

THE MAP DISTANCE BETWEEN THE *e* AND *wlo* LOCI

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The presence of locus *e* on chromosome 6 was first indicated by linkage between *e* and *p* (2). Subsequently, testcross data involving genes *e*, *p* and *pl* revealed the gene order to be *e-p-pl* (4). The observed crossover value of 28% between *e* and *p* placed the *e* locus at the upper extremity of chromosome 6 above the existing uppermost marker, *wlo*. The joint segregation data in Table 1 indicate a distance of 26 units between *e* and *wlo*. The map of chromosome 6 as drawn by Blixt (1) is therefore extended by that amount. The data in Table 1 were obtained by crossing line 60 (*lf E Sn hr Wlo*) to segregates with the genotype *lf e Sn hr wlo* in the F_2 of cross 256 (L31 x L58).

Table 1. Joint segregation data for genes *e* and *wlo* (coupling phase) obtained from cross 284 F_2 .

| Phenotype | | | | Total | Joint seg. χ^2_1 | CrO% \pm S.E. |
|-----------|-------|-------|-------|-------|-----------------------|-----------------|
| E Wlo | E wlo | e Wlo | e wlo | | | |
| 62 | 10 | 14 | 15 | 101 | 17.19*** | 26.3 \pm 5.3 |

Although the position of the *e* locus is now established, the locus itself has certain disadvantages as a marker. Firstly, the E/e segregation is only apparent with certain combinations of the major flowering genes, e.g. it is obscured by the epistatic action of the genes *Lf* and *Lf^a* (2,5). These higher alleles in the *lf* series are in fact very common. Secondly, the variable penetrance of *Sn* (with respect to flowering node) means that with certain polygenic backgrounds genotype *lf e Sn hr* can flower at a low node and may therefore be confused with *lf E- Sn hr* (2). Thirdly, a single dose of *E* is sometimes insufficient to overcome the latening action of *Sn* and segregates of genotype *lf Ee Sn hr* may flower at a high node and thus be confused with the penetrant *lf e Sn hr* plants (5). The last two situations can often be resolved by growing F_3 . Such was the case in Cross 284 where, fortunately, the penetrance of *Sn* was very close to one but about 25% of *Ee* heterozygotes escaped from the early class. These escapes were readily identified in F_3 . Finally, as with the other flowering genes, observation of an E/e segregation is facilitated by the use of controlled environment conditions. However, special conditions are not necessarily obligatory (3).

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4. Murfet, I.C. 1977. *PNL* 9:38.
5. Murfet, I.C. 1978. *PNL* 10:48-52.