

Genome Sequence of *Acinetobacter baumannii* TYTH-1

劉明麗, 劉志俊, Chia-Wei Lu, Ming-Feng Hsieh, Kai-Chih Chang, Han-Yueh Kuo, Chi-Ching Lee, Chun-Tien Chang, Cheng-Yao Yang, Chuan Yi Tang

Bioinformatics

Computer Science and Informatics

ccliu@chu.edu.tw

Abstract

*Acinetobacter baumannii* has emerged recently as a major cause of health care-associated infections due to the extent of its antimicrobial resistance and its propensity to cause large nosocomial outbreaks. Here we report the genome sequence of *Acinetobacter baumannii* TYTH-1 isolated in Taiwan during 2008.

Keyword : *Acinetobacter baumannii*, genomes