

## Cytogenetic Studies of *Trifolium* Spp. Related to Berseem

### IV. The Relationships in the *Alexandrinum* and *Vavilovi* Crossability Groups, and the Origin of the Cultivated Berseem †

E. PUTIYEVSKY, J. KATZNELSON and D. ZOHARY

Division of Pasture and Forage Crops, Agricultural Research Organization, Nve-Ya'ar, P. O. Haifa, and  
Division of Genetics, Hebrew University, Jerusalem (Israel)

**Summary.** The interspecific cytogenetic relationships among *T. alexandrinum* L., *T. berytheum* Boiss., *T. salmoneum* Mout., *T. apertum* Bobr., *T. meironense* Zoh. et Lern. and *T. vavilovi* Eig. were studied in several hundreds of intra-specific (WS) and interspecific (BS) F<sub>1</sub> hybrids in all possible combinations, and in many of their progenies.

Seed germination and development of F<sub>1</sub> BS hybrids were normal, except in crosses involving *T. vavilovi*. Their pollen fertility and seed set was subnormal, with somewhat lower values in *T. vavilovi* progenies. No multivalents were observed in any F<sub>1</sub> BS hybrid, but in some plants, four and also six chromosomes did not pair to form bivalents. B chromosomes were common in hybrids involving *T. berytheum* or *T. salmoneum*.

Some BS F<sub>1</sub> hybrids set ample seeds when selfed, and most of them originated from either one of two self-incompatible *T. berytheum* plants.

The six taxa studied form three groups: (i) *T. vavilovi*, which is quite remote from the other five; (ii) *T. meironense* and *T. apertum*; and (iii) *T. alexandrinum*, *T. berytheum* and *T. salmoneum*. The two last species, especially *T. salmoneum*, seem to be the progenitors of the cultivated berseem.

### Introduction

In studies of various cytogenetic aspects of several *Trifolium* species related to berseem (*T. alexandrinum* L.) published earlier, we described the pattern of crossability of 12 clover species (Putiyevsky and Katznelson, 1973). These species could be divided into five crossability groups, and the relationships within and between the second and the third crossability groups have also been described (Katznelson and Putiyevsky, 1974; Putiyevsky and Katznelson, 1974). Two other crossability groups are more intimately associated with the cultivated berseem; the first one (Group I) includes, besides the cultivated berseem, four wild East Mediterranean taxa, *T. berytheum* Boiss., *T. salmoneum* Mout., *T. apertum* Bobr. and *T. meironense* Zoh. et Lerner. These five species intercross quite readily (Putiyevsky and Katznelson, 1973). Another species, *T. vavilovi* Eig., is morphologically somewhat similar to the five taxa mentioned above, and was recently located with them in subsection *Alexandrina* by Zohary (1972). Hybridization between *T. vavilovi* and the other species, however, failed almost completely and it comprises Crossability Group IV.

The six species, or taxa, concerned here impose several taxonomic problems, and they are often discussed in connection with aspects of the origin of

berseem. Although the East Mediterranean region was exploited botanically quite extensively in the 19th Century, most of the species related to berseem have been described only recently (*T. vavilovi* Eig. in 1934, *T. apertum* Bobr. in 1941, *T. salmoneum* Mout. in 1953, and *T. meironense* Zohary and Lerner in 1972), in contrast to other clover sections and groups from the same region. They were, and still are, often regarded as intraspecific entities of berseem. Even *T. berytheum* Boiss., which was described in 1856 by Boissier, was named as a variety of *T. echinatum*, *T. constantinopolitanum* or *T. alexandrinum*, and intermediates between cultivated berseem and *T. berytheum* have been found in nature (Oppenheimer, 1959; Zohary, 1972). Oppenheimer's interpretation of *T. alexandrinum* var. *berytheum* as wild *T. alexandrinum* definitely focuses the search for the wild progenitor of berseem on *T. berytheum*. Bobrov (1947), on the other hand, claims that his *T. apertum* is the true progenitor of berseem.

Cytogenetic studies may clarify some of the problems mentioned above, but up to now all the interspecific studies with berseem involved crossing it with remote species such as *T. repens* L. (Evans, 1962), *T. resupinatum* (Müller, 1960), and *T. pratense* L. and *T. campestre* Schreb. (Abdel-Tawab, 1968).

In the present study large-scale hybridization was carried out with all the species related to berseem and the results obtained illuminate some of the cytogenetic problems mentioned, including the origin of the cultivated berseem.

† Contribution from the Volcani Center, Agricultural Research Organization, Bet Dagan, Israel, 1974 Series, No. 201-E. The work reported here was financed by Grant FG-IS-222 from the U.S. Department of Agriculture under P.L. 480.

### Materials and Methods

The origin of lines used as parent plants as well as the two methods of obtaining hybrid seed, i.e., manually and by the natural hybridization programme (NHP), have been presented elsewhere (Putiyevsky and Katznelson, 1973; Katznelson, 1971). Altogether, 245 intraspecific and 594 interspecific hybrid seeds of this group were germinated, each seed separately in a petri dish. The seedlings were planted in soil in 3-kg pots and kept in a glasshouse.

The methods employed in the cytological and pollen fertility studies were described earlier (Putiyevsky and Katznelson, 1970). In each  $F_1$  plant, several heads were bagged and triggered for self-pollination. Other heads were back-crossed, usually to the two parental lines (of the same accession number), and the number of flowers treated and the seeds produced was counted. All other heads were collected, threshed, the seed weight was determined, and calculated as percent of total plant dry weight. Some  $F_2$  and  $BC_1$  seeds were germinated and their pollen fertility, cytology and seedset upon selfing and open pollination were determined, as in their parents.

### Results

#### a. Germination and Development

The results of germination and development of intra- and interspecific  $F_1$  hybrids in various combinations among the six species dealt with in this publication are presented in Table 1. These data deserve two

explanations: *First*, the seeds of *T. mei*\* intraspecific (WS) hybrids that turned to albinos resulted from two parent plants which produced many albinos in various intra- and intergroup combinations (Katznelson and Putiyevsky, in press); both plants were albino heterozygotes, probably in the same locus. *Second*, the relatively low germination and high mortality rates in *T. vav* intraspecific combination may stem from the fact that most of these seeds were obtained by the NHP, in which heads and seeds were not selected, and many of them were somewhat shrunken (see Putiyevsky and Katznelson, 1973). This does not exclude some cytogenetic heterogeneity within *T. vav* as another possible cause.

Except for these cases, most of the intraspecific hybrids (86–100%) reached maturity, higher than the percentage in an interspecific (BS) cross combination. Within the latter, those including *T. mei* as a parent usually had better germination and development, while those of *T. vav* were much poorer (43.6%) than hybrids within crossability group I (64.7%).

\* Abbreviations: *T. ale*—*T. alexandrinum* L., *T. apr*—*T. apertum* Bobr., *T. ber*—*T. berytheum* Boiss., *T. meis*—*T. meironense* Zoh. et Lern., *T. sal*—*T. salmoneum* Mout., *T. vav*—*T. vavilovi* Eig.

Table 1. Germination and development of  $F_1$  hybrids

Species combination	Total $F_1$ seed planted	No. $F_1$ seed germinated	No. albino seedlings	No. dead before flowering (albinos excluded)	No. reached maturity	% of germinated seedlings
<i>T. ale</i> — <i>T. ale</i>	15	15		2	13	86.7
<i>T. ber</i> — <i>T. ber</i>	10	10			10	100.0
<i>T. apr</i> — <i>T. apr</i>	8	8		1	7	87.5
<i>T. mei</i> — <i>T. mei</i>	2	2	2		0	0.0
<i>T. vav</i> — <i>T. vav</i>	210	129		32	97	75.2
Total intraspecific combinations	245	164	2	35	127	77.4
<i>T. ale</i> — <i>T. ber</i>	32	32		14	18	56.2
<i>T. ale</i> — <i>T. sal</i>	28	22		9	13	59.1
<i>T. ale</i> — <i>T. apr</i>	30	28		6	22	78.6
<i>T. ale</i> — <i>T. mei</i>	27	27		4	23	85.2
<i>T. ber</i> — <i>T. sal</i>	68	65		27	38	58.5
<i>T. ber</i> — <i>T. apr</i>	75	70		35	35	50.0
<i>T. ber</i> — <i>T. mei</i>	26	26	1	2	23	88.5
<i>T. sal</i> — <i>T. apr</i>	45	39		13	26	64.5
<i>T. sal</i> — <i>T. mei</i>	26	25	4	3	18	72.0
<i>T. apr</i> — <i>T. mei</i>	35	29	1	9	19	65.5
Interspecific within crossability Group I	392	363	6	122	235	64.7
<i>T. ale</i> — <i>T. vav</i>	28	19		8	11	50.8
<i>T. ber</i> — <i>T. vav</i>	76	52		39	13	25.0
<i>T. sal</i> — <i>T. vav</i>	72	36		20	16	44.4
<i>T. apr</i> — <i>T. vav</i>	7	7	3	3	1	14.3
<i>T. mei</i> — <i>T. vav</i>	10	10		2	8	80.0
Between crossability Group I and <i>T. vavilovi</i>	193	124	3	72	49	40.0

b. Pollen Fertility of  $F_1$  Hybrids

Pollen fertility was examined in 127 WS and 293 BS  $F_1$  hybrids (Table 2). Although a few BS hybrids were sterile or semi-sterile, most of the hybrids were

fertile or subfertile (70–80% fertile pollen). There was only a small difference in pollen fertility between WS and BS combinations, and hardly any difference between BS crosses within crossability group I and between them and *T. vavilovi*.

Table 2. Frequency distribution of pollen fertility of intraspecific (W.S.) and interspecific (B.S.)  $F_1$  hybrids

Hybrids combination	Total plants examined	No. of plants with the following of pollen fertility percentage									Average pollen fertility %	
		0–10	10–20	20–30	30–40	40–50	50–60	60–70	70–80	80–90		90–100
T. ale—T. ale	13								1	3	9	90.8
T. ber—T. ber	10							2		3	5	85.6
T. apr—T. apr	7									2	5	91.4
T. vav—T. vav	97									2	95	94.8
Total intra-specific combinations	127							2	1	10	114	93.6
T. ale—T. ber	18					1				5	12	90.2
T. ale—T. sal	13								3	3	7	87.5
T. ale—T. apr	22	1						1	4	9	7	80.5
T. ale—T. mei	23							2	3	8	10	84.4
T. ber—T. sal	38	1				2	3	1	3	8	20	81.7
T. ber—T. apr	35				1	1	2	2	4	17	8	80.3
T. ber—T. mei	23								1	8	14	86.1
T. sal—T. apr	26					2			4	11	8	81.5
T. sal—T. mei	18							1	7	8	2	80.0
T. apr—T. mei	19							1	1	10	7	87.0
Interspecific within crossability Group I	235	2			1	5	6	9	30	87	95	84.5
T. ale—T. vav	11					1	3	4		2	1	63.6
T. ber—T. vav	13						1	2	3	5	2	78.6
T. sal—T. vav	16							6	2		8	79.0
T. mei—T. vav	8							1	2	4	1	81.4
T. apr—T. vav	1									1		81.0
Between crossability Group I and <i>T. vavilovi</i>	49					1	4	13	7	12	12	78.0

Table 3. Chromosome configurations in meiosis of hybrids ( $2n = 16$ )

Hybrids combination	$F_1$				$F_2$					
	No. plants examined	No. cells examined	Cells with		% Cells with univalents	No. plants examined	No. cells examined	Cells with		% Cells with univalents
			7II + 2I	8II				7II + 2I	8II	
T. ale × T. ber	4	110	9	101	8.1	5	84	1	83	1.2
T. ale × T. sal	1	30	—	30	0	3	45	—	45	0
T. ale × T. apr	5	131	11	120	8.4	6	99	5	94	5.0
T. ale × T. mei	5	92	5	87	5.4	8	127	8	119	6.3
T. sal × T. ber	7	192	4	188	2.0	6	101	—	101	0
T. ber × T. apr	10	236	3	233	1.3	5	77	1	76	1.3
T. ber × T. mei	3	55	5	50	9.0	4	68	—	68	0
T. sal × T. apr	5	124	2	122	1.6	2	24	—	24	0
T. sal × T. mei	4	114	1	113	0.7	3	48	—	48	0
T. apr × T. mei	6	164	2	162	1.2	10	172	3	169	1.1
T. ale × T. vav	1	30	4	26	13.3	1	18	1	17	5.5
T. ber × T. vav	3	90	1	89	1.1	2	30	—	30	0
T. sal × T. vav	4	88	9	79	10.2	—	—	—	—	—
T. mei × T. vav	1	30	1	29	3.3	2	28	—	28	0
T. apr × T. vav	1	21	1	20	4.8	—	—	—	—	—

Table 4. Seed set upon selfing, backcrosses and open pollination of *F<sub>1</sub>* hybrids

Hybrid combination	Total No. plants examined		Seed set upon:				Backcross to ♀ parent				Backcross to ♂ parent				Open pollination		
	No. of flowers	No. of seeds	No. of flowers	% seedset	No. of flowers	No. of seeds	No. of flowers	% seedset	No. of flowers	No. of seeds	No. of flowers	No. of seeds	% seedset	No. of flowers	No. of seeds	Seed weight, g	seed weight / plant weight %
<b>Intraspecific</b>																	
T. ale - T. ale	13	2015	464	23.1	—	—	—	—	—	—	—	—	—	—	5.92	1.73	
T. ber - T. ber	10	1659	45	2.1	—	—	—	—	—	—	—	—	—	—	7.96	7.19	
T. apr - T. apr	7	781	0	0	—	—	—	—	—	—	—	—	—	—	0.76	1.69	
<b>Interspecific</b>																	
T. ale - T. ber	18	5688	1089	19.1	243	70	28.8	282	119	42.2	17.11	3.70					
T. ale - T. sal	13	3886	1090	28.0	89	33	37.0	78	35	44.9	16.85	3.50					
T. ale - T. apr	21	5728	227	4.1	318	38	11.9	172	39	22.7	11.82	1.96					
T. ale - T. mei	23	5755	788	13.8	252	63	25.0	230	82	35.6	22.15	1.57					
T. ber - T. sal	37	10712	1192	11.1	206	100	48.5	231	106	45.9	26.45	2.74					
T. ber - T. apr	34	9136	490	5.4	273	75	27.5	337	79	23.4	17.22	1.27					
T. ber - T. mei	23	4398	255	5.1	284	43	15.1	175	50	28.6	12.20	1.79					
T. sal - T. apr	25	5846	5	0.1	117	49	41.9	115	33	28.7	15.59	2.21					
T. sal - T. mei	18	5201	3	0.1	220	59	26.8	213	54	25.3	6.06	0.38					
T. apr - T. mei	19	6401	67	1.0	260	28	10.8	297	47	15.8	12.37	0.75					
T. ale - T. vav	11	1247	48	3.8	—	—	—	—	—	—	9.35	1.91					
T. ber - T. vav	13	4000	1	0.02	25	7	28.0	21	0	0.0	4.35	1.61					
T. sal - T. vav	16	3441	53	1.6	111	17	15.3	45	12	20.8	1.74	0.46					
T. apr - T. vav	1	230	0	0	25	0	0.0	25	0	0.0	0	0					
T. mei - T. vav	8	1369	0	0	25	11	44.0	—	—	—	2.05	2.50					

c. Cytology of *F<sub>1</sub>* Hybrids

Cytology in meiosis was examined in 60 interspecific *F<sub>1</sub>* hybrids and appeared to be quite regular (Table 3). No multivalents were discerned in 1498 cells that were examined, but quite a few of the cells had two univalents instead of a bivalent. These were mainly in T. ale × T. vav, T. sal × T. vav and T. ber × T. mei cross combinations. The more or less normal meiosis corresponds to the normal or subnormal pollen stainability.

An interesting feature found in the majority of *F<sub>1</sub>* hybrids within crossability group I (but not between them and T. vav) is the appearance of B chromosomes. This topic will be reported and discussed elsewhere.

d. Seed Set of *F<sub>1</sub>* Hybrids

The data on seed set in *F<sub>1</sub>* hybrids upon selfing, backcross to both parental populations and on exposure to a large variety of pollen sources, are given in Table 4.

i) *Selfing*. Seeds were formed in many cross combinations. Ample seeds (up to 28% seed set) were produced when T. ale was one of the parents. Good seed set upon selfing, though lower than the T. ale combinations, was obtained in progenies of T. ber — especially T. ber × T. sal, T. ber × T. mei, and reciprocals. However, a closer look at the results of the individual plants traces the good seed set upon selfing to two individual T. ber plants, NYT 206/1 and NYT 206/2. This can be seen in Table 5.

*T. ber × T. sal hybrids*: Of 37 *F<sub>1</sub>* plants, only 12 set any seed when selfed, and 11 of them were progenies of NYT 206/1.

In the *T. ber × T. apr* hybrids, seed set upon selfing may be similarly traced to Plant NYT 206/2. However, one T. apr × T. ber *F<sub>1</sub>* hybrid, a progeny of NYT 206/3, a third plant of NYT 206, did not set any seed. It should be stressed here that seed set upon selfing of the two parental plants, 206/1 and

Table 5. Seed set upon selfing in some *F<sub>1</sub>* hybrids, progenies of *T. ber* (206, 252, 529) crossed with *T. sal* (260) and *T. apr* (1471, 1772)

Parents	Number of plants		Hybrid	
	♀	♂	florets	seeds
206/1	260	6	2015	313
252	260	6	1506	6 (all from one plant)
529/1	260	11	3192	0
260/1	206/1	5	1902	873
260/1	252	6	1130	0
260/1	529	3	967	0
206/2	1471	6	1021	241
206/2	1772	3	1298	200
529/3	1772	3	803	0
529/3	1471	8	2039	0
252	1772	4	1107	0
but 1471	206/3	1	240	0

206/2, was 0.51% and 0.0%, respectively (Putiyevsky and Katznelson, 1970).

Of 23 *T. ber* × *T. mei*  $F_1$  hybrids only two set any seed. Their common parent was *T. ber* NYT 252/1, which had quite good seed set when selfed.

Other cross combinations set hardly any seeds, and those which did form were often shrunken.

ii) *Backcrosses*. Seed set upon backcrossing, though varying considerably in individual cross combinations, was quite high on average, and exceeded 40% in some combinations. There seems to be a positive correlation between seed set in backcrosses to the two parental populations, the only serious deviation being crosses involving *T. vav*.

Also, seed set was higher in the combinations between *T. ale*, *T. ber* and *T. sal* (41%) than in combinations between any of the three taxa and *T. mei*—*T. apr* (22.6%), or between any of all five taxa and *T. vav* (16.9%). Reciprocal differences of the magnitude found in the crossability between species (Putiyevsky and Katznelson, 1973) were quite common.

iii) *Open pollination*. The seed set in WS and BS  $F_1$ s, given in grams seeds and as percent of plant weight, is also presented in Table 4. Outstandingly high amounts of seed were produced in *T. ber* ×

*T. ber* combinations, followed by those of *T. ale*—*T. ber*—*T. sal*. This is in accordance with the success in backcrossing. However, unexpected high seed set was found in the *T. mei* × *T. vav* cross combination, both in backcrossing and in open pollination. The hybrid nature of these plants was ascertained both by morphology and pollen data, and we cannot explain this.

#### e. $F_2$ Plants

As in the case of the second and third crossability groups (Katznelson and Putiyevsky, 1974; Putiyevsky and Katznelson, 1974), germination of  $F_2$  seeds was excellent: 367 out of 381 seeds, which was much better than that of parents or  $F_1$  hybrid seed. One of the seedlings, *T. ber* × *T. mei*, was albino, and 92 plants died before flowering.

Pollen stainability was estimated in the remaining 274 plants; 57  $F_2$  plants, of relatively low pollen stainability, were studied cytologically.

The results on pollen fertility of  $F_2$  and  $BC_1$  hybrids appear in Table 6. Comparison of these results with those of their parents (Table 2) shows that pollen fertilities of  $F_2$  were no higher, and often lower, than those of  $F_1$ s. This is especially obvious in the hybrids involving *T. vav*, which fact helped us to ascertain the hybrid nature of their highly fertile  $F_1$  parents.

Table 6. Pollen fertility of  $F_2$  +  $BC_1$  interspecific hybrids

Combination		No. of plants with the following pollen fertility percentage										Total no. plants examined	Ave. pollen fertility %
♀ = $F_1$	♂	0-10	10-20	20-30	30-40	40-50	50-60	60-70	70-80	80-90	90-100		
<i>T. ale</i> — <i>T. ber</i>	B.C.	1						2	1	7	12	23	85.6
<i>T. ale</i> — <i>T. sal</i>	B.C.							1	1	5	5	12	87.0
<i>T. ale</i> — <i>T. apr</i>	B.C.						4	3	4	8	4	23	80.6
<i>T. ale</i> — <i>T. mei</i>	B.C.					1	3	5	4	4	2	19	71.9
<i>T. ber</i> — <i>T. sal</i>	B.C.								5	9	16	30	88.9
<i>T. ber</i> — <i>T. apr</i>	B.C.						1	1	2	15	9	28	85.6
<i>T. ber</i> — <i>T. mei</i>	B.C.								2	3	3	8	86.2
<i>T. sal</i> — <i>T. apr</i>	B.C.					1			2	6	2	11	82.6
<i>T. sal</i> — <i>T. mei</i>	B.C.								2	2	3	7	86.4
<i>T. apr</i> — <i>T. mei</i>	B.C.								3	6	12	21	89.1
<i>T. ale</i> — <i>T. ber</i>	Selfing						1	1	1	8	4	15	82.9
<i>T. ale</i> — <i>T. sal</i>	Selfing									3	5	8	90.0
<i>T. ale</i> — <i>T. apr</i>	Selfing							1	1	6	8	16	84.8
<i>T. ale</i> — <i>T. mei</i>	Selfing			1	1	2	1	2	2	7	6	22	75.0
<i>T. ber</i> — <i>T. sal</i>	Selfing	1							2	4		7	70.0
<i>T. ber</i> — <i>T. apr</i>	Selfing								1	2		3	84.0
<i>T. ber</i> — <i>T. mei</i>	Selfing					1				2	2	5	81.0
<i>T. sal</i> — <i>T. apr</i>	Selfing									2		2	85.5
<i>T. apr</i> — <i>T. mei</i>	Selfing								1	3	3	7	87.9
Within crossability Group I		2	1	1	5	10	16	34	102	96	267	83.2	
<i>T. mei</i> — <i>T. vav</i>	B.C.					2						2	44.5
<i>T. ale</i> — <i>T. vav</i>	Selfing						1	1	1	1		4	66.0
<i>T. ber</i> — <i>T. vav</i>	Selfing						1					1	50.0
Between crossability Group I and <i>T. vavilovi</i>					2	2	1	1	1			7	60.7

Table 7. Seedset in various progenies of interspecific  $F_2$  hybrids

Combinations			No. of plants examined	Seedset upon selfing			Seedset upon open pollination	
♀ = $F_1$	♂			No. of flowers	No. of seeds	% Seedset	Seed weight, g	seed weight / plant weight %
T. ale	T. ber	B.C. <sub>1</sub>	23	4929	322	6.5	4.38	0.55
		Selfing	15	2159	97	4.5	5.26	1.44
T. ale	T. sal	B.C. <sub>1</sub>	12	2455	250	10.2	10.20	1.71
		Selfing	8	903	11	1.2	3.12	1.92
T. ale	T. apr	B.C. <sub>1</sub>	23	3556	151	4.3	5.33	0.44
		Selfing	16	2997	184	6.1	4.44	1.15
T. ale	T. mei	B.C. <sub>1</sub>	19	4010	29	0.7	1.50	0.19
		Selfing	22	4830	12	0.2	0.12	0.02
T. ber	T. sal	B.C. <sub>1</sub>	30	5915	408	6.9	5.17	0.46
		Selfing	7	1435	14	1.0	0.45	0.34
T. ber	T. apr	B.C. <sub>1</sub>	28	4773	304	6.4	20.97	1.90
		Selfing	3	630	5	0.8	0.20	0.28
T. ber	T. mei	B.C. <sub>1</sub>	8	1318	47	3.6	2.07	0.83
		Selfing	15	1140	110	0.1	0.12	0.13
T. sal	T. apr	B.C. <sub>1</sub>	11	1812	111	6.1	6.72	2.28
		Selfing	2	300	62	20.7	1.18	3.76
T. sal	T. mei	B.C. <sub>1</sub>	7	1000	0	0	0.07	0.04
T. apr	T. mei	B.C. <sub>1</sub>	21	4120	0	0	0.07	0.01
		Selfing	7	1600	0	0	0.07	0.06
T. ale	T. vav	Selfing	4	462	47	10.2	1.80	2.40
T. ber	T. vav	Selfing	1	618	213	34.5	1.82	1.74
T. mei	T. vav	B.C. <sub>1</sub>	2	270	0	0	0.03	0.01

Cytologically, the  $F_2$  were similar to their  $F_1$  parents in the absence of multivalents in meiosis and in the appearance of 7II and 2I cells (Table 3).

Seed set upon selfing of  $F_2$  hybrids varied between 0–34.5% (Table 7), and some seed was obtained in most combinations. Quite obviously, plants that originated from selfed  $F_1$  produced many seeds when selfed. Thus, the single  $F_2$  T. ber × T. vav seed that was obtained from selfing some 4000 florets (Table 4) produced a highly self compatible plant. It is also obvious that selfed flowers formed seeds in many BC plants, in combinations whose  $F_1$  plants were almost completely self incompatible. A good correlation found between seed set upon selfing and seed set by open pollination may suggest that many of the seeds obtained in open pollination may result from selfing.

### Discussion

An obvious conclusion from the data presented is that the six taxa of subsection *Alexandrina* Zoh. (Zohary, 1972) represent a heterogeneous group. First, *T. vavilovi*, though superficially somewhat similar morphologically to the five other taxa, is quite isolated genetically from them, both in terms of crossability (Putiyevsky and Katznelson, 1973) and of lower seed germination and development of  $F_1$  seedlings, much smaller plants with somewhat lower pollen fertility and only few seeds. The effect of isolation through even lower pollen fertility and seed

formation is obvious also in  $F_2$  and BC plants. Thus, *T. vavilovi* is a clear-cut species, well isolated from the five other taxa.

It may be added that there is some evidence that *T. vavilovi* is not homogeneous cytogenetically, as some hybrids between geographically remote populations were sterile and did not set seeds (Katznelson, 1969).

The five other taxa are hardly isolated genetically. Crossability, germination, development, pollen fertility and seed set (BS  $F_1$ s) are similar to WS  $F_1$ s (normal) or only somewhat lower (subnormal).

The mechanism isolating T. ale—T. ber—T. sal from T. mei—T. apr is geographical and ecological. T. apr grows in the Caucasus and Turkey in shrubby habitats (Bobrov, 1950), and the Type Specimen of T. mei was collected in a small clearing within a dense maquis on Mt. Meiron, Israel, *i.e.*, both taxa grow in regions and conditions where T. ale is not grown, and where T. sal and T. ber do not occur naturally. The natural habitats of T. sal and T. ber are wet places, roadsides and waterlogged soils, etc., often border areas sown to berseem. Hybrid swarms between T. ale and T. ber under these conditions are quite common, and they are often regarded as spontaneous berseem. T. ale-like plants are eliminated by natural selection in a process — and due to causes — similar to *Hordeum vulgare* — *H. spontaneum* hybrid swarms (Zohary, 1971). Moreover, in

a study of the domestication process in T. ber we found a clear-cut gene flow from T. ale to T. ber, originating from some 2–3% interspecific hybridization, and the F<sub>1</sub> plants were easily discerned morphologically. By this slow process, lines intermediate between T. ale and T. ber in many traits are being formed (Katznelson, in preparation).

Thus, T. ber and/or T. sal are, or are closest to, the wild progenitor of berseem, the cultivated T. ale. This contrasts with Bobrov's opinion that *T. apertum* is the wild progenitor of berseem. He concluded this on the basis of the open calyx tube of *T. apertum*, without knowing *T. berytheum*, and especially *T. salmoneum*, whose calyx throat is not hairy. The two latter species are much more similar morphologically to berseem than either T. apr or T. mei. In fact, the five species of crossability group I can be separated into two species complexes that differ morphologically: T. mei–T. apr are more delicate, have smaller heads and leaves, and the fruiting head, especially in T. mei, is more cylindrical than in the T. ale–T. ber–T. sal group.

It should be stressed here that as the five taxa are very homogeneous cytogenetically, our conclusion on the origin of berseem is based mainly on morphology, but we have studied a wealth of material at all growth stages. While some taxonomic considerations will be discussed elsewhere, the following are worth discussing here. The three taxa, T. ale, T. sal and T. ber, show morphological intergradation in many characteristics. However, some characteristics of T. ale were not found in the two wild taxa, and vice versa.

a) A very important agronomic trait in the cultivars of the Musgawi group is the very low basal branching of erect branches, which is the basis for the multi-harvest nature of these cultivars. T. sal and some populations of T. ber are erect, but they never branch (and rebranch) at their bases. (This trait was seen in a few plants of T. apr, continuously grazed, from Sinop). On the other hand, the branching pattern of all cultivars of the Fahli and Saidi groups of T. ale is similar, if not identical, to that of some wild types of T. sal and T. ber.

b) The calyx tube in T. ale is completely open, while it is closed by a ring of hairs in T. ber, and by a callus ring without hairs in T. sal. This character affects the easy threshability of the cultivated berseem and much stronger seed retention in the wild taxa. However, the callus ring in T. sal is sometimes hardly seen.

In other morphological characters the three taxa show more or less continuous variation.

c) We found that many populations of T. ber–T. sal have B chromosomes, in almost every plant examined. We, and many other scientists, have never found them in T. ale (Mehta et al., 1963; Almeida

and Carvalho, 1962; Putiyevsky and Katznelson, 1970).

d) While the wild T. ber and T. sal are practically self-incompatible, T. ale varies in its genetic system from selfers to self-incompatibles, even within one population (Putiyevsky and Katznelson, 1970). This change in the genetic system is quite common in many other cases of domestication, such as tomato, barley, cotton and, to some extent, wheat. This may lead to the conclusion that berseem is a newly domesticated plant, but this is erroneous, as berseem was probably the earliest forage crop to be sown from the first Egyptian dynasty, i.e., 3500–3800 B.C.

One may connect the last topic to the breakdown of self-incompatibility in crosses involving T. ber. Such breakdown in interspecific hybrids has already been reported by Pandey (1957) in the F<sub>1</sub> of *T. uniflorum* × *T. repens*. Pandey concluded that although the two quite remote species have the same system of incompatibility, they differ in the location of the S gene in the chromosomal complement.

Our case, however, is quite different. Intrapopulation differences (NYT 206/1 and 206/2 versus NYT 206/3) show that we deal here not with species but with individuals. This apparently rules out differences in location of the S locus, and suggests that the breakdown of the incompatibility system is genetic. Genetical analysis in the future may elucidate this subject.

#### Literature

- Abdel-Tawab, F. M.: Interspecific hybridization in five species of *Trifolium* L. Ph. D. Thesis. Univ. Michigan Press (1968).
- Almeida, J. L. F., Carvalho, H. P.: Trevos autotetraploides. III. O caso de *Trifolium alexandrinum* L. Agron. Lusit. **24**, 45–78 (1962).
- Bobrov, E. G.: Vidi Kleverov U.R.S.S. (in Russian). Acta Inst. Bot. N. Acad. Sci. U.S.S.R. Ser. 1, **6**, 164–336 (1947).
- Bobrov, E. G.: New clovers for culture (in Russian). Akad. Nauk, U.R.S.S. Leningrad (1950).
- Evans, Alice M.: Species' hybridization in *Trifolium*, II. Investigating the prefertilization barriers to compatibility. Euphytica **11**, 256–262 (1962).
- Katznelson, J.: Population studies and selection in berseem clover (*Trifolium alexandrinum* L.) and the closely related taxa. 3rd An. Rep. to the USDA, Project A10-CR-56 (1969).
- Katznelson, J.: Semi-Natural interspecific hybridization in plants. Euphytica **20**, 266–269 (1971).
- Katznelson, J., Putiyevsky, E.: Cytogenetic studies in *Trifolium* spp. related to berseem. II. Relationships within the *echinata* group. Theoret. appl. Genet. **44**, 87–94 (1974).
- Mehta, R. K., Subramanyan, K. W., Swaminathan, M. S.: Studies on induced polyploids in forage crops. III. Growth, cytological behaviour and seed fertility of C<sub>1</sub>, C<sub>2</sub>, C<sub>3</sub> cultures of berseem. Indian J. Genet. Pl. Breed. **23**, 67–81 (1963).
- Müller, Gertrud: Untersuchungen über das Pollenschlauchwachstum bei verschiedenen *Trifolium*-Artkreuzungen. Züchter **30**, 28–30 (1960).

- Oppenheimer, H. R.: The origin of Egyptian clover with critical revision of some closely related species. Bull. Res. Coun. Israel **7D**, 202–221 (1959).
- Pandey, K. K.: A self-compatible Hybrid from a cross between two self-incompatible species in *Trifolium*. J. Hered. **48**, 278–281 (1957).
- Putiyevsky, E., Katznelson, J.: Chromosome number and genetic system in several *Trifolium* species related to *T. alexandrinum*. Chromosoma (Berl.) **30**, 476–482 (1970).
- Putiyevsky, E., Katznelson, J.: Cytogenetic studies in *Trifolium* ssp. related to berseem. I. Intra- and inter-specific hybrid seed formation. Theoret. appl. Genet. **43**, 351–358 (1973).
- Putiyevsky, E., Katznelson, J.: Cytogenetic studies in *Trifolium* spp. related to berseem. III. The relationships between the *T. scutatum*, *T. plebeium* and the *echinata* groups. Theoret. appl. Genet. **44**, 184–190 (1974).
- Zohary, D.: Origin of South-West Asiatic cereals: wheats, barley, oats and rye. In: Plant Life of South-West Asia (Eds. Davis, P. H., Harper, P. C., Hedge, I. C.) pp. 236–260. Bot. Soc. Edinburgh, Univ. Press Aberdeen (1971).
- Zohary, M.: A revision of the species of *Trifolium* sect. *Trifolium* (Leguminosae). II. Taxonomic treatment. Candollea **27**, 99–158, 249–264 (1972).

Received September 12, 1974

Communicated by H. Stubbe

E. Putiyevsky  
J. Katznelson  
Division of Pasture and Forage Crops  
Agricultural Research Organization  
Nve-Ya'ar, P. O. Haifa (Israel)  
D. Zohary  
Division of Genetics  
Hebrew University  
Jerusalem (Israel)