinical trials in patients with ischemic stroke are necessary in order to evaluate the therapeutic the cytokines potential inhibition.

For successful practical applications of inflammatory cytokines-based stroke management, successful implementation of cytokine targeting approaches in heart failure patients is essential. This will require dissection of specific pro-inflammatory pathways with a critical role in dysfunction and progression of adverse remodeling. This patient also includes identification of subpopulations with dysregulated or overactive inflammatory responses that may derive maximal benefit from targeted cytokine or chemokine inhibition in effective stroke treatment.

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# **Biotechnology Advances around the World**

# **Editor's Picks**

Every issue of Biotechnology Kiosk presents select latest research news picked by the editorsin-chief on significant research breakthroughs in different areas of biotechnology around the world. The aim is to promote further R&D in all of these cutting-edge areas of biotechnology. The editors have compiled and included the following innovations and breakthroughs to highlight the latest biotechnology advances.



Dr. Megha Agrawal Co Editor-in-Chief





Dr. Shyamasri Biswas Co Editor-in-Chief

#### **Targeted Cancer Drugs**

# Experimental drug agent pentagamavumon-1 (PGV-1) for stopping cancer growth

Turmeric is known as a popular spice that is used for flavoring. It is also known as a useful medicine for its anti-inflammatory and even anti-cancer benefits. Some previous studies have shown that the medicinal benefits of turmeric come from the compound curcumin, which is sold as an herbal supplement. Curcumin's anti-cancer properties have been studied. These studies have revealed that the high doses of curcumin are required. However, incomplete understanding of the chemical process involving therapeutic actions of curcumin have restricted efforts for its use in cancer therapy.

One way to stop cancer growth is suggested by suppressing the enzyme activity. This leads to the reactive oxygen species that are allowed to cause stress on cells. This ultimately results in cell death. Similar to this process, many anti-cancer drugs operate to control cancer growth. However, these drugs sometimes have with severe side-effects that occur due to stress on healthy cells.

Scientists in Japan and Indonesia reported in a new study how an experimental drug agent stops cancer cells from growing. Their research was published in Scientific Reports (Pentagamavunon-1 (PGV-1) inhibits ROS metabolic enzymes and suppresses tumor cell growth by inducing M phase (prometaphase) arrest and cell senescence. *Scientific Reports*, 2019; 9 (1) DOI: <u>10.1038/s41598-019-51244-3</u>). Pentagamavumon-1 (PGV-1) was first reported some time back as an analogue of a molecule found in turmeric that was later shown to have anti-cancer effects. In this new study, researchers conducted tests on cancer cells and animals. The study revealed that these anti-cancer effects were due to PGV-1 that inhibited a series of enzymes responsible for the metabolism of reactive oxygen species. It is reported that PGV-1 arrests cells in the cell cycle at M phase and it inhibits many ROS-metabolic enzymes. Further, this arrest is believed to prevent the cancer cells from dividing, and the subsequent enzyme inhibition causes the cancer cells to die. This finding is expected to lead to better understanding on the possible modifications to PGV-1 for its use for cancer treatment.

Further, researchers showed that PGV-1 was effective on numerous types of cancers and interestingly, when administered to mice injected with human cancer cells, the mice showed no evidence of the cancer along with no side-effects. Additionally, the anti-cancer effects were shown to persist even after the cessation of PGV-1 administration unlike other anti-cancer drugs.

In view of the high drug efficacy of PGV-1 along with very low amount of side effects in animals, researchers envisioned PGV-1 to be of pharmaceutically relevant therapeutics that could be developed as an orally administered drug for cancer.

#### **Cardiovascular Disease**

#### Potential therapeutic target for clogged arteries

It is known that cardiovascular disease arises mainly as a result of lipids from the blood plasma that are deposited in the walls of blood vessels, which ultimately restricts blood flow. This complex and deadly disease is believed to affect about a third of the world's population. It is critically important to advance the knowledge and understanding of the mechanisms that regulate the levels of blood lipids.

Studies have shown that fat molecules that are also called lipids, for example, cholesterol and triglycerides move back and forth around the circulatory system. This shuttling of fat molecules are caused by a protein called Apolipoprotein-B (ApoB). Further, researchers have named these complexes of lipid and protein as ApoBcontaining lipoproteins. These proteins are considered to be essential for transporting lipids from the intestine and liver to the tissues of the body. However, for the reason that they can also cause cardiovascular disease, they are known as bad cholesterol.

In a new research, researchers in the US reported a protein critical for the synthesis of ApoB-containing lipoproteins. This protein, called MTP, or microsomal triglyceride transfer protein, is shown to be highly conserved in animals, from insects to humans. Researchers have shown that MTP loads lipids onto ApoB, a key initial step in the synthesis of ApoB-containing lipoproteins. Their study was published in PLOS Genetics (A point mutation decouples the lipid transfer activities of microsomal triglyceride transfer protein. *PLOS Genetics*, 2020; 16 (8):

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#### DOI: 10.1371/journal.pgen.1008941).

From previous studies, it is known that MTP can transfer different types of lipids to ApoB, including triglycerides. Thev correspond to a major source of energy. These phospholipids are known as the building-blocks of membranes in the cell. In their study, researchers revealed that a mutation in MTP blocks the loading of triglycerides, does but it not block phospholipids onto ApoB. The separation of these two transfer functions was observed for the first time and was unexpected. This finding is important for the reason that high triglyceride levels in lipoproteins are correlated with bad clinical, which results in diseases such as diabetes and heart disease.

Other identified mutations in MTP preventing both transfer functions of the protein have been reported that can cause a malabsorption syndrome. This makes the intestines to have difficulty absorbing fats and fat-soluble vitamins from the diet, which can result in gastrointestinal distress or more serious problems, such as malnutrition or severe weight loss. However, in this new study that employed zebrafish using the newly identified mutation did not exhibit malabsorption or growth defects. It was attributed to the ability to transfer phospholipids to make ApoB-containing lipoproteins.

MTP has been an important subject of research for its possible role as potential

therapeutic target to help lower triglyceride levels in the blood and prevent cardiovascular disease. But it could not meet the desired goals due to the fact that the existing chemical inhibitors of MTP were proven to be effective and block all MTP function. Such blocking can cause intestinal malabsorption and fat а dangerous accumulation of fat in the liver. This new study paves the way for breakthroughs in the design of more specific MTP inhibitors that can mimic new mutation and selectively block triglyceride transfer to ApoB. The data presented in this study suggests that this type of inhibitor could potentially reduce circulating triglyceride levels without affecting intestine and liver.

Compiled and Edited by Dr. Megha Agrawal & Dr. Shyamasri Biswas.



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