

Seal Analytical AA SOP for Phosphorus Analysis

1.0 Reagent preparation

1.1 Ammonium molybdate solution

Ammonium molybdate	1.8 g	0.9 g	0.45 g
Sulfuric acid	22.3 mL	11.15 mL	5.575 mL
Antimony potassium tartrate	0.05 g	0.025 g	0.0125 g
Sodium dodecyl sulfate	2 g	1 g	0.5 g
DI water	1000 mL	500 mL	250 mL

1.2 Ascorbic acid solution

Ascorbic acid	15 g	7.5 g	3.75 g
DI water	1000 mL	500 mL	250 mL

1.3 SDS solution

Sodium dodecyl sulfate	2 g	1 g	0.5 g
DI water	1000 mL	500 mL	250 mL

1.4 Standards, blanks and check samples

- Prepare standards in the 0.01 – 6 ppm range using KH_2PO_4 . There is room for 10 standards.
- Prepare a blank using just MilliQ water. Run the blank as the first standard, as well as a sample.
- Prepare a check sample using H_3PO_4 CRM.

2.0 Starting the machine

- Turn black power bar at the back of the instrument on.
- Remove dust covers.
- Inspect tubing for signs of damage or dirt. Replace any tubing if needed.
- Place each pump tube in its respective solution. See marks on the sampling lines, as well as Figure 1 in the figures section.
- Tighten the tubes, and make sure they are parallel to one another.
- Attach the platen.
- Turn on the computer (and fill in the log book).
- Open the AACE6.10 software.
- Start charting. (Previous run is ok for now).
- Start the pump by hitting the red button. Let solutions flow through the system for about 10 mins. The setting can be switched to 'fast' during this time. The spectrometer will also warm up during this time.
- Place standards in their racks, preferably from lowest to highest. 901 will be blank.
- Place samples in the sample rack.

3.0 Setting up a new run

While the machine is warming up, you can setup a new run on the software.

- Go to Set Up > Analysis/Run
- Click 'New Analysis'. Choose System 1.
- Fill the Main Page tab. See Figure 2 in the figure section for details. (Method 2 can be deleted.)
- Fill in Channel 1 tab. See Figure 3 in the figure section for details. Note: start standards from 0 ppm, not 50 ppb.
- Set up the Tray Protocol tab. See Figure 4 in the figures section for details.
You can also load a tray from previous run and modify it.
- Save the tray after setting it up.
- Close the screen. Say 'yes' to "Do you want to create a run?"
- Close the old charting screen. Click on 'Charting' again. Make sure the method you saved is the one that is currently active.
- Close Ch.2-1 -.
- Switch pump to 'normal' pace.
- Right click in the chart area and 'Set light power'. This will take a few seconds.
- Right click and 'set base'. You should just be running water as a sample, and reagents during all this.
- Wait for the baseline to become stable. The graph should be a stable continuous line.
- Double click on 'XY2 Sampler 1'. Make the sampling probe go into the highest standard.
After 1 minute, it can be switched back to wash.
- Wait for the highest standard peak to show up. It'll take about 12-15 mins to show up. Don't miss it!
- When you see the peak, right click and 'Set Gain'.
- Now you can hit 'run' and start your analysis. You may start a new charting session if you like.

4.0 Clean-up

- Put all lines in the water and let run for 5 mins on 'fast' speed.
- Inject 3% H₂SO₄ as a sample for 5 mins.
- Then put the probe back in the wash line for 5 mins.
- Turn off the switch on the power bar.
- Take off the platen.
- Loosen the sampling lines
- Put the dust covers on.
- Put all reagents away.

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5.0 Figures

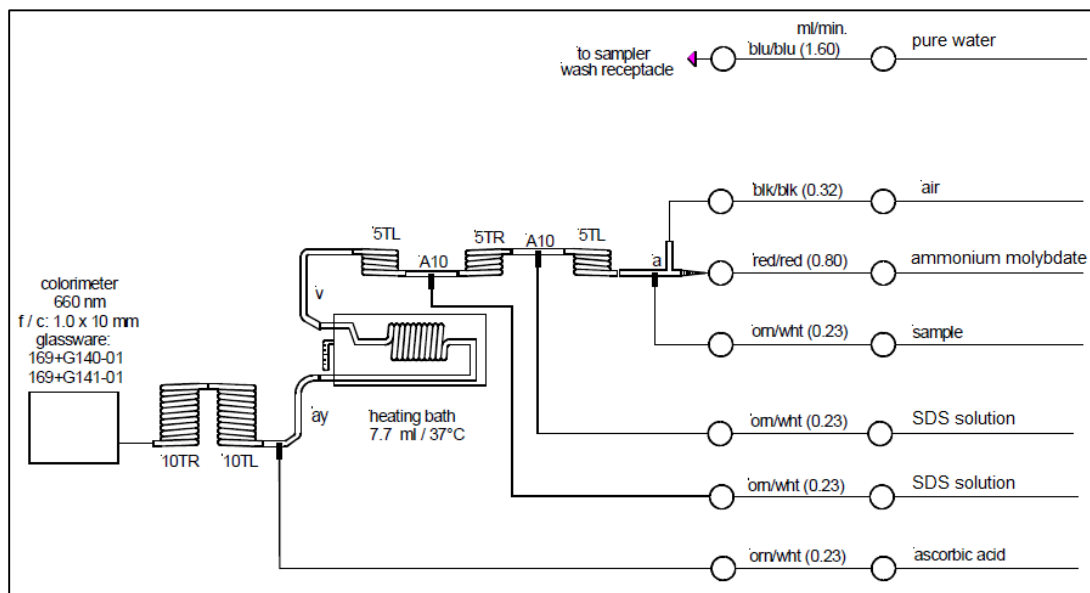


Figure 1. Schematic of the AA instrumentation. Taken and modified from Method No. G-103-93 Rev. 10, page 11.

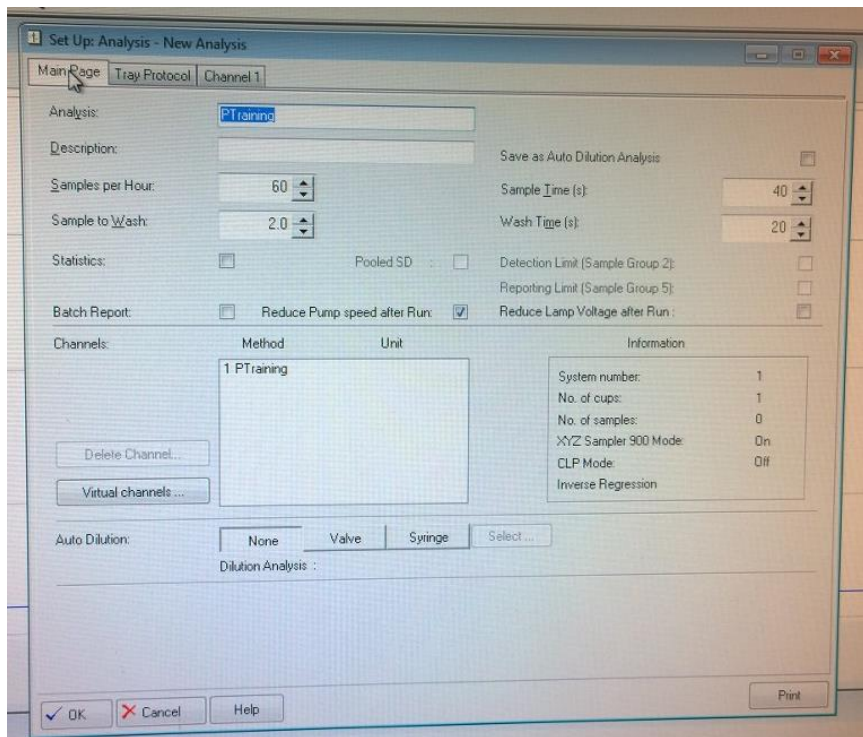


Figure 2. Method Page set-up.

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Set Up: Analysis - New Analysis

Main Page Tray Protocol Channel 1

Method: PTraining Units: µg/L

Baseline Correction: ☒ Calibration Fit: L Q C P Z S

Drift Correction: ☒ Baseline in Calibration: ☐

Carryover Correction: ☒ Baseline Concentration: 0.000

Carryover (%): 0.0 Inverse Chemistry: ☐

Resample: 0 1 2 Autobase reference (%): 5

Use Auto Dilution: ☐ Run Delay Time (min): 0.0

Sample Limits

Set ☐ Low: 0.000 High: 0.000

Peak Window

Start: 50 End: 105 % of sample time: Default

Filter

Smoothing (0 = Off): 0 Spike Recovery Standard conc.: 0.000

Calibrants

1: 50.000

2: 250.000

3: 500.000

4: 750.000

5: 1,000.000

6: 0.000

7: 0.000

8: 0.000

9: 0.000

10: 0.000

Decimals: 3

Quality Control >> Performance Parameters >> Temperature >> Peak Parameters >>

OK Cancel Help Undo Print

Figure 3. Channel 1 set-up.

Set Up: Analysis - New Analysis

Main Page Tray Protocol Channel 1

Peak	Icon	Type	Cup	Sample ID
1	P	P	907	Primer
2	D	D	907	Drift
3	H1	H1	907	High
4	L1	L1	901	Low
5	L1	L1	901	Low
6	N	N	0	Null
7	C	C	901	0
8	C	C	902	5ppb
9	C	C	903	10ppb
10	C	C	904	25ppb
11	C	C	905	50ppb
12	C	C	906	100ppb
13	C	C	907	500ppb
14	S	S	1	Blank Water
15	S	S	6	100ppb
16	S	S	7	50ppb Alex
17	S	S	8	150ppb Vito
18	S	S	9	Blank Water
19	D	D	907	Drift
20	E	E	0	End
21	S	S	10	
22	S	S	11	
23	S	S	12	
24	S	S	13	
25	S	S	14	

OK Cancel Help Load Tray

Figure 4. Tray Protocol set-up.