

plants in F1. According to the results, under a) the genetic cause for fasciation may not be the direct cause of this heterosis either.

Segregation: Wherever a marked heterosis effect was evident in F1, all the segregations studied so far in F2 showed a majority (sometimes more than three-quarters) of plants as long as the F1 hybrids. However, a simple 3:1 segregation is excluded by several hypostatic genes segregating for shorter plants of different length.

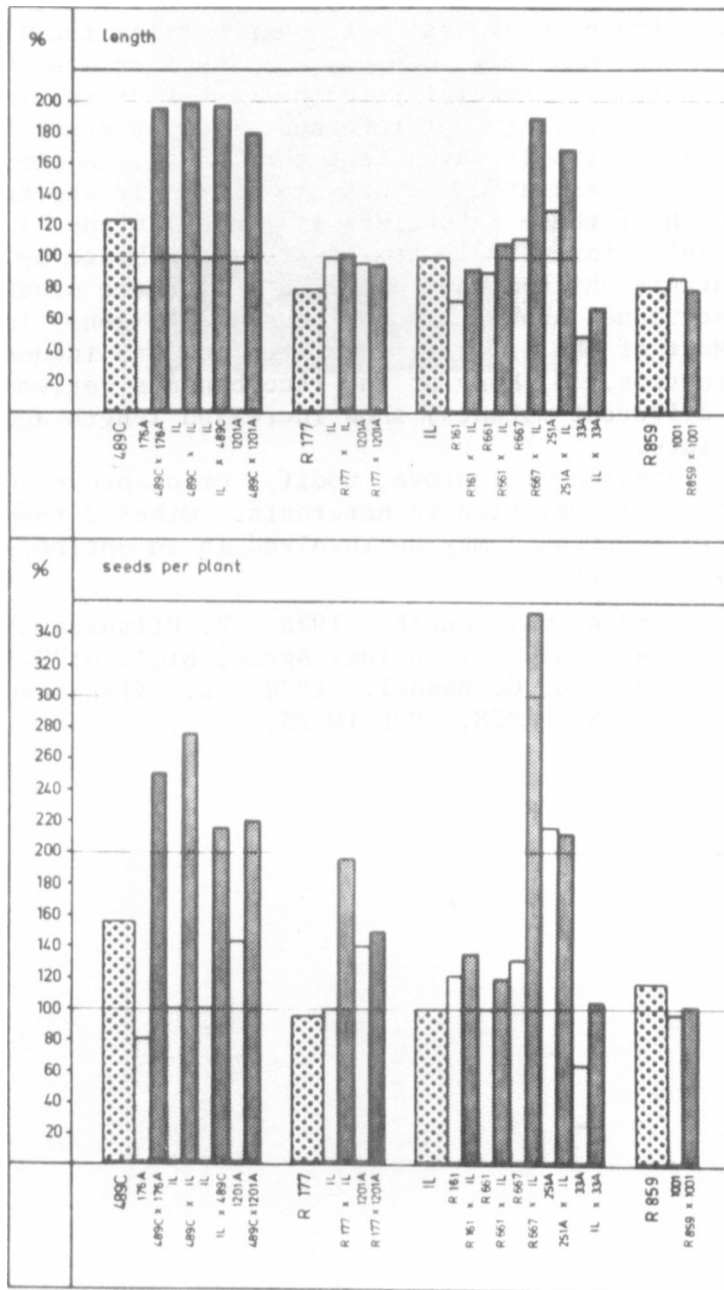


Fig. 2. F1's of the crosses

The following facts seem to support the working hypothesis for special importance of the genes for increased stem length and, perhaps, lateness:

- 1) Where there was no detectable heterosis in length in the F₁, no long plants were found in F₂; unusual length doesn't seem to be a simple hypostatic factor. None of these F₁'s was late.
- 2) The height of the hybrids was, of course, a function of internode number and—even more important in this case—internode length. Keeble and Pellew (1910) found both characters to show dominant inheritance in crosses between non-fasciated lines. The situation is, however, sometimes very complex.
- 3) The fasciated mutants causing heterosis show an increase, especially in the number of internodes (Milutinovic, 1972), but the total length of internodes may be counteracted by recessive genes. Fasciation itself may affect the length. Since the uppermost internodes of strongly fasciated mutants are extremely shortened, one could imagine that the length of these internodes is longer in the F₁ as a result of the action of alleles for normal stem growth contributed by the IL. Strongly fasciated mutants which do not cause heterosis are usually smaller than the IL and, as mentioned above, may not possess the genes for special length and lateness. Most of the linearly fasciated mutants induce weaker heterosis in many characters.
- 4) Many of the recombinants derived from crosses of 489 C x IL (and different mutants) show increased length and yield (3, 4). Most of them are late.

Further research will prove, modify, or disprove some of the above-mentioned points in relation to heterosis. Other determinants (heterozygosity and/or gene interactions) may be involved in an entirely satisfying explanation of this case of heterosis.

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