Geographical patterns of *in vivo* spontaneously emitted volatile organic compounds in *Salvia* spp.



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INTRODUCTION

STATE OF THE ART

Salvia, with its over 900 species, is the largest genus in the Labiatae family, with a remarkable range of **variation**. The specie has undergone a **geographical radiation** in many areas of the world and, since Bentham's classification (1848), no other study has been made on the new (500+) species. Walker et al. investigated the monophily of the genus postulated by Bentham analysing two chloroplast DNA regions (rbcL and trnL-F). This study showed the existence of **at least 3 distinct lineages**, in which much of the diversification fits along **biogeographical lines**:

Salvia Clade I	Salvia Clade II	Salvia Clade III
Largely Europe but with one	America	Independent Asian lineage







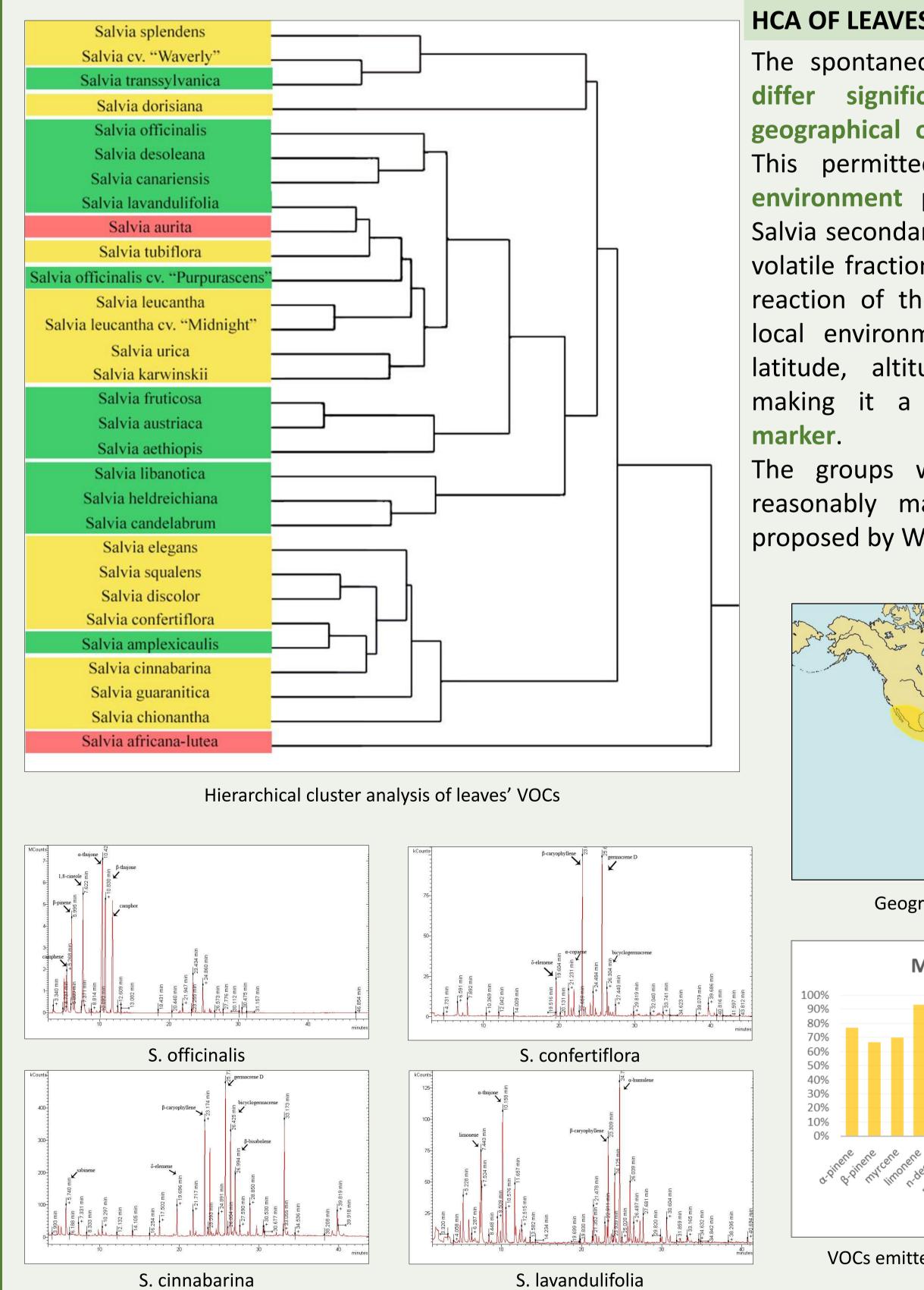


Salvia austriaca Jacq.

American lineage

AIM OF THE STUDY

We investigated 30 species of Salvia by means of HS-SPME-GC-MS to to evaluate the existence of possible **patterns** in the spontaneous emission of VOCs and to find out possible **parameters** that lead to such patterns. We also investigated the collected leaves samples to assess the presence (or the lack) of thujone (α - and/or β -thujone) in the volatile fraction: species with high thujone content are less viable to be used in the food and pharmaceutical industry because of the neurotoxicity of these molecules.

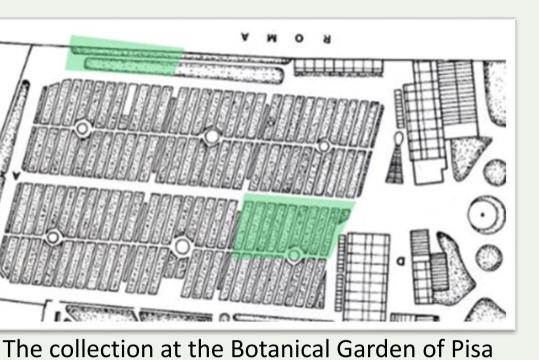


RESULTS AND STATISTICS

HCA OF LEAVES VOCs EMISSION

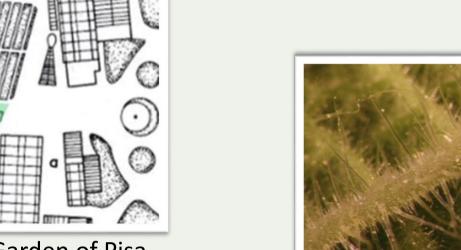
The spontaneous volatile emission profiles differ significantly accordingly to the geographical origin of the various species. This permitted to hypothesize that the environment plays a fundamental role in Salvia secondary metabolites production: the volatile fraction, in particular, represents the reaction of the specimen to the particular local environment (temperature, humidity, latitude, altitude, pollinators, enemies...), making it a possible chemotaxonomical marker.

The groups we identified also seem to reasonably match the clades distribution proposed by Walker et al.





Salvia libanotica Boiss. & Gaill.



Multicellular leaf trichome of *Salvia dorisiana* Standl.

MATERIALS & METHODS

Salvia canariensis L.

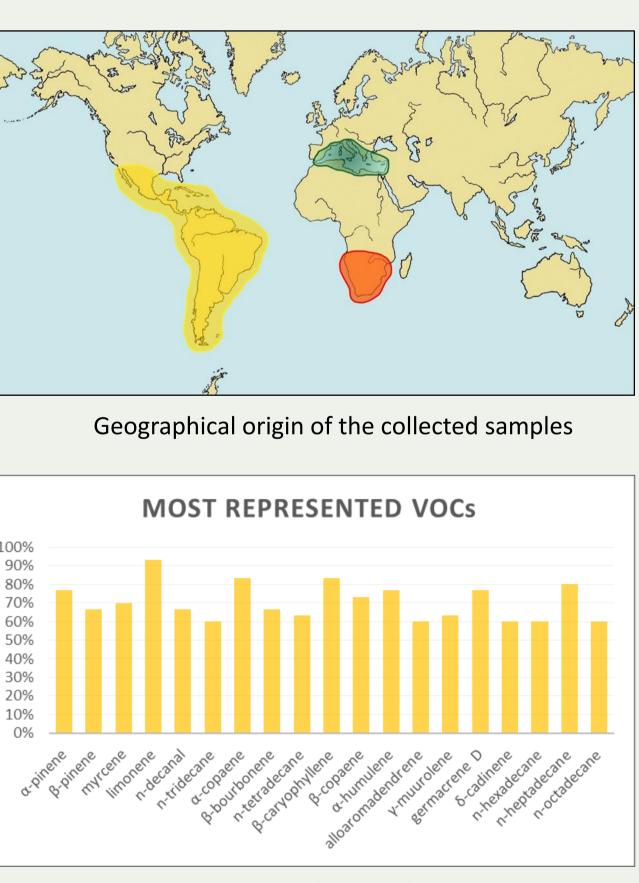
SAMPLES

30 living leaves samples taken from a collection located at the **Botanical Garden of Pisa**.

HS-SPME-GC-MS

Sampling was carried out for a variable time (30 min – 1 h) with a **PDMS coated fibre**. The GC/EI-MS analyses were performed with a **Varian CP-3800** apparatus equipped with a DB-5 cap. column (30 m x 0.25 mm i.d., film thickness 0.25 mm) and a **Varian Saturn 2000** ion-trap mass detector.

MULTIVARIATE STATISTICAL ANALYSIS



VOCs emitted by the 60% (or more) of the analysed specimens

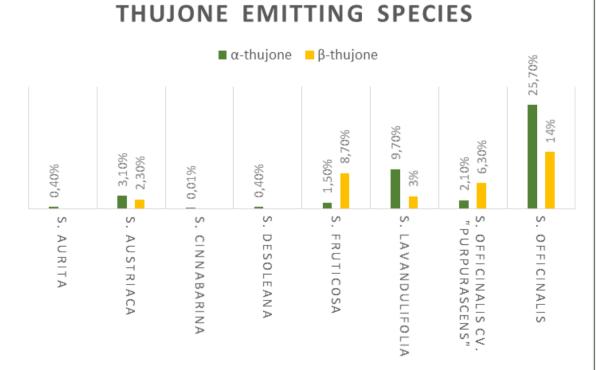
The statistical analyses were carried out with the JMP software package (SAS Institute, Cary, NC, USA). The hierarchical cluster analysis (HCA) was performed using Ward's method with squared Euclidian distances as a measure of similarity.

THUJONE EMITTING SPECIES

Of all the analysed specimens, eight were found to emit α - and/or β thujone. The four species with the highest percentage of these molecules in their VOCs emission profile are:

- 1. S. officinalis
- 2. S. officinalis cv. «Purpurascens»
- 3. S. lavandulifolia
- 4. S. fruticosa

All of them are plants whose origin is located in the Mediterranean area.





OUR FINDINGS

The spontaneous VOCs emission profiles showed a distribution which significantly matches the geographical origin of the analysed

CONCLUSION

specimens. The identified cluster also showed a **correspondence with the clades** identified in Walker et al., whose study seems to be confirmed by our results. The volatile fraction of the specimens is a result of the plant adaptation to the environment, thus making it a possible marker of the plants' origin.

POSSIBLE FUTURE ANALYSIS

- Analysis of a wider number of specimens.
- Analysis of Asian specimens.
- Evaluation of essential oils composition patterns.

Bibliography

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