

Genetic Regulation

1. Staffs and Students (in 2009)

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|-------------------------------------------------------------|---------------------------------------------------------------------|
| Professor | Akinori KIMURA |
| Associate Professor (Graduate School of Biomedical Science) | Toshiaki NAKAJIMA |
| Assistant Professor | Takuro ARIMURA |
| Research Associate | Taeko NARUSE |
| Graduate Student | Daisuke SHICHI, Haruna OHTSUKA |
| Graduate Student (Biomedical Science PhD program) | Taisuke ISHIKAWA, Hitoshi OHTANI, Jianbo ANN, Yusuke SAITO |
| Visiting Graduate Student | Marika TAKAHASHI |
| Visiting Post-doc | Zhiyong CHEN |
| Visiting Student (Nanyang Technology University) | Hui Chiang LAU |

2. Purpose of Education

Genetic factors, i.e. structural and/or functional diversity of human genome, are more or less involved in the etiology and pathogenesis of human diseases. Main objective of Genetic Regulation is to identify the gene mutations or polymorphisms and to decipher the molecular mechanisms involved in the etiology and pathogenesis of intractable diseases, in order to develop new strategies for diagnosis, treatment and/or prevention of the diseases. Current research is focused on the cardiovascular diseases (e.g. idiopathic cardiomyopathy, idiopathic arrhythmia, and coronary heart disease), autoimmune diseases (e.g. Burger disease, Behcet disease, rheumatoid arthritis, and chronic thromboembolic pulmonary hypertension) and infectious diseases (e.g. HIV/AIDS). In addition, genome diversity in immune-related genes is investigated from the view-point of primate evolution.

3. Research Subjects

- 1) Identification and functional analysis of disease-related genes for cardiovascular diseases
- 2) Identification and functional analysis of disease-related genes for autoimmune diseases
- 3) Identification and functional analysis of disease-related genes for infectious diseases
- 4) Structural, functional and evolutional analyses of MHC and immune-related genes in vaccination

4. Publications

Original Article

1. Arimura T, Hayashi YK, Murakami T, Oya Y, Funabe S, Hirasawa EA, Hattori N, Nishino I, Kimura A. Mutational analysis of fukutin gene in dilated cardiomyopathy and hypertrophic cardiomyopathy. *Circ J.* 2009; 73(1): 158-161.
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 9. Romphruk AV, Romphruk A, Naruse TK, Raroengjai S, Puapairoj C, Inoko H, Leelayuwat C. Polymorphisms of NKG2D ligands: diverse RAET1/ULBP genes in Northeastern Thais. *Immunogenetics*. 2009; 61(9): 611-617.
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 11. Arimura T, Inagaki N, Hayashi T, Shichi D, Sato A, Hinohara K, Vatta M, Towbin JA, Chikamori T, Yamashina A, Kimura A. Impaired binding of ZASP/Cypher with phosphoglucomutase 1 is associated with dilated cardiomyopathy. *Cardiovasc Res*. 2009; 83(1): 80-88.
 12. Moulik M, Vatta M, Witt SH, Alora AM, Murphy RT, McKenna WJ, Boriek A, Oka K, Labeit S, Bowles NE, Arimura T, Kimura A, Towbin JA. ANKRD -the gene encoding cardiac ankyrin repeat protein- is a novel dilated cardiomyopathy gene. *J Am Coll Cardiol*. 2009; 54(4):325-333.
 13. Arimura T, Bos MJ, Sato A, Kubo T, Okamoto H, Nishi H, Harada H, Koga Y, Moulik M, Doi YL, Towbin JA, Ackerman MJ, Kimura A. Cardiac ankyrin repeat protein gene (ANKRD1) mutations in hypertrophic cardiomyopathy. *J Am Coll Cardiol*. 2009; 54(4): 334-342.
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 18. Hinohara K, Ohtani H, Nakajima T, Sasaoka T, Sawabe M, Lee BS, Ban J, Park JE, Izumi T, Kimura A. Validation of eight genetic risk factors in East Asian populations replicated the association of BRAP with coronary artery disease. *J Hum Genet*. 2009; 54(11), 642-646.
 19. Sato A, Arimura T, Makita N, Ishikawa T, Aizawa Y, Ushinohama H, Aizawa Y, Kimura A. Novel mechanisms of trafficking defect caused by KCNQ1 mutations found in long QT syndrome. *J Biol Chem*. 2009; 284(50): 35122-35133.

Review Article

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2. Kimura A. Does a gene polymorphism predisposing to the intermediate phenotype predict the risk of disease? *Circ J*. 2009; 73(6): 1016-1017.