[Grant-in-Aid for Scientific Research (S)] Biological Sciences (Medicine, Dentistry, and Pharmacy)



Title of Project : Development of Novel Anti-Infectious Drugs Exhibiting Therapeutic Effects

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Research Project Number: 15H05783 Researcher Number: 90126095

Research Area: Environmental and hygienic pharmacy

The assay systems we develop will then be used to

screen for inhibitors.

Keyword: Microbiology and infectious diseases, Pathogenicity

| Keyword Microbiology and mectious diseases, ratiogenicity | |
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| [Purpose and Background of the Research] | [Expected Research Achievements and |
| The development of novel anti-infectious drugs | Scientific Significance |
| with therapeutic effects is urgently needed to | 1. Understanding bacterial pathogenesis |
| establish effective strategies against | Based on comprehensive analyses of data |
| multidrug-resistant pathogens. Current strategies, | obtained in this project, we will identify novel |
| however, are inadequate and the number of newly | genes responsible for bacterial pathogenesis. |
| discovered drugs has dramatically decreased, | Biochemical analysis of the functions of the |
| resulting in a very limited number of | products of the responsible genes will allow us to |
| anti-infectious drugs with novel mechanisms. One | uncover networks of gene expression involved in |
| possible reason for this is that the behavior of | bacterial pathogenicity. These findings will |
| pathogenic bacteria in test tubes differs | contribute to our understanding of bacterial |
| considerably from that in hosts. In this project, we | pathogenesis in the host. |
| will focus on gene products of pathogens that are | 2. Development of novel anti-infectious drugs |
| necessary for pathogenesis in the host | This project will identify novel genes necessary |
| environment. To achieve this goal, we will identify | for bacteria to survive in the host environment. |
| the genes in pathogens necessary for pathogen | Evaluating the products of these novel genes will |
| proliferation and pathogenesis in host animals. | lead to potential targets for drug development. |
| Based on the findings, we will establish screening | Bacterial growth inhibitors obtained by screening |
| systems to identify inhibitors against the gene | will be useful seed compounds for anti-infectious |
| products and establish a method for developing | treatments. |
| antibacterial agents with novel mechanisms of | |
| action. Our project also aims to elucidate the | [Publications Relevant to the Project] |
| molecular aspects of bacterial pathogenesis. | 1. Hamamoto H, Urai M, Ishii K, Yasukawa J, |
| | Paudel A, Murai M, Kaji T, Kuranaga T, Hamase |
| [Research Methods] | K, Katsu T, Su J, Adachi T, Uchida R, Tomoda H, |
| 1. Screening of pathogenic genes in bacteria using | Yamada M, Souma M, Kurihara H, Inoue M, |
| silkworms | Sekimizu K: <i>Nat Chem Biol</i> , 11, 127-133, 2015 |
| We have established silkworm infectious disease | 2. Kaito C, Saito Y, Ikuo M, Omae Y, Mao H, |
| models with human pathogenic bacteria. Using this | Nagano G, Fujiyuki T, Numata S, Han X, Obata K, |
| model, we will identify deletion mutants of | Hasegawa S, Yamaguchi H, Inokuchi K, Ito T, |
| pathogenic bacteria whose pathogenesis is | Hiramatsu K, Sekimizu K: <i>PLoS Pathog</i> , 9, |
| decreased compared with the wild-type strain. We | e1003269, 2013 |
| will also identify bacterial genes whose expression | 3. Kaito C, Kurokawa K, Matsumoto Y, Terao Y, |
| is appreciably increased in mouse organs compared | Kawabata S, Hamada S, Sekimizu K: Mol |
| with that in test tubes. The decreased pathogenesis | <i>Microbiol</i> , 56, 934-944, 2005 |
| of the gene deletion mutants will be confirmed in | |
| mouse infection models. | Term of Project FY2015-2019 |
| 2. Establishment of an assay system to screen novel | F |
| antibacterial agents | (Budget Allocation) 154,500 Thousand Yen |
| We will then establish methods for biochemical | • |
| analysis of the enzymes encoded by the pathogenic | [Homepage Address and Other Contact |
| genes identified in this study. Using these methods, | Information |
| we aim to elucidate the functions of the enzymes. The assay systems we develop will then be used to | http://www.f.u-tokyo.ac.jp/~bisei/ |
| The assay systems we develop will then be lised to t | |