[Grant-in-Aid for Scientific Research(S)]

Integrated Science and Innovative Science (New multidisciplinary fields)



Title of Project: Mammalian-specific genomic functions

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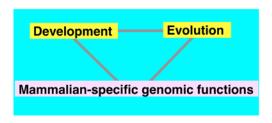
Research Area: New multidisciplinary

Keyword: Genomic function, Genome evolution

[Purpose and Background of the Research]

Viviparity and placentation are representative characters in mammalian reproduction system. We have recently demonstrated that retrotransposon-derived genes specific mammals, Peg10 and Peg11/Rtl1, play essential roles in placenta formation and functions. These are paternally expressed imprinted genes relating early embryonic lethality and late fetal/neonatal lethality, respectively. Genomic imprinting is one of mammalian-specific epigenetic mechanisms and known to essential for mammalian development. this mechanism emerged and how these mammalian-specific genes are acquired from retrotransposons? These questions are related to the origin or evolution of the mammals. The aim of this project is to dissolve these questions.

Know mammals Learn from mammals



Mammalian-specific genes (Newly acquired genes)
 Retrotransposon-derived genes

•Mammalian-specific epigenetic mechanisms (Altered gene expression) Genomic imprinting, X chromosome inactivation

[Research Methods]

Our project comprises four subjects as below:

- 1. Biological function of *Peg10* identification of functional domain of Peg10 protein and production of parthenogenous in mice.
- 2. Role of *antiPeg11/Rtl1*: biological functions of essential miRNAs in *antiPeg11/Rtl1*. Its roles in human pUPD14 and development of new miRNA therapy.
- 3. Biological function of *Sirh* genes: Analyses on *Sirh3*, *4-6*, *7* and *9* knockout mice.
- 4. Reprogramming of genomic imprinting memories in germ cells: molecular mechanism of DNA demethylation in PGCs.

[Expected Research Achievements and Scientific Significance]

Elucidation of evolution of mammals, a part of our history as human beings. Elucidation of the meechanism of epigenetic reprogramming for promoting regenerative medicine.

[Publications Relevant to the Project]

- 1. Kaneko-Ishino T and Ishino F. Retrotransposon silencing by DNA methylation contributed to the evolution of placentation and genomic imprinting in mammals. Develop Growth Differ **52**(6), 533-543 (2010).
- Sekita Y, Wagatsuma H, Nakamura K, Ono R, Kagami M, Wakisaka-Saito N, Hino T, Suzuki-Migishima R, Kohda T, Ogura A, Ogata T, Yokoyama M, Kaneko-Ishino T and Ishino F. Role of retrotransposon-derived imprinted gene, *Rtl1*, in the feto-maternal interface of mouse placenta. Nat Genet 40(2), 243-248 (2008).
- 3. Ono R, Nakamura K, Inoue K, Naruse M, Usami T, Wakisaka-Saito N, Hino T, Suzuki-Migishima R, Ogonuki N, Miki H, Kohda T, Ogura A, Yokoyama M, Kaneko-Ishino T and Ishino F. Deletion of *Peg10*, an imprinted gene acquired from a retrotransposon, causes early embryonic lethality. Nat Genet **38**(1), 101-106 (2006).

Term of Project FY2011-2015

Budget Allocation 165, 200 Thousand Yen

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