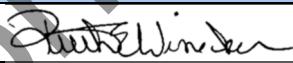
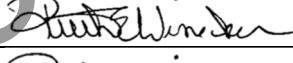
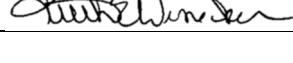


# SOP 053 - Sequence Setup – Thermo LCQuan

## Table of Contents

1.	Principle .....	3
2.	Specimens .....	3
3.	Reagents and Materials .....	3
4.	Instrumentation and Equipment.....	3
5.	Procedure .....	3
6.	References.....	10

# SOP 053 - Sequence Setup – Thermo LCQuan

SOP Name:	SOP #:	
LCQuan Sequence Setup	053	
	Revision:	Revision Date/Initials:
North Carolina Office of the Chief Medical Examiner Toxicology Laboratory		
Approving Authority Name	Approving Authority Signature	Approval Date
Ruth E. Winecker, Ph.D.		04/08/2015
Ruth E. Winecker, Ph.D.		06/10/2016
Ruth E. Winecker, Ph.D.		08/29/2017

# **SOP 053 - Sequence Setup – Thermo LCQuan**

## **1. Principle**

1.1. This method is designed to allow the user to create a sequence and acquire LC/MS/MS data using Thermo LCQuan software.

## **2. Specimens**

2.1. N/A

## **3. Reagents and Materials**

3.1. N/A

## **4. Instrumentation and Equipment**

4.1. Thermo TSQ LC/MS/MS

4.2. LCQuan software

4.3. Data reporting system (PC)

## **5. Procedure**

### **5.1. Create Sequence**

5.1.1. On a networked PC with Thermo LCQuan software installed, open the LCQuan software by double-clicking the ICON.

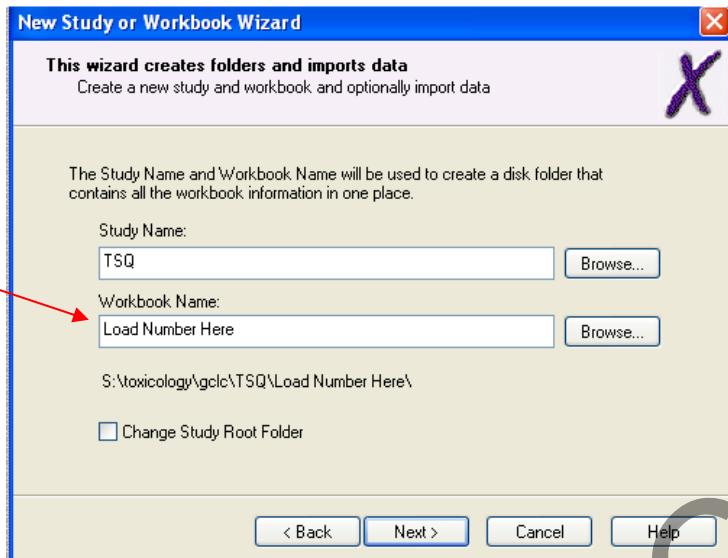


5.1.1.1.

5.1.2. On the welcome screen, click on “Create a new workbook” and click “Next”.

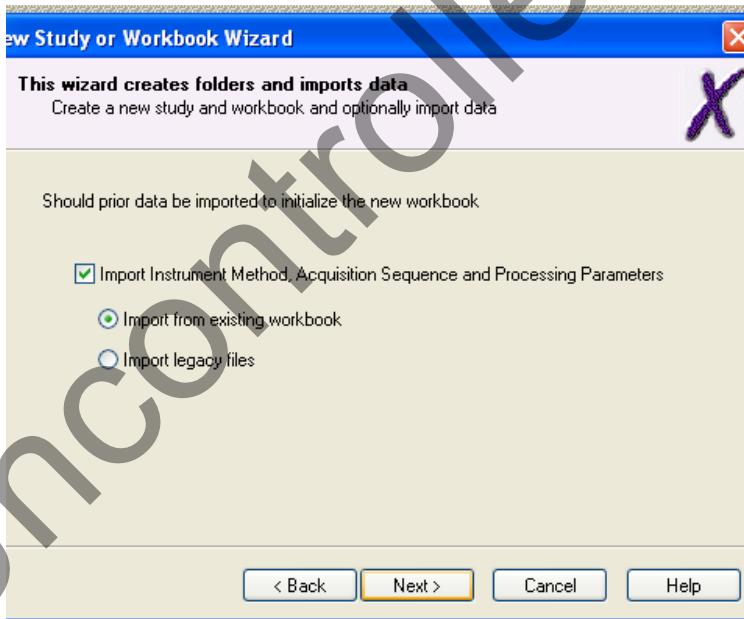
5.1.3. “TSQ” should appear, in the “Study Name” field. In the “Workbook Name” field, enter the load number for which a sequence is to be created, click “Next” (see below).

## SOP 053 - Sequence Setup – Thermo LCQuan



5.1.4. Click “Next” in the following window leaving the box next to “import raw files” un-checked.

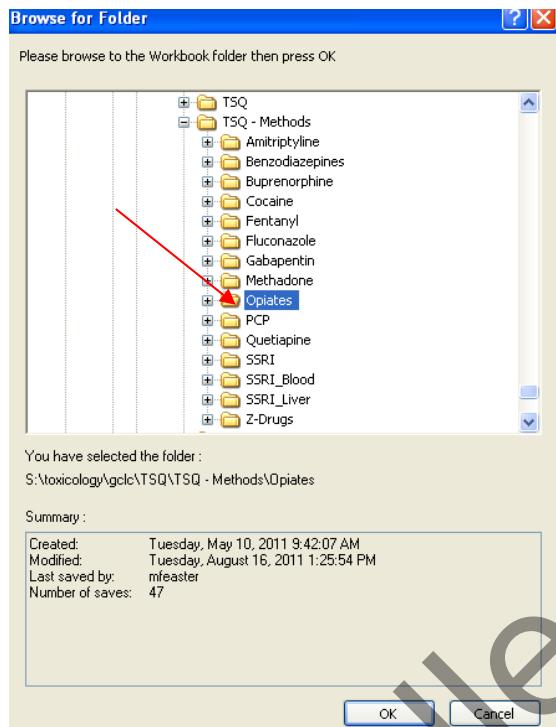
5.1.5. The following window should appear as below, click “Next”.



5.1.6. In the following window, select “Browse” and choose the appropriate method folder corresponding to the assay being performed (Opiates used as example):

## SOP 053 - Sequence Setup – Thermo LCQuan

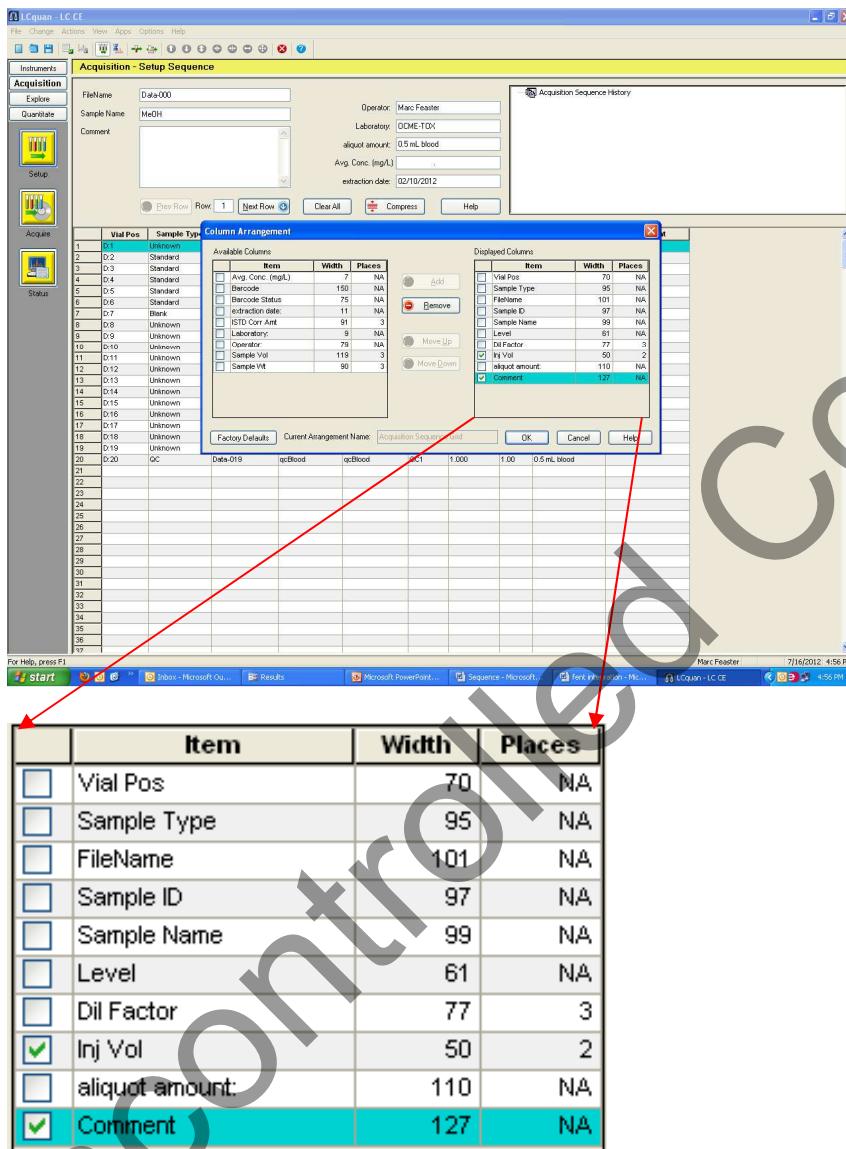
- 5.1.6.1. S:\toxicology\Instrument Methods\LC\Instrument Name). Click “OK”, then click “Next.”



- 5.1.7. The workbook template should now be open, click on the “Acquisition” then “Setup” tabs in the upper left side of the screen.
- 5.1.8. Right click anywhere in the sequence table and choose “Columns” from the dropdown menu. Click to check the boxes of columns to be removed and click “Remove” then “OK” (see below).
- 5.1.8.1. Columns to remain displayed (In the following order):
- 5.1.8.1.1. Vial Pos
  - 5.1.8.1.2. Sample Type
  - 5.1.8.1.3. FileName
  - 5.1.8.1.4. Sample ID
  - 5.1.8.1.5. Sample Name
  - 5.1.8.1.6. Level
  - 5.1.8.1.7. Dil Factor

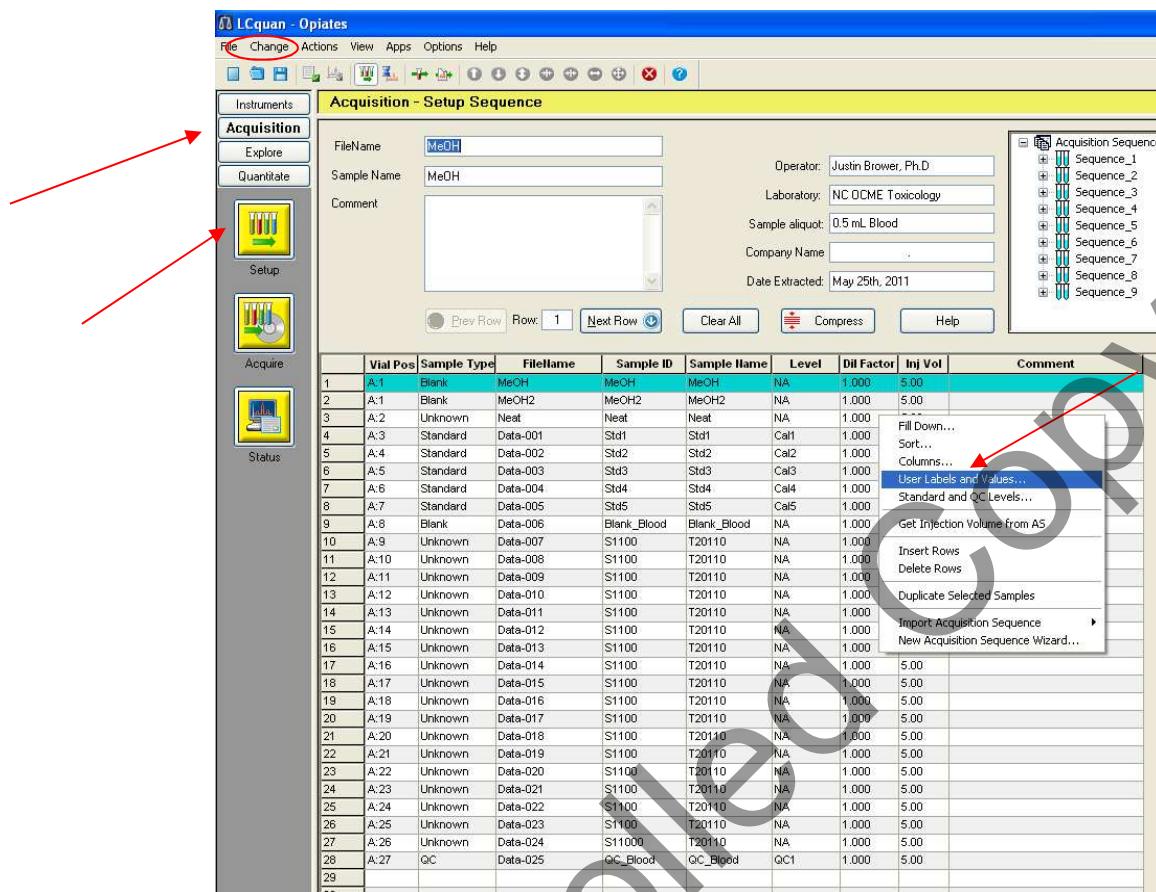
# SOP 053 - Sequence Setup – Thermo LCQuan

## 5.1.8.1.8. Aliquot amount



- 5.1.9. Right click anywhere in the sequence table and choose “User values and labels” from the dropdown menu. Alternatively, click “Change” - “User values and labels”. (see below)

# SOP 053 - Sequence Setup – Thermo LCQuan



5.1.10. In the “User Value 1” and “User Value 5” fields, enter operator name and extraction date respectively. Click “OK”.

5.1.11. In the “User Value 3” field enter default aliquot amount/SOP#/Instrument name. E.G. **0.5mL blood/LC208/TSQ01**

Operator Name  
Extraction Date



# SOP 053 - Sequence Setup – Thermo LCQuan

- 5.1.12. Enter S#'s (Sample ID column) and T#'s (Sample Name Column) as appropriate. Enter dilution factors as needed (Dil Factor column) and enter aliquot amount for each row in the “Sample Aliquot” field at top center of screen. If needed, right click on the row numbers for the option to add or delete rows.

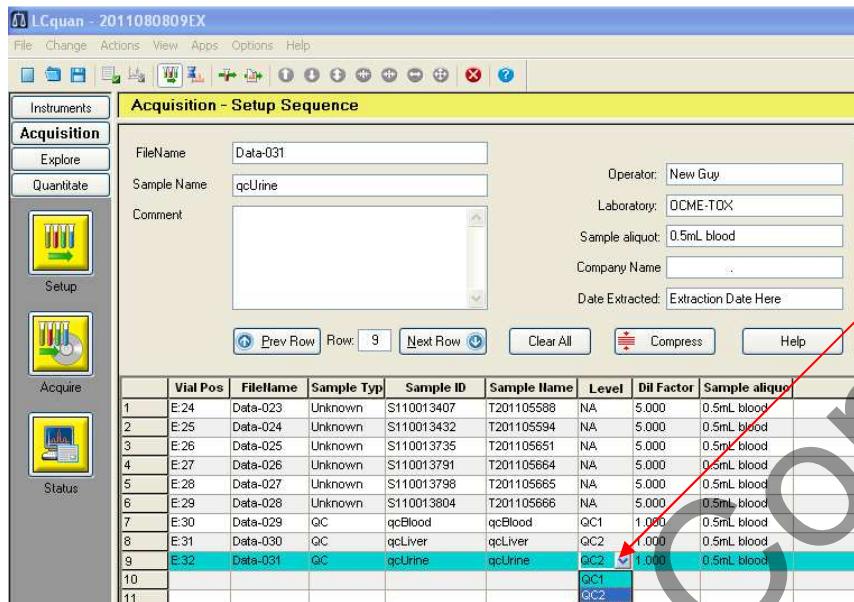
5.1.12.1. Note: Enter “N/A” in “Sample Aliquot” field for MeOH blanks and neat.

- 5.1.13. Shrink each column width to remove any extra spacing. This will allow for the sequence to print on one page.

	Vial Pos	Sample Typ	Filename	Sample ID	Sample Name	Level	Dil Factor	aliquot amount
1	B:1	Unknown	meoh	meoh	meoh	NA	1.000	
2	B:2	Standard	Data-001	Cal1	Cal1	Cal01	1.000	0.1 mL blood
3	B:3	Standard	Data-002	Cal2	Cal2	Cal02	1.000	0.1 mL blood
4	B:4	Standard	Data-003	Cal3	Cal3	Cal03	1.000	0.1 mL blood
5	B:5	Standard	Data-004	Cal4	Cal4	Cal04	1.000	0.1 mL blood
6	B:6	Standard	Data-005	Cal5	Cal5	Cal05	1.000	0.1 mL blood
7	B:7	Blank	Data-006	blankBlood	blankBlood	NA	1.000	0.1 mL blood
8	B:8	Blank	Data-007	blankLiver	blankLiver	NA	1.000	0.1 g liver
9	B:9	Unknown	Data-008	S120011637	T201204482	NA	1.000	0.1 mL blood
10	B:10	Unknown	Data-009	S120011639	T201204482	NA	4.000	0.1 g liver
11	B:11	Unknown	Data-010	S120011989	T201204613	NA	1.000	0.1 mL blood
12	B:12	Unknown	Data-011	S120012033	T201201625	NA	1.000	0.1 g blood
13	B:13	Unknown	Data-012	S120012513	T201204819	NA	1.000	0.1 mL blood
14	B:14	Unknown	Data-013	S120012515	T201204819	NA	4.000	0.1 g liver
15	B:15	Unknown	Data-014	S120012586	T201204818	NA	1.000	0.1 mL blood
16	B:16	Unknown	Data-015	S120012589	T201204610	NA	4.000	0.1 g liver
17	B:17	Unknown	Data-016	S120011637	T201204482	NA	1.000	0.1 mL blood
18	B:18	Unknown	Data-017	S120011639	T201204482	NA	4.000	0.1 g liver
19	B:19	Unknown	Data-018	S120011989	T201204613	NA	1.000	0.1 mL blood
20	B:20	Unknown	Data-019	S120012033	T201201625	NA	1.000	0.1 g blood
21	B:21	Unknown	Data-020	S120012513	T201204819	NA	1.000	0.1 mL blood
22	B:22	Unknown	Data-021	S120012515	T201204819	NA	4.000	0.1 g liver
23	B:23	Unknown	Data-022	S120012586	T201204818	NA	1.000	0.1 mL blood
24	B:24	Unknown	Data-023	S120012589	T201204818	NA	4.000	0.1 g liver
25	B:25	QC	Data-024	qcBlood	QC_Blood	QC01	1.000	0.1 mL blood
26	B:26	QC	Data-025	qcLiver	QC_Liver	QC02	1.000	0.1 g liver

- 5.1.14. If there are multiple QC levels for the assay, click in the “Level” column(s) that correspond to the QC's of interest. From the dropdown menu, select the appropriate QC level.

# SOP 053 - Sequence Setup – Thermo LCQuan

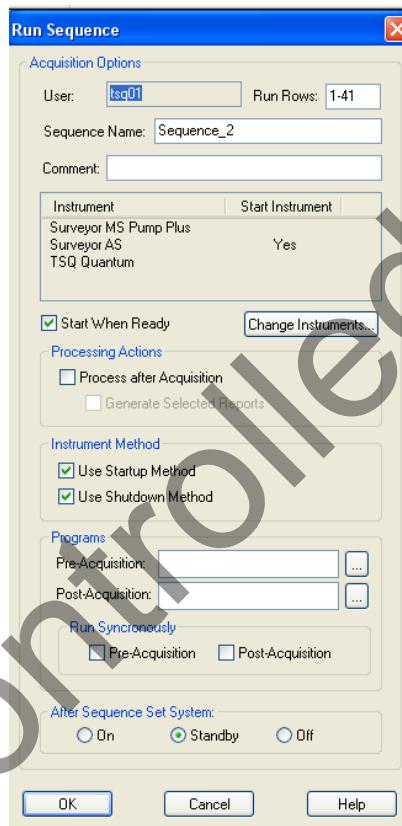


- 5.1.15. Click the “Save” icon to save workbook. If the PC being used is not “attached” to the TSQ, close LCQuan and complete the following steps on the TSQ PC, otherwise skip to step 5.1.17.
- 5.1.16. Open LCQuan and choose “Open an Existing Workbook”. Choose the folder with the corresponding load number and click “OK”.
- 5.1.17. Determine which autosampler rack slot is to be used (A through E) and enter in the first row/Vial Pos column of the sequence table. (If the sequence starts with two MeOH blanks to be sampled from the same vial, enter “rack slot”:vial position” in the first two rows. E.g. row1 - C:1; row2 - C:1.)
- 5.1.18. To fill down the remaining vial locations, left click once and hold on the final sample location using slot1, and drag down to the last sequence entry. Right click on highlighted column and choose “Fill Down”. Click “OK” on the Fill Down dialog box.
- 5.1.19. Repeat step 5.1.17 for the “File Name” column starting with Data-001.
- 5.1.20. Save the workbook (Step 5.1.15).
- 5.1.21. Click “File” – “Print Sequence Info...” to print sequence.
- 5.1.22. Verify sequence vial locations with another analyst, obtaining their initials and date on the sequence sheet, and place the autosampler rack in the appropriate slot in the TSQ autosampler.

## SOP 053 - Sequence Setup – Thermo LCQuan

- 5.1.23. Check the fluid levels of the LC solvents, making sure there is enough to inject the entire load – refill if necessary.
- 5.1.24. Click the “Acquire” button on the left side of the screen. In the dialog box, verify that the appropriate number of rows is displayed in the “Run Rows” field in the upper right. Click “OK” to start acquiring data. It should appear as below.

5.1.24.1. **Note: For TSQ02 & TSQ03, do not check the boxes next to “Use Startup Method” and “Use Shutdown Method”.**



5.1.24.1.1.

## 6. References

- 6.1. Thermo Scientific. *Xcalibur LCquan Quantitative Analysis User Guide*. Aug. 2007. XCALI-97166 Revision D. USA.