

Incorporating standardised drift-tube ion mobility to enhance non-targeted assessment of the wine metabolome (LC × IM-MS)

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Outline







1) Low-field drift tube ion mobility with time-of-flight mass spectrometry (IM-TOFMS)

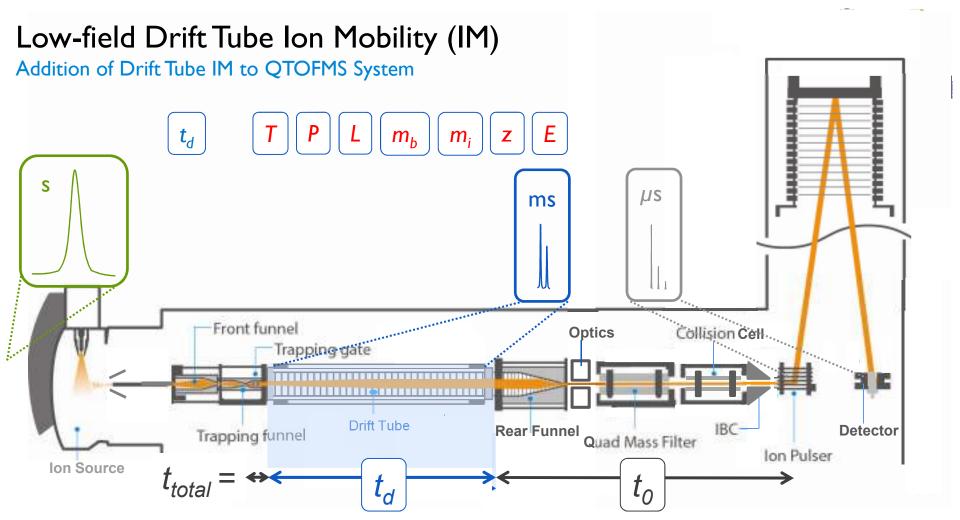
Instrument design overview Collision Cross Section (DTCCS)

2) Macedonia – an emerging wine producer

Climate and wine growing regions Preparation and collection of wine samples

3) Analytical workflow and results

Wine fingerprinting Using DTCCS for identity confirmation



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Ion Mobility: Drift Time (t_d) and DTCCS





LC-compatible Methodology

Single-field Determination [1]:

Transient peaks allow only enough time for one field strength to be used.

Use drift times of calibrant ions (reference ^{DT}CCS values) to perform universal linear calibration.

Universal calibration function applied to IM-MS or LC × IM-MS files.

=> One set of calibrant ions for all metabolite classes and charge states.

[1] Fjeldsted et al, Highly Accurate Collision Cross Section Measurements for Comprehensive High Throughput Applications, 2016, 64th ASMS, San Antonio, USA.

Wine in Macedonia (collaboration with Štip)







Location & Climate



Republic of Macedonia

A sunny, pristine and mountainous country in the heart of the Balkans, Macedonia is a centuries-old producer of fine wines.

Mediterranean and continental climates, with warm summer days and cooler nights. Rich, heavy and clayfilled soils are good base for lengthy ripening of the grapes.

- higher concentrations of sugar, acids and phenolics
- wines with intense aroma, freshness and harmony

Wine in Macedonia



Stanušina





Grape varieties: red

Vranec (most economically important), Stanušina (autochthonous variety), Kratošija (Balkan variety), Pinot Noir, Merlot, Kadarka, Cabernet Sauvignon, Cabernet Franc



Vranec



Kratošija

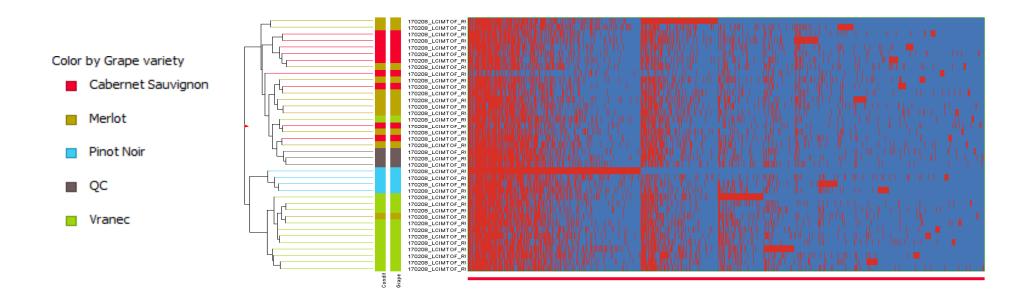


Hierarchical Clustering on Grape Variety





Unsupervised Eucledian (Wards Linkage)



Summary



Current Status and Outlook for Metabolomics

- Wine fingerprinting is maturing, but will allow building of libraries
- Identity confirmation potential with DTCCS is powerful
- General method development for metabolomics workflows is promising, but not yet mature