

**Internode length in *Pisum*: *le*<sup>5839</sup> is a less severe allele than Mendel's *le***

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Mutant alleles at the *le* locus, particularly Mendel's *le* (5, 16), have proved valuable in unravelling the role of gibberellins (GAs) in the control of internode length in *Pisum sativum* (1, 13). The allele *le* partially blocks the conversion of GA<sub>20</sub> to GA<sub>1</sub> the bioactive gibberellin in peas (1, 2), and consequently confers the dwarf phenotype. Quantification of GA<sub>1</sub> levels in the apical portions of isogenic *LeLe* and *lele* lines (using an internal standard and gas chromatography-mass spectrometry) has shown that tall (*Le Le*) plants typically contain 10-18 times more GA<sub>1</sub> than comparable dwarf plants (14). However, similar determinations on another isogenic pair of lines, Torsdag (tall, *LeLe*) and NGB5839 (dwarf, *le*<sup>5839</sup>*le*<sup>5839</sup>) yielded a somewhat smaller difference in GA<sub>1</sub> level (5-6 fold; 11). This suggests that the mutant allele in NGB5839, *le*<sup>5839</sup> (an induced mutation, 3) may be "leakier" than Mendel's *le*. However, this is not supported by the very short internode length of NGB5839 (3). Here we examine this question further at the phenotypic level. The evidence comes from a linkage study in which a gene pair linked to the *le* locus (*V*, normal pods / *v*, sugar pods) was used to monitor the inheritance of the mutant allele present in NGB5839. The *v* and *le* loci are linked (10) with an overall RCV of  $12.6 \pm 0.47\%$  (4). The effect of a photoperiod extension with incandescent light on internode length in *lele* and *le*<sup>5839</sup>*le*<sup>5839</sup> plants is also examined.

**Materials and Methods**

The pure lines used were Nordic Gene Bank line 5839 (NGB5839) (*VV le*<sup>5839</sup>*le*<sup>5839</sup>), NGB463 (*vv lele*) and cv. Dippes Gelbe Viktoria (*VV lele*). NGB5839 and Dippes Gelbe Viktoria carry allele *Lf* (minimum flowering node 11) while NGB463 carries *lf*<sup>a</sup> (minimum flowering node 5; 6). NGB5839 was produced by mutagenesis from cv. Torsdag, by Dr K.K. Sidorova (Novosibirsk, U.S.S.R.).

The plants were grown in a heated glasshouse. The day temperature was usually 20-25°C and the night temperature was 15-18°C. The growing medium was a 1:1 mixture of dolerite chips and vermiculite, topped with 3-4 cm of potting mix. For generations F<sub>1</sub> to F<sub>6</sub> the light regime consisted of natural daylight extended with mixed fluorescent (Thorn 40 W cool white tubes) and incandescent (Mazda 100 W pearl globes) light (intensity ca. 30  $\mu\text{mol m}^{-2} \text{s}^{-1}$  at pot top) to give a photoperiod of 18 h. Certain F<sub>6</sub> plants were grown in either an 8 h photoperiod (8 h natural light) or a 24 h photoperiod (8 h natural light extended to 24 h with weak incandescent light at an intensity of ca. 3  $\mu\text{mol m}^{-2} \text{s}^{-1}$  at pot top). The flowering node is defined as the number of the node bearing the first initiated flower, counting from the cotyledons as zero.

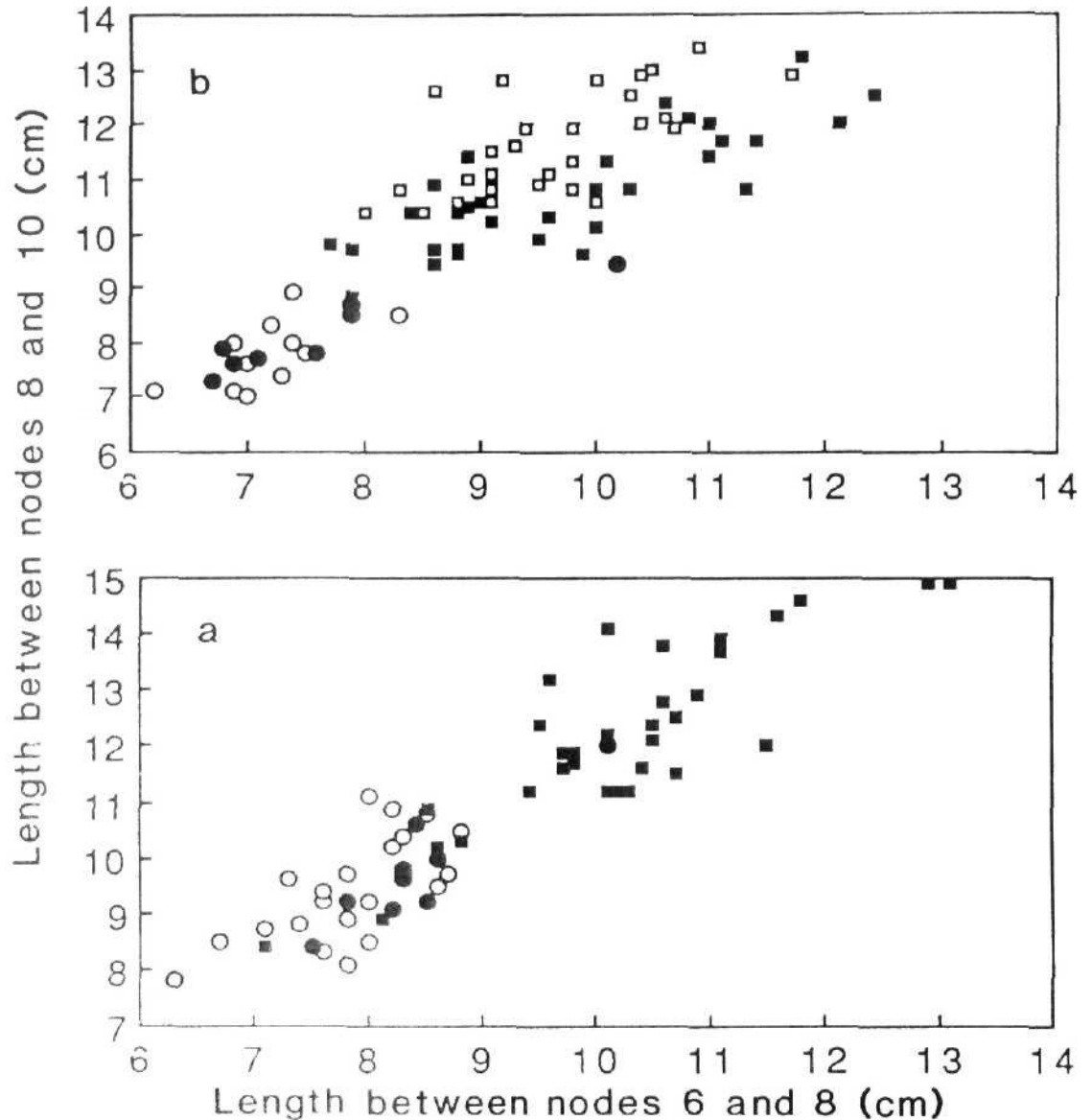


Fig. 1. Stem length between nodes 8 and 10 plotted against stem length between nodes 6 and 8 for F<sub>5</sub> (a) and F<sub>6</sub> (b) plants from cross NGB463 x NGB5839. Key: ○, vv plants from vv parents; ●, vv plants from Vv parents; ■, V- plants from Vv parents; □, VV plants from VV parents. The F<sub>5</sub> was generated from 4 vv and 3 Vv F<sub>4</sub> plants, and the F<sub>6</sub> from 2 vv, 3 Vv and 2 VV F<sub>5</sub> plants.

### Results

The F<sub>1</sub> of cross NGB463 x NGB5839 was dwarf, as were all plants in subsequent generations. Generations F<sub>3</sub> to F<sub>6</sub> were produced by single plant selection beginning with an F<sub>2</sub> segregate of genotype *lf<sup>a</sup> lf<sup>a</sup> Vv*. Plants of genotype vv (sugar pod) possessed, on average, significantly ( $P < 0.001$ ) shorter internodes than V- (normal pod) plants in all generations from F<sub>2</sub> to F<sub>6</sub> (e.g. Table 1, Fig. 1, a and b). This suggests that the mutant allele in NGB5839, *le<sup>5839</sup>* (which entered the cross linked to V), results in longer internodes and is, therefore, a leakier allele than *le*.

Table 1. Stem length (cm) between nodes 6 and 9 for lines NGB463 and NGB5839, and for *Lf*- *V*- and *Lf*- *vv* segregates in the F<sub>2</sub> generation of crosses NGB463 x NGB5839 and NGB463 x Dippes Gelbe Viktoria. The data are shown as mean  $\pm$  SE with n in parentheses. Photoperiod 18 h.

Line or Cross	Genotype	Stem length
NGB463	<i>lf<sup>a</sup> lf<sup>a</sup> vv</i>	16.10 $\pm$ 0.50 ( 6)
NGB5839	<i>LfLf VV</i>	6.32 $\pm$ 0.29 (6)
NGB463 x NGB5839 F <sub>2</sub>	<i>Lf</i> - <i>V</i> -	10.41 $\pm$ 0.35 (50)
	<i>Lf</i> - <i>vv</i>	7.50 $\pm$ 0.35 (11)
NGB463 x Dippes Gelbe Viktoria F <sub>2</sub>	<i>Lf</i> - <i>V</i> -	16.33 $\pm$ 0.46 (43)
	<i>Lf</i> - <i>vv</i>	16.18 $\pm$ 1.28 (10)

Table 2. Stem length (cm) between nodes 6 and 9 for *le*<sup>5839</sup>*le*<sup>5839</sup> and *lele* segregates from cross NGB463 x NGB5839 grown in either an 8 h or a 24 h photoperiod. Data are shown as mean  $\pm$  SE of 10 replicates.

Genotype	Stem length	
	8 h photoperiod	24 h photoperiod
<i>le</i> <sup>5839</sup> <i>le</i> <sup>5839</sup>	11.91 $\pm$ 0.30	18.01 $\pm$ 0.56
<i>lele</i>	8.14 $\pm$ 0.17	10.97 $\pm$ 0.24

While there was some overlap of internode length values between the presumed *lele* and *le*<sup>5839</sup>- plants in earlier generations, this was minimal by F<sub>6</sub> (Fig. 1, b). From F<sub>2</sub> to F<sub>6</sub>, short plants of genotype *vv* produced only short offspring (e.g. Fig. 1, a and b). Plants of genotype *V-* were usually taller, while the vast majority (e.g. 89% in F<sub>5</sub> and 88% in F<sub>6</sub>) of *vv* segregates from *Vv* parents were short. However, several taller *vv* segregates from *Vv* parents were observed; in one such case from the F<sub>4</sub>, the progeny was grown and comprised 5 short and 12 taller types (data not shown). This suggests that this F<sub>4</sub> plant was of genotype *vv le*<sup>5839</sup>*le* (a recombinant). The F<sub>5</sub> generation included 5 short *V-* segregates (Fig. 1, a). The F<sub>6</sub> generation from one of these plants was grown and consisted of 1 short *vv* plant and 13 taller *V-* plants. Thus this F<sub>5</sub> plant was clearly not a recombinant; the reason for its short stature is not clear. However, it is noteworthy that all *VV* F<sub>6</sub> plants possessed considerably longer internodes than all *vv* F<sub>6</sub> plants from *vv* parents (Fig. 1, b).

In contrast to the results from cross NGB463 (*vv lele*) x NGB5839 (*VV le*<sup>5839</sup>*le*<sup>5839</sup>) segregation of the gene pair *V/v* was not associated with differences in internode length in cross NGB463 (*vv lele*) x Dippes Gelbe Viktoria (*VV lele*) (Table 1).

Although the number of individuals available for comparison was small, the internode length of heterozygous *le*<sup>5839</sup>*le* plants was intermediate between that of homozygous *le*<sup>5839</sup>*le*<sup>5839</sup> and *lele* plants. For example, in F<sub>3</sub> the mean values for the stem length between nodes 6 and 9 for *le*<sup>5839</sup>*le*<sup>5839</sup>, *le*<sup>5839</sup>*le* and *lele* plants were (in cm) 17.4 ± 1.40 (n=3), 14.43 ± 0.92 (n=4) and 11.67 ± 0.03 (n=3), respectively. It therefore appears there is very little dominance of either allele over the other.

When *le*<sup>5839</sup>*le*<sup>5839</sup> and *lele* plants with a similar genetic background (F<sub>6</sub> segregates descended from a single F<sub>4</sub> plant from cross NGB463 x NGB5839) were grown in 8 h and 24 h photoperiods, the internodes of both genotypes were longer in 24 h (8 h natural light plus 16 h incandescent light) than in 8 h (Table 2), in accordance with previous results (e.g. 8, 11). However, the response shown by *le*<sup>5839</sup>*le*<sup>5839</sup> plants (a 51 % increase) was greater than that of *lele* plants (a 35% increase, Table 2). There was no evidence that this difference was due to factors other than the genotype at the *le* locus. For example, the flowering behaviour of both *le*<sup>5839</sup>*le*<sup>5839</sup> and *lele* plants was similar. Both groups initiated flowers at nodes 6-8 in both photoperiods and in the 24 h photoperiod flower development ensued either at the node of initiation or at one node higher. In the 8 h photoperiod substantial flower abortion occurred, but to a similar extent in both genotypes. (The flowering genotype of these plants therefore appears to be *lf*<sup>a</sup>*lf*<sup>a</sup> *EE SnSn DneDne*, see 7).

### Discussion

In cross NGB463 (*vv lele*) x NGB5839 (*VV le*<sup>5839</sup>*le*<sup>5839</sup>), *V le*<sup>5839</sup>- segregates, on average, possessed longer internodes than did *vv lele* segregates. In NGB5839, GA<sub>1</sub> levels were not reduced to the same extent as in *lele* lines (compared with isogenic *LeLe* lines; 11,14).

Table 3. Effect of segregation for the  $le^{5839}/le$  pair of alleles on the number of seeds per plant in the F<sub>5</sub> and F<sub>6</sub> generations of cross NGB463 x NGB5839. The data are shown as mean  $\pm$  SE with n in parentheses.

Generation	Genotype	
	$le^{5839}-$	$lele$
F <sub>5</sub>	28.47 $\pm$ 0.78 (30)	23.55 $\pm$ 0.60 (33)
F <sub>6</sub>	18.53 $\pm$ 0.36 (59)	15.63 $\pm$ 0.41 (19)

Considered together, these results strongly suggest that allele  $le^{5839}$  imposes a less severe block on GA<sub>1</sub> biosynthesis than does  $le$ . Clearly  $le^{5839}$  is a different allele from  $le$  and the designation  $le^{5839}$  should remain to indicate this. On the basis of measurements of true-breeding  $le^{5839}le^{5839}$  and  $lele$  F<sub>6</sub> families (descended from a single F<sub>4</sub> plant) allele  $le^{5839}$  increases stem length between nodes 6 and 9 by ca. 40% in an 18 h photoperiod, compared with  $lele$  plants. The paradoxical aspect of the present and previous work is that NGB463 ( $lele$ ) possesses much longer internodes than NGB5839 ( $le^{5839}le^{5839}$ ) (Table 1). Clearly the two pure lines differ with respect to other loci which affect internode length (e.g. possibly at the *Cry* locus). The presence of  $lf^a$  in NGB463 would most likely also result in longer internodes. However, this cannot alone explain the stature of NGB463, since this line was considerably taller than  $lf^a lf^a lele$  (and  $lf^a lf^a le^{5839}-$ ) segregates in F<sub>2</sub>-F<sub>6</sub> (data not shown).

The existence of at least one proven recombinant (genotype  $le^{5839} le vv$ ; phenotype long internodes, sugar pods) in cross NGB463 x NGB5839 strongly indicates that the short stature of  $vv$  plants in F<sub>2</sub> to F<sub>6</sub> of this cross is not due to a pleiotropic effect of  $v$ . This is confirmed by the lack of effect on internode length of segregation for  $V/v$  in cross NGB463 x Dippes Gelbe Viktoria, which, furthermore, is consistent with the presumption that both NGB463 and Dippes Gelbe Viktoria possess the "normal"  $le$  allele.

The identification of allele  $le^{5839}$  increases to four the number of alleles at the  $le$  locus (in order of increasing length,  $le^d$ ,  $le$ ,  $le^{5839}$  and  $Le$ , see 12). It seems possible that allele  $le^{5839}$  may be of some agronomic value since it has the effect of increasing internode length compared with  $le$ . In this context it is of interest that in the F<sub>5</sub> and F<sub>6</sub> generations of cross NGB463 x NGB5839 (on a  $lf^a lf^a EE SnSn DneDne$  genetic background in an 18 h photoperiod, see 9),  $le^{5839}-$  plants produced ca. 20% more seeds than did  $lele$  plants ( $P < 0.001$ ; Table 3). However, this effect cannot for certain be attributed to the difference at the  $le$  locus since the plants also differed at the  $v$  locus.

It is well known that in pea internode elongation is enhanced by photoperiod extensions with incandescent or far-red-rich light (3, 8, 11, 15). It has also been shown (3, 11) that line NGB5839 ( $le^{5839}le^{5839}$ ) responds to such extensions to a lesser extent than does its tall ( $LeLe$ ) progenitor, Torsdag. In contrast, some  $le$  dwarf lines or selections are at least as responsive as Torsdag (Table 1 from 8; Torsdag = L107). However, in the present study  $le^{5839}le^{5839}$  plants responded to an incandescent photoperiod extension to a greater extent than did  $lele$  plants (Table 2). Therefore the difference in responsiveness between NGB5839 and  $le$  lines referred to above is probably attributable to differences in genetic

background. The results shown in Table 2 support the suggestion (3) that on a constant genetic background the response to an incandescent (or far-red-rich) photoperiod extension decreases as the severity of the genetic block in GA<sub>1</sub> biosynthesis increases (see also 11).

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