32nd WPI Seminar

"Next generation sequencing: how you can use it for your own research"

Recent advances in next-generation sequencing (NGS) technologies allow the rapid identification of disease-causing mutations. The examples of NGS experiments in our laboratory, including the successful identification of the location of transgenes in transgenic mice and the use of a bioinformatics approach to identify candidate mutations in a mouse model of atopy-related disorders, will be discussed.



Speaker: Dr. Emiko Noguchi

Department of Medical Genetics, University of Tsukuba

Date: Thursday, June 26, 2014
Time: 12:00-13:00
Venue: Room #402, 4F, Health and Medical Science Innovation Laboratory, University of Tsukuba

☆Light refreshments will be served.

INTERNATIONAL INSTITUTE FOR INTEGRATIVE SLEEP MEDICINE

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