## STUDIES ON CYTOLOGICAL CHARACTERIZATION OF CISSUS QUADRANGULARIS ECOTYPES FOR ASSESSMENT OF PLOIDY LEVEL

## S. Padmapriya\*, K. Vinoth and K. Rajamani

Department of Medicinal and Aromatic Crops, TNAU, Coimbatore Email: spadmapriyaa@yahoo.co.in

## Received-06.11.2021, Revised-20.11.2021, Accepted-27.11.2021

**Abstract:** *Cissus quadrangularis*, commonly known as veldt grape, is one of the medicinally important perennial, climbing succulent, widely distributed in Africa, the Arabian Peninsula, Northern India, and Southeast Asia. The present investigation on cytological characterization of veldt grape was conducted at Department of Medicinal and Aromatics crops, Horticultural College and Research Institute, Tamil Nadu Agricultural University, Coimbatore from 2019 -2020. Fifty veldt grape ecotypes were collected from different geographical locations of Tamil Nadu and five morphologically superior ecotypes identified were subjected to ploidy level estimation bt using flow cytometry method. The mean FL1-H value varied from 1018.05 to 2896.2 for the five superior ecotypes selected. The highest mean value FL1-H were found in TNCq23 and the lowest mean value recorded in TNCq34 from the plot 4.CV value of the veldt ecotypes ranged from 217.38 (TNCq 23) to 334.00 (TNCq9). The histogram of the mean position of G1 phase of the selected veldt grape ecotypes using radish as the reference indicated that the mean position of G1 peak for all the five ecotypes (TNCq32, TNCq34, TNCq29, TNCq23 and TNCq9) exhibited diploid number (2n) of chromosomes.

Keywords: Cissusquadrangularis, Ecotypes, Ploidy level, Flow cytometry

## REFERENCES

**Barow, M. and G. Jovtchev.** (2007). "Endopolyploidy in plants and its analysis by flow cytometry. Flow cytometry with plant cells: analysis of genes, chromosomes and genomes:349-372.

```
[ Google Scholar ]
```

**De Rocher, E.J., Harkins, K.R., Galbraith, D.W. and Bohnert, H.J.** (1990). Developmentally regulated systemic endopolyploid in succulents with small genomes. Science 250 (4977):99-101.

[Google Scholar] Dolezel, J. (1997). Application of flow cytometry for the study of plant genomes. Journal of applied Genetics 3 (38):285-302.

[ Google Scholar ]

Gichuki, D.K., L. Ma, Z. Zhu, C. Du, Q. Li, G. Hu, Z. Zhong, H. Li, Q. Wang, and H. Xin. (2019). Genome size, chromosome number determination, and analysis of the repetitive elements in *Cissusquadrangularis*. Peer Journal 7:8201.

[ Google Scholar ]

**Karkamkar, S.P., Patil, S. and Misra, S.C.** (2010). Cyto–morphological studies and their significance in evolution of family Vitaceae. The Nucleus 53 (1):37-43.

[ Google Scholar ]

Robert, G.W., Qing-feng, W., Yong, W. and Youhao, G. (2001). A taxonomic investigation of variation within *Cissusquadrangularis* L.(Vitaceae) in Kenya. Wuhan University Journal of Natural Sciences 6 (3):715-724.

[ Google Scholar ]

**Soltis, D.E., Soltis, P.S., Bennett, M.D. and Leitch, LJ.** (2003). Evolution of genome size in the angiosperms. American Journal of Botany 90 (11):1596-1603.

[ Google Scholar ]

Wen, J., L.M. Lu, Z.L. Nie, X.Q. Liu, N. Zhang, S. Ickert-Bond, J. Gerrath, S.R. Manchester, J. Boggan, and Z.D. Chen. (2018). A new phylogenetic tribal classification of the grape family (Vitaceae). Journal of Systematics and Evolution 56 (4):262-272.

[ Google Scholar ]