



NATIONAL INSTITUTE OF GENETICS

Research Organization of Information and Systems

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Kosei Tanaka , Christopher S. Henry , Jenifer F. Zinner , Edmond Jolivet ,
Matthew P. Cohoon , Fangfang Xia , Vladimir Bidnenko , S. Dusko Ehrlich,
Rick L. Stevens , and Philippe Noirot : Building the repertoire of
dispensable chromosome regions in Bacillus subtilis entails major
refinement of cognate large-scale metabolic model
Nucleic Acid Res. in press
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