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1. Linkage relationships for two mutants detected in Italian populations.

Further investigations have been accomplished on linkage relationships of two mutants, described in 1967 MNL, with known genetic markers.

For the ij-type mutant, F<sub>2</sub> segregations (repulsion phase) presented the following data (inclusive of 1966 results):

<u>G</u> <sub>1</sub>	<u>I</u> <sub>j</sub>	<u>g</u> <sub>1</sub>	<u>I</u> <sub>j</sub>	<u>G</u> <sub>1</sub>	<u>ij</u>	<u>g</u> <sub>1</sub>	<u>ij</u>
3882		2037		1889		8	

(c.o. 6.5% ± 1.5 st. error).

The data previously reported about close linkage between a shrunken type (bt) mutant and su<sub>1</sub>, have been confirmed by the scoring of ears obtained from backcrossing, to the triple recessive, plants of the constitution Su<sub>1</sub> bt G<sub>1</sub><sub>3</sub> / su<sub>1</sub> Bt g<sub>1</sub><sub>3</sub>, as follows:

<u>Su</u> <sub>1</sub>	<u>Bt</u>	<u>su</u> <sub>1</sub>	<u>Bt</u>	<u>Su</u> <sub>1</sub>	<u>bt</u>	<u>su</u> <sub>1</sub>	<u>bt</u>
113		4124		4157		20	

All the seedlings from the su<sub>1</sub> bt kernels had the G<sub>1</sub> phenotype, while only 26 plants from Su<sub>1</sub> Bt seeds turned out to be g<sub>1</sub>, indicating that part of them derived from contamination. Consequently, considering the bt phenotypes only, the su-bt recombination is 0.5% ± 0.1.

The bt mutant, then, has to be placed on chromosome 4 (probably allelic to bt<sub>2</sub>), between su<sub>1</sub> and g<sub>1</sub><sub>3</sub> and very close to su<sub>1</sub>.

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2. Abnormal segregations (significantly different from a 1:3 ratio) of genetic markers in the F<sub>2</sub> of lines derived from Italian populations.

In the analysis of a number of F<sub>2</sub> progenies derived from crossing lines from Italian populations to some genetic testers bearing recessive mutants, the following abnormal segregations have been observed: