

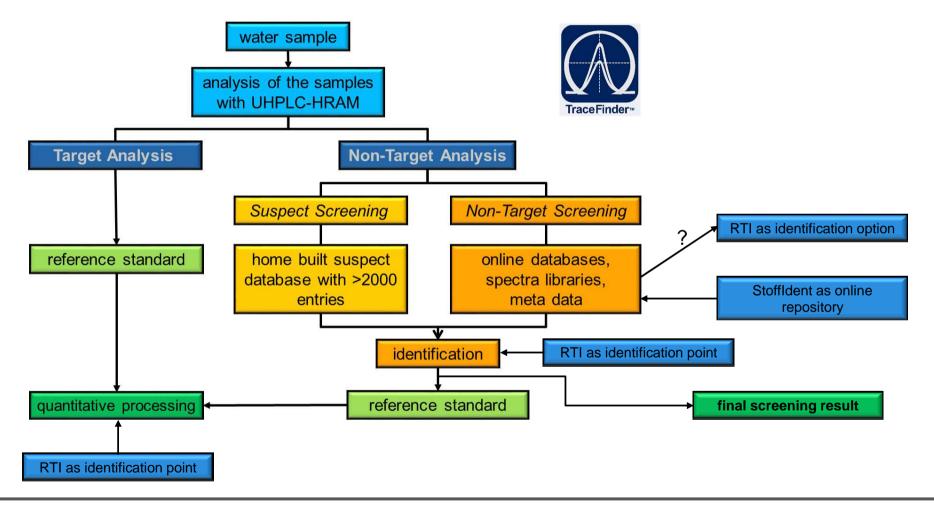


Forldent Workshop Langenau – Water Sample Screening Workflow

Frank Steiner, Anthony Squibb (Slides contributed from Olaf Scheibner, Ryo Komatsuzaki)

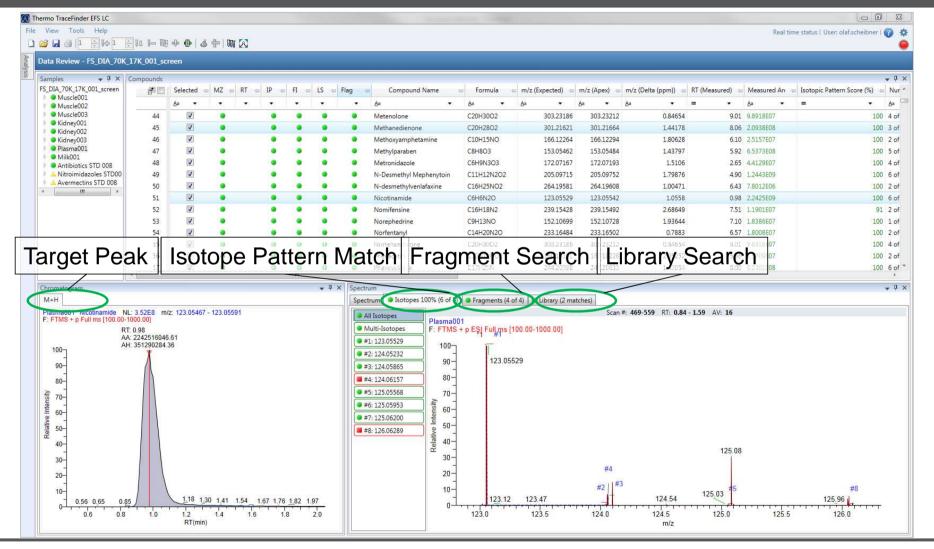
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Quantitative and (Unknown) Screening Workflow



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Result table for suspect screening ("final screening result")



Biologics New Peak Detection

• Non-Targeted MS Peak Detection via Differential Analysis

• Brand new category

Home Tables Frame Table Format	PEP QUAN_DEMO (Sequence) - CI	romeleon Chromatography Studio	- 0
	Annomatogram Plot Start Start All Start A	lent labels	
Non-Targeted MS Processing	Proposition Algoment Detection Use Default Parameters Set as default		Start
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JL, TIC 02- 4 Frames [15 of 3711]) 11.05 min #305 - 495.2870 m/z @ 11.05 min 02- #449 - 613.3168 m/z @ 10.17 min 02-	Peek Internaty Threshold: 1,000,000 Scan Ref(s) If TMS + p ESF Full teck res [300,001 800,00] Peek Internaty Threshold: 1,000,000 Scan Ref(s) If TMS + p ESF Full teck res [300,001 800,00]	•	Norg
1449-4227365 m/c ⊕ 13.40 min ↔ ↔ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓ ↓	Acceleration of the second sec	Pade: 1694 44 20130624_13_Strepty02_500mok - Scan. #308 RT 1594 min NL 7.015-008 FTMS + p ESI Ful lock ms [20 retrory	m 1,750
Data Processing Non-Targeted MS Processing Report Designer	A B C D E F G H I J J K L M 2 Frame ID Component ID Mass PT P	 Frame list with filter mechanism (hidden) Differential ratio threshold Root component for multi-isotope peaks Observed m/z 	1,750
UV Spectral Library	19 Or Frame Table / Frame Flat (S of 3.711) with rule "Frame Flat o> 100.00000 and Frame PatternRecognitor/Element = 0"	Other fields available to display	RKomatsuz

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Biologics New Peak Detection

- Non-Targeted MS Peak Detection
 - Report Variable, Plot, and Table extensions

