Alignment and Data Collection for GISAXS Users

- 1) [optional] Make the name of the spec datafile appropriate for your sample: Spec> newfile filename
- 2) Spec> go_apd
- 3) Change the sample(s), search and interlock the hutch
- 4) Open the photon shutter
- 5) Drop the sample out of the way: Spec> umvr samz -0.2
- 6) Check that the APD is centered on the direct beam. If you see some counts on the APD (>1000), run the macro: Spec> pind3zlup. If not, tweak tth until you see > 1000 counts on the APD, then run the macro: Spec> pind3zlup.
- 7) Adjust the sample height until the sample cuts the beam20-80%: Spec> umvr samz [increment in mm]
- 8) Spec> zcen
- 9) (Optional) if the last sample was on a different substrate: Spec> thcen_coarse
- 10) Spec> thcen
- 11) If scan looks ok, reset the zero value: Spec> set th 0
- 12) Repeat steps (7)-(10) as needed
- 13) Look at a reflection: Spec> an 0.4 0.2
- 14) (Optional) May want to decrease attenuation: Spec> att [att value]
- 15) Spec> lup th -0.05 0.05 20 0.2
- 16) Spec> umv th CEN
- 17) Spec> set th 0.2
- 18) Spec> an 0.2 0.1
- 19) Adjust the attenuation to get APD readout to 50-80K: Spec> att [14]
- 20) Run reflectivity macro. For example: Spec> refgc
- 21) Choose the incident angle you want for gisaxs, thinc: Spec> umv th thinc
- 22) Spec> go_ccd
- 23) Update the filename on the EPICS interface for the Pilatus, update the Next File to 1
- 24) Try a 1-second test image: Spec> pilexp 1
- 25) The detector saturates at 1 million (2^20). The maximum counts seen by any pixel is on the STATS interface, the max for ROI 2. Adjust the exposure time accordingly and record more images
- 26) Record more images at other incident angles, or azimuthal angles, if desired. To help keep track of images, you may want to try Spec> smartnames_on (this puts the incident angle, exposure time, and ccdz state into the filename). To disable this feature, Spec> smartnames_off.
- 27) If you want to patch in the gaps between modules, change the detector height (Spec> pildown)
- 28) Do it all again for the next sample!
- 29) Return the detector to the original height (Spec> pilup)

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