

DELINEATION OF TWO MORPHO-TYPES OF SOLANUM MELONGENA VAR. INSANUM (L.) PRAIN. USING MORPHOLOGICAL, ANATOMICAL AND MOLECULAR TECHNIQUES

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Many of the Solanaceae members, especially the genus *Solanum*, are with taxonomical controversies. There were two morpho-types of *Solanum melongena* var. *insanum*, botanically considered as same. One of it is used in Ayurvedic preparations, locally known as 'Cheruvazhuthina', and the other locally known as 'Punyahachunda', was not. The morphological, macro-morphological, micro-morphological and anatomical features showed similarities and little differences. Micro-morphological characters are unchangeable within a species or variety. Differences in micro-morphological features showed that these two plants may belong to two different varieties. So for confirmation, the advanced molecular techniques are used. DNA barcoding using two genes, ITS and matK, was carried out as preliminary molecular assay. The results showed similarities for both the morpho-types. Success of these two genes, matK and ITS, in Solanaceae family is doubtful. However a detailed molecular profiling is warranted to confirm the recently obtained results to solve the taxonomic position of the two morpho-types of *S. melongena* var. *insanum*.

Key words: Solanum melongena var. insanum, DNA barcoding, ITS, matK

From the time of origin, man began to classify the plants based on the usefulness as poisonous and non-poisonous plants. This system of classification got modified time to time and the recent system of classification is APG system of classification. Taxonomic classification of plant species basically depends on the morphological and anatomical characters. these features are changeable and sometimes difficult to observe, so it is necessary to be supported by molecular techniques, in which molecular markers are used to detect the genetic variability. Molecular markers are biochemical constituents viz. secondary metabolites and macromolecules like proteins and deoxyribo nucleic acids (DNA) that play a very important role in plant taxonomy, physiology, embryology, plant breeding, ecology, genetic engineering, etc.(Joshi et al. 1999). So today taxonomical classification is not merely based on the morphological characters, it uses other biological techniques like spectroscopic studies and the emerging techniques based on molecular biology for classifying plants (Rosselló-Mora et al. 2001), and also for solving the taxonomic ambiguity

among complex groups like the genus Solanum.

The Solanaceae family consists of approximately 98 genera and some 2,700 species, with a great diversity of habitats, morphology and ecology (Kumari et al. 2017.) Plants belonging to this family contain different secondary metabolites coming under alkaloids, terpenoids, phenolics, etc. Tropane alkaloids, such as hyoscyamine and scopolamine, constitute one of the most distinctive group of secondary metabolites of the Solanaceae (Griffin and Lin 2000) and many plants containing them have long been utilised for their medicinal, hallucinogenic and poisonous properties. Withanolides represent a group of steroidal lactones with strong insecticidal properties which appear to be restricted to the Solanaceae (Harborne and Baxter 1993).

Solanum is a largest, cosmopolitan genus with around 1700 species. The genus *Solanum*, however, has certain difficulty in identifying natural species groups. Poor definition of the generic limits, the occurrence of suites of attributes in varying combinations throughout the genus, the phenotypic plasticity and genetic variation in many of the species, and the continual reclassification of parts of genus has ended in nomenclatural confusion (Kumar and Murugan, 2013).

The present investigation focuses on two morpho-types of Solanum to identify its uniqueness if any in Ayurveda. There are two morpho-types available for the genus Solanum and are locally known as 'Cheruvazhuthina' and 'Punyahachunda', considered as separate plants in Ayurveda. The root of 'Cheruvazhuthina' is included in 'Dasamool'. But 'Punyahachunda' is not used in Ayurvedic preparations. In certain places roots of 'Punyahachunda' is used as substitute for roots of 'Cheruvazhuthina'. But taxonomically these plant specimens are treated as ecotypes of a single plant variety - Solanum melongena var. insanum(L.) Prain. (Gamble 2016). Both of these plants show many differences in morphology, but considered as same plant variety of a species. Hence the aim of the present investigation is to assess its difference if any. In the present study the two plants are considered as two morpho-types of S. melongena var. insanum. For convenience 'Cheruvazhuthina' is denoted as morpho-type 1 and 'Punyahachunda' as morpho-type 2 throughout this work.

MATERIALS AND METHODS

The samples for the present investigation were collected from different localities in Kerala such as Nhangattiri, Thirumittacode and Koottanad of Palakkad district, Guruvayur of Thrissur district and Evoor in Alappuzha district of Kerala and voucher specimens were prepared and deposited in the Herbarium of Department of Botany, Sree Neelakanta Govt. Sanskrit College, Pattambi. (Fig. 1 a and b)

Morphological Studies

Macro morphological study:

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Macromorphological characters (trichome and seed coat ornamentation) were studied by close observation using handlens and stereomicroscope (Leica). The morphological characters were analysed using the work of Beentje (2010). Freshly collected leaf and mature dried seed samples were used for investigation. The trichome morphology was analysed according to Payne (1978).

The mature dried seeds of each sample was stained with safranin and observed under compound microscope in 10x magnification. The ornamentation pattern was analysed using the standard work on seed coat ornamentation by Barthlott (1981).

Anatomical Studies: For anatomical studies thin hand sections of stem, leaf and petiole of each sample were observed under compound light microscope after stained with safranin. Stem samples of the third internode of four plants with almost same maturity were selected for the study. For DNA barcoding DNA isolates from both the plant specimens were submitted for barcoding in collaboration with Rajiv Gandhi Centre for Biotechnology, Thiruvananthapuram.

Plant genomic DNA extraction was done as per the method described by Sambrook *et al.* (1989). DNA barcoding was carried out for two genes – *mat K* and ITS. These genes showed sequence polymorphism between the species and constant within a species. (Patwardhan *et al.* 2014)

RESULTS AND DISCUSSION

Morphological Studies

Micro morphological study (Table. 1): The micro morphological peculiarities of the two morpho-types were studied using the collected samples. The results obtained are summarised in the Table 2.

The two morpho-types were analysed for macro morphological as well as micro morphological characters. Macro-

Table 1. The macro morphological characters analysed for the two morpho-types (Bold words denote the characters in difference)

Characters	'Cheruvazhuthina' (Morpho-type 1)	'Punyahachunda' (Morpho-type 2)
Habit	Highly branching shrubs, ~1m height	Shrub, less branched, ~1 m height
Habitat	Terrestrial, slightly wet	Terrestrial, slightly wet
Stem	Highly branching; dense, long purple coloured spines; young parts dark purple coloured, less tomentose	Less branching; sparsely distributed, short, yellow coloured spines; young parts dark purple coloured, densely tomentose
Leaves	Alternate, simple, undulate, ~8 spines on adaxial surface, 3-6 on abaxial surface	Alternate, simple, sinuate, ~2 spines on adaxial surface, 0-3 spines on abaxial surface
Inflorescence	Racemose, 2-3 flowers from single point, in sub- axillary position	Solitary cyme, sub-axillary position
Flower	Bisexual, regular, hypogynous, complete	Bisexual, regular, hypogynous, complete
Calyx	5 sepals, gamosepalous, persistant, spiny	5 sepals, gamosepalous, persistant, not very spiny
Corolla	5 petals, gamopetalous, corolla lobes prominent	5 petals, gamopetalous, corolla lobes not prominent, umbrella shaped
Androecium	5 stamens, epipetalous, porous dehiscence, alternipetalous	5 stamens, epipetalous, porous dehiscence, alternipetalous
Gynoecium	Superior ovary, bicelled, axile placentation on swollen placenta, capitate stigma	Superior ovary, bicelled, axile placentation on swollen placenta, capitate stigma
	Leaf Characters	
Leaf length (Mean \pm SD)	13 ± 0.6 cm	7.5 ± 0.37 cm
Leaf width (Mean \pm SD)	$11 \pm 1.5 \text{ cm}$	6 ± 1.75 cm
Intermodal length (Mean \pm SD)	$8\pm0.5~\mathrm{cm}$	$4\pm0.5~\mathrm{cm}$
	Flower and Inflorescence Chan	racters
Petiole length (Mean \pm SD)	$4.5 \pm 0.6 \text{ cm}$	$2.25\pm0.58~cm$
Pedicel length (Mean \pm SD)	$2.25\pm0.49~\mathrm{cm}$	1.5 ± 0.5 cm
Diameter of flower (Mean ± SD)	$3.25\pm0.5\ cm$	$1.5\pm0.52~\mathrm{cm}$
Stamen length (Mean ± SD)	$0.85 \pm 0.3 \text{ cm}$	$0.45\pm0.12~\text{cm}$
Style length (Mean \pm SD)	1.3 ± 0.32 cm	$0.75\pm0.11~\mathrm{cm}$
Corolla colour	Dark violet	Dark violet with pale striations in the petal junctions
	Spine characters	
Spine length (Mean \pm SD)	0.6 ± 0.1 cm	$0.25 \pm 0.1 \text{ cm}$
Spine distribution	Upper – 6- 8 Lower – 3-6	Upper – 1-3 Lower – 0-3
Spine morphology	Dimorphic spines	Simple spines

Delineation of two morphotypes of Solanum melongena

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Α

В





Figure. 1(A-J): Macro morphological characters of two morpho-types of *Solanum melongena*var. *insanum.* 1A. Inflorescence of morpho-type 1, **B**. Inflorescence of morpho-type 2, **C**. Fruit of morpho-type 1, **D**. Fruit of morpho-type 2, **E**. Leaf of morpho-type 2 and 1, **F**. Dimorphic spines of morpho-type 1, **G**. Simple spines of morpho-type 2 **H**. Leaf with numerous spines in morpho-type 1, **I**. Leaf with less spines in morpho-type 2, **J**. Ripe fruit of morpho-type 1 and 2 respectively.

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Table 2: T	he micro mo	orphol	ogical	pecul	liarities o	fthe	two morr	bho-tr	vnes
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characters	'Cheruvazhuthina' (Morphotype 1)	'Punyahachunda' (Morphotype 2)
	Trichome characters (Fig.	3a and b)
Morphology	Stellate trichome	Stellate trichome
Number of arms	6 - 7 arms	8 arms
Number of trichomes in adaxial surface	2-3	Numerous
	Seed characters (Fig3. c	and d)
Size	2 – 3 mm	2 – 3 mm
Seed coat ornamentation	Thick muri with wide lumen	Thick and compact muri



Figure 2 (A-D). Micro morphological characters of two morpho-types of *Solanum melongena* var. insanum **A**. Trichomes of morpho-type 1 **B**. Trichomes of morpho-type 2 **C**. Seed coat ornamentation of morpho-type 1 **D**. Seed coat ornamentation of morpho-type 2



Figure 3 (a-c): a. The epidermal and cortical regions morpho-typ e 1 and 2 respectively, **b.** Stem T.S. of morpho-type 1 and 2 respectively, after secondary thickening, **c.** The secondary xylem of morpho-type 1 and 2 respectively.



Figure. 4: Isolated DNA from two morpho-types after gel electrophoresis – A: morpho-type 1 and B: morpho-type 2.

SR593-A-MATK	1	CTTTTTTAGAGGACAATTTGTCACATCTAAATTATGTATTAGATATACTA	50
SR593-B-MATK	1	CTTTTTTAGAGGACAATTTGTCACATCTAAATTATGTATTAGATATACTA	50
SR593-A-MATK	51	ATACCCTACCCGTTCATCTGGAAATCTTGGTTCAAACTCTTCGTTA	100
SR593-B-MATK	51	ATACCCTACCCGTTCATCTGGAAATCTTGGTTCAAACTCTTCGTTATTG	100
SR593-A-MATK	101	GGTAAAAGATGCCTCTTCTTTACATTTATTACGATTCTTTCT	150
SR593-B-MATK	101	GGTAAAAGATGCCTCTTCTTTACATTTATTACGATTCTTTCT	150
SR593-A-MATK	151	ATTGTAATTTGAATAGTCTTATTACTTCAAAGAAGCCCGGTTACTCTTTT	200
SR593-B-MATK	151	ATTGTAATTTGAATAGTCTTATTACTTCAAAGAAGCCCGGTTACTCTTTT	200
SR593-A-MATK	201	ТСААААААААТССААААТТСТТСТТСТТСТТАТАТААТТСТТАТGТАТА	250
SR593-B-MATK	201	ТСААААААААТССААААТТСТТСТТСТТСТТАТАТААТТСТТАТGTATA	250
SR593-A-MATK	251	TGAATGCGAATCCACTTTCGTCTTTCTACGGAAACAATCTTTTCATTTAC	300
SR593-B-MATK	251	TGAATGCGAATCCACTTTCGTCTTTCTACGGAAACAATCTTTTCATTTAC	300
SR593-A-MATK	301	GATCAACATCTTTTGGAGCCCTTCTTGAACGAATATATTTCTATGGAAAA	350
SR593-B-MATK	301	GATCAACATCTTTTGGAGCCCTTCTTGAACGAATATATTTCTATGGAAAA	350
SR593-A-MATK	351	ATAGAACGTCTTGTAAAAGTCTTTGCTAAGGATTTTCAGGTTACCCTATG	400
SR593-B-MATK	351	ATAGAACGTCTTGTAAAAGTCTTTGCTAAGGATTTTCAGGTTACCCTATG	400
SR593-A-MATK	401	GTTATTCAAGGATCCTTTGATGCATTATGTTAGGTATGAAGGAAAATCAA	450
SR593-B-MATK	401	GTTATTCAAGGATCCTTTGATGCATTATGTTAGGTATGAAGGAAAATCAA	450
SR593-A-MATK	451	TTCTGGCTTCAAAAGGGACGTTTCTTTTGATGAATAAATGGAAATTTTAC	500
SR593-B-MATK	451	TTCTGGCTTCAAAAGGGACGTTTCTTTTGATGAATAAATGGAAATTTTAC	500
SR593-A-MATK	501	CTTGTCAATTTTTGGCAATGTCATTTTTCTATGTACTTTCACACAGG	550
SR593-B-MATK	501	CTTGTCAATTTTTGGCAATGTCATTTTTCTATGTACTTTCACACAGGAAG	550
SR593-A-MATK	551	GATCCATATAAACCAATTATCCAACCATTCCCGTGACTTTATGGGCTATC	600
SR593-B-MATK	551	GATCCATATAAACCAATTATCCAACCATTCCCGTGACTTTATGGGCTATC	600
SR593-A-MATK	601	GTTCAAGTGTGCGACTAAATCATTCAATGGTACGTAGTCAAATGTTCGAA	650
SR593-B-MATK	601	GTTCAAGTGTGCGACTAAATCATTCAATGGTACGTAGTCAAATGTTCGAA	650
SR593-A-MATK	651	AATTCATTTCTAATCAATAATCCAATTAAGAAATTCGATACCCTTGTTCC	700
SR593-B-MATK	651	AATTCATTTCTAATCAATAATCCAATTAAGAAATTCGATACCCTTGTTCC	700
SR593-A-MATK	701	AATTATTCCTTTGATTGGATCATTAGCTAAAGCACACTTTTGTACCGTAT	750
SR593-B-MATK	701	AATTATTCCTTTGATTGGATCATTAGCTAAAGCACACTTTTGTACCGTAT	750
SR593-A-MATK	751	TAGGGCATCCCATTAGTAAACCGGTTTGGTCCGATTTATCAGATTCTGAT	800
SR593-B-MATK	751	TAGGGCATCCCATTAGTAAACCGGTTTGGTCCGATTTATCAGATTCTGAT	800
SR593-A-MATK	801	ATTATTGACCGATTTGGGCGTATATGCAGAAATCTTTTTCATTATTATAG	850
SR593-B-MATK	801	LATTATTGACCGATTTGGGCGTATATGCAGAAATCTTTTTCATTATTAT	
SR593-A-MATK	851	CGGATCTTCCAAAAAAAGAC 871	
SR593-B-MATK	851	CGGATCTTCCAAAAAAAAGAC 871	

Figure 5: The pair-wise sequence alignment of *matK* using BLAST

8				
A	ITS	1	TTTCCGTAGGTGAACCTGCGGAAGGATCATTGTCGAAACCTGCACAGCAGAACGACCCGC	60
В	ITS	1	TTTCCGTAGGTGAACCTGCGGAAGGATCATTGTCGAAACCTGCACAGCAGAACGACCCGC	60
A	ITS	61	GAACACGTTCAAACACCGGGGGAGCCGCGCGCGGGGGGGG	120
В	ITS	61	GAACACGTTCAAACACCGGGGGAGCCGCGCGCGCGGGGGGCGCCGCCCCGCGC	120
A	ITS	121	GTCTGCCCCTCGCCCCCTCTTCGGGGGGGCCAAACGAACCCCGGCGGGAAAGCGCCAAG	180
В	ITS	121	GTCTGCCCCTCGCCCCCTCTTCGGGGGGGCCAAACGAACCCCGGCGCGGAAAGCGCCAAG	180
A	ITS	181	GAATACTCAAACGAGAGCCCTCCGCCGTGCCCGTCCGCGGGGGGGG	240
В	ITS	181	GAATACTCAAACGAGAGCCCTCCGCCCGTGCCCGTCCGCGGGGCGTGCGGGCGG	240
A	ITS	241	TGCTTCTTTCGAAACCAAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATG	300
В	ITS	241	TGCTTCTTTCGAAACCAAAAACGACTCTCGGCAACGGATATCTCGGCTCTCGCATCGATG	300
A	ITS	301	AAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTC	360
В	ITS	301	AAGAACGTAGCGAAATGCGATACTTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTC	360
A	ITS	361	TTTGAACGCAAGTTGCGCCCGAAGCCGTCAGGCCGAGGGCACGTCTGCCTGGGCGTCACG	420
В	ITS	361	TTTGAACGCAAGTTGCGCCCGAAGCCGTCAGGCCGAGGGCACGTCTGCCTGGGCGTCACG	420
A	ITS	421	CATCGCGTCGCCCCCGCACGCCGCGCGCGCGCGGGGGGGG	480
В	ITS	421	CATCGCGTCGCCCCCCGCACGCCGCTCGGCGTCGCGGGGGGGG	480
A	ITS	481	GCCTCGCGCCGGCCGGCCTAAATGCGAGTCCACGTCGACGGACG	540
В	ITS	481	GCCTCGCGCCCGCGGCCGGCCTAAATGCGAGTCCACGTCGACGGACG	540
A	ITS	541	TGGTTGTAGCTCAACTCTCTTGGTGCCGCGGCCACAGCCCGTCGCGCGTGCGCGCTCCAC	600
В	ITS	541	TGGTTGTAGCTCAACTCTCTTGGTGCCGCGGCCACAGCCCGTCGCGCGCG	600
A	ITS	601	GGCCCCTCCGGCGCTAGCGCGCTCCGACCGCGACCCCAGGTCAGGCGGGATTACCC 656	
В	ITS	601	GGCCCCTCCGGCGCTAGCGCGCTCCGACCGCGACCCCAGGTCAGGCGGGATTACCC 656	

Figure 6: The pair-wise sequence alignment of ITS using BLAST

morphological characters include qualitative and quantitative characters. The qualitative macro morphological characters including habit, habitat, characters of stem, leaf, inflorescence and flower were investigated. About 10 quantitative characters viz., leaf length and width, inter-nodal length, petiole length, pedicel length, diameter of flower, stamen, style, length and number of spines were examined. The qualitative as well as quantitative characters investigated the two

morpho-types showed variations in leaf characters, flower morphology, spine characters, seed and fruit morphology (Fig. 1). In leaf morphology the leaf shape of the morpho-type 1 was undulate and that of morpho-type 2 was sinuate (Fig. 1 E). The length and width of the leaf of morpho-type 1 $(13 \pm 0.6 \text{ cm})$ was higher than that of morphotype 2 ($7.5 \pm 0.37 \text{ cm}$) (Table 1). Morpho-type 1 is spinier than morpho-type 2 and morpho-type 1 has dimorphic spines, but morpho-type 2 possess only one type of spine (Fig. 5 f and g).

Colour of spine was also showing considerable difference ie. morpho-type 1 has dark purple coloured spines whereas morpho-type 2 has small yellow coloured spines (Fig. 1 F and G). The seed coat ornamentation also showed some differences (Fig. 1 C and D). Morpho-type 1 has thick muri with wide lumen seed coat ornamentation, whereas the morpho-type 2 has thick, compact muri seed coat ornamentation. Morpho-type 1 has racemose inflorescence, whereas morpho-type 2 has solitary cyme inflorescence. Trichome distributions on leaf also showed variations. Morpho-type 2 has densely distributed stellate trichomes on the upper surface of the leaf and that of morphotype 1 has scarcely distributed stellate trichomes on the upper leaf surface (Fig. 2A and B). The collected samples showed distinguishable differences (Table 1 and 2). Using these characters the collected samples can be distinguished as morpho-type 1 and morpho-type 2. So it might be pointed out that the characters showing differences were conserved in each morpho-type. The significant variations in the micro morphological characters such as trichome characters (Fig. 2 A and B) and seed coat ornamentation (Fig. 2 C and D) were conserved in each morpho-types. Krishnakumar et al. (2018) reported that majority of the accessions of Abelmoschus esculentus could be easily distinguishable using trichome characters even in the early vegetative stage of the plants. This variability in the micro morphological characters were supported by RAPD studies of Abelmoschus esculentus (L.) Moench. (Christeena, 2014). Zhigila et al. (2015) also used trichome characters to distinguish five varieties of Capsicum annuum L. Fareedkhan et al. (2012) conducted a study on the morphological characterisation of the wild varieties of Solanum tuberosum L. and the variations found were confirmed by RAPD.

Anatomical Studies: The cross section of the leaf and petiole of two morpho-types had no significant differences. However cross section of stem showed some differences among the

morpho-types studied. The anatomy of the two morpho-types were almost same; a single epidermal layer followed by 1 - 2 layers of chlorenchyma, then by 4 layers of collenchyma in morpho-type 2 and 8 layers of collenchyma in morpho-type 1. Then 4 layers of paranchyma followed by the stelar region (Fig. 3 a). The stelar region contain single layered endodermis and stone cells above the secondary phloem in discontinuous pattern in morpho-type 2. The stone cells were absent or poorly developed in morpho-type 1. Morpho-type 1 contain more xylem vessels than morpho-type 2. Morphotype 2 started its secondary thickening earlier than morpho-type 1. Both morpho-types contain intra-xylary phloem in pith region and cell inclusions like crystals in pith cells (Fig. 3 b and c).

Anatomical studies of the stem, leaf and petiole showed no significant differences. The secondary growth in morpho-type 2 is started earlier than that of morpho-type 1 (Fig. 3 b). One notable difference observed was the number of xylem vessels in stem. Morph-type 1 possessed more xylem vessels than that of morpho-type 2. It may be due to the increased girth of the stem of morpho-type 1. Further detailed anatomical studies are warranted to confirm the anatomical difference. Earlier stuies revealed that anatomical differences were observed in different varieties of *Artemisia absinthium* (Konowalik and Kreitschitz, 2012).

DNA Barcoding of the two Morpho-types:

The isolated DNA from the two morphotypes were subjected for polyacrylamide gel electrophoresis. The results showed that the isolated DNA from two morpho-types were of similar equal size ie.1 kb (Fig.4).

Two genes- the chloroplast gene *mat* K and the nuclear gene ITS – were sequenced. Both sequence were submitted to NCBI GenBank using BankIt submission tool. ITS sequence got accession number MG572181 and *mat* K sequence got accession number MG543677.

The sequence of *mat K* and ITS of two morphotypes were then pair-wise aligned using NCBI BLAST (Fig. 5 and 6). The pair-wise sequence alignment results obtained from BLAST showed that the sequences of both *mat K* and ITS genes were 100% similar in both morphotypes.

In the present investigation the morphological as well as anatomical analysis showed many differences between the two morpho-types. In another study these two morpho-types showed a characteristic difference in phytochemical constituents present in root and fruit (Thulasi and Krishnakumar, 2018). In such cases of taxonomical controversies molecular level analysis, like DNA barcoding is relied for further clarification. Hence in the present study DNA barcoding of the two morpho-types were performed. The sequence of the two genes mat K and ITS were compared. These two genes are frequently used for DNA barcoding studies. Mat K, the chloroplast gene, coded maturase kinase and ITS, the nuclear gene, is the internal transcribed spacer region which present between the small rRNA subunit gene and large rRNA subunit gene. These genes were selected because these two genes showed sequence polymorphism between species and constant within a species (Patwardhan et al., 2014). Pair wise sequence alignment using the NCBI tool BLAST was carried out. But this analysis showed 100% similarity. Based on the DNA barcoding results, the two morpho-types were with same taxonomic status. The difference seen in the morphology, anatomy and phytochemical constituents might be due to the environmental differences. However there were reports that the use of DNA barcoding for differentiating two verities is a failure in the case of complex groups like wild potatoes (Spooner, 2009) where whole chloroplast genome sequencing was used. So for further confirmation the complete chloroplast genome sequencing is to be performed. We could suggest that one of the morpho-type (morphotype1) might be the wild alley of the other morpho-type.

In conclusion, morphologically the two morpho-types were different in many macro

morphological characters like spine morphology, leaf shape, inflorescence type and fruit morphology, as well as micro morphological characters like seed coat ornamentation and trichome characters. Micro morphological characters are less changed within a species or variety as the influence of ecological factors. So micro morphologically the two morpho-types are different varieties of Solanum melongena. Anatomically two morpho-types does not showed much variations. Except in the number of xvlem vessels and early starting of secondary thickening in morpho-type 2 than in morphotype 1. Finally the DNA barcoding studies showed that the two morpho-types are same. The success of the two genes – mat K and ITS – used in barcoding is doubtful. So a further confirmation is needed, by whole chloroplast genome sequencing, to confirm the taxonomical position of the two morpho-types.

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