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Division of Cardiology

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General Summary

We have 6 research groups for covering the broad field of cardiology. In respective study groups, we have been studying the problems that face us in clinical practice. Our research is based on clinical studies that use the large database we have been developing. In specific, we recently used covariance structure analysis as a new solution for action assignments. Basic research is also performed to solve clinical questions.

Research Activities

Ischemic Heart Disease Research Group

We have converted patients' data, including risk factors and coronary lesion morphology, hemodynamic data, from cardiac catheterization examinations and treatments in patients with ischemic heart disease, into our large, precise database. Using this database, we have been performing a study comparing risk factors, clinical outcomes, and other data. We have recently reported the differences of risk factors between coronary organic stenosis and acute coronary syndrome by using covariance structure analysis. We have been using a similar method to study a contributing pattern of obesity to ischemic heart disease. In the analysis, we reported a possible risk of low-reactivity of natriuretic peptide. Furthermore, since fractional flow reserve is reportedly a good method for evaluating significant coronary stenosis, we have been collecting and analyzing information about the clinical data of fractional flow reserve (FFR).

Arrhythmia Research Group

In our arrhythmia team, we have been focusing on the curative treatment of atrial fibrillation. Our research activities include the comparison of efficiency and safety among different ablation methods (radiofrequency vs. cryoballoon ablation), factors associated with the occurrence of pulmonary vein stenosis following balloon ablation, and the association of ablation methods and asymptomatic cerebral ischemia.

Heart Failure Research Group

1. Study group on heart failure

Since last year, we have constructed and updated a database of approximately 3,000

patients who have been hospitalized for cardiac catheterization and treatment. Regarding this database, analysis is conducted using the Structural Equation Modeling (SEM) or Covariance Structure Analysis by adjusting the statistical analysis method, AMOS (Analysis of Moment Structures), which has been previously known but the use of which has been rarely reported in the cardiovascular field. In particular, we analyze the interrelationships of clinical factors that cannot be expressed by multivariate statistical analysis alone, focusing on statistical analysis research on clinical data including plasma BNP. Using a path diagram, it is easier to have a visual understanding of the relationships between each factor, and to perform multiple regression analysis and path analysis (repetition of multiple regression analysis) using factors and confirmatory factor analysis. Last year, with respect to the relationship between obesity and BNP, we reported the relationship between the change in BNP concentration before and after treatment along with the change in body weight. This spring, we published an article on the influence of remodeling changes in the left ventricular cavity on BNP concentration. By publishing articles on wide-ranging analysis results including detailed data analysis of the disease state of chronic heart failure along with the relationships between various valvular diseases and atrial fibrillation, we will continue to promote a wide range of clinical studies based on the experience gained from daily clinical practice. We are also continuing our efforts to clarify the mechanism of these findings via fundamental research.

Imaging Research Group

1. Study group on imaging

With the increasing number of cases involving transcatheter aortic valve replacement (TAVR) since last year, cardiac CT and echocardiograms are also vital as preoperative examinations for evaluating the aortic valve. From this valuable case information, we are seeking research agendas for clinical studies. Through other imaging modalities such as cardiac MRI and myocardial isotope tests, we are continuing to seek research agendas for clinical studies and conduct analyses on cardiomyopathy and arrhythmias.

Molecular Biology Research Group

Glucose becomes an important preferential substrate for cardiac metabolism and ATP generation during ischemia-reperfusion injury (IRI). Therefore, acceleration of glucose uptake and its metabolism is critical for myocardium to develop ischemic tolerance. Although insulin plays a pivotal role in this process, we have recently reported that insulin resistance increases during ischemic attack of acute coronary syndrome (ACS). The study also suggested that there are endogenous mechanisms of promoting glucose metabolism. One of the potential mechanisms is sodium-glucose co-transporter 1 (SGLT1). A study of Langendorff murine heart perfusion demonstrated that the inhibition of SGLT1 during IRI reduces glucose uptake into the myocardium, leading to a decrease in the cardiac tissue ATP content. As a consequence, cardiac functional recovery after IRI was impaired by SGLT1-inhibition. The present findings provide the significant role of SGLT1 in optimizing cardiac energy metabolism during IRI.

Cardiac Physiology Research Group

We have demonstrated that thrombin, the final product of the coagulation cascade, is present in the heart. Coagulability is increased in patients with dilated cardiomyopathy (DCM). Using knock-in mice that have a cardiac troponin T deletion mutation that causes human DCM ($\Delta K210$ knock-in mouse) (B6; 129-Tnnt2^{tm2Mmt0}). We assessed the effects of a direct thrombin inhibitor, dabigatran, in $\Delta K210$ knock-in mice. Dabigatran significantly improved fractional shortening in echocardiographic findings and survival outcomes. In conclusion, tissue thrombin is involved in the pathogenesis of DCM, and thrombin inhibition can be beneficial for the treatment of DCM.

To investigate this mechanism, we practiced microarray analysis, which has demonstrated that Casq1·Postn·Myh7 may be involved in the mechanism. We further practiced Western blot analysis, but there were no significant differences between hearts of DCM mice and Wild type mice in these three gene protein products. We next investigated the apoptosis by the TUNEL assay. The apoptotic index, the percentage of TUNEL-positive nuclei, was significantly increased in the DCM group in comparison to that observed in the Wild group. Treatment with dabigatran significantly reduced the apoptotic index. The apoptosis may be involved in the mechanism.

Publications

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