Cladistic analysis of the genera: *Trifolium, Trigonella* and *Melilotus* (Fabaceae: Papilionaceae) in Egypt.

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ABSTRACT

The morphological characters of the three genera (*Trifolium*, *Trigonella* and *Melilotus*) of the tribe *Trifolieae* were analysed and used to construct their phylogenetic relationships. The cladistic analysis reveals strong similarities between the species of genus *Trifolium* which clustred in one clade, and also for species of genus *Trigonella*. For *Melilotus*, all species are strongly related to each other except for *M. albus* which is evolved early and its taxonomic position is unclear. This species is very rare in Egypt (recorded once), the number of flowers per raceme and length of inflorescence in relation to subtending leaves play an important role in the identification of the species. Leaflets are oblong obovate to lanceolate, pods are obliquely ovate to elliptic with sharp acute beak. The seed protein pattern of the most common species of the previous genera was analyzed using Sodium Dodecylsulphate –Polyacrylamide Gel Electrophoresis (SDS-PAGE) technique. The seed protein data showed that all species of genus *Trifolium* form one clad, the second clad includes all species of *Trigonella*, the third clad contains all species of *Melilous*. These data are compatible with taxonomic position of species of the three genera.

KEYWORDS: Cladistics, Trifolium, Trigonella and Melilotus, Papilionaceae, Egypt

INTRODUCTION

The genera *Trifolium, Trigonella, Melilotus* and *Medicago* are in the tribe Trifolieae of subfamily Papilionoideae (Family Fabaceae). These three genera are closely related to each other and are belonging to tribes *Trigonelleae* and *Trifolieae*, (Schulz 1901). The *Trifolieae* was described by Hyen (1981). Trautvetter (1841) stated that it is difficult to establish clear cut limits between the genera of *Trifolieae*. Linnaeus (1753) classified *Melilotus* as one group under *Trifolium*. Seringe (1825) placed the two genera *Melilotus* and *Trigonella* in a special section known as *Grammocarpus* of genus *Trigonella*. Battacharyya (1958) concluded that *Trigonella* and *Melilotus* might be treated as two subgenera. Heyn (1966) reported the presence of intermediate species between the three genera *Medicago*, *Trigonella* and *Melilotus*. Therefore, some species of *Medicago* viz. *M.ruthenica* and *M.polycarpa* are considered by most botanists to be intermediates between *Trigonella* and *Medicago*.

The tribe *Trifolieae* seems to be natural units. Its four genera do not constitute a phyletic group; though, there are still some morphological links between its genera (Zohary & Heller 1984). Molecular phylogenetic relashionships of species of genus *Medicago* was reviewed by Bena (1998), who concluded that genus *Medicago* is closely related to the other studied three genera, and it was chosen by this study as outgroup. Small *et al.* (1987) stated that the highly evolved syndrome of floral adaptations provides in *Medicago* an appropriate means of circumscribing the genus. Accordingly, 23 *Trigonella* species were transfered to *Medicago*. Some intermediate characters such as the horn, the stigma and the flower tripping mechanism characterize these species. Moreover, Greuter *et al.* (1989) regarded *Trigonella polyceratia* as a member of genus *Medicago*, which is recognized by Tächkolm (1974) as *Trigonella polyceratia*. Endo & Ohashi (1997) carried out a cladistic analysis of tribes Cicereae, Trifolieae and Vicieae; of which Cicereae and Vicieae were found to be a monophyletic group.

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and Trifolieae was its sister group. Steel *et al* (1997) studied the phylogenetic relationships of the tribes Trifolieae and Vicieae. They pointed out that *Medicago* supported a monophyletic group distinct from that formed by *Melilotus*, which is nested within *Trigonella* and *Trifolium* basal to the remainder of *Trifolieae*.

Due to the existing overlap in identifying the genera of the tribe *Trifolieae* on morphological basis, more accurate and precise criterion; viz electrophoretic analysis of storage proteins, was applied to achieve more convenient generic identification (Robnson & Megarrity 1975; Dallying et *al*: 1979; Wrigley *et al*: 1982; Ferguson & Grobe 1986). The present work represents the first attempt to study the phylogenetic relationship between the three genera in Egypt using morphological characters and variability of seed storage proteins.

MATERIALS AND METHODS

Morphology: A total of 34 species belonging to the genera *Trifolium, Trigonella* and *Melilotus* of tribe Trifolieae were the subject of systematic revision by Kamel (1992), Hassan (1992), and El-Bous (1995). The morphological characters of their species (31charaters) are of taxonomic importance and were scored (Table 1). Many of these characters are used for the first time to construct the phylogenetic relationships.

Seed protein analysis: seeds belonging to sixteen species belonging to. Trifolium, Trigonella and Melilotus of different origins were obtained from herbarium of Suez Canal University. Characterization and molecular mass determination of seed storage proteins was carried out using one-dimensional SDS-polyacrylamide gel electrophoresis. Samples were prepared for electrophoresis by extracting protein from 0.5g seed powder in 1cm³ of 0.2M Tris/Hcl, pH 8, and 1mM phenylmethylsulphonyl fluoride (PMSF) for 2h. in a refrigerator. The extract was centrifuged at 5000g for 20 min. and proteins in the supernatant were precipitated with 5 volumes of cold acetone at -20 ^oC for 2h. Pellets obtained after centrifugation were dissolved in 0.02 ml of sample buffer (0.125 M Tris/Hcl, pH6, 8, 2% m/v SDS, 10% m/v sucrose, 1% v/v β -mercaptoethanonl, 0.1% m/v bromophenol blue) and denaturated by heating at 80 $^{\circ}$ C for 2-3 min. 17% SDS-polyacrylamide gel slabs were prepared as described by Laemmli (1970). Equal amounts of protein were loaded per lane. Electrophoresis was carried out in Tris/glycine. SDS running buffer (0.25M Tris, 1.88M glycine 0.1% SDS) using vertical gel electrophoresis units (Mini-protein cell, BioRad, USA) at 140 V for the first 15 min followed by 150 V until the indicator dye reached the bottom of the gel. Gels were stained over night in 20 cm³ of 0.25% acetic acid, 50% (v/v) methanol, 7% (v/v) glacial acetic acid and destained by shaking overnight in 50% methanol and 7% glacial acetic acid. The relative mobility of the band was calculated and the presence or absence of each band was treated as a binary character (Table 2).

The cladistic analysis of the protein and the morphological characters are followed the principles of phylogenetic reasoning elaborated by Hennig (1966). The results were analyzed by computer using Henning 80 program of Farris. The data were entered in computer with the assumption that species *Medicago sativa* was considered to be the outgroup because it is a native species to the Old World as mentioned by Small (1987).

RESULTS

Analysis of the morphological characters (Table 1)

- 1. Life Span: annual (0), perennial (1).
- 2. Stem: procumbent (0), erect or decumbent (1), ascending (2), prostrate (3).
- **3. Indumentum:** Hairy (0), and glabrous (1).
- 4. Stipules shape: Triangular oblong (0), ovate (1), and lanceolate (2).
- 5. Stipules margin: Entire (0), dentate (1), and incised (2).

6. Leaf shape: obovate (0), Elliptical (1), and lanceolate (2).

7. Leaf margin: Entire (0), dentate to denticulate (1), and serrate to serrulate (2)

8. Leaf apex: acute to acuminate (0), rounded to truncate (1), emarginate or mucrounate (2).

9. Calyx teeth: shorter than calyx tube (0), as long as calyx tube (1), longer than calyx tube (2).

10. Corolla colour: white -to cream (0), yellow (1), and pink to mauve (2).

11. Stander length: longer than keel and wing (0), shorter than keel and wing (1).

12. Stander: distinct into limb and claw (0), not distinct into limb and claw (1).

13. Apex: notched (0), and obtuse (1). **14. Wing shape:** oblong (0),

oblanceolate (1). **15. Wing claw:** shorter than Limb (0),

or claw as long as limb (1), also claw longer than limb(2).

16. Wing horn: present (0), absent (1).

17. Keel shape: oblong (0), ovate (1).

18. Keel apex: obtuse or rounded (0), Acute (1).

19. Filament of Stamens: connate for nearly; 2/3 of its length (0), connate for 1/2 of its length (1), or 3/4 of its length (2).

20. Shape of Ovary: Linear-oblong (0), or subglobular to ovate (1), cylindrical (2).

21. Number of ovules: from one to two (0), from three to eight (1), or from five to twenty (2).

22. Ovary: glabrous (0), hairy (1).

23. Style length: longer than ovary (0), or shorter (1).

24. Style: curved (0), straight at apex (1).

25. Pod shape: or linear (0), ovate to elliptical (1), obovate (2), curved oblong-circular (3).

26. Pod surface: membranous (0), with concentric striation (1).

27. Pod apex: obtuse (0), sharp acute (1), mucronate (2), or tapering (3).

28. Seed shape in outline: ovate (0), oblong to circular (1), or oblong ovate (2).

29. Seed colour: brown (0), or yellow (1), orange red (2).

30. Surface of seed coat: smooth (0), wrinkled (1).

31. Length of peduncle: from 0-4 cm (0), from 4-10 cm (1), of from 10-33 cm (2).

Cladogram (Figure 1)

The cladogram is based on 31 morphological characters, each number at a node or branching point representing a hypothetical ancestor and with the numbers of lines correspond to the characters tabulated in table 1.

Only one tree resulted as shown in (Figure 1), with two main clades. The first clade includes all *Trifolium* species. *Trifolium argutum* evolved early based on characters: leaf margine serrate to sirrulate, calyx teeth is shorter than calyx tube and ovary hairy. Other *Trifolium* species are separated in two groups or two subgenera separated at nodes 59, 60. The first node at node 59 uses the characters: leaf obviate, calyx teeth as long as calyx tube and style up curved at apex, while node 60 carries different characters: leaf lanceolate, calyx teeth longer than calyx tube and style straight. The first subgenus can also be separated into two

Table 1: Data matrix of the 31 morphological characters used to construct the phylogeny of the three genera (*Trifollium*, *Trigonella & Melilotus*) of family Papilionaceae of Egypt

Species	1	1	2	3			
- -		0	0	0			
Medicago sativa	1202	011202011	.000001001	103100000			
Trifolium alexandrinum	0202	011210000	020010100	001020211			
Trifolium tomentosum	0201	101112000	020110000	002032112			
Trifolium repens	1311	002210001	.011001010	013031012			
Trifolium stellatum	0201	101220001	.010011100	001022111			
Trifolium resupinatum	0200	001112000	000010000	001030011			
Trifolium campestre	0201	001121011	.100010101	103020000			
Trifolium argutum	0200	020222000	000010100	011030000			
Trifolium incarnatum	0201	101222001	.020010100	011032112			
Trifolium nigrescens			000000010				
Trifolium glanduliferum	0211	201222000	000000100	002032110			
Trifolium angustifolium	0310	002202101	.011000101	001030010			
Trifolium fragiferum	1312	221110000	111000000	002032012			
Trifolium philistaeum	0211	011022011	.120010101	103022111			
Trifolium scabrum	0201	201110001	.020000000	001030210			
Trifolium lappaceum	0210	010222001	.020010100	011030010			
Trifolium purpureum	0202	021222001	.000010100	011030110			
Trifolium dasyurum	0201	210222101	.020000100	011022011			
Trifolium dichroanthum	0200	011122001	.020010100	011021011			
Trigonella maritima	0111	201201000	000110021	000112100			
Trigonella arabica	0211	200200000	000100021	000012010			
Trigonella stellate	0010	001201100	000010010	103212100			
Trigonella cylindracea	0102	001201000	000001201	103212100			
Trigonella glabra	0111	201201000	000002210	103132000			
Trigonella laciniata	0102	201211010	000000111	101120000			
Trigonella occulta	0000	101211000	120100101	111130000			
Trigonella foenum-graecum	0110	011201000	000102121	100232010			
Trigonella polyceratia	0110	001001000	020102010	100132000			
Trigonella anguina	0312	201211110	010102121	003122000			
Melilotus indicus			.000000101				
Melilotus segetalis			000001101				
Melilotus sulcatus			000000101				
Melilotus messanensis			000002101				
Melilotus albus Medicus			.000001010				
<i>Melilotus serratifolius</i>	0200	212101100	000102101	002211000			

subgroups at node 55, including *Trifolium alexandrinum*, *T. scabrum*, *T. stellatum*, *T. incarnatum*, *T. tomentosum* and *T. glanduliferum*. *Trifolium alexandrinum* considered as basal group to the other members in this group by having stipule shape ovate and its margin dentate, *Trifolium glanduliferum* is a sister to *Trifolium tomentosum*. *Trifolium glanduliferum* having hairy plants, margin of stipules incised, leaf apex rounded, calyx teeth as long as calyx tube, claw of wing longer than limb, keel shape ovate, keel apex acute and length of peduncle from 0-4 cm. *Trifolium tomentosum* is a glabrous plant or glabrescent or with sub apprised to patent hairs, stipules margin dentate, leaf apex rounded, and length of peduncle 10 - 33 cm. The second subgroup at node 56, includes *T. resupinatum*, *T. nigrescens*, *T. repens*,

T. frgiferum. Trifolium resupinatum in this group is basal to others since it has stem ascending, plant hairy, corolla yellow, keel apex acute and seed shape is ovate. *T. repens* is sister to *T.fragiferum*. The first has stipules ovate, with entire margine, leaf obovate, with serrate to serrulate margine, leaf apex retuse, stander apex obtuse, staminal tube of androecium connate for half of its length, style straight, pod curved oblong and seed shape oblong to circular. *Trifolium fragiferum* has lanceolate stipules, and its margin incised, leaf shape lanceolate, with dentate margin and apex rounded, stander apex notched, staminal tube of androecium connate for 2/3 of its length, style curved at apex, pod shape obovate and seed shape oblong ovate.

The second group of genus *Trifolium* is separated at node 60 to including *T. purpureum*, *T. angustifolium*, *T. lappaceum*, *T. dasyurum*, *T. dichroanthum*, *T. campestre and T. philistaeum*. *Trifolium purpureum* is basal group having calyx teeth longer than calyx tube, ovary glabrous and style straight. *Trifolium campestre and T. philistaeum*. Are closely related species, *Trifolium campestre* is hairy plants, obovate leaf and a rounded apex; the claw of wing is shorter than limb, seed shape ovate, seed colour brown, and surface of seed coat smooth and length of peduncle 0-4cm. *Trifolium philistaeum*, glabrous, leaf shape is elliptic, and its apex is acute, the claw of wing is longer than limb, seed shape oblong ovate, seed colour yellow, surface of seed coat wrinked and length of peduncle 4-10 cm.

The second largest clade includes *Trigonella* and *Melilotus* species, which is separated from other clade at node 65 which has the characters: stem erect to decumbent, corolla yellow, stander are not distinct into limb and claw, pod ovate elliptical. At node 61, *Trigonella occulta* has evolved early based on having: stipules margin dentate, wing shape oblanceolate, style straight, seed shape ovate and is regarded as basal species for other *Trigonella* and *Melilotus* species.

At node 58, *Trg. cylindracea, Trg. stellata* and *Trg. maritima, Trg. arabica* clustered together based on the characters: staminal tube connected for nearly 2/3 for its length, gynecium linear oblong, pod surface *reticulate. Trg. cylinderacea* and *Trg. stellata* are closely related to each other, while *Trg. maritima* and *Trg. arabica* are forming another sister species. At node 54, *Trg. polycerata, Trg. foenum – graecum and Trg. glabra* clustered together on: number of ovules 3 - 8, style shorter than ovary, pod apex tapering. On the other hand, *Trg. foenum _ graecum and Trg. polyceratia* are sister species having stipules margine dentate and pod shap linear

For genus *Melilotus, M. segetalis* is considered as sister species to other three species (*M. messanensis, M. sulcatus,* and *M. serratifolius*) which has the characters: stem erect to decumbent, leaf apex retused emarinated mucronate and pod apex tapering. *Melilotus segetatis* is considered as basal species at node 35 having standard length shorter than keel and wing, seed shape oblong to circular. *Melilotus albus* comes early in this tree due to its biennial, mature leaflets are oblong obovate to lanceolate, the number of flowers per raceme and length of inflorescence in relation to subtending leaves play an important role in the identification of species (30-60 flowers), corolla is white, pods are obliquely ovate to elliptic with sharp acute beak.

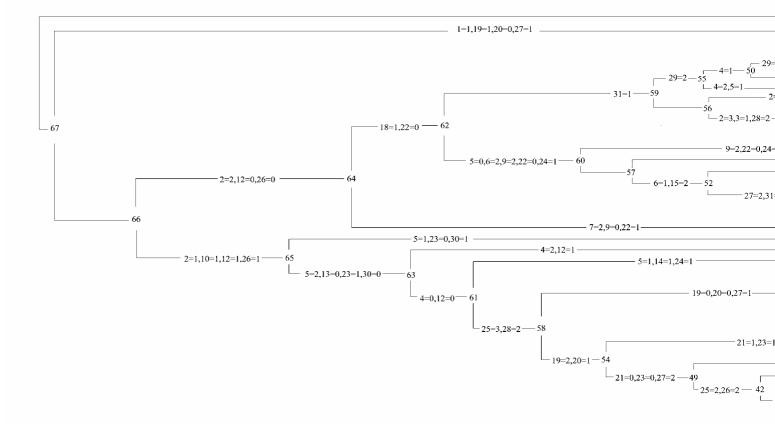


Figure. 1: Morphological cladogram for the Egyptian three genera (Subfamily: Papilionidae) showing the relationship between them (Length=241, Ci= Me. Medicago, Mel. Melilotus, Tri. Trifolium, Trg. Trigonella

Protein analysis (Table 2 and Figure 2)

1- Band with molecular weight 160 KD: in all species of *Melilotus*, two *Trigonella* (*stellata* and *cylindraceae*), and in two *Trifolium* (*stellatum* and *resupinatum*).

2- Band with molecular weight 130 KD: in all Trigonella and Trifolium.

3- Band with molecular weight 110 KD: in two *Trifolium* (*stellatum* and *angustifolium*), in all *Melilotus* except *messanensis*

4- Band with molecular weight 95 KD: in *Trifolium carnatum* and in three *Trigonella* (*maritime*, *stellata* and *cylindraceae*).

5- Band with molecular weight 90 KD: in all Trigonellae and two Trifolium (stellatum and incarnatum).

6- Band with molecular weight 80 KD: in all *Melilotus* except species *messanensis*, and in five *Trifolium* (*alexandrinum*, *tomentosum*, *repens*, *resupinatum* and *campestre*).

7- Band with molecular weight 75 KD: in three *Trifolium* (*stellatum*, *angustifolium* and *incarnatum*), also in two *Trigonella* (*maritima* and *stellata*), and in all *Melilotus* except *messanensis*.

8- Band with molecular weight 73 KD: in three *Trifolium* (*stellatum*, *angustifolium* and *incarnatum*), also in *Trigonella stellata*, and in three *Melilotus* (*indicus*, *segetalis* and *messanensis*).

9- Band with molecular weight 71 KD: only in three *Trifolium* (tomentosum, repens and campestre)

10- Band with molecular weight 69 KD: in two Trifolium (tomentosum and repens), and in Melilotus indicus.

11- Band with molecular weight 67 KD: in three Melilotus (segetalis, sulcatus and messanensis)

12- Band with molecular weigh 65 KD: only in Melilotus indicus.

13- Band with molecular weight 62 KD: in two *Trifolium* (*angustifolium* and *incarnatum*), also in two *Trigonella* (*arabica* and *cylindraceae*)

14- Band with molecular weight 60 KD: in all Trifolium and in two Trigonella (maritima and cylindraceae)

15- Band with molecular weight 56 KD: in all Trigonella

16- Band with molecular weight 52 KD: in three Trigonella (maritima, arabica and stellata)

17- Band with molecular weight 48 KD: only in Trigonella cylindraceae.

- 18- Band with molecular weight 46 KD: in all Trifolium and in two Melilotus (indicus and sulcatus)
- 19- Band with molecular weight 44 KD: only in Trigonella maritime.
- 20- Band with molecular weight 42 KD: in all Trigonella.
- 21- Band with molecular weight 40 KD; in all Trigonella.
- 22- Band with molecular weight 38 KD: in two Melilotus (sulcatus and messanensis).
- 23- Band with molecular weight 36 KD: in two Trifolium (angustifolium and incarnatum)
- 24- Band with molecular weight 35 KD: in Trifolium angustifolium and incarnatum.

25- Band with molecular weight 33 KD: in Melilotus segetalis and sulcatus.

- 26- Band with molecular weight 31 KD: in Melilotus segetalis and sulcatus.
- 27- Band with molecular weight 30 KD: in all Melilotus.
- 28- Band with molecular weight 29 KD: in all Melilotus.

29- Band with molecular weight 28 KD: in five *Trifolium (alexandrinum, tomentosum, repens, stellatum, and resupinatum)*.

- 30- Band with molecular weight 26 KD: only in Trifolium compestre.
- 31- Band with molecular weight 24 KD: in all Trifolium.
- 32- Band with molecular weight 22 KD: in all Trifolium except (repens and angustifolium)
- 33- Band with molecular weight 21 KD: in three Trifolium (alexandrinum, angustifolium and incarnatum)
- 34- Band with molecular weight 20 KD: in Trifolium alexandrinum and stellatum.
- 35- Band with molecular weight 19 KD: in Trifolium and Trigonella, also Melilotus indicus.
- 36- Band with molecular weight 18 KD: in Trifolium and Trigonella, also in Melilotus except indicus.

37- Band with molecular weight 17 KD: in four *Trifolium (alexandrinum, stellatum, angustifolium, and incarnatum).*

- 38- Band with molecular weight 16 KD: in all Trifolium species.
- 39- Band with molecular weight 15 KD: in all Trifolium species.
- 40- Band with molecular weight 14 KD: in all Melilotus species.
- 41- Band with molecular weight 13 KD: in all Melilotus species.
- 42- Band with molecular weight 12 KD: in all Trigonella species.
- 43- Band with molecular weight 10 KD: in all Trigonella species.

Species		Characters					
	1	1	2	3	4		
		0	0	0	0		
Medicago sativa	00000	000000000000000000000000000000000000000	000000000	0000000000000000	00000000		
Trifolium alexandrinum	01000	100000010	001000000	00001011111	.11110000		
Trifolium tomentosum	01000	1001100010	001000000	00001011001	.10110000		
Trifolium repens	01000	1001100010	001000000	00001010001	.10110000		
Trifolium stellatum	11101	.0110000010	001000000	00001011011	.11110000		
Trifolium resupinatum	11000	100000010	00100000	00001011001	10110000		
Trifolium campestre	01000	1001000010	001000000	00000110001	.10110000		
Trifolium argutum	01100	0110000110	001000011	00000011101	.11110000		
Trifolium incarnatum	01011	.0110000110	001000011	00000011101	.11110000		
Trigonella maritima	01011	.0100000011	100111000	00000000001	10000011		
Trigonella arabica	01001	.0000000101	100011100	00000000001	10000011		
Trigonella stellata	11011	.0110000011	100011100	00000000001	10000011		
Trigonella cylindraceae	11011	.0000000101	.010011000	00000000001	10000011		
Melilotus indicus	10100	1110101000	00100000	00110000001	.00001100		
Melilotus segetalis	10000	0100010000	000000000	1111000000	10001100		
Melilotus sulcatus	10100	111001000	001000000	1111000000	11001100		
Melilotus messanensis	10100	111001000	000000000	0011000000	11001100		

Table 2: Data matrix of the 43 protein pattern characters used to construct the phylogeny of the three genera (*Trifollium, Trigonella & Melilotus*) of family Papilionaceae of Egypt. (0= band absent, 1= band present).

Cladogram (Figure 3)

The electrophoretic pattern of the most common species of the three genera is shown in Fig.2. Protein marker with wide range (2-212 KD) which contains 13 bands as analyzed by SDS-PAGE in which 3 bands (97,36, and 20) have high intensity. A great homology of protein patterns within range 160-73 KDa can be recognized amongst all studied genera except *Melilotus* and *Trifolium* which showed the absence of one band 95 KD and two extra bands at the range of 110-80 KD, where *Melilotus* shows the absence of the two bands at 90 and 130 KD respectively. Another area of great similarity in protein profiles was observed in the range 19-18 KD, where are agreed similarity in protein profiles was observed, between *Melilotus* and *Trigonella* in 11 bands ranged from 71-15 KD. The similarity between *Trigonella* and *Trifolium* is also observed within the range 130-13 KD, where 7 bands were shared between *Melilotus* and *Trifolium* ranged from 56-38 KD

Electrophoretic protein profiles corresponding to different accessions of the different species in *Melilotus* showed 7 indentical patterns (67, 33, 31, 30, 29,14 and 12 KD). There are 5 bands that characterized *Trigonella* species at 56, 46, 42, 40, 38 KD, while in *Trifolium* the electrophoretic protein profiles showed 6 identical patterns (71,28,24,20, 16,15 KD).

On the basis of 43 protein storage characters, only one tree resulted from the cladogram of the 16 Egyptian species (Figure 3), which it distinguished into two main monophyletic groups based on nodes 30, 29.

The first clade includes genus *Melilotus*; *M. segetalis* is a sister group to the monophyletic group due to the absence of bands 3, 6 and 8. These bands with molecular weight 110 KD, 80 KD and 37 KD which are present in all other member of the genus. *Melilotus sulcutus* with character 36 has a band with 23 KD is a sister to species *M. indicus*.

The second largest clade includes two sub-clades: the first includes all *Trigonella* group, *Trigonella arabica* is considered as a basal species by the absence band with molecular weight 62 KD. *Trigonella stellata* is sister to *T. maritima* because they share bands of molecular weight 60 & 44 KD.

On the other hand, *T. cylinderica* separated having band with molecular weight 60 KD. The other sub-clade which nested all *Trifolium* species, *T. campestre* is considered as a basal group which has bands of molecular weight 50, 46 26, 24, 16 & 15 KD. *Trifolium argutum* is sister to *T. incarnatum* having bands with molecular wight 35 & 21KD.

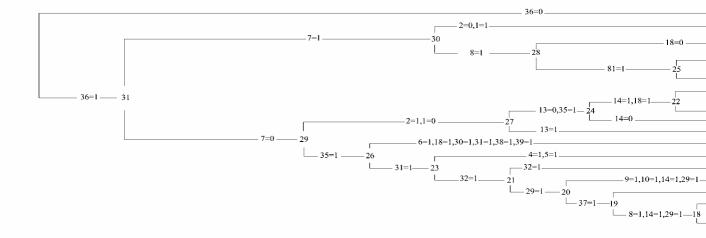


Figure 3 : Cladogram of the Egyptian genera (*Trif., Trg., Mel.*) of the tribe Trifolieae, Subfamily: Papilionideae, showing the relationship between the *Medicago, Mel. Melilotus, Trg. Trigonella, Tri. Trifolium.*

DISCUSSION

The present study aimed to analyze 31 morphological characters and 43 seed storage protein characters. The first analysis based on morphological characters suggests that there are two clades in one tree among all species of the three genera examined. The first clade reveals strong similarities between *Melilotus* and *Trigonella* (Sering 1825; Battacharyya 1958; Heyn 1966). These two sister genera based on sharing the characters: stipules margin incised, stander apex notched, style longer than ovary and surface of seed coat smooth. In this clade *Melilotus indicus* comes out early as a basal group to the other members at the node 65 which has the characters: stipule margin entire, stander apex obtuse, style longer than ovary and surface of seed coat wrinked. *Melilotus albus* evolved early depending on the characters: perennial plant, staminal tube connate for half of its length, gynoecium linear oblong and pod apex sharp acute. Our results agree completely with those of Shulz (1901) and Survorov (1950) who classified *Melilotus* into two subgenera; one includes *M. albus* and the other

subgenus included two sections; one including *M. indicus* and the other including all the other species investigated.

The second clade includes all species of the genus *Trifolium* sharing these characters: stem ascending, corolla pinks to mauve, stander distinct into limbe, claw and pod surface membranes. According to Steel *et al.* (1997) *Trifolium* is basal to the remainder of Trifolieae.

The protein analysis suggests one tree with 3 clades. The first clade includes genus *Melilotus* sharing bands with molecular weight 160 and 75 KD, *M. segetalis* evolved early based on the presence of the band with molecular weight 18 KD and the absence of bands with molecular weight 110, 80 and 73 KD. The second clade at the node 27 comprised all species of *Trigonella*, *T. stellata*, *maritima* and *cylindracea* which are evolved together depending on the presence of the bands 95, 75, 60 KD.

Trigonella arabica is basal group in this clade by having a band with molecular weight 62 KD. While *T. stellata* and *maritima* are sister species depending on having bands with molecular weights 95, 80, 71, 60, 24, 16 and 15 KD.

The third clade, at the node 26 is then separated on the basis of bands with molecular weight 80, 24, 16, 15 KD. *Trifolium campestre* has evolved from the hypothetical ancestor number 23 based on having the band with molecular weight 80, 46, 26, 24 KD. *Trifolium angustifolium* and *incarnatum* are sister species, due to sharing the intensity of bands with molecular weight 130 and 21 KD.

In conclusion, the morphological tree reflects that *Melilotus* is nested within *Trigonella* and that *Trifolium* is basal in this tree Steel *et al.* (1997) while *M. albus* has a special taxonomic position and collected only once from one locality El-Hadidi (1956). *Melilotus albus* is believed adventive in Egypt (El-Bous 1995).

The seed protein profiles are powerful discrimination tool for the identification of the three genera. In addition, seed protein profiles may provide evidence of the genetic relationships between *Trifolium*, *Trigonella*, and *Melilotus*.

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