

CLASSICAL TO MODERN GENETICS

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Summary

The history and the research results obtained by the recent research by use of genetic approach were briefly summarized in the organisms, *Bacillus subtilis*, *Agrobacterium tumefaciens*, *Neurospora crassa*, *Aspergillus nidulans*, and *Lotus corniculatus* L. var. *japonicus* Regel, with special focus on *Neurospora crassa*, since the organism played important roles in the development of genetics and molecular biology. In *Bacillus subtilis* the molecular analysis of the process of sporulation is described. In *Agrobacterium tumefaciens*, which forms crown gall in the plant, the nature of Tumor inducing (Ti) plasmid is reported. In *Neurospora crassa* history and relatively precise

research results of genetics and epigenetics are summarized. Not only classical genetics but the recent development of epigenetics including repeat induced point mutation (RIP) mediated by the specific methylation of cytosine will provide new fields in the genetic analysis of higher eukaryote. *Aspergillus nidulans* shows specific parasexual (mitotic) genetic cycle, and *Lotus corniculatus* L. provided excellent systems to analyze the molecular mechanism of *Rhizobium*-Legume symbiosis

1. Introduction

During the development of the research on genetics and molecular biology, *Neurospora crassa* played very important roles. The process of development of early stages of research are well described in the preface of "The *Neurospora* Compendium, Chromosomal Loci, eds. Perkins, D. D. *et al.*, 2001." In it Dodge, B.O., who analyzed sexual cycle of *Neurospora crassa* in 1927, pioneered to establish the life cycle. He was very much impressed by the advantages for genetic analysis, and he tried to persuade Morgan, T.H. and his *Drosophila* group to adopt the organism. Morgan encouraged a graduate student, Lindegren, C. to examine potential nature of *Neurospora*. Ten years later the research results issued from Dodge, and Lindegren led Beadle, G.W. and Tatum, E.L. to adopt *Neurospora* in their research for nutritional mutants. The results endowed *Neurospora* to gain wide recognition, and to be the fungal counterpart of *Drosophila*. The work "Control of Biochemical Reactions in *Neurospora*" broke down the barriers that had separated biochemistry and genetics. The work of *Neurospora* went on to those in *Drosophila*, *maize*, and other higher eukaryotes. The *Neurospora* work was also expanded to other micro organisms, such as *Escherichia coli*, *Chlamidomonas*, *Aspergillus*, *Sordaria*, *Ustilago*, *Saccharomyces*, *Schizosaccharomyces*, *Coprinus*, *Podospora*, *Ascobolus*, and so on. In the present review the author would like to focus on *Neurospora crassa*.

2. *Bacillus subtilis*

Bacillus subtilis was characterized in 1876, which is Gram positive and grow in an aerobic condition. Under nutritionally rich condition it grew forming typical rod-shaped vegetative cells. However, under nutritionally deficient condition *B. subtilis* forms endospores.

At the beginning of 1960, it was recognized as a cellular differentiation system from vegetative cells to endospores, and provided excellent system to analyze the development of endospores by genetical methods.

Further it was discovered to be transformed by DNA, which supported the genetic analysis of sporulation.

2.1 Life cycle of *Bacillus subtilis*

As shown in Fig. 1, the life cycles of *Bacillus subtilis* are composed of two alternative growth phases, vegetative cell growth (Stage 0 and I) and entering into the process of sporulation (Stage, II~VII). Limitation for the source of carbon, nitrogen or phosphorous made the vegetative cells to entering into the developmental phase of

sporulation.

The life cycle of *Bacillus subtilis*, showing the separate phases of vegetative growth and sporulation. The vegetative cycle is proliferative and comprises alternating round of growth by elongation and central division. On starvation, the sporulation cycle is chosen. It begins with a modified cell division, in which an asymmetric septum is produced. A series of intermediate morphological states then ensues, leading to the production of a mature spore. Completion of the septum is defined as *Stage II*. The edges of the septal disc, which comprises a pair of lipid bilayers, migrate towards the pole of the cell, engulfing the smaller cell, known now as the prespore, within the cytoplasm of the larger “mother cell”. During *Stage IV*, a layer of cell-wall-like material, known as the cortex is synthesised in the space between the membranes surrounding the prespore. In *Stage V*, a multilayered proteinaceous coat is deposited on the outside surface of the spore. During the final period of development, known as “maturation”, the extreme resistance and dormancy properties that typify the mature spore develop, with little over change in morphology. Finally, the completed spore is released by lysis of the mother cell (*Stage VII*). The cycle is completed by spore germination and outgrowth of a new vegetative cell, which occurs when conditions suitable for the resumption of growth recur.

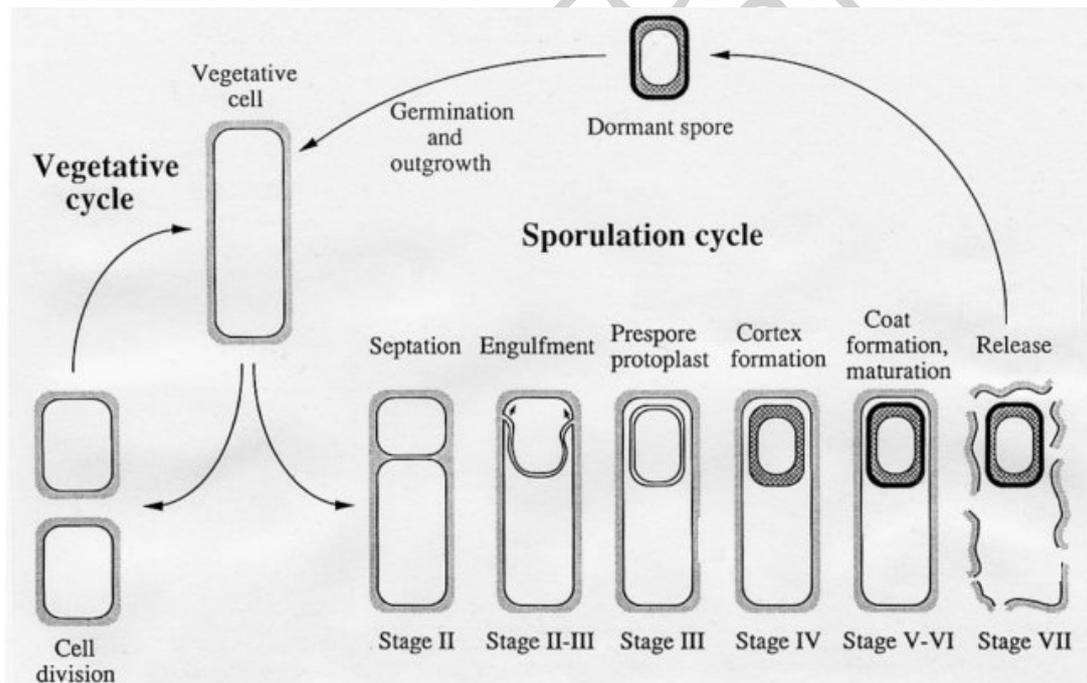


Figure1: The life cycle of *Bacillus subtilis* (Errington, J. 1992)

The processes of sporulation initiated on starvation are comprised of seven distinct morphological stages including 0, II, III, IV, V and VI as shown in Fig. 1. About 50 genes (*Spo*) regulating the sporulation are identified to drive the process.

3. *Agrobacterium tumefaciens*

As plant pathogens *Agrobacterium* was detected and isolated from soil. *Agrobacterium*

infected to wounded plant deliver the plasmid DNA into host plants, where a part of plasmid T-DNA is integrated to the DNA of host plants. *Agrobacterium tumefaciens* causes crown gall on the infected stem of the plants. *Agrobacterium rhizogenes* infects to plants, and proliferates in the root inducing the roots. The former harbor large (~200 kb) plasmids called Ti for tumor-inducing and the latter harbor Ri for root-inducing. A part of Ti plasmid transferred into plants is integrated in plant chromosome, which was designated as T-DNA (transferred DNA). The T-DNA expressed in the plant causes over-production of or enhanced-sensitivity of plant hormones resulting in the formation of crown gall tumors or in the induction of roots, and causes the production of opines, which is catabolized by the inciting bacteria.

The processes of infection of *Agrobacterium* are as follows:

Wounded plants produce acetosylingone, which chemotactically attract bacteria. The bacteria attach to the plant cell, the process of which is controlled by the chromosomal genes, *att chvA*, *B* and *exoC* controlling virulence in bacteria. The acetosylingone stimulate sensor kinase, *VirA*, which phosphorylates response regulator, *VirG* constituting a two-component system. *VirG* transcriptionally activate *virB*, *virC*, *virD*, *virE* and *virG* on Ti plasmid by attaching to *vir*-boxes of hexanucleotide motives in the promoter regions of *vir* genes. The concentrations of acetosylingone to attract bacteria to plant is two orders of magnitude lower than those to induce *vir* genes.

The proteins *VirD1* and *VirD2* control the processing of the T-DNA. *VirC1* protein controls the efficiency of T-DNA processing. *VirE2* protein is a single stranded DNA binding protein, and *VirE1* protein was suggested to support the stability of *VirE2* protein. The *virB* operon is composed of 11 genes, the proteins coded by the operon are essential for virulence.

4. *Neurospora crassa*

A filamentous fungus *Neurospora crassa* is an ascomycete, and heterothallic microorganism. The organism was first described by Payen, A. (1843) in France. "One gene-one enzyme" hypothesis was provided after genetic and biochemical analysis of mutants in *Neurospora crassa* by Beadle, G. W. and Tatum, E. L. (1941). The process of gene conversion was also successfully analyzed in *Neurospora crassa* by Mitchel, M. B. (1955). The history of pioneering genetics to modern biology has been reviewed by Perkins, D. D. (1992).

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Biographical Sketches

Kohji Hasunuma, graduated from Tokyo University, Faculty of Science, Department of Biology (Plant Science) at 1966, and Graduate School of Biology (Plant Science) at 1971. Research associate of Tokyo University, Faculty of Arts and Culture 1971~1979. Associate Professor of National Institute for Basic Biology, 1979-1990. Visiting researcher at Carnegie Institution of Washington at Stanford at 1990. From 1990 professor of Yokohama City University, Kihara Institute for Biological Research. The Hirase Prize was awarded to Prof. K. Hasunuma by Japanese Society of Plant Morphology in 2000 in the success to prove molecular mechanism of light signal transduction in *Neurospora crassa* and *Pisum sativum*.

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