



グローバル COE セミナーのご案内

Mutational Profiling in Cancer by Next Generation Sequencing

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Associate Professor

David A. Wheeler, Ph.D. is an Associate Professor in Molecular and Human Genetics at Baylor College of Medicine. His interests are in the discovery, analysis and interpretation of genetic variation using high throughput DNA sequencing methodologies. He has worked extensively in the genome sequencing field for 10 years, and in computational biology for 20 years. He has expertise in the production of DNA sequence, the storage, processing and distribution of the sequence data, and the analysis and interpretation of the data. He has been a key participant in many large-scale multi-institutional DNA sequencing projects such as the Human Genome Project, the ENCODE and HapMap projects, and The Cancer Genome Atlas (TCGA) project. Dr. Wheeler is currently principal investigator of a Genome Characterization Center within the TCGA.

December 7th, Monday 16:00~

医学研究科 F 棟 1 階 セミナー室

Building F, 1F, Seminar Room

連絡先 病態生物医学分野

(Department of Pathology and Biology of Diseases)

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