## Positively Heterochromatic Segment in Mitotic Metaphase Chromosomes of Vicia faba

It is well established that heterochromatic segments in the metaphase chromosomes of Vicia-faba are revealed by cold treatment or other methods. But all the reports¹ available so far concerned the understained, i.e. negatively heterochromatic (-H) segments except the one by Ockev² who recognized the existence of more densely stained, i.e. positively heterochromatic (+H) segments in addition to the -H segments. The present report deals with the +H segments of metaphase chromosomes which are disclosed by a new method.

Primary root-tips of *Vicia faba* (Nagasaya Soramame) were treated according to the procedure shown in the Table, and squashed after Feulgen stain. +H segments were clearly observed as the segments which remained dense in stainability in the metaphase chromosomes which swelled slightly and were more weakly stained than usual (Figure 1).

It is well known that DNA and/or histones are selectively extracted by TCA<sup>3,4</sup>, HCl<sup>5</sup> or acetic acid<sup>6</sup>. Therefore, it is supposed that the present procedure caused the preferential extraction of chromosomal constituents and consequently the differential staining along chromosomes. This implies that the + H segment revealed by the present method is a representation of heterogeneity in the chemical make-up of chromosomes. The + H segment was not detected when the sequence of TCA and HClacetic acid treatments was reversed, or when the HClacetic acid treatment was omitted. Relevant to the in situ selective extraction of chromosomal constituents, it was already suggested by Holtzman<sup>7</sup> that the sequence or combination of the treatments was also important.

The number and position of the +H segment in the present experiment are different from those plotted by  $Ockev^2$ . The site of +H was confined only to a very tiny segment at the satellite of the M chromosome directly adjacent to the satellite stalk where the nucleolus is

Scheme outlining the procedure for demonstration of + H segment

Primary root-tips

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Pretreatment with colchicine

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Tap-water rinse

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Fixation by acetic alcohol (1:3) or by 4% neutral formalin

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Tap-water rinse

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Treatment with 5% trichloroacetic acid (TCA) at 90 °C for about 3 min

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Tap-water rinse

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Treatment with 1N hydrochloric acid (HCl) and 45% acetic acid mixture (1:9) at 60 °C for about 3 min

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Tap-water rinse

Feulgen stain

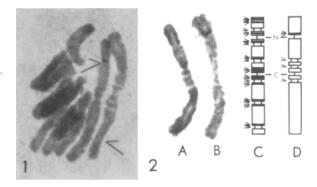


Fig. 1. The metaphase chromosome complement of  $Vicia\ faba$  after the application of the procedure shown in the Table. Arrows indicate + H segments in the single pair of M chromosomes.

Fig. 2. M chromosomes at metaphase showing -H and +H segments after various treatments. A, treatment with HCl-acetic acid<sup>9</sup>; B, present method; C and D, schematic representation of M chromosome. The position of the centromere (C), nucleolus organizing region (N), -H (gaps indicated by single arrows) and +H segments (black parts indicated by double arrows) as described by Ockey<sup>2</sup> (C) and myself (D).

formed (Figures 1 and 2). McClintock<sup>8</sup> reported that in Zea mays the part of the SAT chromosome immediately proximal to the satellite stalk is responsible for the organization of the nucleolus and this part shows positively heterochromatic, deep staining characteristic at pachytene. Therefore, the  $+\,\mathrm{H}$  segment in the present case may be a nucleolus organizing body in Vicia faba  $^{10}$ .

Zusammenfassung. Nach sukzessiver Behandlung der Wurzelspitze von Vicia faba mit 5% iger Trichlorsäure und dem Gemisch von einem Teil 1 N HCl und 9 Teilen 45% iger Essigsäure treten bei Feulgenfärbung die schmalen, tief gefärbten + H-Segmente der Trabanten beider metaphasischen M-Chromosomen direkt an der Astbasis hervor. Das + H-Segment ist als «nucleolus organizing body» nach McClintock<sup>8</sup> anzusehen.

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