EDAX EBSD

EBSD - Start Up

- 1. Open the 'IR' camera program to view specimen movement. (F4)
- 2. Tilt specimen holder to a total of 70°

Note: Avoid using carbon tape to secure specimen

- 3. Focus the sample (With SEM software) so that **WD** is at a height of 15mm.
- 4. Open 'TSL OIM Data Collection' program.
- 5. Insert the EBSD 'Camera' (camera pull down menu)

Note: Caution – Make sure the detector does not hit the holder.

Camera Adjustment (Instrument Console)

- 6. Select a 'Binning' image resolution. (4x4 is recommended for most samples)
- 7. Close the IR camera
- 8. From the 'Image Processing' tab select 'Image Processing Mode' (Standard or Enhanced)

Standard

- a. Unclick 'Background Subtraction' (No check mark)
- b. Adjust the 'Gain' and 'Exposure'

Note: 100 fps - EBSD, 30 fps -EDS/EBSD

- c. Press the 'Capture Bkd' button.
- d. From the 'Inage Processing' tab select the 'Background Subtraction' (Red check mark will appear)

Enhanced

- a. Click 'On' (Red check mark will appear Affect will be seen immediately in the live camera window above)
- b. Press 'Modify' to change enhanced image processing mode.

Collecting a Scan

- 9. On main toolbar select the correct WD ('*' will appear if calibrated) from the pull down menu
- 10. Click **'Ext XY'** icon to get SEM control back.
- 11. Select the imaging detector (SED or FSD)
- 12. Go to the 'Phase page' tab
- 13. Press the 'Load' button to select the phase(s) of interest.
- 14. Go to the 'Interactive page' tab
- 15. Press the 'Capture' button to see EM image.

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- 16. Click on EM image (green X will appear)
- 17. Press 'Index' button and verify quality of diffraction pattern. (Ideal: CI = 1.0, fit = 0.0, Minimum: CI = 0.1)
- 18. Go to the 'Scan" tab.
- 19. Draw a Box (scan properties will pop-up)
- 20. Set all properties (File directory, scan dimensions, step size etc.)

Note: EDS/Chemistry is set up from the chemistry tab in the 'Scan Properties'

21. Start Scan

EBSD - Shut Down

- 22. Retract the EBSD 'Camera' (camera pull down menu)
- 23. Click 'Exit' from the file pull down menu to close program.

Analysis (Clean Up)

- 24. From the desk top click on 'TSL OIM Analysis' ion
- 25. Open your file
- 26. Select 'Clean Up' from the right click on the mouse
- 27. Check off 'Grain CI Standardization' and press 'OK'
- 28. Select 'Clean Up' from the right click on the mouse
- 29. Check off 'Grain Dilation' with single iteration and press 'OK'
- 30. Select 'Clean Up' from the right click on the mouse.
- 31. Check off 'Neighbor Phase Correlation' and press 'OK'. (If more than one phase are selected)

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