



Effect of genetic diversity on adaptation of endangered Sinai Thyme in Saint Katherine Protectorate, Egypt

Asmaa A. Zaghloul^{1*}, Om Mohamed A. Khafagi¹, Mohamed M. Moursy², Hany A. Marghany²

¹Department of Botany, Faculty of Science (Girls branch), Al-Azhar University, Cairo, Egypt

²Department of Botany, Faculty of Science (Boys branch), Al-Azhar University, Cairo, Egypt

ARTICLE HISTORY

Received: February 18, 2023

Accepted: July 1, 2023

INDEXING: Arab International Journal of Environment Science – (AIJES), published by General Association for Studies, Research and Technology is covered by Egyptian Knowledge Bank

<https://aijes.journals.ekb.eg/>

*CORRESPONDING AUTHOR

Asmaa A. Zaghloul Department of Botany, Faculty of Science (Girls branch), Al-Azhar University, Cairo, Egypt. Email:

asmaazaghloul2828.2@gmail.com

CITE THIS ARTICLE

Zaghloul, A.A., Khafagi, A.O., Moursy, M.M., Marghany, H.A. (2023). Effect of genetic diversity on adaptation of endangered Sinai Thyme in Saint Katherine Protectorate, Egypt. Arab International Journal of Environmental Sciences, 2(1): 4. <http://doi.org/10.21608/aijes.2023.1478>

ABSTRACT Saint Katherine Protectorate (SKP) is located in South Sinai, Egypt. It harbors many endemic and endangered medicinal plants that are overexploited and facing a serious threat of extinction. From these important medicinal plant species, we select Sinai Thyme, *Thymus decussatus* (a near-endemic and endangered plant) to investigate the genetic diversity level among some populations using both random amplified polymorphic DNA (RAPD) and inter-simple sequence repeat (ISSR) molecular markers. RAPD and ISSR represent a level of polymorphism of 86.70%. Assessing genetic polymorphisms among the *Thymus* populations at varying microhabitats can help in their genetic improvement and conservation programs.

Key words: Sinai Thyme, genetic diversity, Saint Katherine Protectorate.

INTRODUCTION

The Sinai Peninsula forms a land bridge between Africa and Asia and its flora and fauna have been influenced by both continental masses; and contains the Saint Katherine Protectorate which covers the mountainous region of Southern Sinai. St. Katherine Protectorate is characterized by the highest mountains in Egypt, a dense wadi system and an arid climate, it was declared as a protected area in 1996 due to its biological and cultural interest (SKP-MP, 2003). St. Katherine Protectorate is located between 33° 30' to 34° 30' E, and 28° 50' to 29° 50' N (Zaghloul *et al.*, 2022)

Saint Katherine Protectorate (SKP) located at Southern Sinai is considered one of the largest protected areas in Egypt which included the highest mountains rich with about 44 % of endemic plant species in Egypt. In Southern Sinai, Mountains flora varies from the other areas because of the higher variability in ecosystems, climatic conditions, water environment, and elevations (Omar, 2014; Soliman *et al.*, 2022). Saint Katherine Protectorate harbors many rare endangered and native plants that are overexploited and facing severe threat of extinction. *Thymus* genus, belonging to family Lamiaceae, integrates 928 species distributed all over the world

and among them about 215 species were mainly distributed in Mediterranean area (Minarchenko *et al.*, 2019). Owing to its aromatic nature, thyme plant has good antimicrobial, antiparasitic, antispasmodic and antioxidant properties (Iftikhar *et al.*,

2023). Therefore, their medicinal properties made it one of the most essential and common medicinal herbs (Iftikhar *et al.*, 2023). Three species known as *T. capitatus*, *T. bovei*, and *T. decussatus* have been reported for *Thymus* genus in Egypt.



Figure 1: Photo of endangered Sinai Thyme (*Thymus decussatus* Benth.) in Saint Katherine Protectorate, Egypt

Effect of genetic variation on plant adaptation

Knowledge of genetic variation among *T. decussatus* populations is critical for their survival and also for better use of their genetic resources. Sinai thyme is distributed in scattered and patches microhabitats (Khedr, 2021), where it was restricted to mountains heads at altitudes above 1500 m (Soliman *et al.*, 2022) and it was growing well in the bases of cliffs and wades. Thyme will be regularly a significant species for corrupted ecosystems particularly arid environments for longer grazing pressure (Thompson and Gilbert, 2013).

Genetic polymorphism refers to the degree of genetic variation that can occurs at different scales including populations, subpopulation or even among neighboring individuals and it plays an important role in adaptation of a plant species to environmental stimuli (Caliebe *et al.*, 2022). Ecogeographical patterns affect the distribution of genetic diversity in a species. Association of genetic diversity with variation in ecogeographical patterns has been detected in several species of aromatic plants (Youssef and Mahgoub, 2015).

Determination genetic variation in plant species

DNA markers are used in determining genetic variability in many plant species. Random amplified polymorphic DNA (RAPD), inter simple sequence repeats (ISSRs), simple sequence repeats (SSRs), and amplified fragment length polymorphism (AFLP) are the most widely used DNA markers (Seehalak et al., 2006). Guanine-Cytosine content is the percentage of nitrogenous bases in a DNA content has been reported to be a factor deciding a primer's efficiency, since GC content is correlated with annealing temperature and is related to the generation of more DNA fragments (Cervantes-Salguero, 2022).

Genetic variability of aromatic and medicinal plant populations has been analyzed using either RAPD or ISSR or both (Akçali Giachino, 2020). The genetic diversity among the different accessions or populations of thyme plant using either RAPD or ISSR or both molecular markers was evaluated in different studies. Genetic variation within and among eight populations of the endemic Tunisian *Thymus algeriensis* plant species collected from variant geographical habitats was assessed using seven RAPD primers (Caliebe et al., 2022). Their results recorded presence of high genetic variation within and among the studied populations.

RAPD markers were also used in genetic identification of five accessions of Palestinian *Thymus* for their ability to produce polymorphisms (Solyman and

Alkowni, 2014). Moreover, genetic diversity of ten *Thymus kotschyanus* populations obtained from various ecosystems in Iran was analyzed using RAPD markers (Khoshshokhan et al., 2014). Their findings showed high level of genetic diversity among the ten studied populations of *T. kotschyanus*. Even, twenty ISSR primers were applied for evaluation of the genetic relationships among fourteen *Thymus* accessions belonging to three *T. daenensis*, *T. kotschyanus* and *T. vulgaris* species (Yousefi et al., 2015). Their findings showed high level of genetic diversity among populations.

The establishment of efficient conservation strategies for growing plants (Salgotra and Chauhan, 2023). Thus, great efforts should be carried out on preserving the rare endangered species as well as the endemic species with restricted geographical distribution (Zhu et al., 2023). From these near-endemic and endangered plant species, *T. decussates* plant species growing in Saint Katherine Protectorate was selected in this study to assess the degree of genetic variation among their populations using both of RAPD and ISSR molecular markers. RAPD and ISSR markers have been successfully applied in assessing the extent of genetic diversity among eight *T. decussatus* populations collected in Saint Katherine Protectorate, Southern Sinai, Egypt from eight microhabitats (Mahgoub et al., 2020). Mahgoub et al. (2020) suggested all primers with variable percentages of polymorphism with an average of 86.21 % were found to be polymorphic. Mahgoub

et al. (2020) were entirely compatible with results on five Palestinian *Thymus* species, *T. kotschyanus*, *T. caespititius*. In addition, numerous research studies on *Thymus* species from several countries showed that the rate of polymorphism ranges from 62% to 92% (Saboh *et al.*, 2019). Nevertheless, the polymorphism level obtained in Mahgoub *et al.* (2020) is rather lower than the recorded average polymorphism (94.31%), which examined genetic polymorphism in 13 *Thymus* accessions collected from different habitats in Iran using 20 RAPD primers (Yousefi *et al.*, 2015).

Mahgoub *et al.* (2020) showed that the number of DNA fragments produced per primer ranged from seven to ten which are close to the result obtained in their analysis on populations of *Thymus vulgaris*. High level of polymorphism recorded in our study reflects the great degree of genetic variation among *Thymus* populations collected from various microhabitats in Saint Katherine Protectorate (SKP). Such heterogeneity may be due to the high degree of gene exchange between the studied populations (Khoshokhan *et al.*, 2014).

Numerous studies have documented that the RAPD method is an effective tool for thyme taxa genome. The analysis of a large number of DNA fragments can be considered as a good representation of the *Thymus* taxa genome and the ability to distinguish all the samples studied by RAPD banding profiles indicates that this technique may provide a faster and cheaper way for the identification of the *Thymus*

taxa (Mahgoub *et al.* 2020). Mahgoub *et al.* (2020) showed high polymorphic among the Thyme populations surveyed (87.18 %) by ISSR markers. The polymorphism has an average of 88.9% in *Thymus daenensis* (Rahimmalek *et al.*, 2009).

In another study also it was recorded that the number of polymorphic bands ranged from 75% to 100%, with an average of 96.76% between fourteen *Thymus* accessions belonging to three species (*T. daenensis*, *T. kotschyanus*, and *T. vulgaris*) (Sarfaraz *et al.*, 2020)

ISSR markers can be a highly insightful, efficient, and reliable method for genetic diversity trials (Iqbal *et al.*, 2023). The efficacy of the molecular marker techniques depends primarily on the degree of polymorphism produced by the primers used. To characterized the degree of genetic variation among species of *T. decussatus* populations (Mahgoub *et al.*, 2020).

RAPD and ISSR markers were appeared genetic variation among the populations of *Thymus decussates* with an average of 86.70% (Mahgoub *et al.*, 2020). RAPD and ISSR markers were used in similar studies to assess the genetic variation among *Nepeta septemcrenat*, *Thymus decussates* and *Phlomis aurea* plant species collected from Saint Katherine Protectorate, South Sinai, Egypt (EL-Sadek *et al.*, 2017). The primers of ISSR produced more polymorphic bands than primers of RAPD, hence ISSRs were found to be more efficient than RAPD in

estimating genetic diversity (**EL-Sadek et al., 2017**).. Similar results also have been obtained for other plants including *Salvia* (**Javan et al., 2012**)

(**Mahgoub et al., 2020**) support the fact that there are SKP region-specific variations in relation to soil and climatic adaptation cycle. Also **Catană et al. (2013)** support this fact in their study on endemic plant species (*Draba dorneri* Heuff and *Commiphora wightii*), in their study on three medicinal plant species belongs to the Lamiaceae family (*Nepeta septemcrenata*, *Ballota undulate* and *Teucrium polium*) which grows naturally in Saint Catherine Mountain at three different altitudes. All of these result indicated that there are relationship between geographical distributions and the genetic diversity.

There are high genetic variation occurs between *T. decussatus* populations as recorded in **Mahgoub et al. (2020)** contrasts with mountain plants. For endemic plants from Tibet and Central Asian desert plants and several other studies of outcrossing endemic species, it was found that there was high genetic variability in colonies than in habitats (**Chen et al., 2009**). Differences in polymorphism among restricted species may be linked to the degree to which their populations, in accordance with occur in heterogeneous habitats (**Wang and Zhang, 2022**). Despite the short geographical distances which separate the current *T. decussatus* populations, high genetic variation was observed between the

populations studied using both RAPD and ISSR markers (**Mahgoub et al., 2020**).

Also, the extent and distribution of genetic diversity is significantly influenced by the mating mechanism and reproduction mode. Higher homozygosity results in low levels of genetic variation in self-compatible species, while species with primarily outcrossing mating systems experience higher rates of genetic variation (**Pluhár et al., 2012**).

CONCLUSION

Both RAPD and ISSR analyses showed the same percentage polymorphism, as 86.21 percent for RAPD markers and 87.18 percent for ISSR markers It can be concluded that amongst the *Thymus* populations rising in the Saint Katherine region there is a high genetic diversity. Specific geographical and ecological conditions permit certain potential genetic modifications or DNA changes such as translocation, deletion, point mutation and so on. The molecular analysis using markers RAPD and ISSR showed that both markers were able to differentiate between the different populations of species as *T. decussatus* and can thus be used to research the degree of genetic variation between species as *T. decussatus*.

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