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Miniaturization and Automation of Dutch mini Luke extraction of Pesticides in Fruits and Vegetables

Klaudia Dyrda
17th June 2025

The reasons for carrying out the analysis



- To provide a monitoring service in accordance with the harmonised EU monitoring programme.
- REGULATION (EC) No 396/2005 OF THE EUROPEAN PARLIAMENT AND OF THE COUNCIL
- on maximum residue levels of pesticides in or on food and feed of plant and animal origin and amending Council Directive 91/414/EEC

Sampling and sample plans - 2024



Routine samples

F & V

TAT 4 weeks

817 samples

FAO

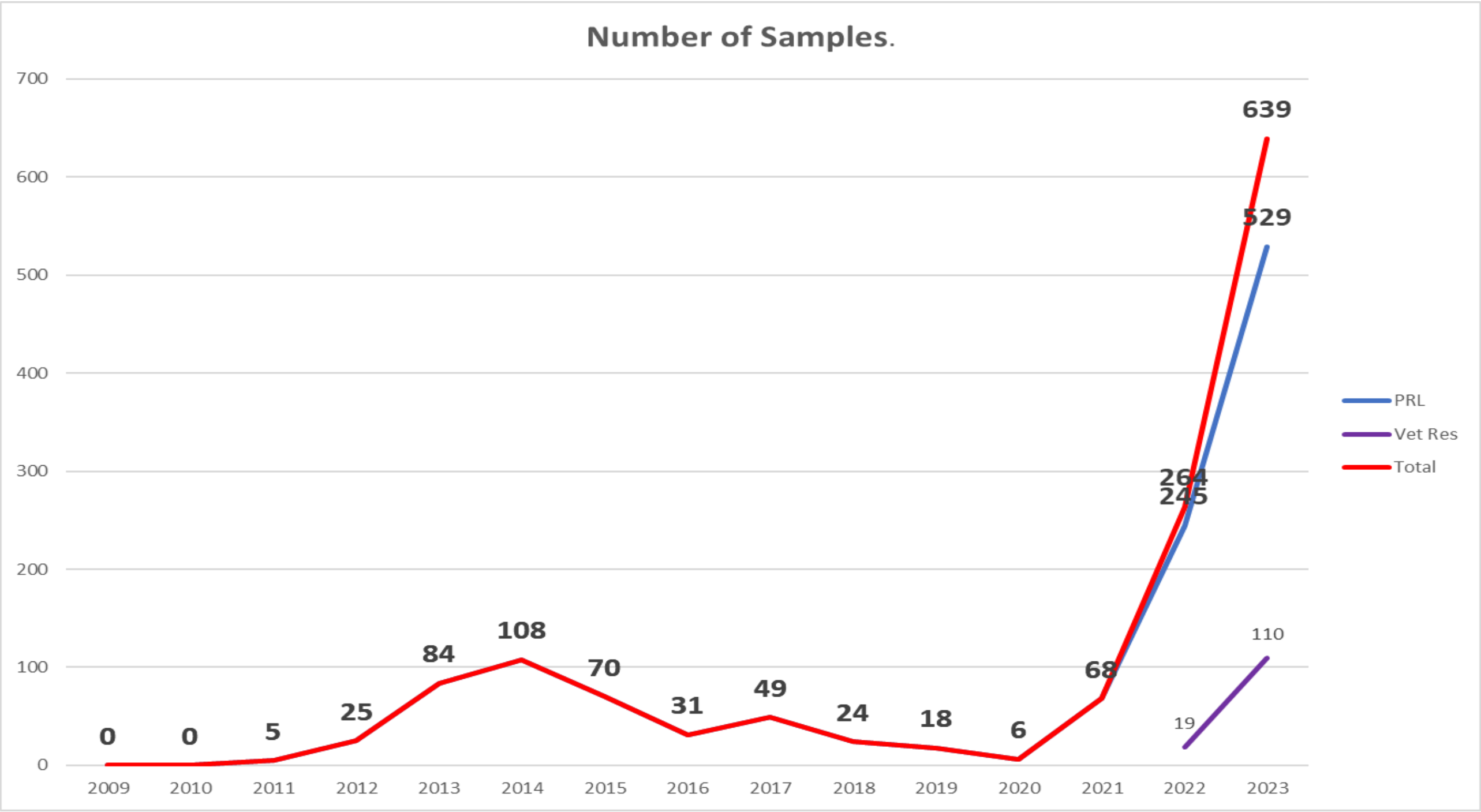
TAT 4 weeks

290 samples

BCP Samples

TAT 48 hours

146 samples

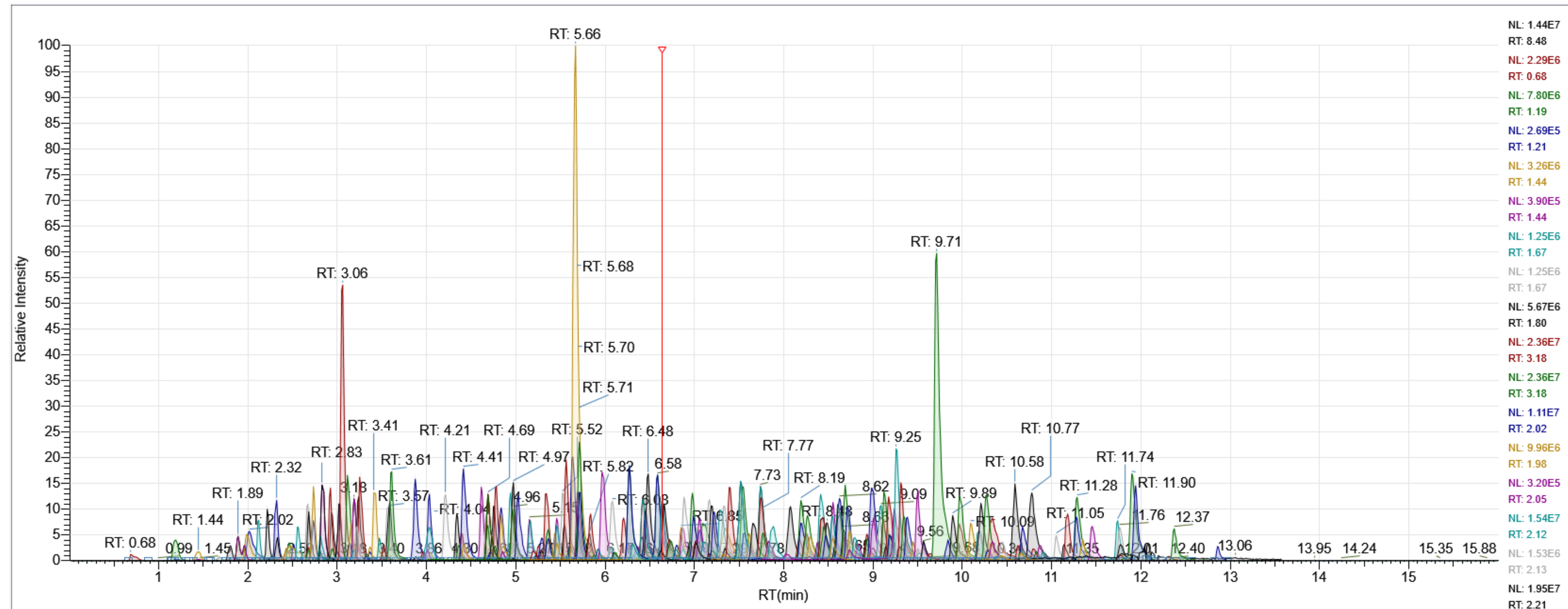


Scope

501 analytes

397 analytes accredited

79% of Fruit and vegetable accredited



Dutch mini-Luke Extraction



Advantages

- Well established
- Sensitive
- Low matrix effects
- Broad Spectrum Extraction
- Amenable with GC-MS and LC-MS

Disadvantages



Dutch mini-Luke Extraction



Samples are
homogenised
15g aliquots

Diluted 1in20 in
Methanol
(LC analysis)



s
to

Centrifuged

yl
(s)

Evaporated down to low
volume

Dutch mini-Luke Extraction

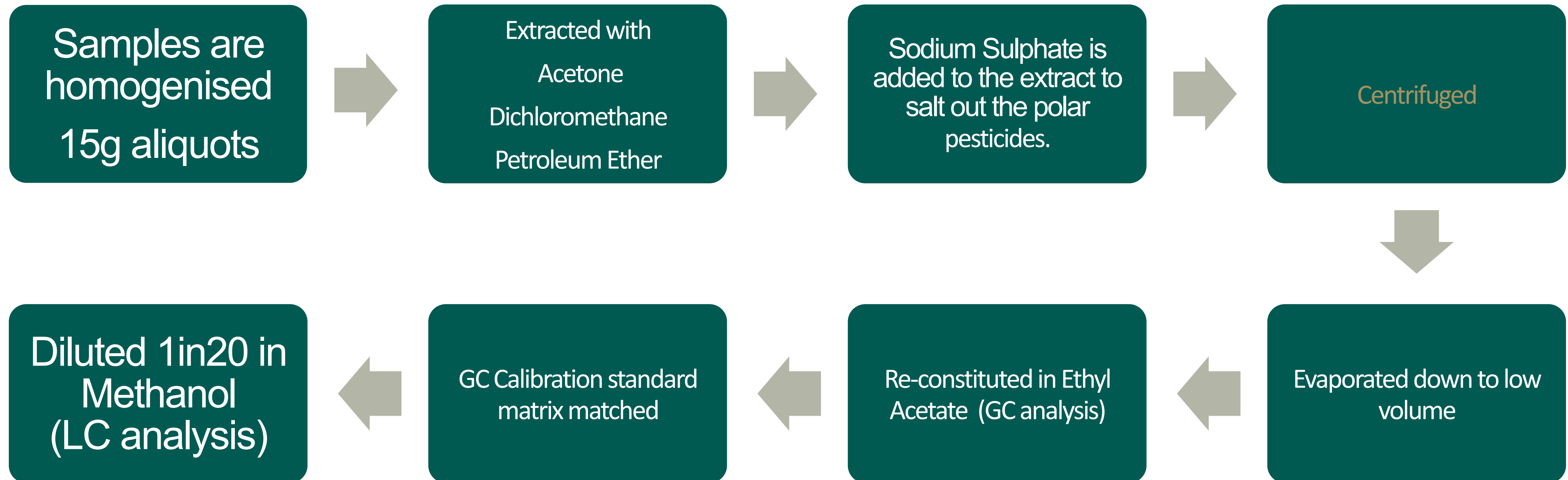


Sodium
added to
salt o
p

Re-cons
Acetate



Dutch mini-Luke Extraction

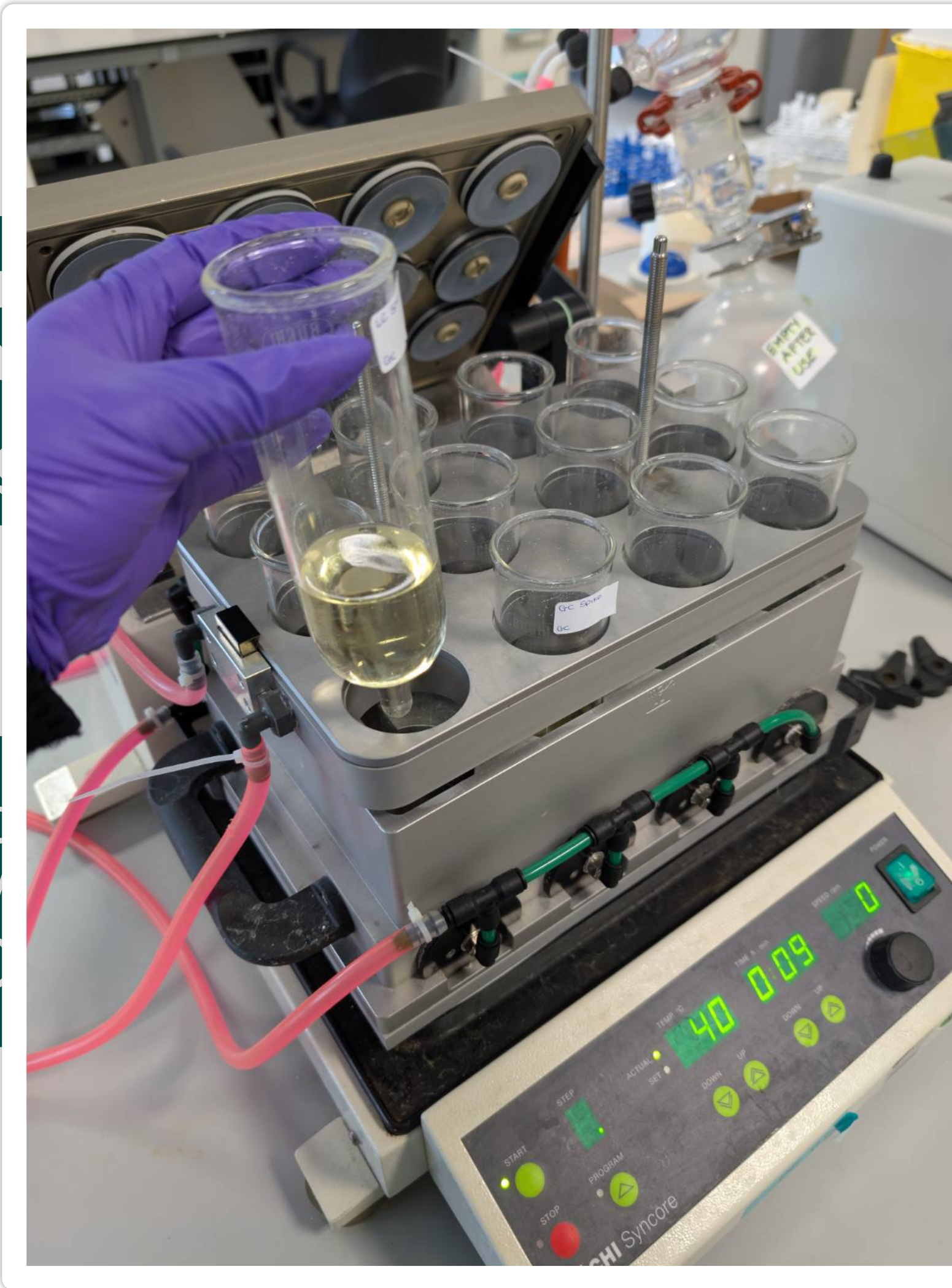


Dutch mini-Luke Extraction



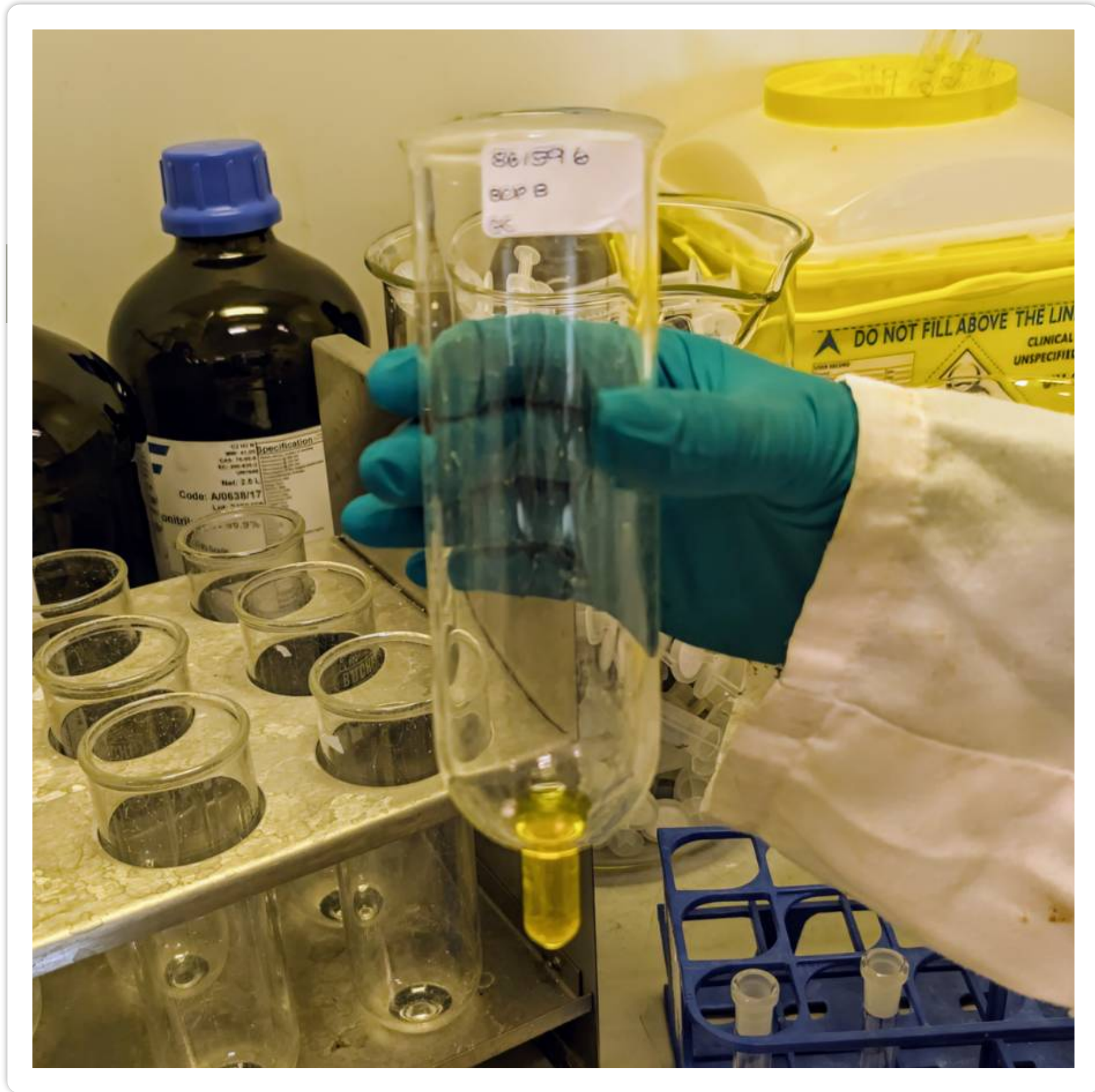
Sample
homogenized
150

Diluted
M
(LO



Standard

Standard



Centrifuged

Reduced down to low
volume

Dutch mini-Luke Extraction



Samples are
homogenised
15g aliquots

Diluted 1in20 in
Methanol
(LC analysis)



Centrifuged

Evaporated down to low
volume

Solvent usage



Acetone	30ml + extra for cleaning
Dichloromethane	30ml
Petroleum Ether	30ml
Ethyl Acetate	25ml
Methanol	9.5ml
Total Solvent usage per sample	>124.5ml

2024 Samples



- 817 Fruit and Vegetables analysed
- Over 101 L of solvent consumed
- 5 water coolers



How do we validate methods?



ANALYTICAL QUALITY CONTROL AND METHOD VALIDATION PROCEDURES FOR PESTICIDE RESIDUES ANALYSIS IN FOOD AND FEED SANTE 11312/2021 v2

Supersedes Document No. **SANTE/11312/2021**. Implemented by 01/01/2024

Initial Full Validation



Tomato
Avocado
Potato

Validation needs to be performed

For all analytes within the scope of the method

For at least 1 commodity from each of the commodity groups

- *High water content – Apple, onion, broccoli*
- *High acid content and high water content – Lemon, strawberry, grape*
- *High sugar content and low water content – dried fruit, honey, fruit jam*
- *High oil content and very low water content – walnut, sunflower seed, peanut butter*
- *High oil content and intermediate water content – olives, avocado*
- *High starch and/or protein content and low water and fat content – lentils, barely, pasta*

Experimental set up

10ppb
20ppb
50ppb
100ppb



Sample set

- Reagent blank
- 1 blank sample
- 5 Spiked samples at target LOQ
- 5 Spiked samples at 2-10 x target LOQ

Instrumental sample sequence

- Conditioning blanks
- Calibration standards
- Reagent blank
- Sample blank
- 5 spiked samples at target LOQ
- 5 spiked samples at 2-10 x Target LOQ
- Calibration standards

Validation parameters and criteria



Parameter	What/How	Criterion
Sensitivity/Linearity	Linearity check from 5 levels	Deviation of back-calculated concentration from true concentration $\leq +20\%$
Matrix effect	Difference of response from standard in matrix extract and standard in solvent	In case of more than 20% signal suppression or enhancement, matrix effects need to be addressed in Calibration
LOQ	Lowest spike level meeting the identification and method performance criteria for recovery and precision	$\leq \text{MRL}$
Specificity	Response in reagent blank and blank control samples	$\leq 30\%$ of Reporting Limit
Recovery	Average recovery for each spike level tested	70 – 120%
Precision (RSD_r)	Repeatability RSD_r for each spike level tested	$\leq 20\%$
Precision (RSD_{wR})	Within-laboratory reproducibility, derived from on-going method validation / verification	$\leq 20\%$
Robustness	Average recovery and RSD_{wR} derived from on-going method validation / verification	
Ion ratio	Check compliance with identification requirements for MS techniques	
Retention time		$\pm 0.1 \text{ min}$

Identification requirements for different MS techniques



MS Detector/Characteristics		Acquisition	Requirements for identification	
Resolution	Typical systems (Examples)		Minimum number of ions	Additionally
Unit Mass resolution	Single MS	Full scan, limited m/z range, SIM	3 ions	S/N ≥ 3
	Quadrupole, ion trap, TOF			Analyte peaks from both product ions in the extracted ion chromatograms must fully overlap.
	MS/MS	Selected or multiple reaction monitoring. Mass resolution for precursor-ion isolation equal to or better than unit mass resolution	2 products ions	Ion ratio from sample extracts should be within $\pm 30\%$ (relative) of average of calibration standards from same sequence
	Triple Quadrupole, ion trap, Q-trap, Q-TOF, Q-Orbitrap			
Accurate mass measurement	High Resolution MS: (Q-)TOF (Q-)Orbitrap	Full scan, limited m/z range, SIM fragmentation with or without precursor-ion selection, or combination thereof	2 ions with mass accuracy ≤ 5 ppm Preferably include the molecular ion Include at least 1 fragment ion	S/N ≥ 3 Analyte peaks from precursor and/or products ion(s) in the extracted ion chromatograms must fully overlap.



Is Miniaturization possible?

Protocol



1. Aliquot 1g sample into 50ml tube
2. Spike
3. Add 2ml Acetone shake with ceramic bead
4. Add 2ml Petroleum ether, 2ml Dichloromethane and 2g of sodium sulphate
5. Shake using shaker
6. Centrifuge
7. Transfer 3ml into glass vial, evaporate to dryness using Turbovap
8. Reconstitute with 0.5ml of Ethyl Acetate
9. Filter the extract through a 0.2 μm filter – GC Fraction
10. Transfer 50 μl of extract into 2ml vial and add 950 μl Methanol – LC Fraction

Solvent usage



Acetone	2ml
Dichloromethane	2ml
Petroleum Ether	2ml
Ethyl Acetate	0.5ml
Methanol	0.95ml
Total Solvent usage per sample	7.45ml

Solvent usage

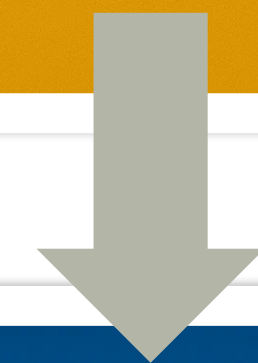
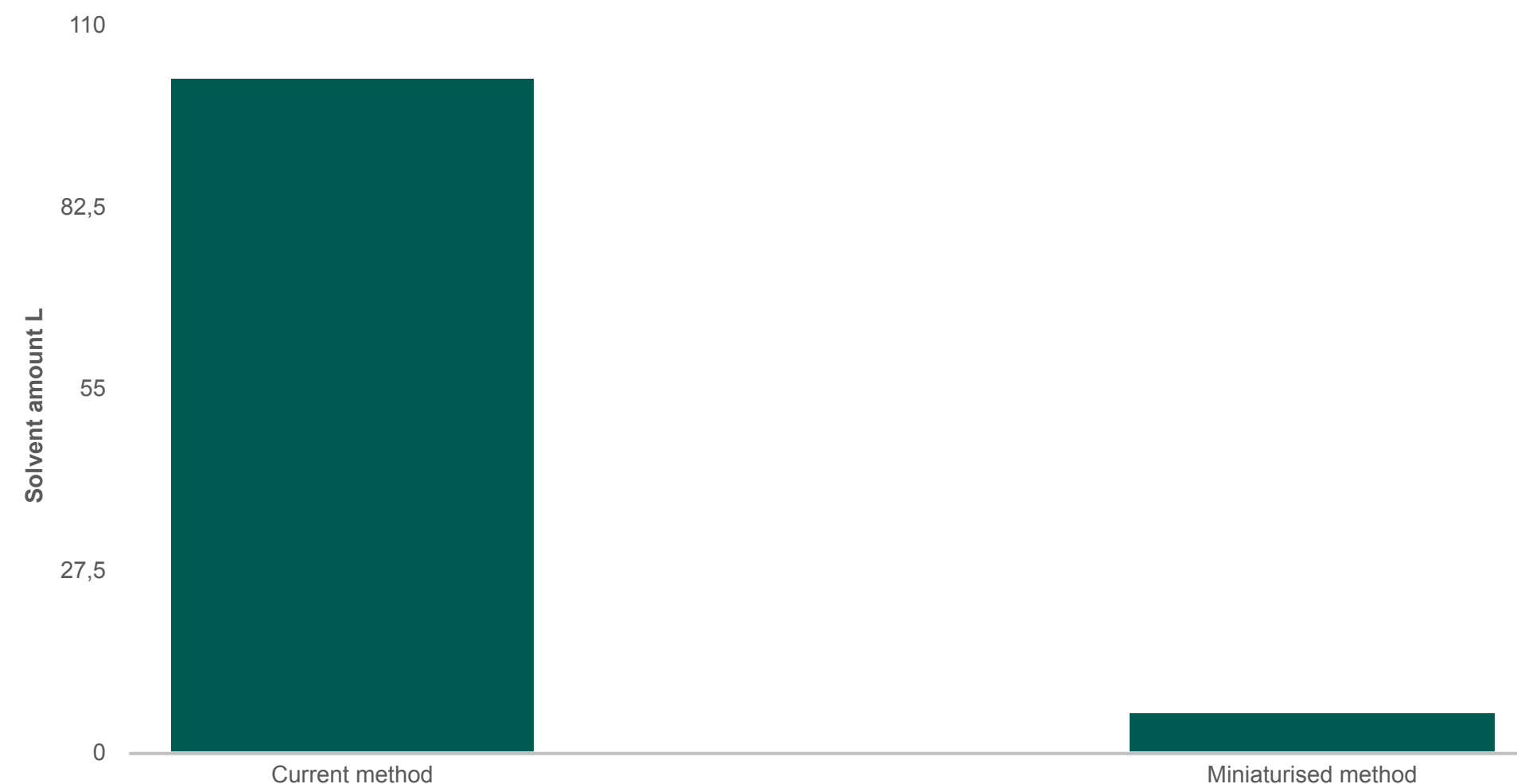


Acetone	15ml + extra for cleaning
Dichloromethane	15ml
Petroleum Ether	15ml
Ethyl Acetate	25ml
Methanol	9.5ml
Total Solvent	>124.5ml



2024 Samples

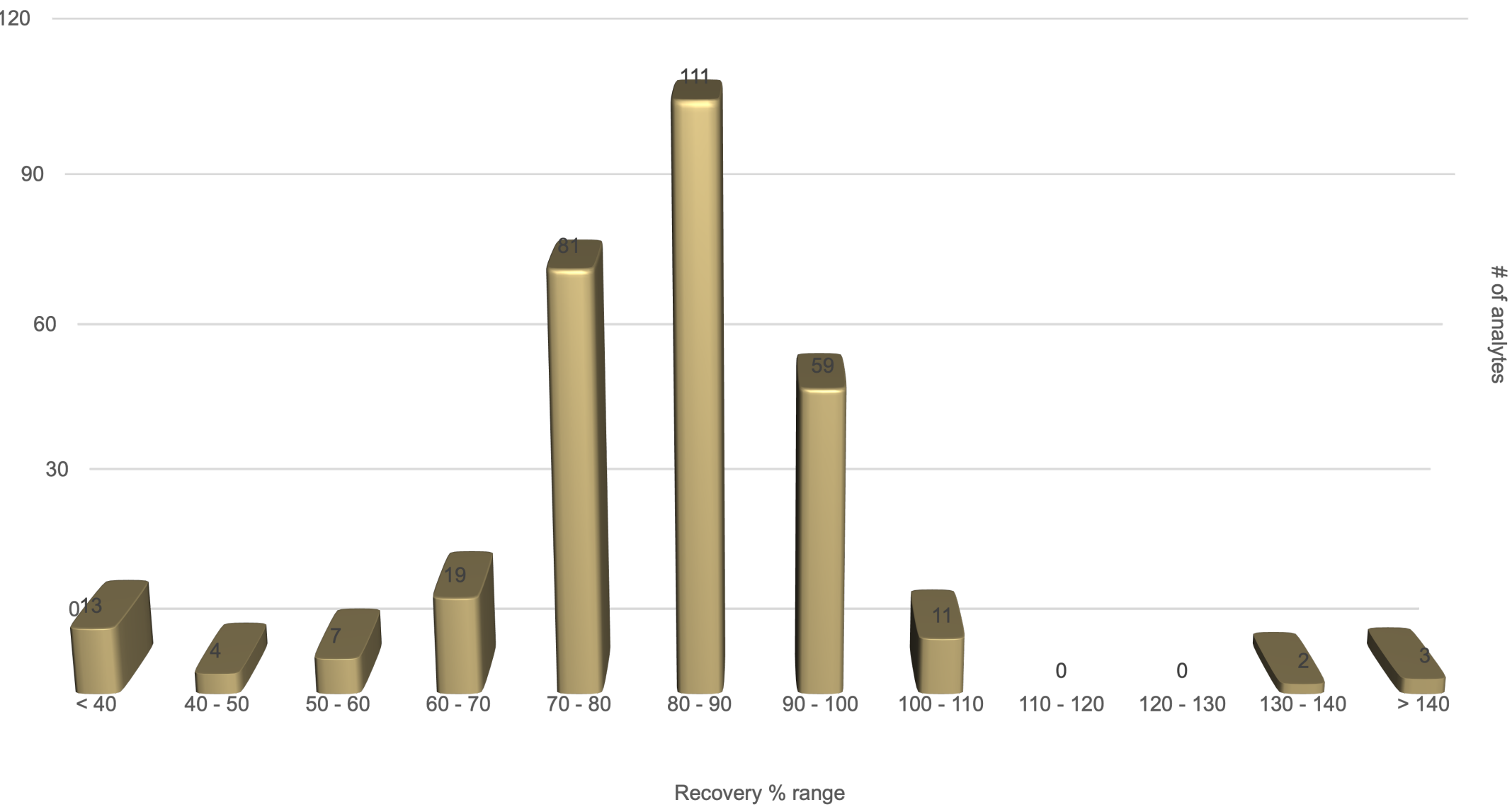
- 817 Fruit and Vegetables analysed
- Over 101L of solvent consumed
- Miniaturised method only uses 6 L
- ~17 times less solvent would be consumed



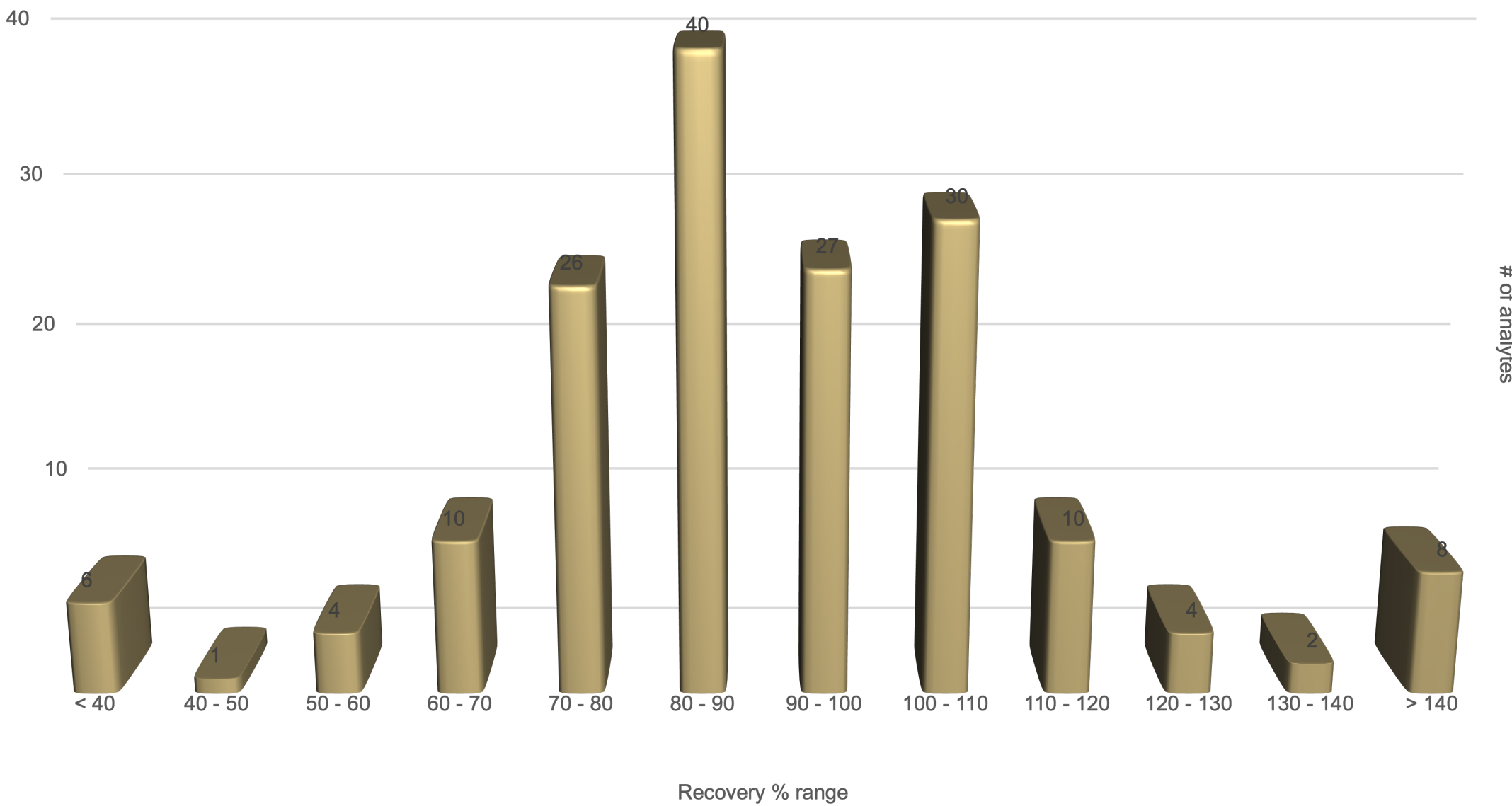
Results



LC HRMS



GC HRMS



310 analytes
262 analytes acceptable recovery

85% acceptable



Can it be automated?

Collaboration with Da Vinci Laboratory Solutions UK and Ireland Ltd.



Colin Hastie – Application Chemist

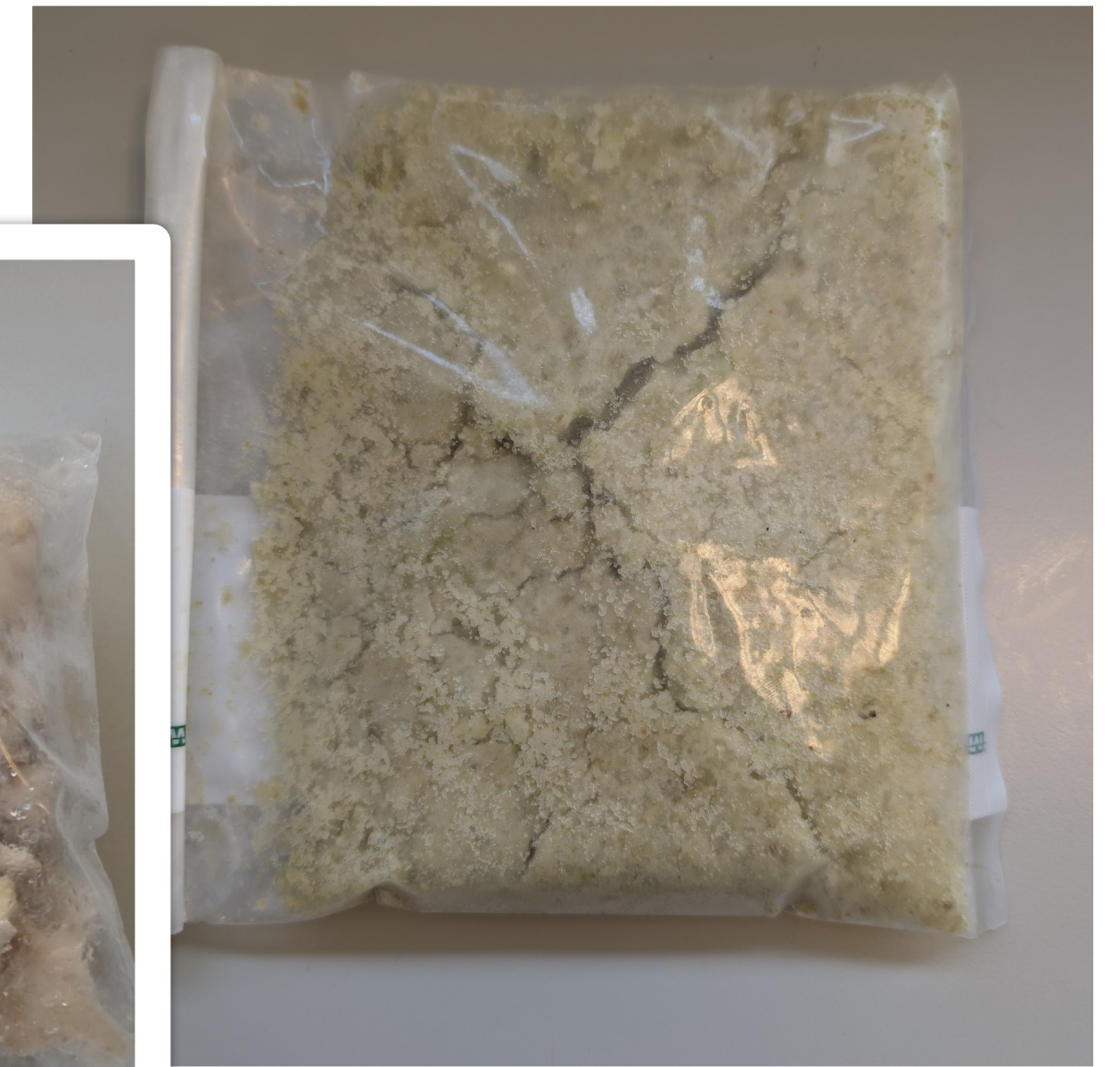
Use of Gerstel Dual head Robotic Pro Multipurpose sample (MPS)



Project Plan

Provide Colin with:

- Standards mixes
- Matrix – tomato, avocado & potato
- GC MS/MS methods



Protocol



Into a 10 mL glass vial 1 g of samples weigh out and 2 g of sodium sulphate, this is then be loaded on to the MPS which is programmed to carry out the following actions on the samples.

Sample is spiked at 4 levels with n=6 replicates at each level.

1. Spike with 1 mg/L Std mix (0, 10 ,50 or 100 µL as appropriate)
2. Add 2 mL of acetone to the sample
3. Move the sample vial to the quick mix and mix at 2000 rpm for 30 seconds
4. Add 2 mL of Petroleum ether to the sample
5. Add 2 mL of Dichloromethane to the sample
6. Mix the sample at 2000 rpm for 30 seconds
7. Move the sample to the centrifuge and centrifuge at 2000 G for 3 minutes.
8. Transfer 3 mL of sample extract to a 4 mL vial
9. Add 50 µL of **nonane** to the 4 mL vial as a keeper solvent
10. Evaporate the extract in the MVar
11. The vial is reconstituted in 0.45 mL of Ethyl acetate and is ready for GC analysis
12. Transfer 50 µL of sample extract to separate 2 mL vial
13. Add 950 µL of methanol to the new vial ready for LC-MS/MS analysis



Experimental



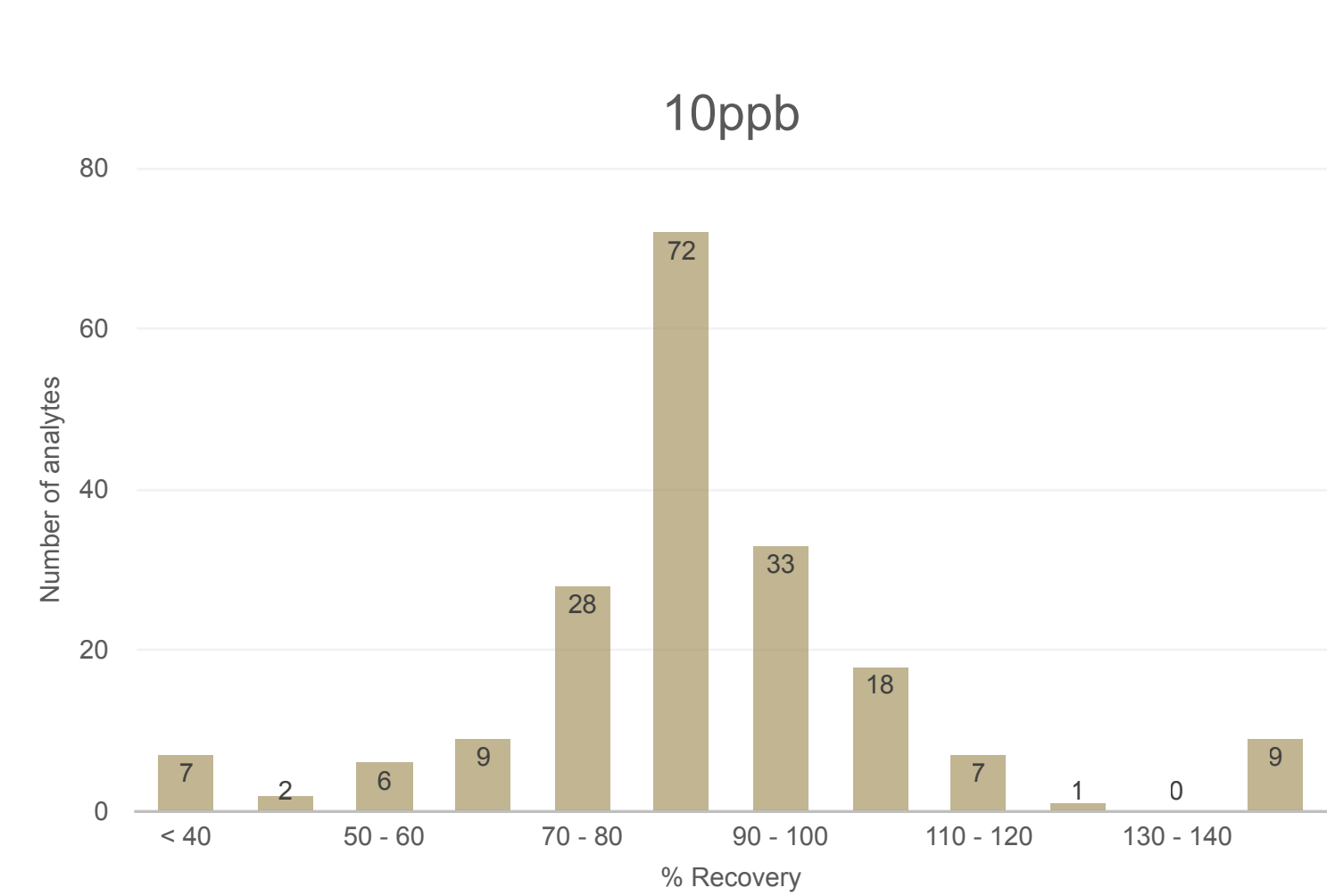
Validation Batch of 30 samples ~7hrs

Approx 15 min per sample

Routine Batches ~ 18 samples

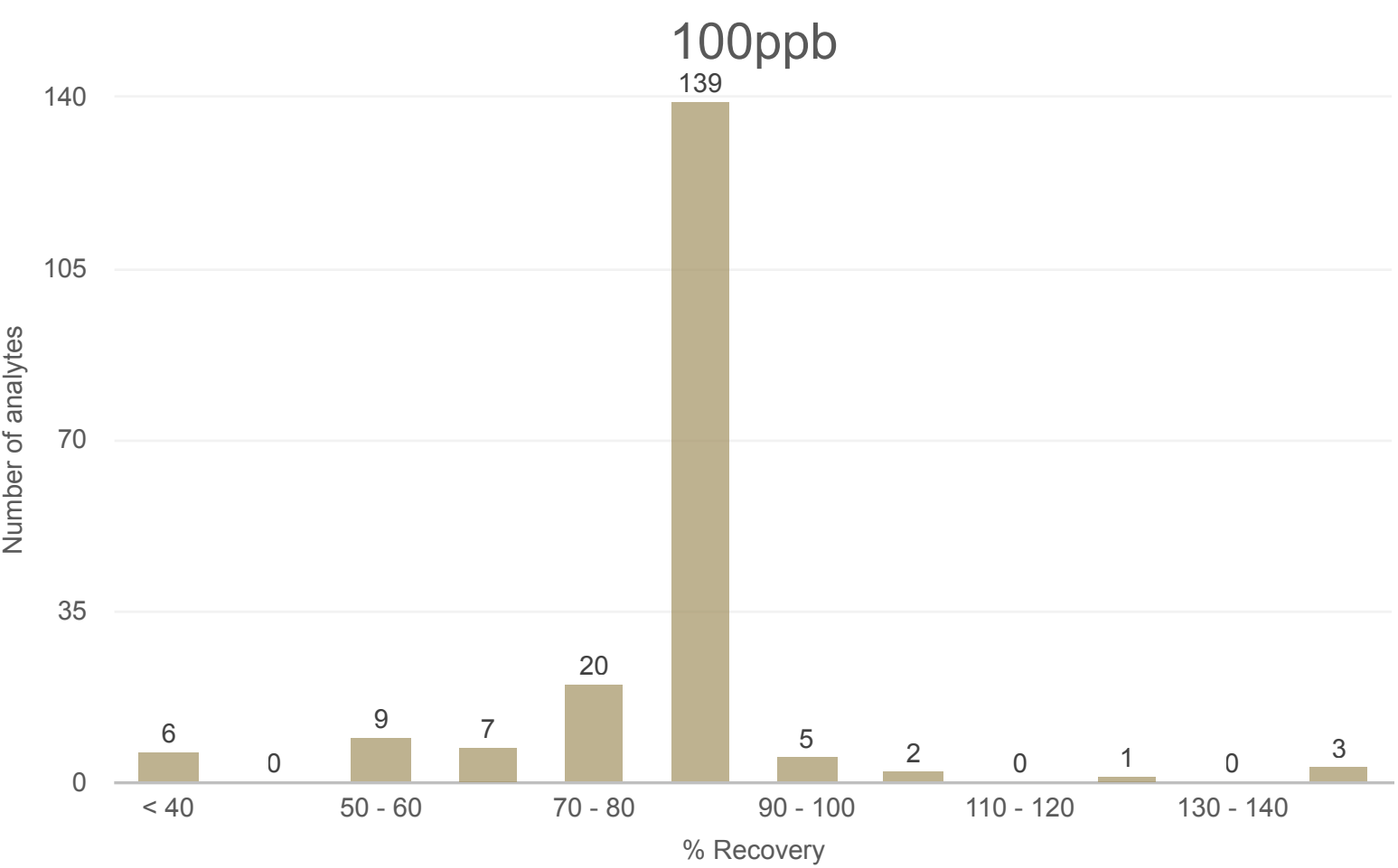
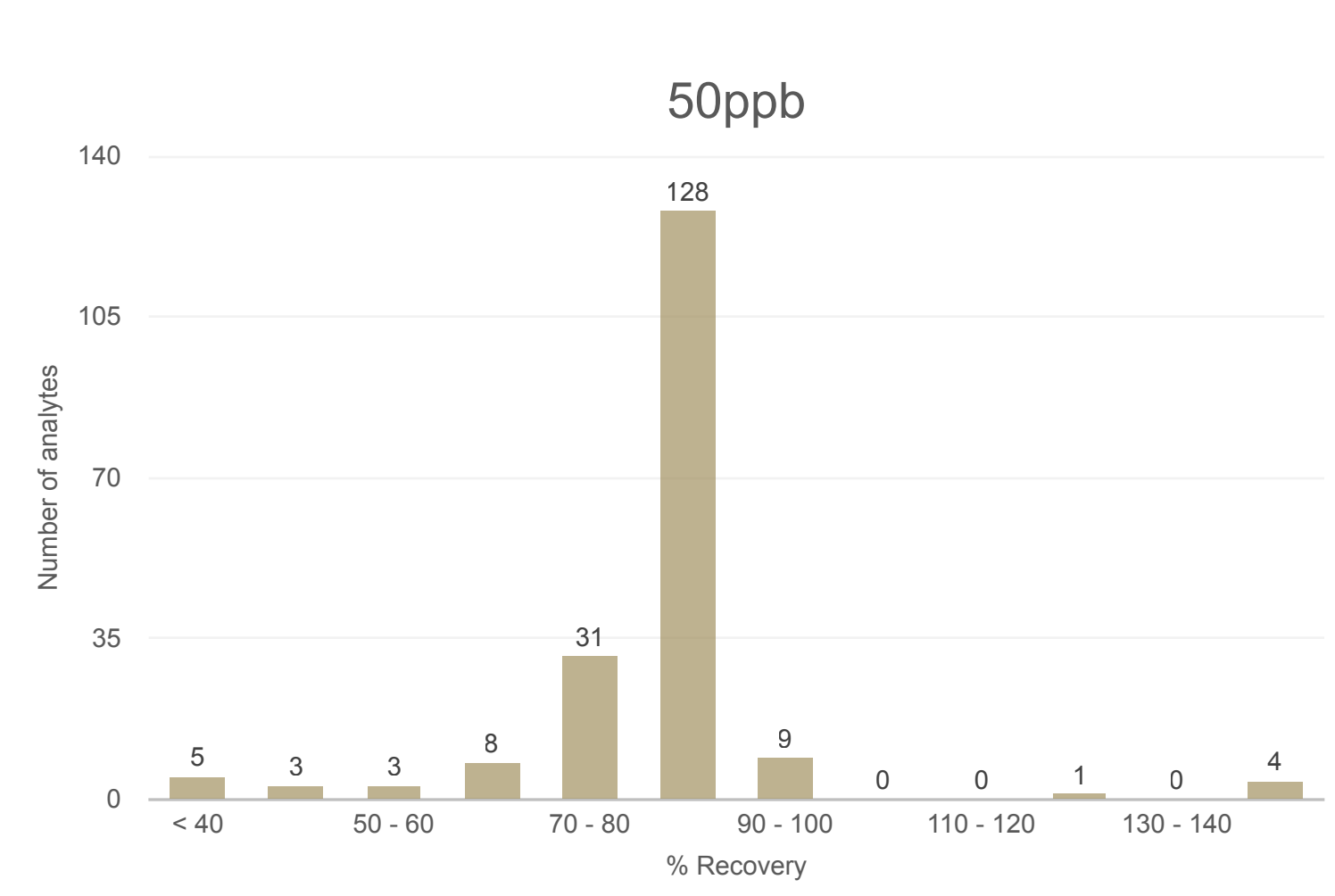
Approx 4.5hrs

Results – tomato GC



192 analytes
160 analytes acceptable recovery

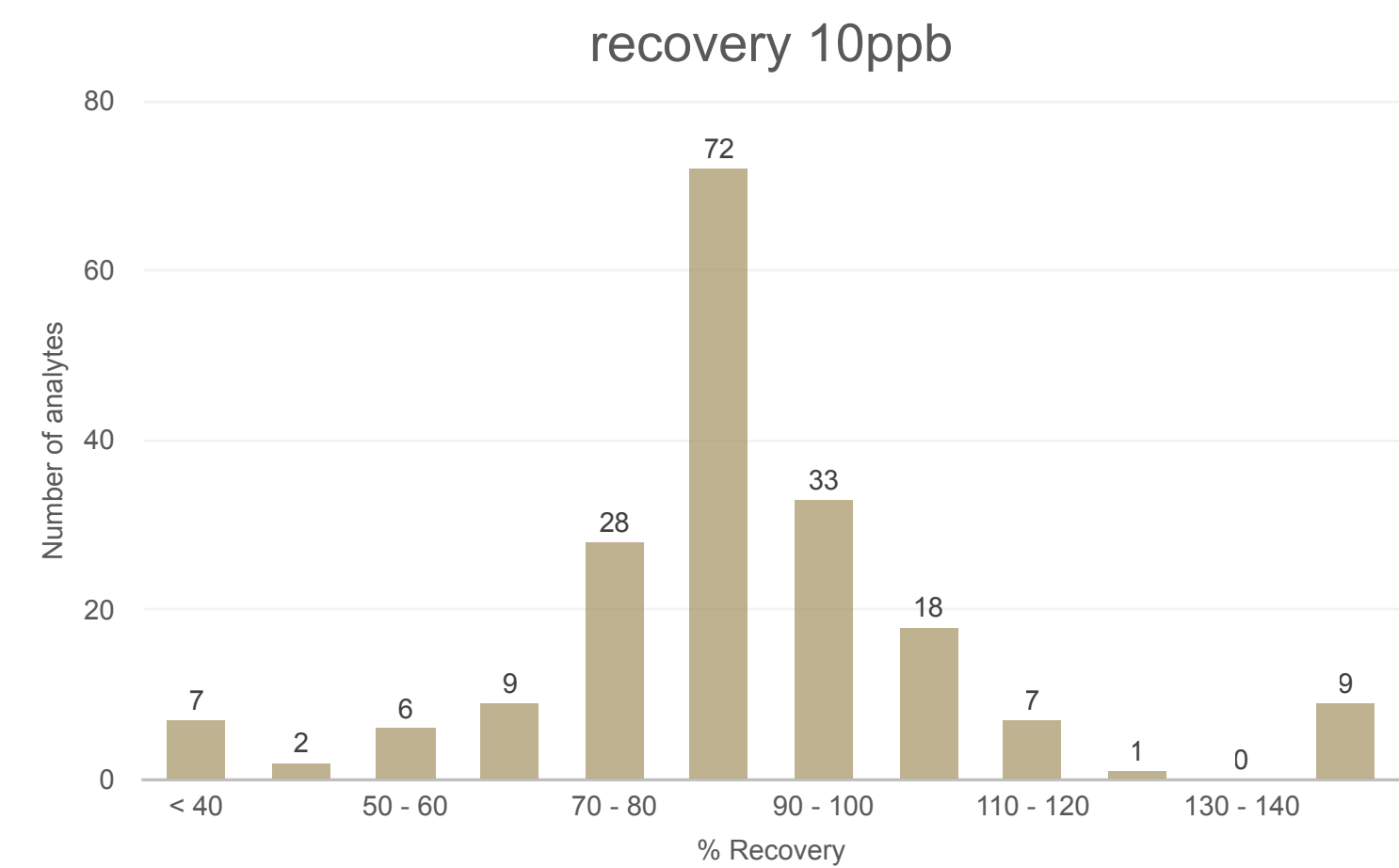
83% acceptable



192 analytes
167 analytes acceptable recovery

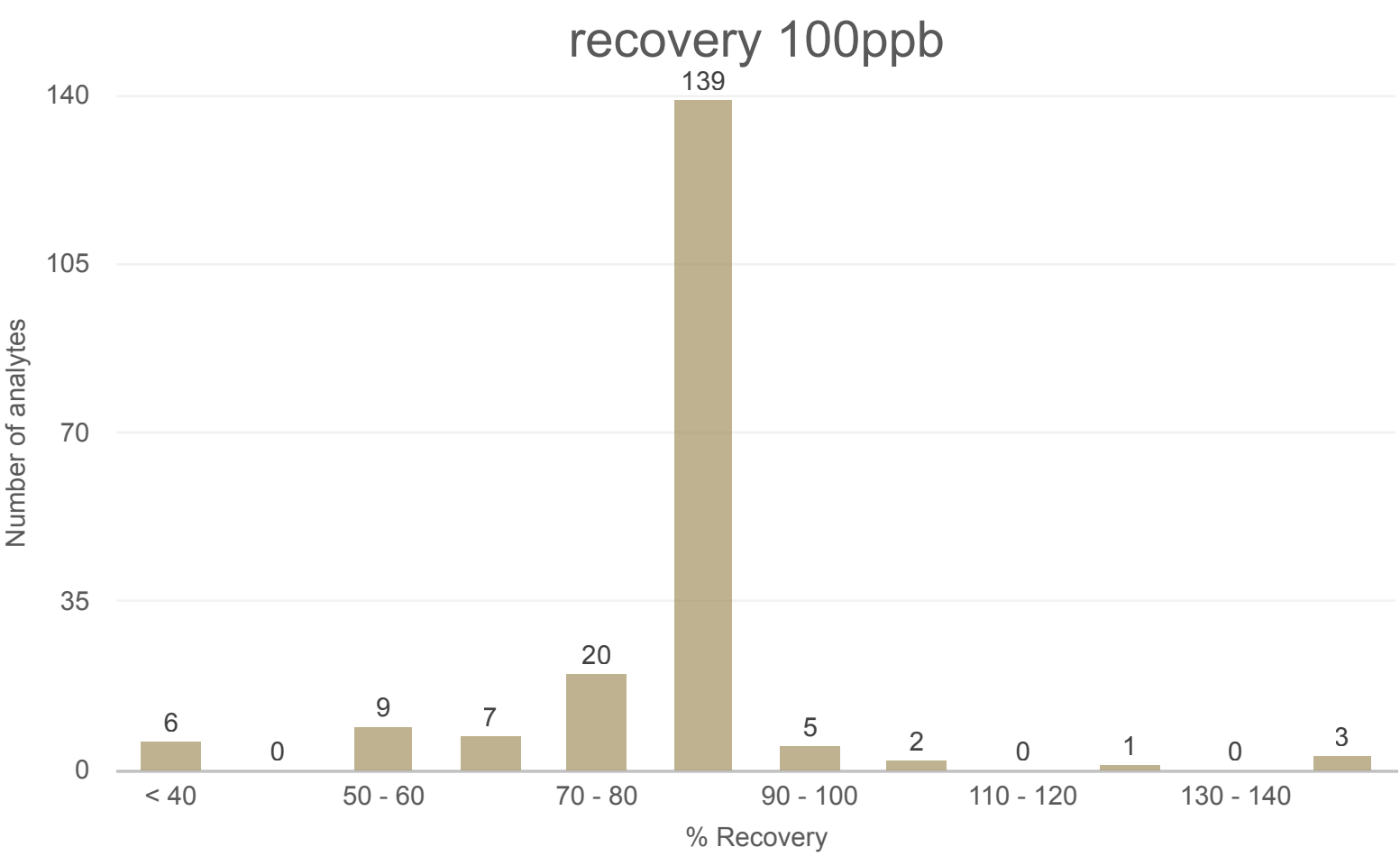
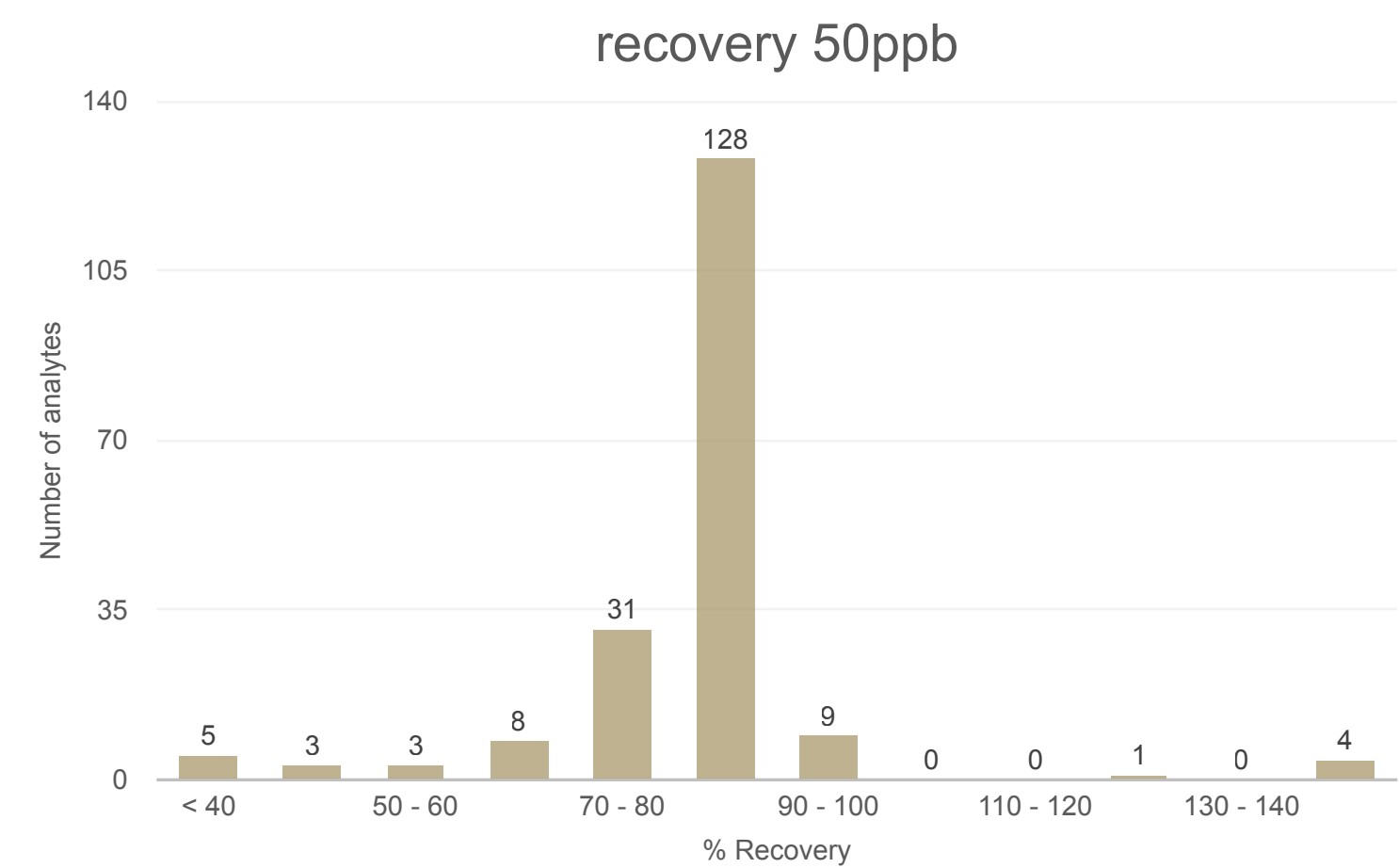
87% acceptable

Results – potato GC



192 analytes
158 analytes acceptable recovery

82% acceptable



192 analytes
166 analytes acceptable recovery

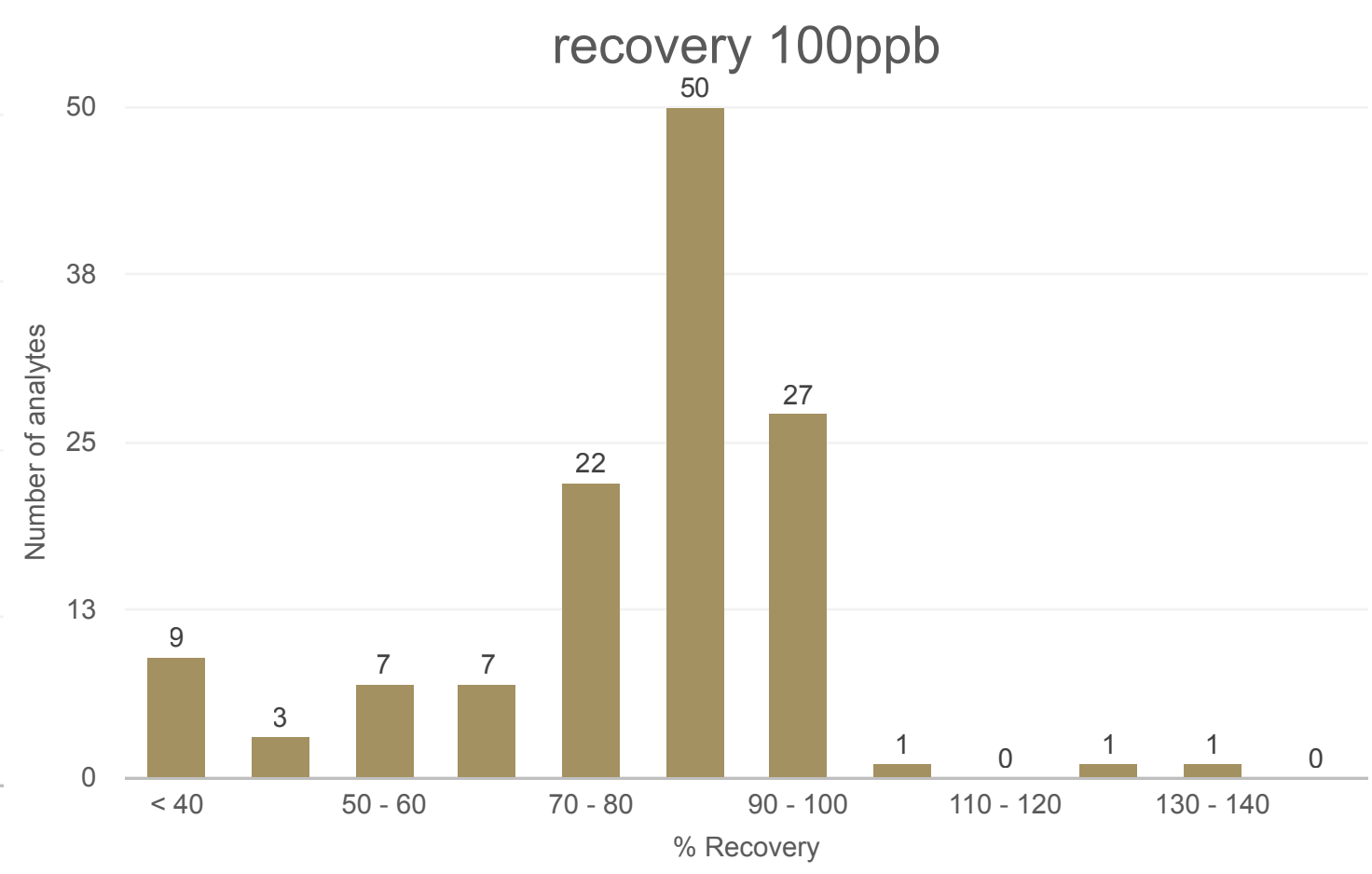
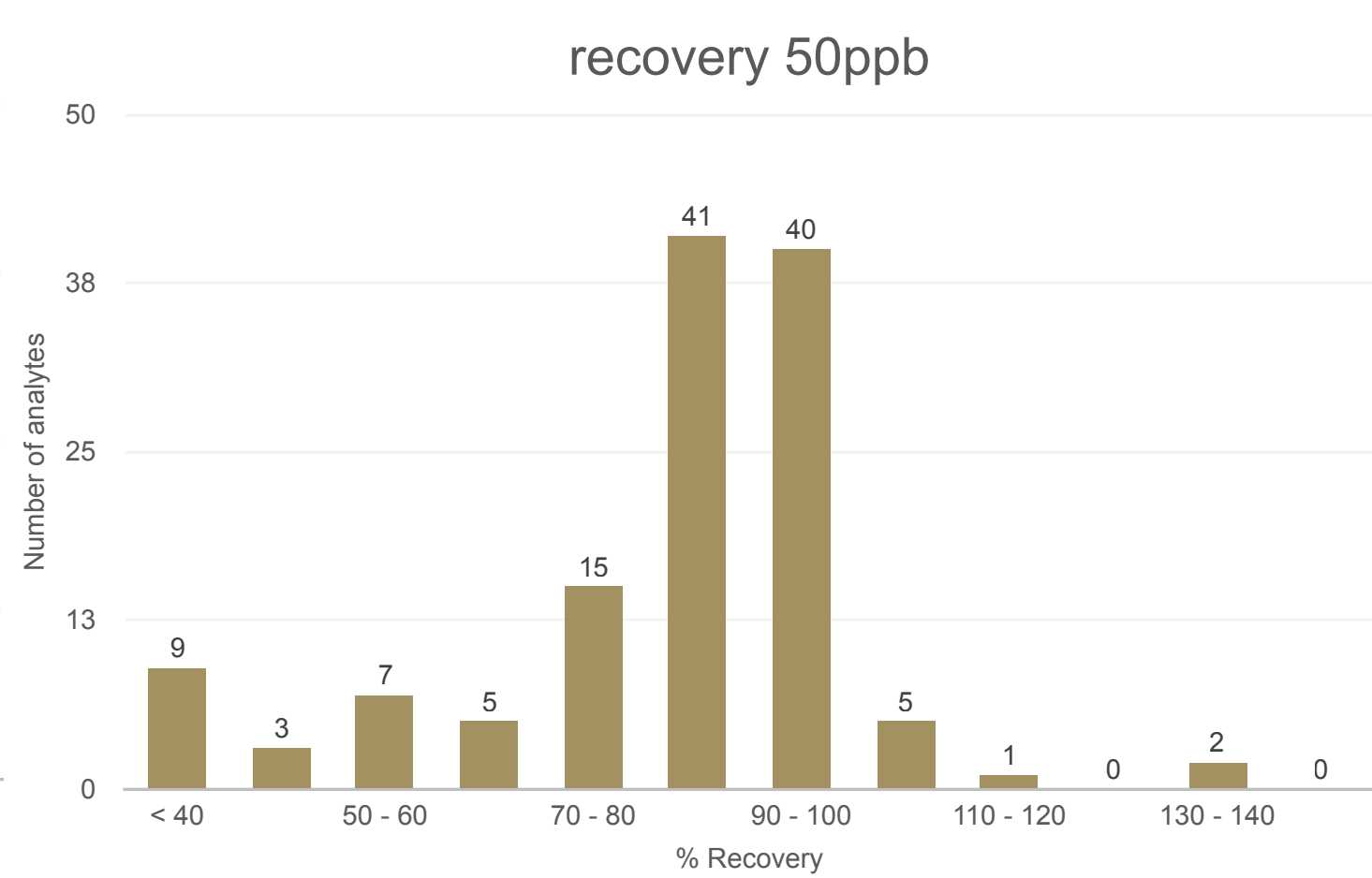
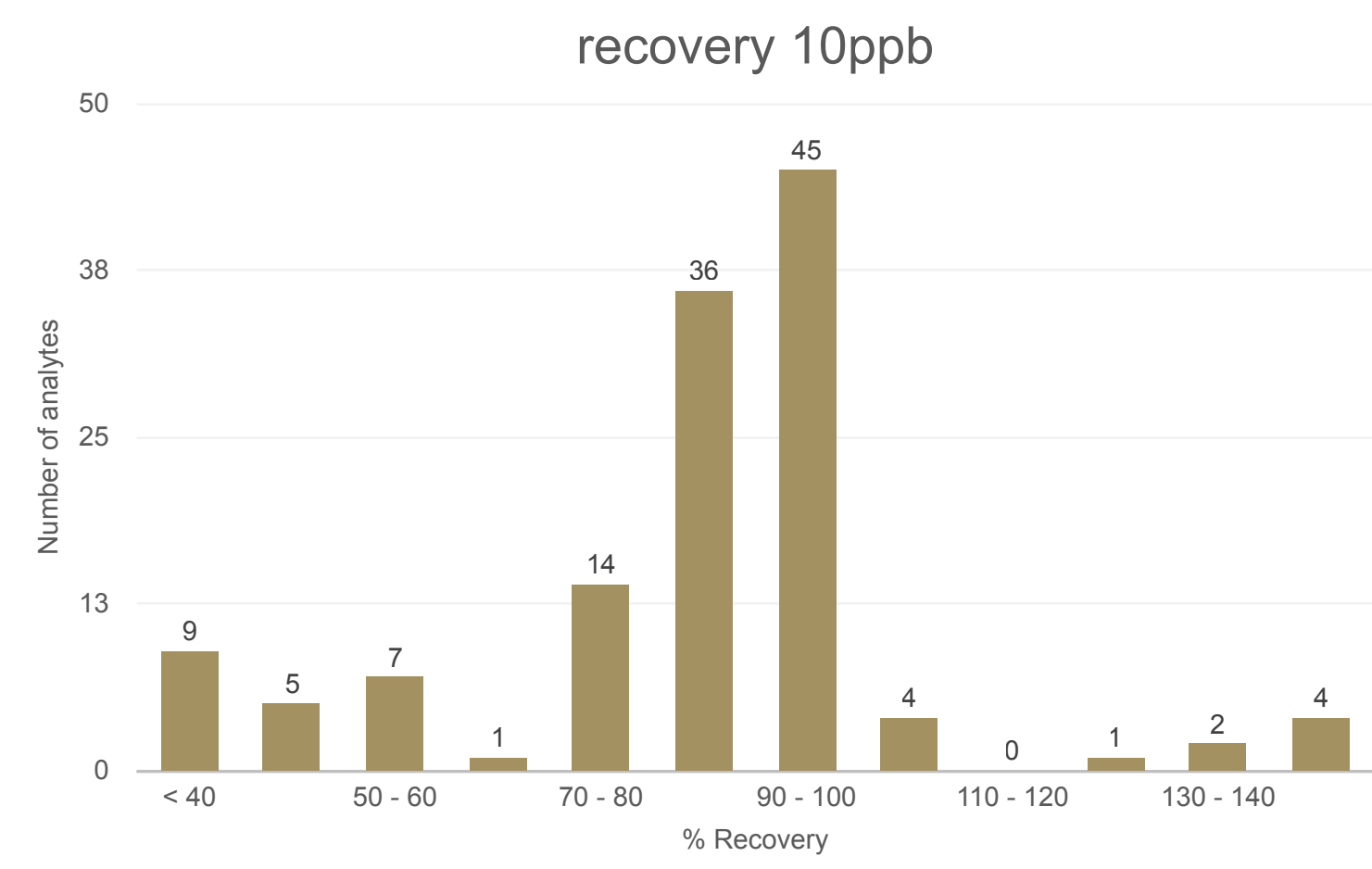
86% acceptable

Problematic compounds



Not currently accredited	Tomato	Potato	Not on GC Scope
1,4-Dimethylnapthalene	1,4 dimethylnapthalene	1,4 dimethylnapthalene	2,4,6-Trichlorophenol
Anthraquinone	Binapacryl	Acephate	3,5-Dichloroaniline
Captan	biphenyl	Aclonifen	3-chloroaniline
Dicofol	Biteranol-II	Aldrin	Methamidophos
Dimoxystrobin	Captan	Anthraquinone	Molinate
Endosulfan-alpha	Carbofuran	Azaconazole	Simazine
Folpet	Chlorothalonil	Biteranol-II	Terbutylazine
Formothion	Dichlobenil	Bromophos-ethyl	
Heptachlor endo-epoxide,trans	Dichlorvos	Chlorbufam	
Isofenphos-oxon	Etridazole	Cyanofenphos I	
Nitrofen	methacrifos	Diazinon	
Oxadixyl	O-phenylphenol	Diphenylamine	
Paraoxon methyl	Phorate	Heptachlor exo epoxide	
PCB28	Propham	hexachlorobenzene	
PCB52	Tecnazene	Iprovalicarb II	
PCB101	Triadimenol	Omethoate	
PCB118		Procymidone	
PCB138		Propachlor	
PCB153			
PCB180			
Pentachloroaniline			
Phorate			
Pirimicarb desmethyl			
Resmethrin			
Silthiofam			
Tefluthrin			

Results – Avocado LC



128 analytes
99 analytes acceptable recovery

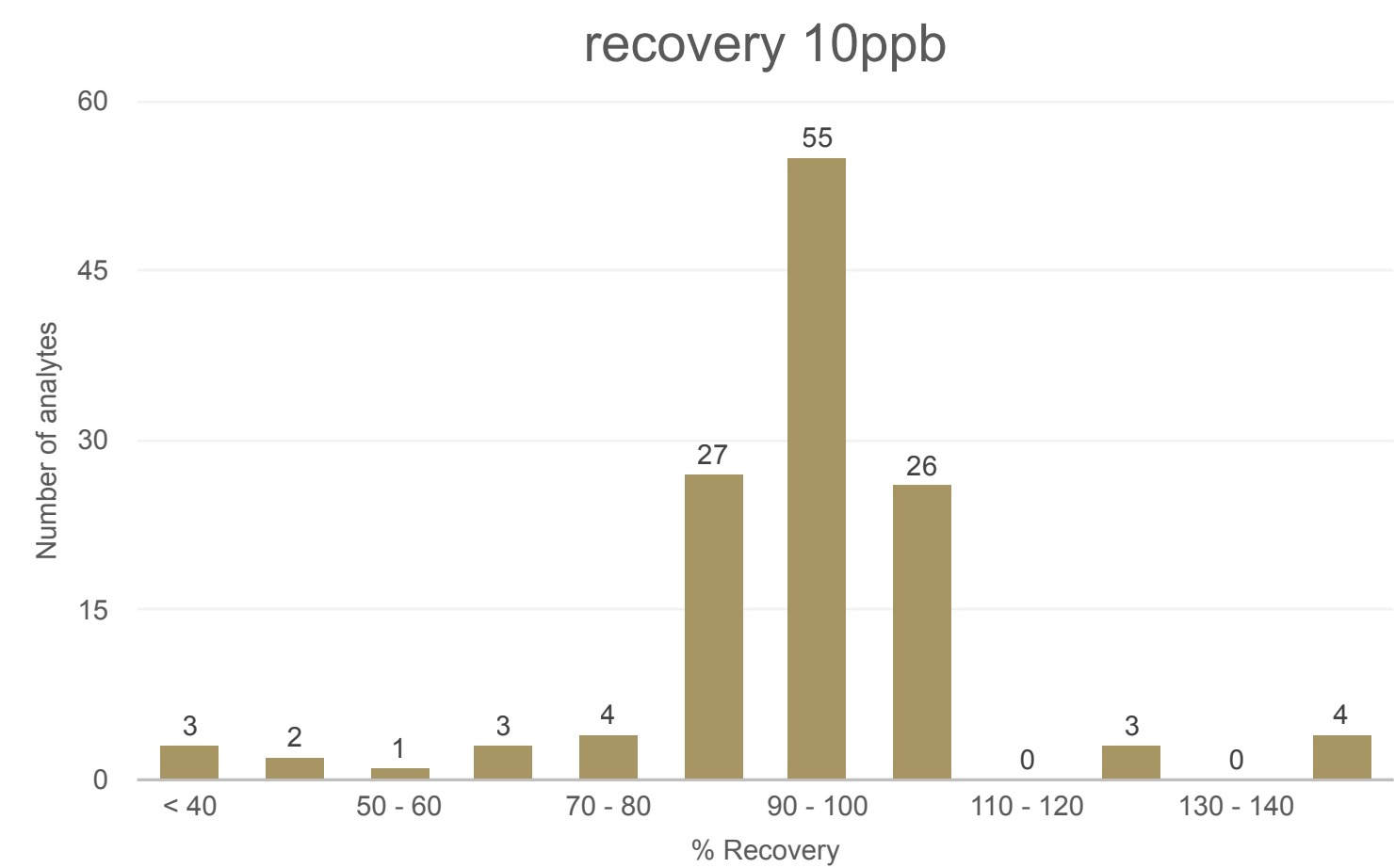
77% acceptable



128 analytes
100 analytes acceptable recovery

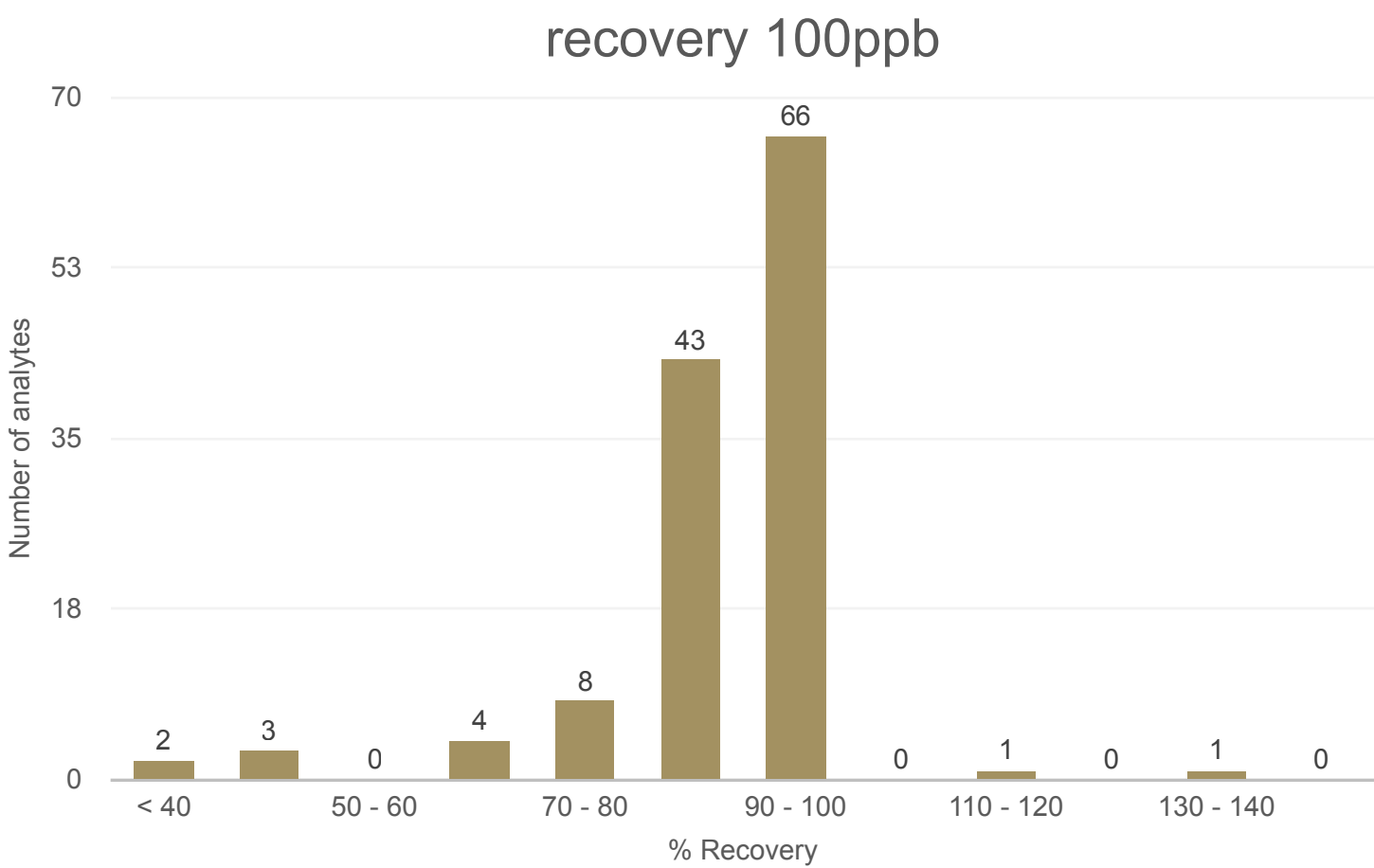
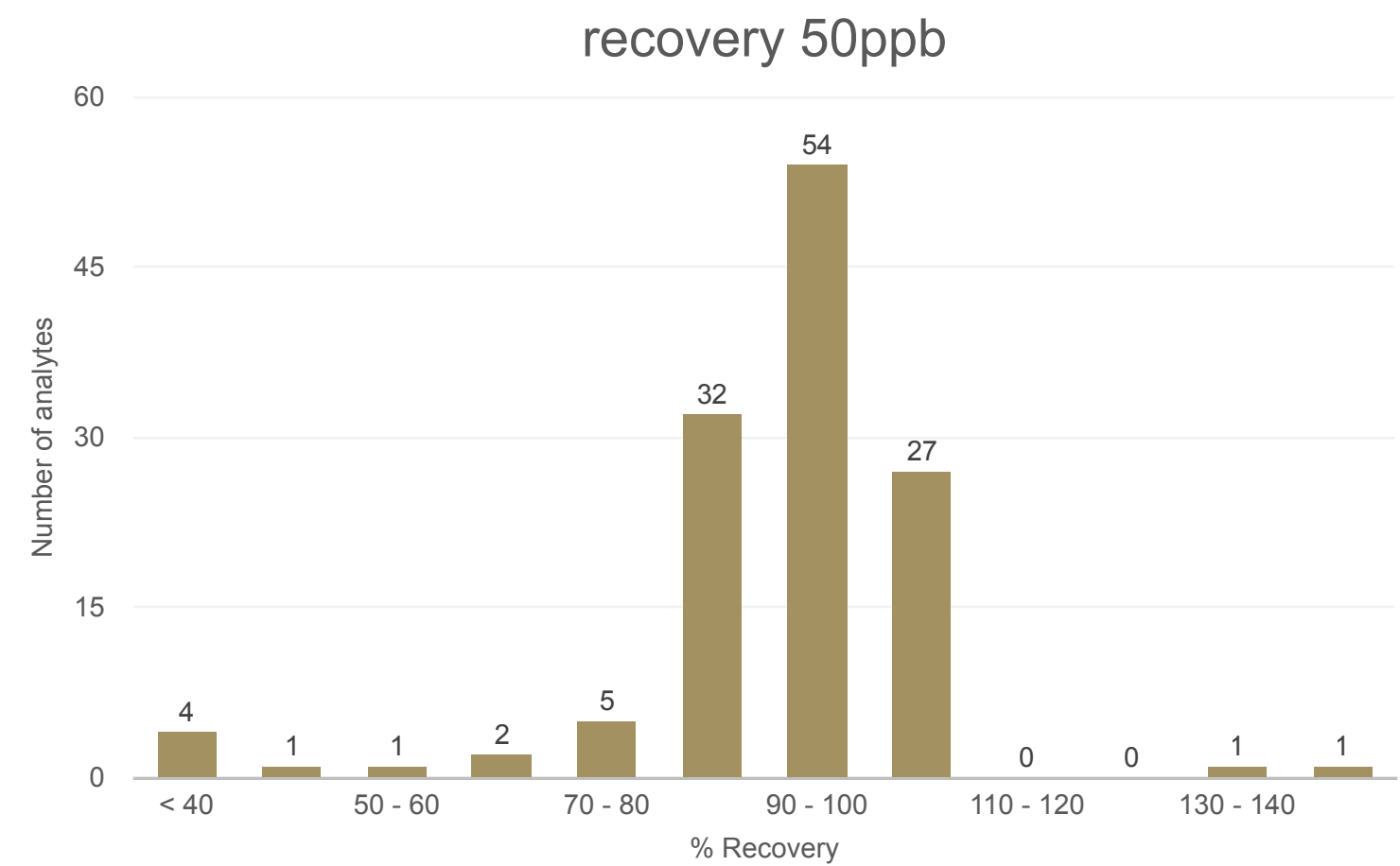
78% acceptable

Results – Tomato LC



128 analytes
112 analytes acceptable recovery

88% acceptable



128 analytes
118 analytes acceptable recovery

92% acceptable



Success!



>79% of assessed analytes passed
recovery criteria

Potential Issues



Homogenisation – current protocol proves difficult to obtain small enough particles to have representative sample





Next steps?

Next steps



1. Purchase Gerstel Dual head
Robotic Robotic Pro
Multipurpose sample (MPS)

3. Validation and Accreditation
of Fruit and Vegetables
method



Thank you for your attention!



Thank you

Da Vinci Laboratory Solutions UK and Ireland Ltd.

Colin Hastie

Jim Garvey

Sadbh Healy



Any questions?