

# Lettuce Genetics: Inheritance, Linkage, and Epistasis

Edward J. Ryder

U.S. Department of Agriculture-Agricultural Research Service, U.S. Agricultural Research Station,  
1636 East Alisal Street, Salinas, CA 93905

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**Abstract.** Two new lettuce (*Lactuca sativa* L.) genes are described and named *truncated leaf* (*tn*), and *sickly* (*si*). A gene for reflexed involucre is identical to that previously described in wild lettuce (*L. serriola* L.). Mosaic reaction (*me*) and light green (*lg*) are linked, with  $P = 0.448$ . Six gene pairs tested for linkage are independently inherited. *Sickly* is epistatic to *light green*.

This paper continues the description of major genes, linkages, epistasis, and other interactive relationships in lettuce. Eventually, the accumulated information of this type will be suitable to be combined with information from RFLP and similar analyses to provide a comprehensive lettuce genome map. I report here two new genes, one linkage and one epistatic combination.

## Materials and Methods

Eight experiments were conducted. The study of truncated leaf began with the observation of off-type plants in a field planting. These plants were in a group of BC<sub>2</sub>F<sub>3</sub> families from the cross M631 × 'Salinas', which were segregating for mosaic resistance. The off-type plants were darker green and smaller than the normal plants' and had entire leaf margins and less-blistered leaves. The reduced size was partially due to truncation of the leaves. The truncated plant, 78-162-1, was crossed to 'Vanguard 75', a crisphead cultivar. Plant type was observed and recorded in F<sub>1</sub>, F<sub>2</sub>, and F<sub>3</sub> families.

A slow-growing, sickly appearing plant (85-1174cd) was observed in the F<sub>2</sub> progeny of a cross 82-1061-1 × 'Vanguard 75'. The plant was severely stunted, chlorophyll deficient, and had necrotic areas on the leaves. It survived to produce flowers and was crossed by 'Vanguard 75' and 'Prizehead', a red leaf lettuce. Plant type was observed in F<sub>1</sub> and F<sub>2</sub> populations of both crosses and in F<sub>3</sub> families of the 'Prizehead' cross. The latter cross also segregated for dark vs. light green, which led to a study of epistasis with the sickly character.

'Australian', a light-green leaf lettuce, produces reflexed mature involucre. A study of shattering (reflexed) vs. nonshattering (erect) -was reported by Whitaker and McCollum (1954) in a cross between wild (*L. serriola*) and cultivated lettuce. A single gene controlled the trait, with shattering dominant. The gene was designated as *b* and later renamed *er* (Robinson et al., 1983). Whitaker and McCollum footnoted a personal communication from R.C. Thompson that 'Australian' was the only known cultivated form with "... some degree of shattering. . . ." To study the inheritance of reflexed as manifested in 'Australian', an F<sub>2</sub> population from a cross 'Calmar' × 'Australian', planted for a linkage study, was also observed for involucre type. To confirm the involucre observations, a random

sample of 80 F<sub>3</sub> families was grown. A reflexed plant from another 'Australian' cross was crossed with *L. serriola* to test for gene identity.

A possible linkage relationship between lettuce mosaic (LMV) reaction and dark- vs. light-green leaves was studied in F<sub>2</sub> populations of three crosses: 'Australian' × 'Vanguard 75', 'Chung Chuk Myun' × 'Vanguard 75', and 'Vanguard 75' × 'Prizehead'. 'Vanguard 75' is dark green and LMV resistant (*Lg-Lgmomo*), while the three other parents are light green and susceptible (*lglgMoMo*). 'Chung Chuk Myun' is a Korean leaf lettuce.

Several other linkage studies were based on F<sub>2</sub> populations of the crosses: M400-27-23 × 'Golden Bibb', yellow vs. salmon, normal vs. virescent, LMV reaction; and 'Calmar' × 'Australian', yellow vs. pale, black seed vs. white, dark vs. light green, reflex vs. erect.

All plantings were in a greenhouse. Plants were grown in flats or plastic pots of various sizes, in soil or sand-soil mixtures. The containers were fertilized and hand-watered as needed. Crosses were made by the method of Oliver (1910), modified to maximize the number of hybrid seeds (Ryder and Johnson, 1974; E. J. R., unpublished).

Genetic ratios were compared by standard  $\chi^2$  procedure. Maximum likelihood tables were used to calculate recombination values (Allard, 1956). Recombination values are denoted by *p* and probability values by *P*.

## Results and Discussion

### Inheritance studies

**Truncated leaf.** Plants of this type grow normally for 45 to 55 days. The 15th or 16th leaf continues to expand normally latitudinally but at a slower rate at the marginal area around the apex, resulting in a truncated or foreshortened appearance. The F<sub>1</sub> was normal. The F<sub>2</sub> segregated ≈3 normal : 1 truncated (Table 1). F<sub>3</sub> families from normal F<sub>2</sub> plants segregated 1 normal : 2 segregating, and segregation within the latter group was again 3:1. The evidence indicates that truncation is recessive and the trait is inherited as a single gene. The trait is named *truncated leaf* and designated *tn*.

The gene will probably be of limited use as a marker because the phenotype appears late in the vegetative growth period. It also has incomplete penetrance. The differential leaf expansion may, however, prove interesting in a developmental study.

**Sickly.** In the greenhouse, these plants emerge as early as normal ones. Cotyledons are smaller than normal and pale green. The first true leaf is yellow-green, slightly malformed, and small.

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Table 1. Inheritance of truncated leaf in lettuce. Segregation in F<sub>2</sub> population and F<sub>3</sub> families of the cross 78-162-1 × Vanguard 75.

| Population                    | No. observed (expected) |             |           | $\chi^2$   | <i>P</i> |
|-------------------------------|-------------------------|-------------|-----------|------------|----------|
|                               | Normal                  | Segregating | Truncated |            |          |
| F <sub>2</sub>                | 70 (68)                 |             | 21 (23)   | 0.24 (3:1) | 0.5-0.7  |
| Among F <sub>3</sub> families | 15 (13)                 | 24 (26)     |           | 0.47 (1:2) | 0.3-0.5  |
| Within seg. families          | 199 (207)               |             | 77 (69)   | 1.24 (3:1) | 0.2-0.3  |

Table 2. Inheritance of sickly in lettuce. Segregation in F<sub>2</sub> populations and F<sub>3</sub> families.

| Population              | No. observed (expected) |             |            | $\chi^2$     | <i>P</i>  |
|-------------------------|-------------------------|-------------|------------|--------------|-----------|
|                         | Normal                  | Segregating | Sickly     |              |           |
| F <sub>2</sub>          |                         |             |            |              |           |
| 85-1174cd x Prizehead   | 127 (126.75)            |             | 42 (42.25) | 0.003 (3:1)  | 0.95-0.98 |
| 85-1174cd x Vanguard 75 | 166 (165)               |             | 54 (55)    | 0.03 (3:1)   | 0.8-0.9   |
| F <sub>3</sub>          |                         |             |            |              |           |
| 85-1174cd x Prizehead   |                         |             |            |              |           |
| Among families          | 43 (36.5)               | 80 (73)     | 33 (36.5)  | 1.38 (1:2:1) | 0.5-0.7   |
| Within seg. families    | 1417 (1407)             |             | 459 (469)  | 0.28 (3:1)   | 0.5-0.7   |

Table 3. Inheritance of reflexed involucre in lettuce. Segregation in F<sub>2</sub> population and F<sub>3</sub> families of the cross 'Calmar' × 'Australian'.

| Population                    | No. observed (expected) |             |           | $\chi^2$     | <i>P</i>  |
|-------------------------------|-------------------------|-------------|-----------|--------------|-----------|
|                               | Reflex                  | Segregating | Erect     |              |           |
| F <sub>2</sub>                | 537 (530)               |             | 170 (177) | 0.35 (3:1)   | 0.5-0.7   |
| Among F <sub>3</sub> families | 21 (20)                 | 43 (40)     | 16 (20)   | 1.08 (1:2:1) | 0.5-0.7   |
| Within seg. families          | 425 (449)               |             | 174 (150) | 5.12 (3:1)   | 0.02-0.05 |

Table 4. Evidence for linkage between genes for leaf color and mosaic reaction in lettuce, repulsion phase. S = susceptible, R = resistant.

| F <sub>2</sub> population            | No. observed (expected) |              |              |              | Linkage<br>$\chi^2$ | P         |
|--------------------------------------|-------------------------|--------------|--------------|--------------|---------------------|-----------|
|                                      | Dark green              |              | Light green  |              |                     |           |
|                                      | S                       | R            | S            | R            |                     |           |
| Australian x Vanguard 75             | 361<br>(364)            | 134<br>(121) | 125<br>(121) | 26<br>(40)   | 5.43                | 0.01-0.02 |
| Chung Chuk Myun x Vanguard 75        | 227<br>(248)            | 91<br>(83)   | 97<br>(83)   | 26<br>(28)   | 2.54                | 0.1-0.2   |
| Vanguard 75 x Prizehead              | 517<br>(526)            | 181<br>(175) | 185<br>(175) | 52<br>(59)   | 1.72                | 0.1-0.2   |
| Pooled                               | 1105<br>(1138)          | 406<br>(379) | 407<br>(379) | 105<br>(126) | 8.67                | <0.01     |
| Test for heterogeneity <sup>a</sup>  |                         |              |              |              |                     |           |
| Total $\chi^2$ (9 df) = 13.38        |                         |              |              |              |                     |           |
| Pooled $\chi^2$ (3 df) = 8.79        |                         |              |              |              |                     |           |
| Heterogeneity $\chi^2$ (6 df) = 4.59 |                         |              |              |              |                     |           |
|                                      |                         |              |              |              |                     | 0.5-0.7   |

<sup>a</sup>Based on  $\chi^2$  values for 9:3:3:1 ratio.

Three weeks after seeding, the first true leaf on a sickly plant is ≈1 cm long and on a normal plant 2 to 3 cm long. The sickly died during 90 days. The F<sub>1</sub> was normal. The F<sub>2</sub> populations produced a leaf every 6 days, about half the rate for the normal plants. Sickly type is partially lethal: 15 of 54 plants segregated ≈3 normal :1 sickly (Table 2). A sample of 156 F<sub>3</sub>

Table 5. Linkage detection tests for lettuce showing no evidence for linkage. F<sub>2</sub> phenotypes: 1 = A-B-, 2 = A-bb, 3 = aaB-, 4 = aabb<sup>z</sup>.

| Loci <sup>y</sup> | Observed no.<br>in each phenotype |     |     |    | Total | Linkage<br>$\chi^2$ | P       |
|-------------------|-----------------------------------|-----|-----|----|-------|---------------------|---------|
|                   | 1                                 | 2   | 3   | 4  |       |                     |         |
| sa;mo             | 513                               | 170 | 156 | 64 | 903   | 1.34                | 0.2-0.3 |
| vi;mo             | 492                               | 168 | 183 | 69 | 912   | 0.45                | 0.5-0.7 |
| lg;sa             | 525                               | 175 | 170 | 70 | 940   | 1.59                | 0.2-0.3 |
| pa;er             | 408                               | 120 | 129 | 50 | 707   | 1.89                | 0.1-0.2 |
| w;er              | 404                               | 129 | 133 | 41 | 707   | 0.02                | 0.8-0.9 |
| lg;er             | 421                               | 129 | 116 | 41 | 707   | 0.62                | 0.3-0.5 |

<sup>y</sup>Comparison based on 9:3:3:1 F<sub>2</sub> ratio. Expected numbers for each test: sa;mo (508,169, 169, 56); vi, mo(513, 171, 171, 57); lg,sa(529, 176, 176, 59); pa,er; w,er, lg,er (398, 133, 133, 44).

<sup>z</sup>Gene symbols: sa = salmon flower; mo = lettuce mosaic resistance; vi = virescent leaf; lg = light green leaf; pa = pale flower; er = erect involucre; w = white seed.

families from the cross with 'Prizehead' segregated  $\approx$ 1 normal :2 segregating :1 sickly. Within the segregating families, segregation was again 3:1. The sickly type is recessive, and the trait is inherited as a single gene, which is named *sickly* and symbolized *si*.

This trait may be useful as a marker, as the effect is easily observed in the cotyledon-first true-leaf stage. It is also interesting as a variant to the chlorophyll-deficient genes previously identified in lettuce (reviewed in Robinson et al., 1983; Ryder, 1983). Among those genes, albino genes are lethal, and golden is partially lethal. Expressivity of the golden genotype varies in the seedling stage with regard to the chlorophyll-deficiency aspect; those plants with severe reduction of chlorophyll, approaching albinism, usually die. Those with less-severe reduction survive, grow and develop normally, and produce golden flow-

ers. The other chlorophyll-deficient genes restrict growth and cause varying degrees and patterns of yellowing. Sickly plants all survive the seedling stage but become progressively weaker, severely stunted, and partially necrotic. This trait would also be of interest in a study of growth and development.

**Reflexed involucre.** Segregation in F<sub>2</sub> was  $\approx$ 3 reflexed : 1 erect (Table 3). The F<sub>3</sub> families segregated 1 reflexed : 2 segregating :1 erect. Segregation within segregating families was significantly different from 3:1, with an excess of erect plants. This result probably is due to a misreading. There was no visible difference between homozygous and heterozygous reflexed plants. The preponderance of evidence is that reflexed is dominant and controlled by a single gene. These are the same results as found by Whitaker and McCollum (1954). A reflexed plant from another experiment with 'Australian' as a parent was crossed with a reflexed plant of *L. serriola*. The F<sub>1</sub> was also reflexed; therefore, the genes are the same.

Linkage studies

**Leaf color-mosaic reaction.** The F<sub>2</sub> population from the cross 'Australian'  $\times$  'Vanguard 75' segregated for several traits, including dark-green vs. light-green leaf and reaction to lettuce mosaic virus. 'Australian' is light green and susceptible (*lglgMoMo*), while 'Vanguard 75' is dark green and resistant (*LgLgmomo*). There was an excess of parental types and a shortage of recombinant types (Table 4). Linkage in the repulsion phase was indicated. Two additional F<sub>2</sub> populations were grown: 'Chung Chuk Myun'  $\times$  'Vanguard 75' and 'Vanguard 75'  $\times$  'Prizehead'. 'Chung Chuk Myun' and 'Prizehead' are also light green and mosaic susceptible. F<sub>1</sub> plants from crosses between 'Australian' and the other two light-green cultivars were light green; therefore, the three cultivars have the same light-green allele.

The other two F<sub>2</sub> populations also showed a shortage of re-

Table 6. Segregation for dark vs. light green and normal vs. sickly in F<sub>2</sub> and F<sub>3</sub> families of the cross 85-1174cd  $\times$  Prizehead lettuce.

| Population                    | No. observed (expected)        |           |                 | $\chi^2$ (9:3:4) | P       |
|-------------------------------|--------------------------------|-----------|-----------------|------------------|---------|
|                               | Dark                           | Light     | Sickly          |                  |         |
| F <sub>2</sub>                | 93(92)                         | 34(32)    | 42(42)          | 0.21             | 0.5-0.7 |
| Among F <sub>3</sub> families | <i>Observed</i>                |           | <i>Expected</i> |                  |         |
| All dark                      | 11                             |           | 9.75            |                  |         |
| Seg. dark : sickly            | 15                             |           | 19.5            |                  |         |
| Seg. 9:3:4                    | 40                             |           | 39.0            |                  |         |
| Seg. dark : light             | 22                             |           | 19.5            |                  |         |
| Seg. light : sickly           | 25                             |           | 19.5            |                  |         |
| All light                     | 10                             |           | 9.75            |                  |         |
| All sickly                    | 33                             |           | 39.0            |                  |         |
|                               |                                |           |                 | 4.03             | 0.5-0.7 |
|                               | <i>No. observed (expected)</i> |           |                 |                  |         |
| Within segregating families   | Dark                           | Light     | Sickly          |                  |         |
| Dark : sickly (3:1)           | 209 (214)                      |           | 76 (71)         | 0.47             | 0.3-0.5 |
| 9:3:4 (A)                     | 426 (440)                      | 184 (147) | 173 (196)       | 12.46            | <0.01   |
| 9:3:4 (B)                     | 192 (193)                      | 62 (64)   | 89 (86)         | 0.30             | 0.8-0.9 |
| Dark : light (3:1)            | 317 (330)                      | 123 (110) |                 | 2.03             | 0.1-0.2 |
| Light: sickly (3:1)           |                                | 344 (349) | 121 (116)       | 0.29             | 0.5-0.7 |

combinant types, but the  $\chi^2$  for linkage was not significant in either test (Table 4). However, the consistent shortage of recombinant in all three populations indicates linkage. When the observed numbers are pooled,  $\chi^2$  for linkage is highly significant. Heterogeneity  $\chi^2$  for the 9:3:3:1 ratio is not significant, indicating the populations segregate in the same manner. I conclude that there is loose linkage in the repulsion phase. The recombination value was calculated as  $P = 0.44 \pm 0.018$ .

*Other linkage detection tests.* Linkage  $\chi^2$  and probabilities were calculated for six other combinations. Independent assortment was shown for all pairs (Table 5).

### Epistasis

*Light green and sickly.* The  $F_2$  of the cross 85-1174cd  $\times$  'Prizehead' segregated for dark- vs. light-green as well as for the sickly trait. Only three phenotypes were distinguishable—dark green, light green, and sickly—and they occurred in the ratio 9:3:4 (Table 6). This ratio suggests that sickly is epistatic to light green. There were seven categories among 156  $F_3$  families, as expected, if *LgLgsisi*, *Lglgsisi*, and *lglgsisi* are all sickly,

but otherwise not distinguishable. Segregation among families was as expected (Table 6). Segregation within  $F_3$  families was also as expected, except for those segregating for all three phenotypes. In this group, there was an excess of light green plants and a shortage of the other phenotypes. A second planting of seven three-phenotype families was grown and segregated 9:3:4, as expected. Therefore, *Sisi* is epistatic to *Lglg*.

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