

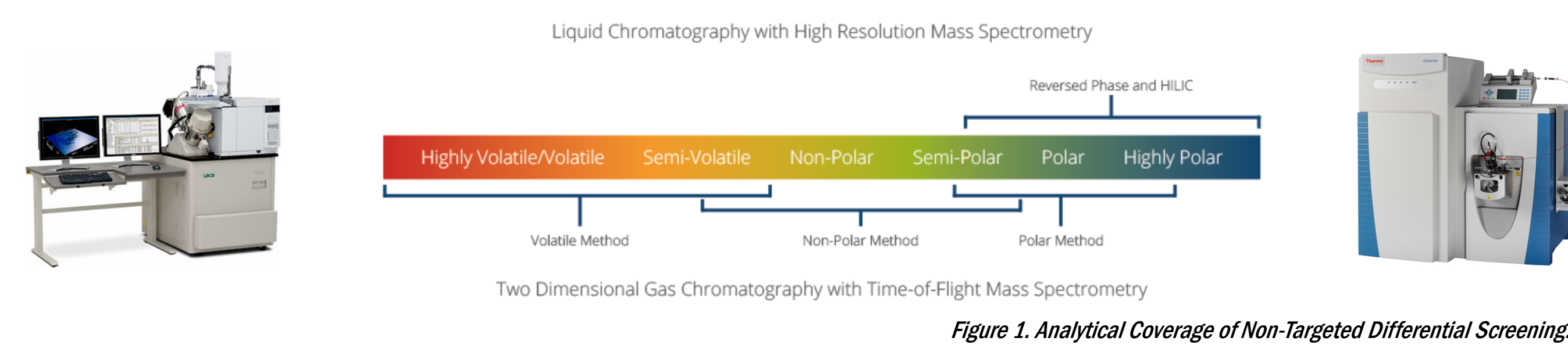
Non-targeted differential screening of complex matrices using GC×GC-TOFMS for comprehensive characterization of the chemical composition and determination of significant differences

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Introduction

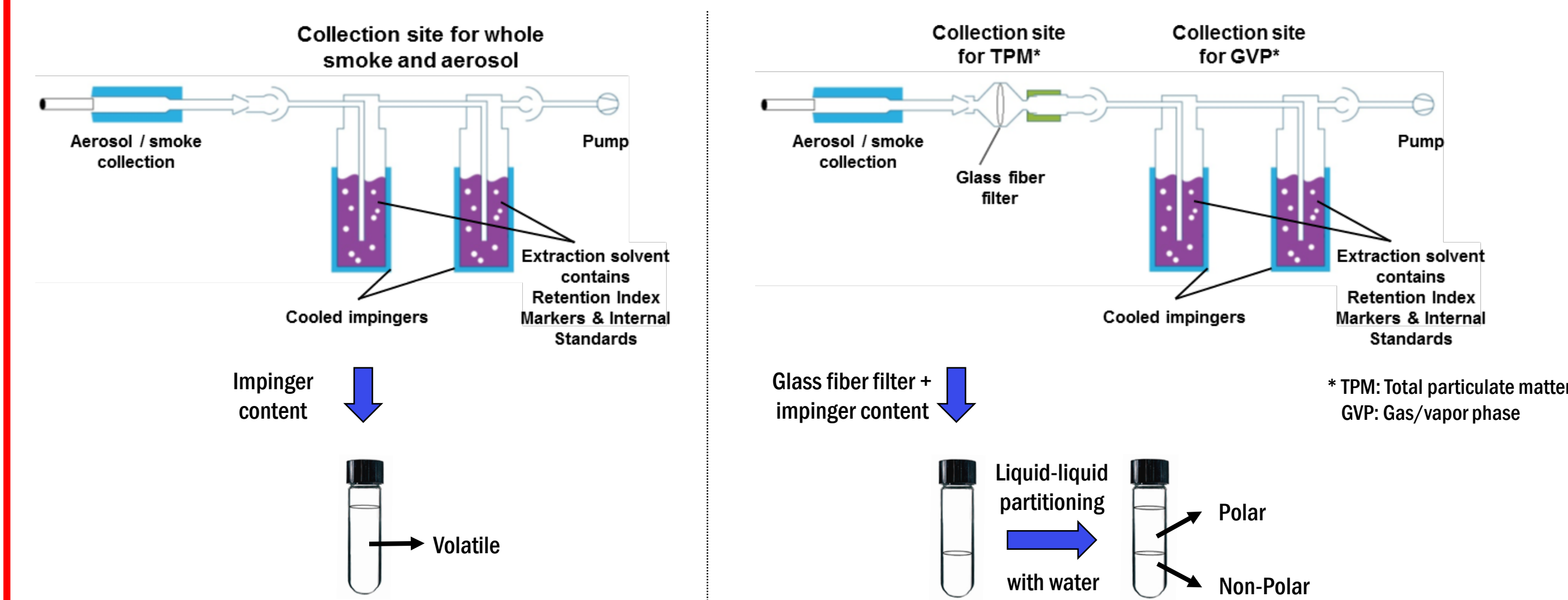
Non-Targeted Differential Screening (NTDS) represents a key methodology to not only comprehensively characterize the chemical composition of aerosol derived from different tobacco products, but also to determine the most relevant differences between complex matrices. Therefore, a complementary NTDS workflow has been developed using two-dimensional gas chromatography coupled to time-of-flight mass spectrometry (GC×GC-TOFMS) in parallel with liquid chromatography coupled to high resolution accurate mass spectrometry (LC-HRAM-MS, see Poster TP 549, Arndt et al.). The approach maximizes the chemical space coverage with multiple analytical methods and determines significant differences based upon semi-quantitative information for all compounds. Together with a Computer-Assisted Structure Identification platform (CASI; in-house development¹), which enhances the accuracy of compound structure identification and accelerates and standardizes the identification process, this approach facilitates the detection of novel compounds and differences in concentration.



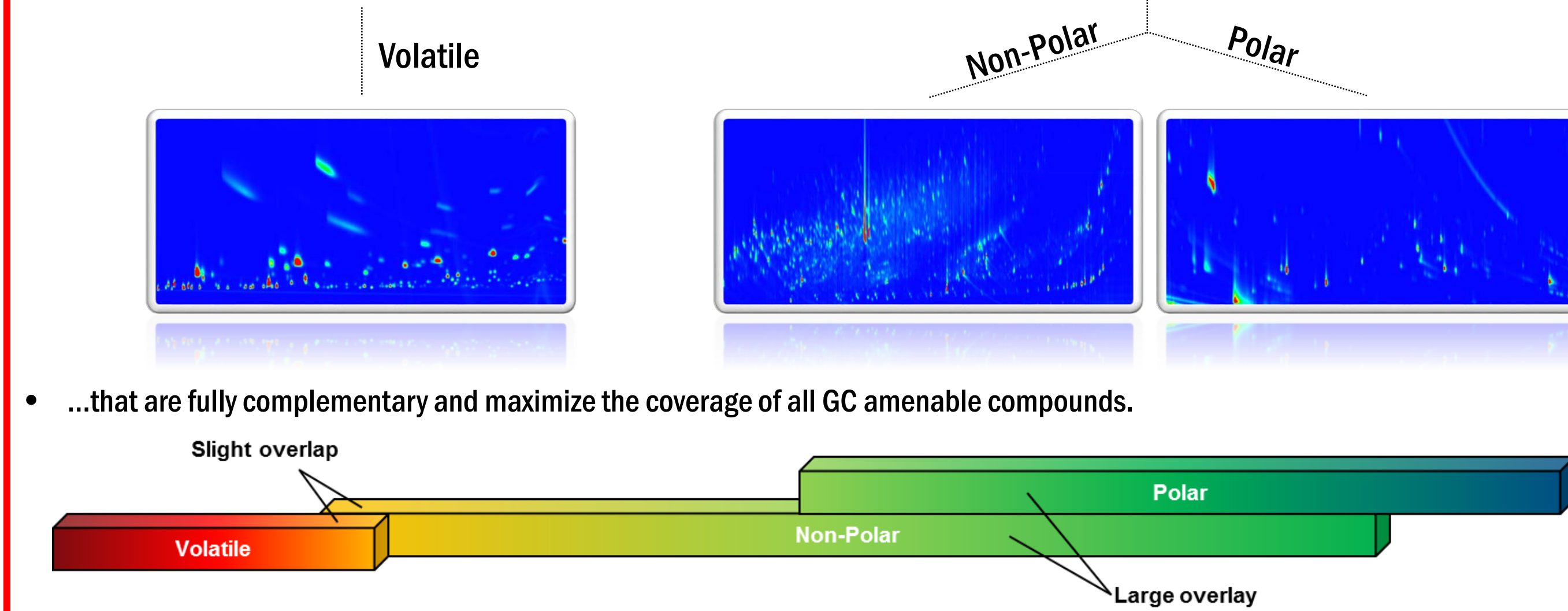
Experimental & Analytical Design

The experimental NTDS workflow comprises:

- Two type of sample collections for constituents that are highly volatile/volatile and non-polar/polar...



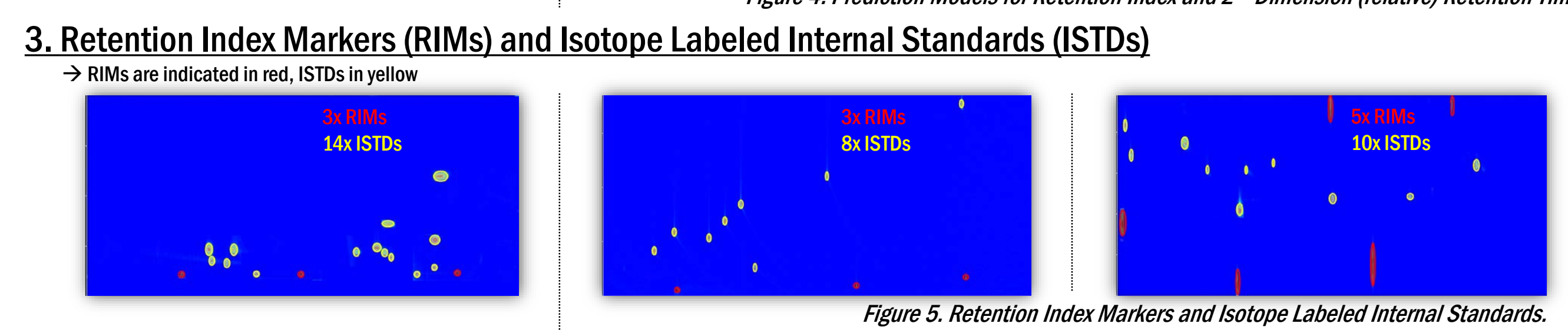
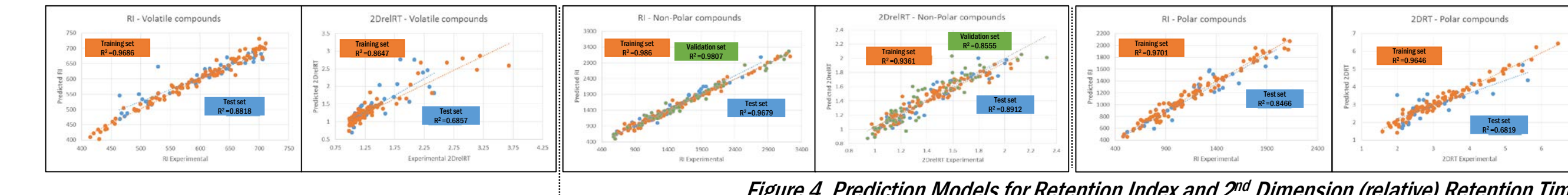
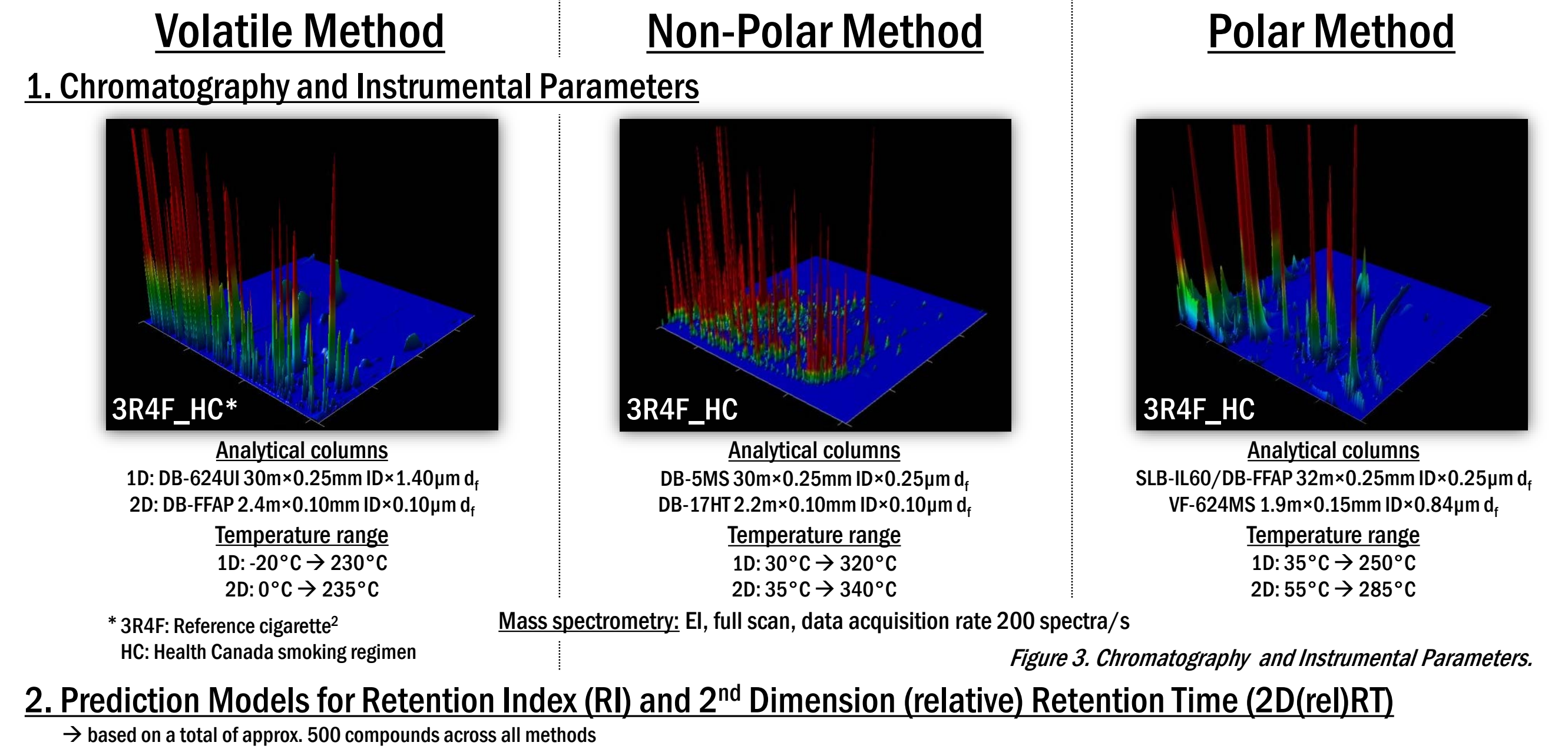
- and three analytical methods with direct cool on-column injection...



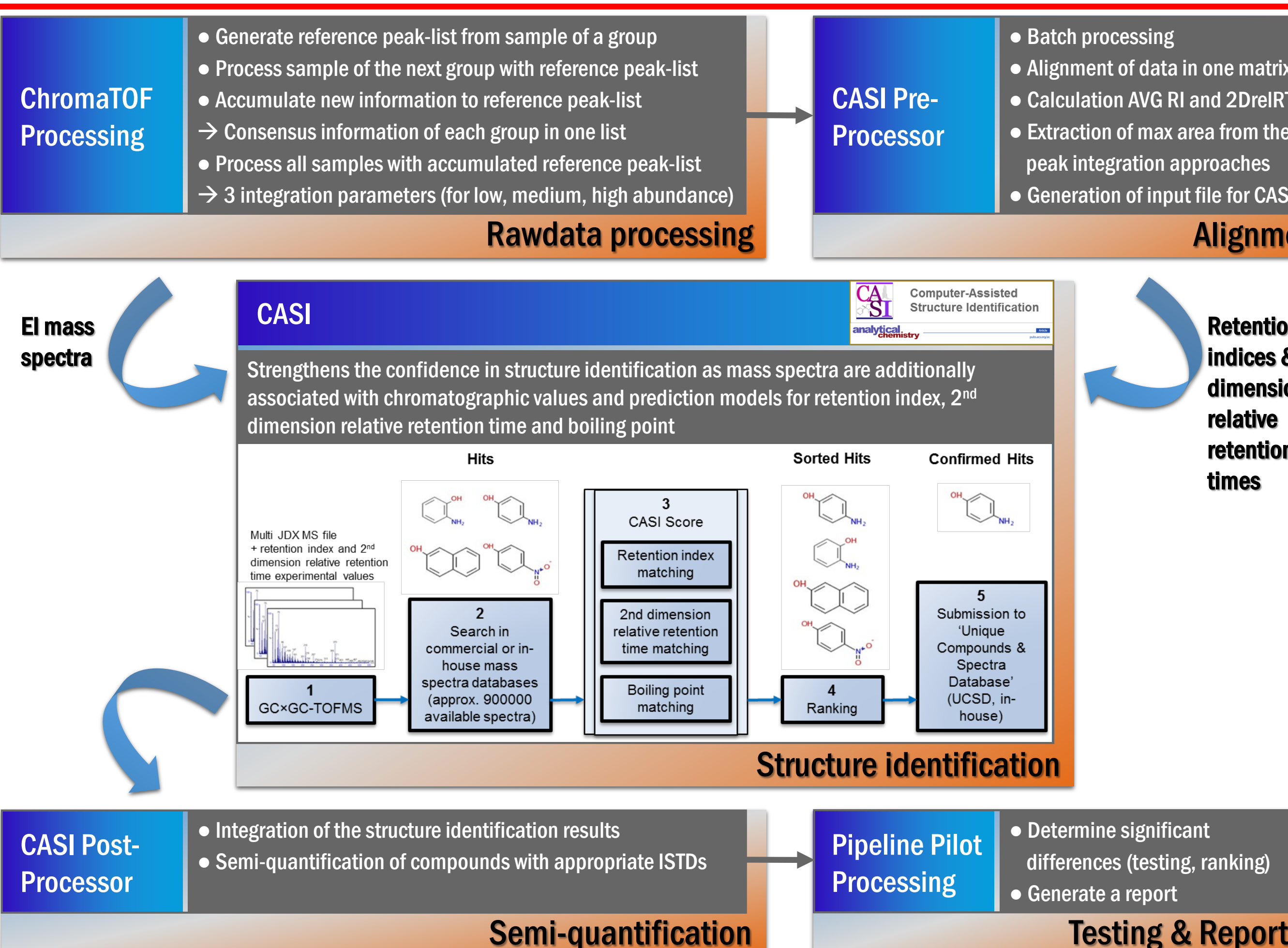
- Advantages:
- Sample preparation is minimized to prevent changes in chemical composition
- Derivatization of polar constituents is avoided by introduction of a novel approach that enables direct injection of aqueous phases
- Compounds that are present in the non-polar and polar fractions can be summed as they derive from the same sample
- Each method contains a dedicated set of retention index markers and stable isotope-labeled internal standards
- Extensive coverage of the chemical functionalities and separation space

Figure 2. Experimental & Analytical Design.

Analytical Methods



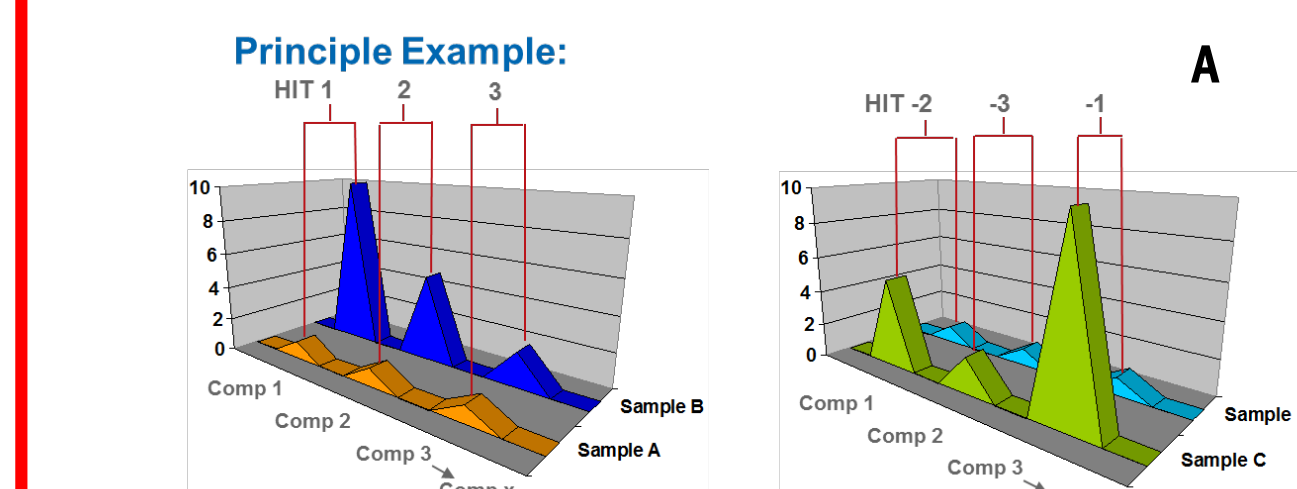
Automated Data Processing Workflow



Validation

Concept

- Addition of selected compounds in different amounts to TPM of a cigarette
- NTDS of fortified TPM vs. non-fortified TPM
- Comparison of experimentally generated HIT-list with theoretical HIT-list
- HIT-list is sorted by RANK formula³ → considers relative differences in abundance of each compound as well as the absolute abundance



Validation Acceptability Criterion

- Correlation coefficient for the reciprocal theoretical HIT numbers and the experimentally found reciprocal HIT numbers > 0.98

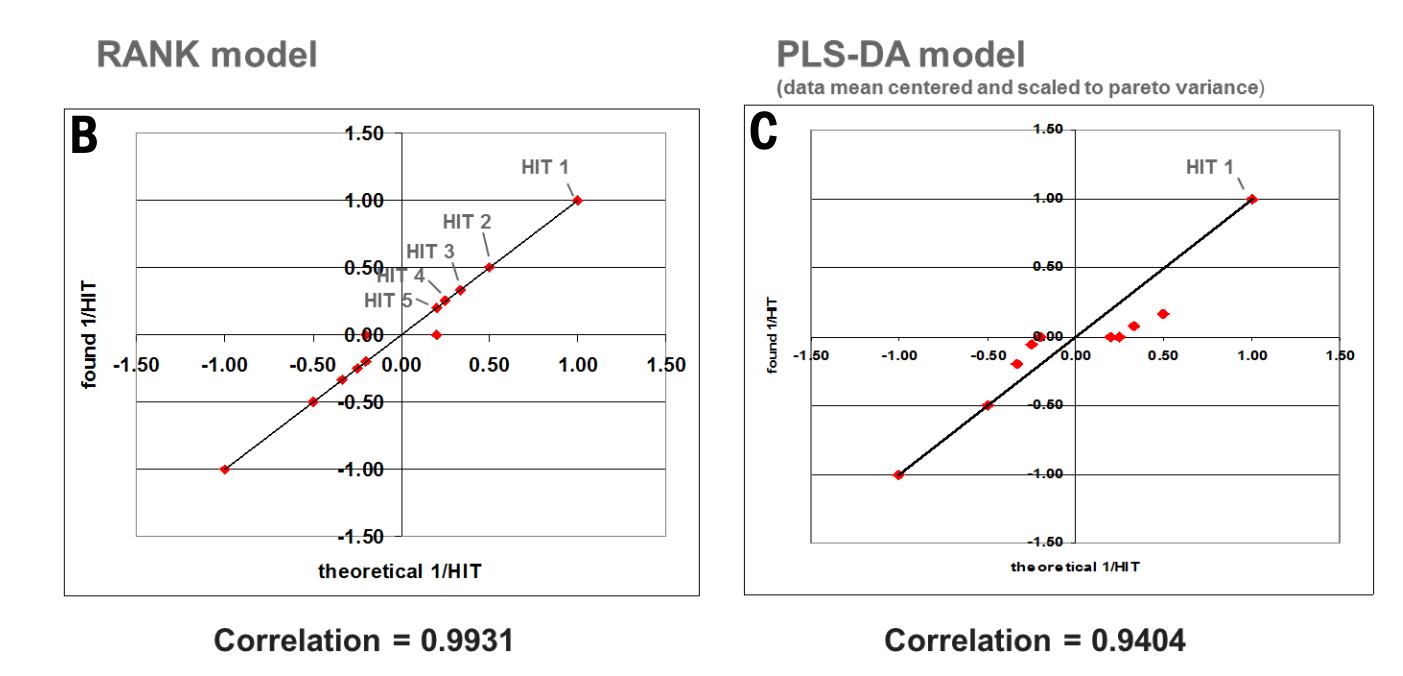


Figure 6. Principle of the Spike-in Experiment (A). Results of an in-house developed RANK Model (B) compared to a conventional PLS-DA Model (C).

Results



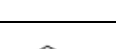
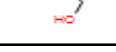
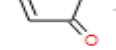
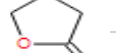

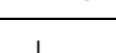
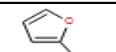
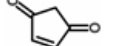
Hit numbers, proposed compound names and structures found in the respective methods				CASI score, probability of the structure proposals and final confidence level upon review of the scientist			Semi-quantitative (N=3) values based on predefined rules, where internal standards are allocated to corresponding compound classes			Unfoldable columns for more details on CASI → RI/2DreIRT/BP deviation (pred vs exp/calc) Structure → PMICODE/CAS/MW/Smiles/Formula Significance → T-test, RANK Compound present in Blank → Exclusion Constituent Origin → Aerosol, Material, Smoke, Plant, Flavor					
Hit #	Proposed compound name	Structure	Method	CASI Score	Probability	Confidence	SEMI-QUANTIFICATION			Identification Details CASI	Structure Identifier	Ranking / T-Test	BLANK	Origin / Probable Source	
							Tobacco Product 1 Sum conc. (μg/gst item)	Tobacco Product 2 Sum conc. (μg/gst item)	X-fold change						
1	Propylene glycol		Polar	927.53	HIGH	IDENTIFIED	Confirmed by REF STD	23.73	174.92	7.4	996.70	PM0001952	43786.26	-	X
2	trans-4-hydroxyphenyl-2-methyl-1,3-dioxane		Polar	834.74	HIGH	HIGH	Agreed	0.00	2.09	unique	1008.05		961.67	-	X
		Non-Polar	846.2	HIGH	HIGH	Agreed	81.28					83.00	-	X	
3	2-Furamethanol		Non-Polar	861.1	HIGH	IDENTIFIED	Confirmed by REF STD	1.12	8.79	7.8	814.86	PM0000003	1602.12	-	X
		Polar	862.38	HIGH	IDENTIFIED	Confirmed by REF STD	1053.41				PM0000003	703.97	-	X	
4	2(5H)-Furanone		Non-Polar	885.1	HIGH	IDENTIFIED	Confirmed by REF STD	1.99	5.32	2.7	898.40	PM0000431	272.99	-	X
		Polar	931.19	HIGH	IDENTIFIED	Confirmed by REF STD	1180.30				PM0000431	83.86	-	X	
5	Butyrolactone		Non-Polar	917.2	HIGH	IDENTIFIED	Confirmed by REF STD	0.73	4.08	5.6	898.62	PM0000396	764.63	-	X
		Polar	942.23	HIGH	IDENTIFIED	Confirmed by REF STD	1066.73				PM0000396	53.08	-	X	
6	2-Furancarboxaldehyde, 5-methyl-		Non-Polar	922.9	HIGH	IDENTIFIED	Confirmed by REF STD	2.94	11.10	3.8	963.74	PM0000001	1393.93	-	X
		Polar	885.66	HIGH	IDENTIFIED	Confirmed by REF STD	984.98				PM0000001	3.07	-	X	
7	1-Hydroxy-2-propanone		Non-Polar	858.1	HIGH	IDENTIFIED	Confirmed by REF STD	6.96	26.65	3.8	589.87	PM00000221	3274.74	-	X
8	Furfural		Non-Polar	883.7	HIGH	IDENTIFIED	Confirmed by REF STD	26.86	47.21	1.8	787.36	PM0000002	767.93	-	X
9	2-Cyclopentene-1,4-dione		Non-Polar	723.0	MEDIUM	IDENTIFIED	Confirmed by REF STD	0.76	3.80	5.0	856.11	PM0000429	670.29	-	X
10	2-Propanone, 1-(acetyloxy)-		Non-Polar	891.2	HIGH	HIGH	Agreed	8.01	16.92	2.1	833.48	PM00003012	569.52	-	X

Figure 7. Example and Description of a Fused Data Result Table.

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Conclusion & Outlook

- The high quality EI mass spectra generated by GC×GC-TOFMS, combined with an enhanced and automated structural identification and semi-quantification workflow, presents a powerful and robust platform for non-targeted assessment of complex matrices
- In total, the NTDS workflow has tentatively identified and semi-quantified more than 3000 constituents in smoke of a cigarette across the different methods
- NTDS is mainly used to compare aerosol of tobacco products, but can also be applied to a variety of other matrices

Next steps:

→ Transfer existing methodologies to GC×GC-HR-TOFMS and establish a workflow for structure elucidation of unknown compounds

References

- Knorr, A., Monge, A., Stueber, M., Stratmann, A., Arndt, D., Martin, E., Pospisil, P., Computer-Assisted Structure Identification (CASI) - An Automated Platform for High-Throughput Identification of Small Molecules by Two-Dimensional Gas Chromatography Coupled to Mass Spectrometry, Anal Chem. 2013, 85(23):11216-24.
- University of Kentucky (www2.ca.uky.edu/refcig/3R4F%20Preliminary%20Analysis.pdf)
- Knorr, A., A method of analyzing data from chemical analysis, International Patent WO 2013098169 A1, PCT/EP2012/076244, 2013, Jul 4.