Reducing AOT SWS01 Data in IA[31]

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NOTE: This document is subject to change due to ongoing IA/ISAP software development.

Before you begin:

This guide is intended to give a basic description of how to use IA tasks to reduce AOT1 data: it assumes that the FITS files containing the raw data have already been retrieved from the archive, and gives only a representative use of each command. This document provides background information for the accompanying IA script. For a full description of input parameters and alternate uses of a particular command, consult the IA help tool, accessed by typing a "?" on the command line after starting an IA session. There are many possible help topics, so for each command described below, we have included, to the right hand side, the help topic where it can be found.

All IDL command syntax in an IA data reduction session is shown in **boldface** below,

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with variable and file names indicated in *italics*. All variable names are meant to be representative, but of course can be chosen by the user. These instructions suggest saving the output SPD after particularly time-consuming steps; the user should weigh the value of additional saves against the constraints of disk space. We also recommend that you work in a separate directory for each of your datasets, to prevent confusion.

Before starting IA, you should close all applications which use color (e.g., Netscape, Mosaic, or additional IDL sessions), as they will interfere with the IA color table. After IA has been started, you may restart these applications without affecting the IA session. In addition, for some tools a better performance is achieved if your window manager is set such that a window is active when the pointer is moved into it, but it is not automatically brought to the foreground. In general, it is advisable to keep the text window from which you started IA accessible, as some error messages from the interactive tools will be displayed in this window (note that some error messages will appear in pop-up windows of their own).

To begin: Start your IA session by typing **ia** at the system prompt. Then follow the numbered instructions in this guide.

1) **cal_select** (help: IA UTILITIES)

This tool is used to select calibration files. See Appendix B.4 of the IA User's Manual for descriptions of each calibration file. The cal_select command opens a window which lists all current calibration files, and there are two possible settings for each CAL file: 'BASE' or 'TEST'. The choice will depend on the current status of the calibration files, so ask an IA guru.

It is possible to load a cal_select.dat file (a binary file saved out of a previous run of cal_select) by clicking the 'load' button. NOTE: if using the 'load' option, the 'path' box must be filled in or cal_select will crash.

After either setting the appropriate CAL files manually or loading your calibration file,

click 'OK' to exit the cal_select tool.

2) *erd*=**read_ferd**('xxxYYYzz') (help: IA INPUT OUTPUT)

This command reads in the ERD from the raw data files. The 8 digits in parentheses refer to the revolution number (xxx), target number (YYY), and observation sequence number (zz) from the archive listing. You may also use the command "read_fspd" to read in the SPD directly and skip to step 5; however, the pipeline reduction is ever-changing, and starting with the ERD allows you to take advantage of the most recent version of the reduction software.

3) *spd*=**dspd**(*erd*,**aotlr**=*aotlr*) (help: IA DERIVE SPD)

This command converts the ERD to an SPD, and uses the following CAL files: (1; 2; 2a,b; 3; 4; 5; 6; 12; 16a,b,c,e; 18; 23; 24). The aotlr (grating position data) will be needed later in updown1_inter.

4) *spd_orig*=**read_fspd**('*SWSPxxxYYYzz.FITS*') (help: IA INPUT OUTPUT)
spd.header=*spd_orig.header*
save,*spd,aotlr*,**file**='*spd.save*',/**xdr**

Unfortunately, the full header is not properly saved by dspd; the two commands in this step correct this problem by reading in the SPD (spd_orig) and copying its header into your recently derived SPD (spd). The filename given to read_fspd is in the format of files coming directly from the archive, where 'xxxYYYzz' are the same digits entered in step 2 above. The third command saves the SPD and the aotlr into an IDL save set, which can be restored in later sessions of IA if necessary. In that case, typing

restore,'*spd.save*' (help: ROUTINES)

will restore the IDL structures *spd* and *aotlr* to your IA session.

NOTE: An SPD is rather large, and the use of the option /xdr makes the saving process

significantly faster. The /xdr (for eXternal Data Representation) option is not necessary to save smaller quantities such as the sctbl or aotlr alone (e.g., step 6), but it is needed if you wish to export any save sets between VMS and UNIX machines (see the IDL help file on "save" for more information.)

5) **list_header,***spd* (help: IA UTILITIES)

set_plot,'ps
device,file='*aper_plot.ps***',/land**
aper_plot,*spd*,'1A',60. (help:IA UTILITIES)
aper_plot,*spd*,'3A',60.
aper_plot,*spd*,'3E',60.
aper_plot,*spd*,'4',60.
device,/close
set_plot,'x

These tasks give you more information about your dataset. The list_header command lists the FITS header, and in particular, you should note the position angle of your observations: the real position angle is (roll angle) + 90 degrees. The remaining commands in this step create a postscript file containing plots of the four SWS apertures on the sky (for bands 1 & 2; 3ACD; 3E; and 4). The value of 60. above is the declination scale of the plot in arcseconds. Consult your local system manager for printer options to determine how to print the postscript file locally.

6) *sctbl*=**make_sctbl(***spd***)** (help: IA UTILITIES)
print_sctbl,*sctbl*,**file='***sctbl.txt***',/spawn**
save,*sctbl*,**file='***sctbl.save***'**

These commands create, print out, and save the scan table, which gives such information as the sequence of observations (darks, photometric checks, and science scans) and

wavelengths covered by each band. The print_sctbl command prints the scan table to the screen as well as to the specified file name. The /spawn option is a local MPE command to print the specified text file to the default printer; again, consult your local system manager for printer options.

7) *spd*=**antimem**(*spd*) (help:IA AUTO ANALYSIS)

This routine is designed for reducing memory effects, as is done in the standard pipeline. The use of antimem is mostly historical, but should be kept in the reduction chain because this task may in the future include a more complicated method for dealing with memory effects.

8) *spd_D*=**dark_inter**(*spd,sctbl*,/**multia**) (help:IA UTILITIES)
save,*spd_D*,**file**='*spd_D.save*',/**xdr**

NOTE: Usually the server or window system is configured to retain plot window after scrolling or the start of a screen saver. The /multia option is only needed if the draw window is not retained (as on a Multia terminal); it causes IDL to handle the plot storage directly.

The dark_inter tool facilitates dark current correction. It can be the most time-consuming of all data reduction steps, but is also one of the most important in producing reliable spectra. Make sure to save the SPD after completing this task. Before you begin, you should note the sequence of darks and other scans in the scan table. Note that the reset time ("tr") and gain columns in the scan table must be the same for a dark and the scan from which it is being subtracted; dark_inter automatically makes available all darks with matching tr and gain. Also check the scan table to determine which (band+scan) combinations hold actual data: those which do not will show "0.000" for both wave_min and wave_max (this information is also shown in the message box in dark_inter as "w_range"). It is not necessary to dark subtract these (band+scan) combinations. For each of the four bands (starting at detectors 1 [SW1:], 13 [SW2:], 25 [LW1:], and 37

[LW2:]), the following sequence should be followed:

A.Select a detector. This detector will represent the whole band.

B.Revise Darks. Click "Revise Darks". In the "Revise Darks" window which pops up:

→ Select the first set of darks. This may be done either by clicking "next", which will automatically choose the first set of darks if no dark has been previously selected in the given band, or by clicking the "select" button first and then clicking the dark you want in the middle left panel.

→ choose clipping options–here are the most general suggestions:

▷ first 1 point and clipping(mean) with kappa=3 (i.e., 3σ clipping around the mean).

▷After strong photometric checks, you may want to use clipping (linear) instead of mean to deal with obvious memory effects (i.e., jump in signal followed by a decline in signal over time).

→ refine dark to be subtracted for each detector of the band, for each scan:

▷"clip" shows the points to be clipped from the first detector (top right window): two vertical lines show the kappa-clipping limits you have specified, and all points above or below these lines are clipped.

▷"apply to whole band" shows the effect of the current choice for all detectors in the band.

▷many iterations may be tried (use "undo" between each to cancel effects of previous try), but in the end, the combination of "apply to whole band" and finally "apply" actually applies the clipping to the current set of dark measurements. NOTE: the use of "clip" and "apply to whole band", without clicking the "undo" button in between, results in the first detector being clipped twice.

→ Click on "next" to select the next dark, and repeat the "revise darks" steps for all darks in the requested band. If you did not "apply" the latest clipping to the dark before moving to the next dark, you will get an error message in the text window reminding you to do so.

→ Click "done" in the "Revise Darks" window when all darks in the band have been revised.

C. Subtract darks from photometric checks and science scans:

→ Select the photometric check or science scan to be dark subtracted by indicating its number in the "phot selection" or "scan selection" box. Your selection will be indicated in green in the upper two windows of the left panel. Make sure that the entire selected check or scan can be seen in the left column of windows before proceeding. If necessary, click on "plot" to access a window where you can adjust the plot parameters, or use the slider bar at the bottom of the left panel to adjust the horizontal range. Note that the plot window may stay open during the entire use of dark_inter, so that you may readjust as necessary.

→ Choose the dark subtraction method: click on "Method".

▷In the "method" menu, choose which darks to use. The darks with matching tr and gain will be highlighted in yellow-orange. If you only have one dark before the check or scan which can be used, you should use "previous" and "mean" for the method. If you have valid dark measurements (that is, with the same value of "tr" and gain) before and after a scan, you can choose to do a linear interpolation between the two. If the two darks come immediately before and after the scan, select them with the "previous" and "next" buttons, and click "linear interpolation" for the method.

▷When you have successfully marked the dark in question, it will be shown in red. If you get strange fits or error messages when trying to subtract a dark, it may be because the dark was not properly selected; go back to the "Method" menu and toggle your dark choices until they appear in red in the left plot windows.

▷ Choosing other darks: If you wish to choose a dark which comes after an intermediate dark for a linear interpolation fit, you must click on "cursor" and then on the appropriate dark measurement in the middle window of the left panel to select it–the "next" button only points to the very next dark, regardless of whether or not it can be used. The "cursor" option is very location-sensitive, so you may have to click a few times (and get a few warning messages). If you change your mind and decide not to use it, click "cursor" and the dark again to unselect it. Be sure that the dark points are colored yellow-orange again to be sure you were successful.

▷If the available darks do not provide a good dark measurement to be subtracted, you may do a spline fit instead of a mean or linear interpolation fit. For now, just choose "spline". NOTE: The spline fit must be done for each detector individually, and is much more time consuming; use only when necessary.

▷ Click "Done" in the Method menu.

→ Subtract fitted dark current.

▷Click on "fit", which will show the fit for the selected detector in the bottom left plot.

▷If the method selected is "mean" or "linear interpolation", all that remains is to click on the "apply to whole band" button. You may iterate through the "method", "fit", and "apply to whole band" sequence if desired. If the dark subtraction is satisfactory for all detectors in this band/scan combination, you have completed the subtraction.

▷However, often "jumps" are seen in the data. If these are simply horizontal steps, then it is advisable to use either the previous or next dark to fit the longest section and then mask out the other section later in the mask_inter tool. To make a correction to the dark subtraction for one detector, use the slider at the top of the tool to select the detector, and then set the scan number in the window below. Using the "fit" button writes a new solution for that detector, and doesn't affect the other detectors.

▷If the method selected is "spline", clicking the "fit" button produces instruction messages in the IA text window which explain how to define spline points, fit the spline, iterate if the first fit is not optimal, and apply the fit to the data. Spline points should be defined on the center left plot. Repeat the "fit" step for each individual detector in the band.

▷Additional notes for spline fits: Previous fits are left on the plot, and can be used to guide a better spline fit, if necessary. Closely spaced spline points are likely to cause a highly structured fit; widely spaced spline points are better for this purpose. You will need to enter the scan number again each time you increment the detector number and want to perform a new spline fit. You will also need to replot each time if you used special plot parameters to get the selected scan to fit in the left windows.

→ Repeat Step C for each photometric check and science scan in this band.

D. Set slider on top to the first detector number of the next band and repeat the process, starting with Step B, Revise Darks. Complete Steps B and C for all four bands. It is a good idea to save the data set periodically during the use of dark_inter, to prevent loss of efforts in the event that the tool crashes. The easiest way to do this is to use the "save current" button in the dark_inter tool, which creates a file called "dark_inter_save.xdr" containing the current data file. This file is overwritten in successive uses of this button, so you may make incremental saves without using additional disk space.

9) *spd_DM*=**mask_inter**(*spd_D,sctbl*) (help: IA UTILITIES)

save,*spd_DM*,**file**='*spd_DM.save*',/**xdr**

This command puts you into the mask_inter tool, which can be used to inspect your SPD and mask out bad data. Sometimes a whole detector is significantly noisier than the others in the band, or there may be significant memory effects after a strong glitch. In addition, you can mask out the jump segments first detected with dark_inter (see step 8C). See the on-line help for this tool for a detailed account of all the utilities of this tool.

Enter a band number in the "band" box, click on the scan number you wish to see, and click "disp_spd" to display a subset of the data. Note that to examine a different scan, you must click on the previous scan number again to "unset" it; otherwise there will be two scans selected and the tool will complain (multiple scans can be selected for producing an AAR within mask_inter, but only one scan can be displayed at a time). If you click the "plotmode" button, a background window will pop up in which you can set the plot scale; we recommend that you choose a fixed y-scale, as this is a good way of finding detectors for which dark subtraction didn't work properly, or for finding noisy detectors or other strange features. Each time you change the plot scale, click "disp_spd" again to display the data with the new plot values.

To mask out a specific section of a detector, click "select" and the detector of interest, which will cause the data from that detector to show in the bottom left window. Then click "mask" and use the left mouse button to select the region to be masked out (e.g., in the case of a memory effect). If you see the same glitch in all 12 detectors of a band, you can use the "do to all" button after masking out the glitch in one of the detectors. If you judge the input from an entire detector to be bad, you can click on the detector number in the list to the right-hand side of the display, and then click "faulty" to set it all bad.

IMPORTANT NOTE: Using the "faulty" button to set a detector "bad" is NOT reversible at the current time, and sets the specified detector bad for all scans. Setting the detector

bad by masking all points out using the "mask" button IS reversible (using "unmask"), and only sets the detector bad for the specified scan.

For now, set 'order' to zero, thus showing all the data in that band; the order is used to separate different AOT bands, which are of interest in later steps. Saving the masked SPD should be considered if a significant amount of masking has been done.

10) *spd_U*=**updown1_inter**(*spd_DM,sctbl,aotlr*) (help: IA UTILITIES)
save,*spd_U*,**file='***spd_U.save*'**,/xdr**

This tool corrects for differences between the 'up' and 'down' scans of the instrument. Observations are taken in pairs of "up" and "down" scans, where wavelength decreases with increasing time in the "up" scan, and increases with time in the "down" scan. Due to memory effects, the shapes of the "up" and "down" scan may be different. The updown1_inter tool takes a smoothed version of each of the up and down scans separately, and determines a "gain" or "offset" correction to make the up and down scans consistent with each other. The top plot of the tool shows a section of the scan including the region you are correcting; you will see darks or photometric checks on either end of this plot. The center plot shows the up and down scan laid over each other, and the bottom of the three long plots shows either the ratio or difference between the up and down scans (depending on whether you have chosen the "gain" or "offset" option). For updown1_inter (as opposed to updown_inter, the version used for other observing modes), the middle and bottom plots are in wavelength-like units, but due to the overlapping AOT bands they are not in actual wavelengths. Note that in these plots, wavelength *decreases* to the right.

The "gain" option is really only useful for observations with very strong signal, where the uncertainties due to dark subtraction can be neglected with respect to gain effects. In addition, the choice of the smoothing radii should depend on speed, choosing smaller values for higher speeds; it is also dependent on resolution. Table 1 summarizes the general recommendations for median and smoothing radii as well as correction method

for the different bands. The key is to look at the resultant correction that the tool wants to make and be sure that it doesn't follow the noise too closely or introduce artifacts into the time spectrum.

NOTE: Ignore "Floating divide by zero" errors upon starting updown1_inter, as they are spurious.

The general procedure:

→Set "det" to the first detector number of the band you want to correct, and select the scan.

→Set the correction mode and median and smoothing radii (see Table 1).

→Click "Do Adjustment"; the bottom of the three long plots shows the correction determined from the smoothed data overlaid as a dotted curve.

→You may iterate on choosing the correction mode and doing the adjustment if the solution is not satisfactory; after finding an acceptable updown correction, click on "Update full array" to perform the up-down correction on all detectors in the band for that scan.

Repeat this process for all scans in each band, and for all four bands. After completing the updown correction, save the SPD.

Table 1: Recommended Up-Down Correction Settings

Band	Scan	R(median)	R(smooth)	method
1	1	50-100	50-200	offset
1	2	”	”	”
2	1	”	”	”
2	2	”	”	”
3	1	”	”	offset or (gain)
3	2	100-200	200-400	”
3	3	”	”	”
4	1	”	”	”

11) **plot_phot,***spd_U,sctbl,*[1],13**,/all** (help:IA UTILITIES)

plot_phot,*spd_U,sctbl,*[1],25**,/all**

plot_phot,*spd_U,sctbl,*[1],37**,/all**

The routine plot_phot can be used to look at photometric check data. The command lines given above request IA to plot the first photometric check for all 12 detectors of bands 2, 3, and 4 (which start with detector numbers 13, 25, and 37); the band 1 photometric check is not used. If you have more than one photometric check, you can replace the [1] above with a range of values (e.g. [1,3]) and compare them on the same plot. By specifying a particular detector and removing the /all flag, you can look at the photometric checks for that detector.

Data points already flagged as bad by the system will be marked (see help file for color code). If these are the only bad points and the photometric checks look sensible, the standard routines will handle the bad points appropriately. Note that the default is to always use the first photometric check, so if it is corrupted it should be replaced with another. If you suspect your photometric data has problems which are severe enough to

affect the ultimate calibration, such as a corrupted first phot. check or large spikes not already flagged, consult an IA guru.

12) *spd_U*=**respcal**(*spd_U*) (help: IA AUTO ANALYSIS)
spd_U=**fluxcon**(*spd_U*)
spd_U=**velcor**(*spd_U*)

The routines respcal and fluxcon apply various calibration files: respcal uses CAL files 13, 19, and 25; fluxcon uses CAL file 13. The output of fluxcon looks like the following:

```
Warning(SPCP): some phot. check data might be doubtful       Det:26 Uv: 2
Warning(SPCP): some phot. check data might be doubtful       Det:39 Uv: 2
Warning(SPCP): some phot. check data might be doubtful       Det:46 Uv: 3
Warning(SPCP): some phot. check data might be doubtful       Det:48 Uv: 2
```

If the number following "Uv:" (the number of outliers) is greater than 3, the photometric check data for that detector indicated may be suspect.

The velcor task performs a correction to the heliocentric frame based on the velocity of the satellite.

13) *spd_F*=**flat_inter**(*spd_U,sctbl*) (help: IA UTILITIES)
save,*spd_F*,**file**='*spd_F.save*',/**xdr**

The first command starts up the flatfield correction tool flat_inter. This task corrects the flatfield of each detector in the band with respect to the others. As with "updown1_inter", the correction can be a "gain" or "offset" correction, and is made on the basis of smoothed data. It is generally recommended to use the "median" filter, and to exclude poor points. As with updown1_inter, you should make sure that the fit you have chosen does not introduce spurious features, and vary the smoothing or correction parameters accordingly,

and the 'gain' option should only be used with strong signal. You should apply flatfields to each AOT separately by choosing the appropriate combination of band, scan, and order. Table 2 shows the suggested settings for correction mode and smoothing radii for each AOT band. NOTE: be sure to press "return" after changing each entry; otherwise, strange effects may result if new values are not registered by the tool.

Table 2: Flatfield correction settings

AOT	Band	scan	order	mode	res1	res2
1A	1	1	4	offset	50-500	50-200
1B	1	1	3	"	"	"
1D	1	2	3	"	"	"
1E	1	2	2	"	"	"
2A	2	1	2	"	"	"
2B	2	1	1	"	"	"
2C	2	2	1	"	"	"
3A	3	1	2	offset or (gain)	25-400	50-100
3C	3	2	2	"	"	"
3D	3	2	1	"	"	"
3E	3	3	1	"	"	"
4	4	1	1	"	"	"

14) *spd_FM*=**mask_inter**(*spd_F,sctbl*) (help: IA UTILITIES)

save,*spd_FM*,**file**='*spd_FM.save*',/**xdr**

Now that all calibrations have been applied, you can go back into the interactive masking tool used in step 9 to check for lines which have been marked as glitches (in low-resolution SWS01 data, lines can be confused as glitches). There can be a significant number of such glitches, even in the slowest-speed SWS01 observations, and they may mark the line wings as well as the line peak. You may also wish to mask out noise spikes or outliers;

for instance, a signal which is clearly spurious, such as a strong spike detected only in the up or down scan of all 12 detectors, will bias the resulting spectrum. This masking step may be best done after following the rest of the steps described below and looking at an unmasked version of the output spectrum to see where lines have been detected, and returning to this step to produce a refined dataset.

See Table 3 below for the wavelength ranges covered by each AOT. Click the "w-mode" button to plot in wavelength units, and specify the wavelength range of interest in the "plotmode" pop-up window. Glitched points are marked by yellow asterisks. To remove an unwanted glitch, click "select" to specify the detector, and if needed click "zoom" and use the cursor to select the section of the scan holding the offending point (both up and down scans will be shown, with wavelength units along the upper x axis of the plot). Then click "unglitch", and click the mouse cursor on either side of the glitched point. The unglitched point will then appear as a white cross, like the rest of the good data. You can also use the "fluxclip" button to mask out data by hand. When you click this button, the data from all detectors is plotted in the lower left hand window, and you can define a series of boxes with the left mouse button; the points inside these boxes are then clipped. Clicking the right mouse button returns you to the main mask_inter tool.

You can see the effects of your masking by using the "extract_aar" and "aar_f" buttons, which will produce a filtered AAR (with the resolution specified in the "resol" window) of the specific section of spectrum that you have selected with band, scan, order, and x_range. Further, you can output your final, masked AAR result to a file by clicking on the "save_aar" button. This action will pop up a window where you can specify the name of the file you wish to write, and what format: XDR, IDL SAVE, or FITS. The first two options both write IDL save files which can be restored in an IA session (the first using the XDR format described in step 4).

If you choose this method of producing AARs, then step 15 below is not necessary; skip to step 16 to complete your reduction.

15) *spd_FM*=**killwait**(*spd_FM*) (help: IA UTILITIES)

aar=**extract_aar**(*spd_FM*) (help: IA AUTO ANALYSIS)

save,*aar*,**file**='*aar.save*',/**xdr**

status=**sap_wfits**('*aar.fits*',*aar*) (help: ISAP)

aar_filt=**aarfilt1**(*aar*,1,/**noglitch**,/**noover**) (help: IA UTILITIES)

The first task flags "wait" points as "bad"; it is useful on SWS01, SWS02, and SWS06 data. Note that running the killwait task is not necessary if AARs are produced from the mask_inter tool (see step 14). The next task extracts an AAR from the reduced SPD, and the following two commands save the AAR as an IDL save set and a FITS file. Note that all the masked and glitched points are still in the AAR, but are ignored by aarfilt1.

The last task produces a final, filtered AAR (bad points removed, order overlaps resolved, each AOT set to a particular resolution). The second parameter of aarfilt1 is the speed factor, which determines the effective resolution. A speed factor of 1 should be used only for the slowest of the 4 AOT1 modes (S01 S=4; see the scan table for scan_type). Be advised that aarfilt1 was optimized for the slowest AOT1 mode, and that the optimum resolution will depend on the source distribution as well as the AOT Band. For S=1 (fastest) data, a speed factor of $\sim$0.3 is appropriate, with intermediate values for S=2 and S=3 data. Consult an IA guru for more advice. Note that glitches are not automatically neglected; you must use the /noglitch option to do so. This is most significant in bands 3 and especially 4, where there are many points marked as glitches. Table 3 shows the band width and resolution settings in the current version of aarfilt1 (version 1.1, 20 May 1996).

Table 3: AOT Bandwidths set by AARFILT1

AOT	(Band,Scan,Order)	(λ_1,λ_2)	R($\lambda/\delta\lambda$) [a]
1A	(1,1,4)	2.3,2.6	1700
1B	(1,1,3)	2.6,3.02	1400
1D	(1,2,3)	3.02,3.52	1650
1E	(1,2,2)	3.52,4.08	1200
2A	(2,1,2)	4.08,5.30	(0.8)*1600
2B	(2,1,1)	5.30,7.03	(0.8)*1000
2C	(2,2,1)	7.03,12.35	(0.8)*1600
3A	(3,1,2)	12.35,16.4	1500
3C	(3,2,2)	16.4,19.5	1900
3D	(3,2,1)	19.5,27.5	1200
3E	(3,3,1)	27.5,29.5	1450
4	(4,1,1)	29.5,45.2	1400

[a] Resolution assumes a speed factor of 1.

16) *aar_fringe*=**aarfringe**(*aar_filt*) (help:IA UTILITIES)

After this step, the basic reduction of SWS01 data is complete, but there is likely significant fringing, especially for lines lying in band 3 (and to some extent in band 2). These may be dealt with using the interactive tool "aarfringe".

The documentation for aarfringe is quite detailed, and we recommend you read it thoroughly to determine your options; only a brief description of tool usage is given here. When you start the tool, you will see buttons across the top of the tool to select the data and method of fringe fitting. Select each band, one at a time, from the pull-down AOT Band menu at the top left. Then you must choose the fringe fitting method: Fast Fourier Transform (FFT), sinusoidal (SIN) or Do Kester's method (DOK) of iteratively fitting sine components. You may also choose gain or offset; gain is the default for this tool, as you usually see fringes most clearly in stronger sources.

Before attempting a fringe fit, you should enter the radial velocity of your source as well as the wavelengths of strong lines in the bottom center panel. These lines will be excluded ¿from the fit; choose the resolution such that the entire line gets masked out. The bottom right panel is where you specify parameters specific to each model; consult the help file for how best to fill in this block. Do Kester's method has been seen to produce the best results in general, but certainly try multiple methods to see what works best for your data.

Clicking the "Do it" button at top right performs a fit; you can change parameters/methods and click this button again. When you think you have a good fit that you want to keep, click "Apply". Repeat for each AOT band that you want to correct. When you are done with all, click "Done" to return to the IA prompt. At this point, you should save your AAR again if you did any significant de-fringing.

Finishing up:

Type "help" to look at the list of IDL structures you have generated in this data reduction

section. Make sure to save all files which were produced from the most time-consuming steps (i.e., any of the interactive tools) as IDL save sets and FITS files before exiting from IA. Then you can restore the SPD or AAR at any stage and redo some of the reduction if you find it necessary at a later date.

Reducing AOT SWS02 Data in IA[31]

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and

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and

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and

Dieter Lutz

NOTE: This document is subject to change due to ongoing IA/ISAP software development.

Before you begin:

This guide is intended to give a basic description of how to use IA tasks to reduce AOT2 data: it assumes that the FITS files containing the raw data have already been retrieved from the archive, and gives only a representative use of each command. This document provides background information for the accompanying IA script. For a full description of input parameters and alternate uses of a particular command, consult the IA help tool. This help tool is accessed by typing a "?" on the command line after starting an IA session. There are many possible help topics, so for each command described below, we have included, to the right hand side, the help topic where it can be found.

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All IDL command syntax in an IA data reduction session is shown in **boldface** below, with variable and file names indicated in *italics*. All variable names are meant to be representative, but of course can be chosen by the user. These instructions suggest saving the output SPD after particularly time-consuming steps; the user should weigh the value of additional saves against the constraints of disk space. We also recommend that you work in a separate directory for each of your datasets, to prevent confusion.

Before starting IA, you should close all applications which use color (e.g., Netscape, Mosaic, or additional IDL sessions), as they will interfere with the IA color table. After IA has been started, you may restart these applications without affecting the IA session. In addition, for some tools a better performance is achieved if your window manager is set such that a window is active when the pointer is moved into it, but it is not automatically brought to the foreground. In general, it is advisable to keep the text window from which you started IA accessible, as some error messages from the interactive tools will be displayed in this window (note that some error messages will appear in pop-up windows of their own).

To begin: Start your IA session by typing **ia** at the system prompt. Then follow the instructions below.

1) **cal_select** (help: IA UTILITIES)

This tool is used to select calibration files. See Appendix B.4 of the IA User's Manual for descriptions of each calibration file. The cal_select command opens a window which lists all current calibration files, and there are two possible settings for each CAL file: 'BASE' or 'TEST'. The choice will depend on the current status of the calibration files, so ask an IA guru.

It is possible to load a cal_select.dat file (a binary file saved out of a previous run of cal_select) by clicking the 'load' button. NOTE: if using the 'load' option, the 'path' box must be filled in or cal_select will crash.

After either setting the appropriate CAL files manually or loading your calibration file, click 'OK' to exit the cal_select tool.

2) *erd*=**read_ferd**('xxxYYYzz') (help: IA INPUT OUTPUT)

This command reads in the ERD from the raw data files. The 8 digits in parentheses refer to the revolution number (xxx), target number (YYY), and observation sequence number (zz) from the archive listing. You may also use the command "read_fspd" to read in the SPD directly and skip to step 5; however, the pipeline reduction is ever-changing, and starting with the ERD allows you to take advantage of the most recent version of the reduction software.

3) *spd*=**dspd**(*erd*) (help: IA DERIVE SPD)

This command converts the ERD to an SPD, using the following CAL files: (1; 2; 2a,b; 3; 4; 5; 6; 12; 16a,b,c,e; 18; 23; 24).

4) *spd_orig*=**read_fspd**('*SWSPxxxYYYzz.FITS*') (help: IA INPUT OUTPUT)
spd.header=*spd_orig.header*
save,*spd*,**file=**'*spd.save*'**,/xdr**

Unfortunately, the full header is not properly saved by dspd; the two commands in this step correct this problem by reading in the SPD (spd_orig) directly and copying its header into your recently derived SPD (spd). The filename given to read_fspd is in the format of files coming directly from the archive, where 'xxxYYYzz' are the same digits entered in step 2 above. The third command saves the SPD into an IDL save set, which can be restored in later sessions of IA if necessary. In that case, typing

restore,'*spd.save*' (help: ROUTINES)

will restore the IDL structure *spd* to your IA session.

NOTE: An SPD is rather large, and the use of the option /xdr makes the saving process

significantly faster. The /xdr (for eXternal Data Representation) option is not necessary to save smaller quantities such as the sctbl or aotlr alone (e.g., step 6), but it is needed if you wish to export any save sets between VMS and UNIX machines (see the IDL help file on "save" for more information.)

5) **list_header,***spd* (help: IA UTILITIES)
set_plot,'ps
device,file='*aper_plot.ps***',/land**
aper_plot,*spd*,'1A',60. (help:IA UTILITIES)
aper_plot,*spd*,'3A',60.
aper_plot,*spd*,'3E',60.
aper_plot,*spd*,'4',60.
device,/close
set_plot,'x

These tasks give you more information about your dataset. The list_header command lists the FITS header, and in particular, you should note the position angle of your observations: the real position angle is (roll angle) + 90 degrees. The remaining commands in this step create a postscript file containing plots of the four SWS apertures on the sky (for bands 1 & 2; 3ACD; 3E; and 4). The value of 60. above is the declination scale of the plot in arcseconds. Consult your local system manager for printer options to determine how to print the postscript file locally.

6) *sctbl*=**make_sctbl(***spd***)** (help: IA UTILITIES)
print_sctbl,*sctbl*,**file='***sctbl.txt***',/spawn**
save,*sctbl*,**file='***sctbl.save***'**

These commands create, print out, and save the scan table, which gives such information as the sequence of observations (darks, photometric checks, and science scans) and

wavelengths covered by each band. The print_sctbl command prints the scan table to the screen as well as to the specified file name. The /spawn option is a local MPE command to print the specified text file to the default printer; again, consult your local system manager for printer options.

7) *spd_mem*=**antimem**(*spd*) (help:IA AUTO ANALYSIS)

This routine is designed for reducing memory effects, as is done in the standard pipeline. The use of antimem is mostly historical, but should be kept in the reduction chain because this task may in the future include a more complicated method for dealing with memory effects.

8) *spd_D*=**dark_inter**(*spd,sctbl*,**/multia**) (help:IA UTILITIES)
save,*spd_D*,**file**='*spd_D.save*',**/xdr**

NOTE: Usually the server or window system is configured to retain plot window after scrolling or the start of a screen saver. The /multia option is only needed if the draw window is not retained (as on a Multia terminal); it causes IDL to handle the plot storage directly.

The dark_inter tool facilitates dark current correction. It can be the most time-consuming of all data reduction steps, but is also one of the most important in producing reliable spectra. Make sure to save the SPD after completing this task. Before you begin, you should note the sequence of darks and other scans in the scan table. Note that the reset time ("tr") and gain columns in the scan table must be the same for a dark and the scan from which it is being subtracted; dark_inter automatically makes available all darks with matching tr and gain. Also check the scan table to determine which (band+scan) combinations hold actual data: those which do not will show "0.000" for both wave_min and wave_max (this information is also shown in the message box in dark_inter as "w_range"). It is not necessary to dark subtract these (band+scan) combinations. For

each of the four bands (starting at detectors 1 [SW1:], 13 [SW2:], 25 [LW1:], and 37 [LW2:]), the following sequence should be followed:

A.Select a detector. This detector will represent the whole band.

B.Revise Darks. Click "Revise Darks". In the "Revise Darks" window which pops up:

→ Select the first set of darks. This may be done either by clicking "next", which will automatically choose the first set of darks if no dark has been previously selected in the given band, or by clicking the "select" button first and then clicking the dark you want in the middle left panel.

→ choose clipping options–here are the most general suggestions:

▷ first 1 point and clipping(mean) with kappa=3 (i.e., 3σ clipping around the mean).

▷After strong photometric checks, you may want to use clipping (linear) instead of mean to deal with obvious memory effects (i.e., jump in signal followed by a decline in signal over time).

→ refine dark to be subtracted for each detector of the band, for each scan:

▷"clip" shows the points to be clipped from the first detector (top right window): two vertical lines show the kappa-clipping limits you have specified, and all points above or below these lines are clipped.

▷"apply to whole band" shows the effect of the current choice for all detectors in the band.

▷many iterations may be tried (use "undo" between each to cancel effects of previous try), but in the end, the combination of "apply to whole band" and finally "apply" actually applies the clipping to the current set of dark measurements. NOTE: the use of "clip" and "apply to whole band", without clicking the "undo" button in between, results in the first detector being clipped twice.

→ Click on "next" to select the next dark, and repeat the "revise darks" steps for all darks in the requested band. If you did not "apply" the latest clipping to the dark before moving to the next dark, you will get an error message in the text window reminding you to do so.

→ Click "done" in the "Revise Darks" window when all darks in the band have been revised.

C. Subtract darks from photometric checks and science scans:

→ Select the photometric check or science scan to be dark subtracted by indicating its number in the "phot selection" or "scan selection" box. Your selection will be indicated in green in the upper two windows of the left panel. Make sure that the entire selected check or scan can be seen in the left column of windows before proceeding. If necessary, click on "plot" to access a window where you can adjust the plot parameters, or use the slider bar at the bottom of the left panel to adjust the horizontal range. Note that the plot window may stay open during the entire use of dark_inter, so that you may readjust as necessary.

→ Choose the dark subtraction method: click on "Method".

▷In the "method" menu, choose which darks to use. The darks with matching tr and gain will be highlighted in yellow-orange. If you only have one dark before the check or scan which can be used, you should use "previous" and "mean" for the method. If you have valid dark measurements (that is, with the same value of "tr" and gain) before and after a scan, you can choose to do a linear interpolation between the two. If the two darks come immediately before and after the scan, select them with the "previous" and "next" buttons, and click "linear interpolation" for the method.

▷When you have successfully marked the dark in question, it will be shown in red. If you get strange fits or error messages when trying to subtract a dark, it may be because the dark was not properly selected; go back to the "Method" menu and toggle your dark choices until they appear in red in the left plot windows.

▷ Choosing other darks: If you wish to choose a dark which comes after an intermediate dark for a linear interpolation fit, you must click on "cursor" and then on the appropriate dark measurement in the middle window of the left panel to select it–the "next" button only points to the very next dark, regardless of whether or not it can be used. The "cursor" option is very location-sensitive, so you may have to click a few times (and get a few warning messages). If you change your mind and decide not to use a selected dark, click "cursor" and the dark again to unselect it. Be sure that the dark points are colored yellow-orange again to be sure you were successful.

▷If the available darks do not provide a good dark measurement to be subtracted, you may do a spline fit instead of a mean or linear interpolation fit. For now, just choose "spline". NOTE: The spline fit must be done for each detector individually, and is much more time consuming; use only when necessary.

▷ Click "Done" in the Method menu.

→ Subtract fitted dark current.

▷Click on "fit", which will show the fit for the selected detector in the bottom left plot.

▷If the method selected is "mean" or "linear interpolation", all that remains is to click on the "apply to whole band" button. You may iterate through the "method", "fit", and "apply to whole band" sequence if desired. If the dark subtraction is satisfactory for all detectors in this band/scan combination, you have completed the subtraction.

▷However, often "jumps" are seen in the data. If these are simply horizontal steps, then it is advisable to use either the previous or next dark to fit the longest section and then mask out the other section later in the mask_inter tool. To make a correction to the dark subtraction for one detector, use the slider at the top of the tool to select the detector, and then set the scan number in the window below. Using the "fit" button writes a new solution for that detector, and doesn't affect the other detectors.

▷If the method selected is "spline", clicking the "fit" button produces instruction messages in the IA text window which explain how to define spline points, fit the spline, iterate if the first fit is not optimal, and apply the fit to the data. Spline points should be defined on the center left plot. Repeat the "fit" step for each individual detector in the band.

▷Additional notes for spline fits: Previous fits are left on the plot, and can be used to guide a better spline fit, if necessary. Closely spaced spline points are likely to cause a highly structured fit; widely spaced spline points are better for this purpose. You will need to enter the scan number again each time you increment the detector number and want to perform a new spline fit. You will also need to replot each time if you used special plot parameters to get the selected scan to fit in the left windows.

→ Repeat Step C for each photometric check and science scan in this band.

D. Set slider on top to the first detector number of the next band and repeat the process, starting with Step B, Revise Darks. Complete Steps B and C for all four bands. It is a good idea to save the data set periodically during the use of dark_inter, to prevent loss of efforts in the event that the tool crashes. The easiest way to do this is to use the "save current" button in the dark_inter tool, which creates a file called "dark_inter_save.xdr" containing the current data file. This file is overwritten in successive uses of this button, so you may make incremental saves without using additional disk space.

9) *spd_DM*=**mask_inter**(*spd_D,sctbl*) (help: IA UTILITIES)

save,*spd_DM*,**file**='*spd_DM.save*',/**xdr**

This command puts you into the mask_inter tool, which can be used to inspect your SPD and mask out bad data. Sometimes a whole detector is significantly noisier than the others in the band, or there may be significant memory effects after a strong glitch. In addition, you can mask out the jump segments first detected with dark_inter (see step 8C). See the on-line help for this tool for a detailed account of all the utilities of this tool.

Enter a band number in the "band" box, click on the scan number you wish to see, and click "disp_spd" to display a subset of the data. Note that to examine a different scan, you must click on the previous scan number again to "unset" it; otherwise there will be two scans selected and the tool will complain (multiple scans can be selected for producing an AAR within mask_inter, but only one scan can be displayed at a time). If you click the "plotmode" button, a background window will pop up in which you can set the plot scale; we recommend that you choose a fixed y-scale, as this is a good way of finding detectors for which dark subtraction didn't work properly, or for finding noisy detectors or other strange features. Each time you change the plot scale, click "disp_spd" again to display the data with the new plot values.

To mask out a specific section of a detector, click "select" and the detector of interest, which will cause the data from that detector to show in the bottom left window. Then click "mask" and use the left mouse button to select the region to be masked out (e.g., in the case of a memory effect). If you see the same glitch in all 12 detectors of a band, you can use the "do to all" button after masking out the glitch in one of the detectors. If you judge the input from an entire detector to be bad, you can click on the detector number in the list to the right-hand side of the display, and then click "faulty" to set it all bad.

IMPORTANT NOTE: Using the "faulty" button to set a detector "bad" is NOT reversible at the current time, and sets the specified detector bad for all scans. Setting the detector

bad by masking all points out using the "mask" button IS reversible (using "unmask"), and only sets the detector bad for the specified scan.

Saving the masked SPD should be considered if a significant amount of masking has been done.

10) *spd_U*=**updown_inter**(*spd_DM,sctbl*) (help: IA UTILITIES)
save,*spd_U*,**file**='*spd_U.save*',/**xdr**

This tool corrects for differences between the 'up' and 'down' scans of the instrument. Observations are taken in pairs of "up" and "down" scans, where wavelength decreases with increasing time in the "up" scan, and increases with time in the "down" scan. Due to memory effects, the shapes of the "up" and "down" scan may be different. The updown_inter tool takes a smoothed version of each of the up and down scans separately, and determines a "gain" or "offset" correction to make the up and down scans consistent with each other.

When you start the tool and select a detector and scan number, you will see to the left side four plots: the section of the stripchart that holds your scan, the up and down scans overplotted, the difference between the up and down scans, and the stdev of the difference. It is recommended to do the updown correction on a detector in the middle of the band, for if there is strong signal it will show up in the middle of this detector, and the effects of the updown correction will be easier to interpret. The recommended settings are the defaults: use the "fit" method, correcting "both" sections, using the "offset" correction mode, and "exclude" bad points. Choose kappa=2-3, and use the default "linear ramp" function for fitting. The "gain" option is really only useful for observations with very strong signal, where the uncertainties due to dark subtraction can be neglected with respect to gain effects. In addition, other fit functions may be used, though this is generally not advised. For example, "Fouks (SPIE)"[2]and "PHT-IST26" are functions

[2]SPIE 2268, 283 (1994).

which are found to fit transients of infrared detectors, but they are not tuned to the SWS detectors and make no claim on reproducing a physical model of them.

At this point, you only need to click "Do Fit" to see the fit; the third plot ¿from the top will then overplot the fit on the "difference up-down" plot. You may iterate on choosing the correction mode, kappa, etc. and doing the fit if the solution is not satisfactory. After finding an acceptable updown correction, click on "Update full array" to perform the up-down correction on all detectors in the band for that scan. The results for all 12 detectors in the band are shown to the right of the menu panel.

Repeat this process for all scans in each band, and for all bands where you have data. NOTE: updown_inter should **not** be used to correct band 4 (LW2:) for very faint sources, where glitch tails can dominate the solution. Consult your IA guru for more information.

11) **plot_phot,***spd_U,sctbl,*[1],13**,/all** (help:IA UTILITIES)
plot_phot,*spd_U,sctbl,*[1],25**,/all**
plot_phot,*spd_U,sctbl,*[1],37**,/all**

The routine plot_phot can be used to look at photometric check data. The command lines given above request IA to plot the first photometric check for all 12 detectors of bands 2, 3, and 4 (which start with detector numbers 13, 25, and 37); the band 1 photometric check is not used. If you have more than one photometric check, you can replace the [1] above with a range of values (e.g. [1,3]) and compare them on the same plot. By specifying a particular detector and removing the /all flag, you can look at the photometric checks for that detector.

Data points already flagged as bad by the system will be marked (see help file for color code). If these are the only bad points and the photometric checks look sensible, the standard routines will handle the bad points appropriately. Note that the default is to always use the first photometric check, so if it is corrupted it should be replaced with another. If you suspect your photometric data has problems which are severe enough to

affect the ultimate calibration, such as a corrupted first phot. check or large spikes not already flagged, consult an IA guru.

12) *spd_U*=**respcal**(*spd_U*) (help:IA AUTO ANALYSIS)
spd_U=**fluxcon**(*spd_U*)
spd_U=**velcor**(*spd_U*)

The routines respcal and fluxcon apply various calibration files; respcal uses CAL files 13, 19, and 25; fluxcon uses CAL file 13. The output of fluxcon looks like the following:

```
Warning(SPCP): some phot. check data might be doubtful       Det:26 Uv: 2
Warning(SPCP): some phot. check data might be doubtful       Det:39 Uv: 2
Warning(SPCP): some phot. check data might be doubtful       Det:46 Uv: 3
Warning(SPCP): some phot. check data might be doubtful       Det:48 Uv: 2
```

If the number following "Uv:" (the number of outliers) is greater than 3, the photometric check data for that detector indicated may be suspect.

The task velcor performs a correction to the heliocentric frame based on the velocity of the satellite.

13) *spd_F*=**flat_inter**(*spd_U,sctbl*) (help: IA UTILITIES)
save,*spd_F*,**file**='*spd_F.save*',/**xdr**

The first command starts up the flatfield correction tool flat_inter. This task corrects the flatfield of each detector in the band with respect to the others. As with "updown_inter", the correction can be a "gain" or "offset" correction, and is made on the basis of smoothed data. It is generally recommended to use the "median" filter, and to exclude poor points. As with updown_inter, you should make sure that the fit you have chosen does not

introduce spurious features, and vary the smoothing or correction parameters accordingly. The default values are Res1=500 and Res2=100; try lower values, especially for band 4 (LW2:) for very faint sources, and larger values may be useful for very bright sources. In general, it is a good idea to keep the relation Res1∼5 Res2. As before, the 'gain' option should only be used with strong signal. NOTE: be sure to press "return" after changing each entry; otherwise, strange effects may result if new values are not registered by the tool.

14) *spd_FM*=**mask_inter**(*spd_FM,sctbl*) (help: IA UTILITIES)

save,*spd_FM*,**file='***spd_FM.save*',**/xdr**

Now that all calibrations have been applied, you can go back into the interactive masking tool used in step 9 to mask out noise spikes or outliers; for instance, a signal which is clearly spurious, such as a strong spike detected only in the up or down scan of all 12 detectors, will bias the resulting spectrum. This masking step may be best done after following the rest of the steps described below and looking at an unmasked version of the output spectrum to see where lines have been detected, and returning to this step to produce a refined dataset.

Click the "w-mode" button to plot in wavelength units, and specify the wavelength range of interest in the "plotmode" pop-up window. You can use the "fluxclip" button to mask out outlying data by hand. When you click this button, the data from all detectors is plotted in the lower left hand window, and you can define a series of boxes with the left mouse button; the points inside these boxes are then clipped. Clicking the right mouse button returns you to the main mask_inter tool. If you want to remove an unwanted glitch (glitches are marked with yellow asterisks), click "select" to specify the detector, and if needed click "zoom" and use the cursor to select the section of the scan holding the offending point (both up and down scans will be shown, with wavelength units along the upper x axis of the plot). Then click "unglitch", and click the mouse cursor on either side

of the glitched point. The unglitched point will show up as a white cross, like the rest of the good data.

You can see the effects of your masking by using the "extract_aar" and "aar_f" buttons, which will produce a filtered AAR (with the resolution specified in the "resol" window) of the specific section of spectrum that you have selected with band, scan, and x_range. It is recommended to use kappa=2.5-3 and resol>2000, especially for faint sources. Further, you can output your final, masked AAR result to a file by clicking on the "save_aar" button. This action will pop up a window where you can specify the name of the file you wish to write, and what format: XDR, IDL SAVE, or FITS. The first two options both write IDL save files which can be restored in an IA session (the first using the XDR format described in step 4). Repeat for each line.

15) *spd_FM*=**killwait**(*spd_FM*) (help: IA UTILITIES)

aar=**extract_aar**(*spd_FM*) (help: IA AUTO ANALYSIS)

save,*aar*,**file**='*aar.save*',/**xdr**

status=**sap_wfits**('*aar.fits*',*aar*) (help: ISAP)

aar=**shiftscans**(*aar*) (help: IA UTILITIES)

line=**aarfilt**(*aar*,[**7.29**,**7.32**],**1200.**,**kappa**=**3.0**,/**noglitch**)

The first task flags "wait" points as "bad"; it is useful on SWS01, SWS02, and SWS06 data. The next task extracts an AAR from the reduced SPD, and the following two commands save the AAR as an IDL save set and a FITS file. Note that all the masked and glitched points are still in the AAR, but are ignored by shiftscans and aarfilt.

If you have an SWS02 dataset which has multiple scans on the same line, you should use the routine "shiftscans". This command does an additional flatfielding of common scans with respect to each other; it is an "offset" mode correction and is particularly valuable for faint sources. This task is not necessary if the wavelength ranges of your scans do not overlap.

The last task produces a final, filtered AAR for a particular SWS02 scan; you should repeat this step for each individual line scan. In this example, the wavelength range is 7.29-7.32 microns, the resolution is 1200, and kappa is set to 3 (3σ). Save the individual filtered files as the whole AAR was saved above. If you have faulty detectors that were not masked out before, you can do so with this command as well–see the help file.

After this step, the basic reduction of SWS02 data is complete, but there is likely significant fringing, especially for lines lying in bands 2 and 3. These can be difficult to deal with for SWS02 data, as there are only very short baselines over which to measure the fringes. There are different tools you can use to try to deal with fringes, either by trying to remove the fringes or fitting a line shape; these include **linefit** (in ISAP), **aarfringe**, or **multifit**. Consult your IA guru for strategies for recovering as much information as

possible.

Finishing up:

Type "help" to look at the list of IDL structures you have generated in this data reduction section. Make sure to save all files which were produced from the most time-consuming steps (i.e., any of the interactive tools) as IDL save sets and FITS files before exiting from IA. Then you can restore the SPD or AAR at any stage and redo some of the reduction if you find it necessary at a later date.

Reducing AOT SWS06 Data in IA[31]

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NOTE: This document is subject to change due to ongoing IA/ISAP software development.

Before you begin:

This guide is intended to give a basic description of how to use IA tasks to reduce AOT6 data: it assumes that the FITS files containing the raw data have already been retrieved from the archive, and gives only a representative use of each command. This document provides background information for the accompanying IA script. For a full description of input parameters and alternate uses of a particular command, consult the IA help tool. This help tool is accessed by typing a "?" on the command line after starting an IA session. There are many possible help topics, so for each command described below, we have included to the right hand side the help topic where it can be found.

All IDL command syntax in an IA data reduction session is shown in **boldface** below,

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with variable and file names indicated in *italics*. All variable names are meant to be representative, but of course can be chosen by the user. These instructions suggest saving the output SPD after particularly time-consuming steps; the user should weigh the value of additional saves against the constraints of disk space. We also recommend that you work in a separate directory for each of your datasets, to prevent confusion.

Before starting IA, you should close all applications which use color (e.g., Netscape, Mosaic, or additional IDL sessions), as they will interfere with the IA color table. After IA has been started, you may restart these applications without affecting the IA session. In addition, for some tools a better performance is achieved if your window manager is set such that a window is active when the pointer is moved into it, but it is not automatically brought to the foreground. In general, it is advisable to keep the text window from which you started IA accessible, as some error messages from the interactive tools will be displayed in this window (note that some error messages will appear in pop-up windows of their own).

To begin: Start your IA session by typing **ia** at the system prompt. Then follow the instructions below.

1) **cal_select** (help: IA UTILITIES)

This tool is used to select calibration files. See Appendix B.4 of the IA User's Manual for descriptions of each calibration file. The cal_select command opens a window which lists all current calibration files, and there are two possible settings for each CAL file: 'BASE' or 'TEST'. The 'BASE' system contains the CAL file configuration as in the current pipeline version. The 'TEST' system provides CAL files, which were updated since the last pipeline delivery. The individual choice depends on the current status of the CAL file system, so ask an IA guru.

If you run the cal_select tool for the first time, save the recommended CAL file selection as file 'cal_select.dat' in your top directory. To do this, press the 'save' button after the

CAL file selection, edit, if necessary, the 'path' box in the popping up window and confirm path and filename by clicking 'OK'.

If you have already saved the cal_select.dat file in a previous run of cal_select, you can restore the saved setting by clicking the 'load' button. NOTE: the 'path' box must be filled in properly, otherwise cal_select will crash.

After either setting (and saving) the appropriate CAL file selection manually or loading your calibration file, click 'OK' to exit the cal_select tool. All CAL files selected from the 'TEST' system are then listed in your test window.

2) *erd*=**read_ferd**('xxxYYYzz',**path**=[.data]) (help: IA INPUT OUTPUT)

This command reads in the ERD from the raw data files. The 8 digits in parentheses refer to the revolution number (xxx), target number (YYY), and observation sequence number (zz) from the archive listing. You may also use the command "read_fspd" to read in the SPD directly and skip to step 5; however, the pipeline reduction is ever-changing, and starting with the ERD you allows you take advantage of the most recent version of the reduction software and calibration files.

3) *spd*=**dspd**(*erd*) (help: IA DERIVE SPD)

This command converts the ERD to an SPD, using the following CAL files: (1; 2; 2a,b; 3; 4; 5; 6; 12; 16a,b,c,e; 18; 23; 24).

4) *spd_orig*=**read_fspd**('*SWSPxxxYYYzz.FITS*') (help: IA INPUT OUTPUT)

spd.header=*spd_orig.header*

spd.data.det.flag=*spd_orig.data.det.flag*

save,*spd*,**file**='*spd.save*',/**xdr**

Unfortunately, the SPD header and detector flags (esp. the reference scan flag of SWS06) are not properly filled by dspd; the three commands in this step correct this problem by reading in the SPD (spd_orig) directly and copying its header and the detector flags into your recently derived SPD (spd). The filename given to read_fspd is in the format of files coming directly from the archive, where 'xxxYYYzz' are the same digits entered in step 2 above. The fourth command saves the SPD into an IDL save set, which can be restored in later sessions of IA if necessary.

In that case, typing

restore,'*spd.save*' (help: ROUTINES)

will restore the IDL structure *spd* to your IA session.

NOTE: An SPD is rather large, and the use of the option /xdr makes the saving process significantly faster. The /xdr (for eXternal Data Representation) option is not necessary to save smaller quantities such as the sctbl or aotlr alone (e.g., step 6), but it is needed if you wish to export any save sets between VMS and UNIX machines (see the IDL help file on "save" for more information.)

5) **list_header**,*spd* (help: IA UTILITIES)

set_plot,'**ps**

device,**file**='*aper_plot.ps*',/**landscape**

aper_plot,*spd*,'1A',60. (help: IA UTILITIES)

aper_plot,*spd*,'3A',60.

aper_plot,*spd*,'3E',60.

aper_plot,*spd*,'4',60.

device,/close

set_plot,'x

These tasks give you more information about your dataset. The list_header command lists the FITS header, and in particular, you should note the position angle INSTROLL of the y-axis of the ISO satellite during your observation (see ISO Observer's Manual for the definition of the ISO coordinate system). The position angle of the long side of the SWS aperture is then defined by: INSTROLL + 90 degrees.

The remaining commands in this step create a postscript file containing plots of the four SWS apertures on the sky (for bands 1 & 2; 3ACD; 3E; and 4). The value of 60. above is the declination scale of the plot in arcseconds. Consult your local system manager for printer options to determine how to print the postscript file locally.

6) *sctbl*=**make_sctbl(***spd***)** (help: IA UTILITIES)

print_sctbl,*sctbl*,**file='***sctbl.txt***',/spawn**

save,*sctbl*,**file='***sctbl.save***'**

These commands create, print out, and save the scan table, which gives such information as the sequence of observations (darks, photometric checks, reference scans and science scans) and wavelengths covered by each scan.

The reference scans are specific for SWS06. They are short, pseudo-science scans, which were thought for additional calibration purposes, but never used in SWS06 reduction. Since summer 1997 they are no longer part of SWS06 observations. Usually reference scans are not passed into the AAR, unless the keyword /all is used in the 'daar' or 'extract_aar' command. Unfortunately, due to software problems, this keyword has currently to be used. The reference scans have therefore to be sorted out on AAR level. WARNING: for high fluxes, where memory effects play an important role, the continuum level of reference

scans after dark measurements is too low. In combination with proper science scans they therefore might create fake features, if they would be included in the AAR.

The print_sctbl command prints the scan table to the screen as well as to the specified file name. The /spawn option is a local MPE command to print the specified text file to the default printer; again, consult your local system manager for printer options.

7) *spd_mem*=**antimem**(*spd*) (help: IA AUTO ANALYSIS)

This routine is designed for reducing memory effects, as is done in the standard pipeline. The use of antimem is mostly historical, but should be kept in the reduction chain because this task may in the future include a more complicated method for dealing with memory effects.

8) *spd_D*=**dark_inter**(*spd,sctbl*,**/multia**) (help:IA UTILITIES)
save,*spd_D*,**file**='*spd_D.save*',**/xdr**

NOTE: Usually the server or window system is configured to retain plot window after scrolling or the start of a screen saver. The /multia option is only needed if the draw window is not retained (as on a Multia terminal); it causes IDL to handle the plot storage directly.

The dark_inter tool facilitates dark current correction. It can be the most time-consuming of all data reduction steps, but is also one of the most important in producing reliable spectra. Make sure to save the SPD after completing this task. Before you begin, you should note the sequence of darks and other scans in the scan table. Note that the reset time ("tr") and gain columns in the scan table must be the same for a dark and the scan from which it is being subtracted; dark_inter automatically makes available all darks with matching tr and gain. Also check the scan table to determine which (band+scan) combinations hold actual data: those which do not will show "0.000" for both wave_min

and wave_max (this information is also shown in the message box in dark_inter as "w_range"). It is not necessary to dark subtract these (band+scan) combinations. For each of the four bands (starting at detectors 1 [SW1:], 13 [SW2:], 25 [LW1:], and 37 [LW2:]), the following sequence should be followed:

A.Select a detector. This detector will represent the whole band.

B.Revise Darks. Click "Revise Darks". In the "Revise Darks" window which pops up:

→ Select the first set of darks. This may be done either by clicking "next", which will automatically choose the first set of darks if no dark has been previously selected in the given band, or by clicking the "select" button first and then clicking the dark you want in the middle left panel.

→ choose clipping options–here are the most general suggestions:

▷ first 1 point and clipping(mean) with kappa=3 (i.e., 3σ clipping around the mean).

▷After strong photometric checks, you may want to use clipping (linear) instead of mean to deal with obvious memory effects (i.e., jump in signal followed by a decline in signal over time).

→ refine dark to be subtracted for each detector of the band, for each scan:

▷"clip" shows the points to be clipped from the first detector (top right window): two vertical lines show the kappa-clipping limits you have specified, and all points above or below these lines are clipped.

▷"apply to whole band" shows the effect of the current choice for all detectors in the band.

▷many iterations may be tried (use "undo" between each to cancel effects of previous try), but in the end, the combination of "apply to whole band" and finally "apply" actually applies the clipping to the current set of dark measurements. NOTE: the use of "clip" and "apply to whole band", without clicking the "undo" button in between, results in the first detector being clipped twice.

→ Click on "next" to select the next dark, and repeat the "revise darks" steps for all darks in the requested band. If you did not "apply" the latest clipping to the dark before moving to the next dark, you will get an error message in the text window reminding you to do so.

→ Click "done" in the "Revise Darks" window when all darks in the band have been revised.

C. Subtract darks from photometric checks and science scans:

→ Select the photometric check or science scan to be dark subtracted by indicating its number in the "phot selection" or "scan selection" box. Your selection will be indicated in green in the upper two windows of the left panel. Make sure that the entire selected check or scan can be seen in the left column of windows before proceeding. If necessary, click on "plot" to access a window where you can adjust the plot parameters, or use the slider bar at the bottom of the left panel to adjust the horizontal range. Note that the plot window may stay open during the entire use of dark_inter, so that you may readjust as necessary.

→ Choose the dark subtraction method: click on "Method".

▷In the "method" menu, choose which darks to use. The darks with matching tr and gain will be highlighted in yellow-orange. If you only have one dark before the check or scan which can be used, you should use "previous" and "mean" for the method. If you have valid dark measurements (that is, with the same value of "tr" and gain) before and after a scan, you can choose to do a linear interpolation between the two. If the two darks come immediately before and after the scan, select them with the "previous" and "next" buttons, and click "linear interpolation" for the method.

▷When you have successfully marked the dark in question, it will be shown in red. If you get strange fits or error messages when trying to subtract a dark, it may be because the dark was not properly selected; go back to the "Method" menu and toggle your dark choices until they appear in red in the left plot windows.

▷ Choosing other darks: If you wish to choose a dark which comes after an intermediate dark for a linear interpolation fit, you must click on "cursor" and then on the appropriate dark measurement in the middle window of the left panel to select it–the "next" button only points to the very next dark, regardless of whether or not it can be used. The "cursor" option is very location-sensitive, so you may have to click a few times (and get a few warning messages). If you change your mind and decide not to use a selected dark, click "cursor" and the dark again to unselect it. Be sure that the dark points are colored yellow-orange again to be sure you were successful.

▷If the available darks do not provide a good dark measurement to be subtracted, you may do a spline fit instead of a mean or linear interpolation fit. For now, just choose "spline". NOTE: The spline fit must be done for each detector individually, and is much more time consuming; use only when necessary.

▷ Click "Done" in the Method menu.

→ Subtract fitted dark current.

▷Click on "fit", which will show the fit for the selected detector in the bottom left plot.

▷If the method selected is "mean" or "linear interpolation", all that remains is to click on the "apply to whole band" button. You may iterate through the "method", "fit", and "apply to whole band" sequence if desired. If the dark subtraction is satisfactory for all detectors in this band/scan combination, you have completed the subtraction.

▷However, often "jumps" are seen in the data. If these are simply horizontal steps, then it is advisable to use either the previous or next dark to fit the longest section and then mask out the other section later in the mask_inter tool. To make a correction to the dark subtraction for one detector, use the slider at the top of the tool to select the detector, and then set the scan number in the window below. Using the "fit" button writes a new solution for that detector, and doesn't affect the other detectors.

▷If the method selected is "spline", clicking the "fit" button produces instruction messages in the IA text window which explain how to define spline points, fit the spline, iterate if the first fit is not optimal, and apply the fit to the data. Spline points should be defined on the center left plot. Repeat the "fit" step for each individual detector in the band.

▷Additional notes for spline fits: Previous fits are left on the plot, and can be used to guide a better spline fit, if necessary. Closely spaced spline points are likely to cause a highly structured fit; widely spaced spline points are better for this purpose. You will need to enter the scan number again each time you increment the detector number and want to perform a new spline fit. You will also need to replot each time if you used special plot parameters to get the selected scan to fit in the left windows.

→ Repeat Step C for each photometric check and science scan in this band.

D. Set slider on top to the first detector number of the next band and repeat the process, starting with Step B, Revise Darks. Complete Steps B and C for all four bands. It is a good idea to save the data set periodically during the use of dark_inter, to prevent loss of efforts in the event that the tool crashes. The easiest way to do this is to use the "save current" button in the dark_inter tool, which creates a file called "dark_inter_save.xdr" containing the current data file. This file is overwritten in successive uses of this button, so you may make incremental saves without using additional disk space.

9) *spd_DM*=**mask_inter**(*spd_D,sctbl*) (help: IA UTILITIES)

save,*spd_DM*,**file='***spd_DM.save*'**,/xdr**

This command puts you into the mask_inter tool, which can be used to inspect your SPD

and mask out bad data. Sometimes a whole detector is significantly noisier than the others in the band, or there may be significant memory effects after a strong glitch. In addition, you can mask out the jump segments first detected with dark_inter (see step 8C). See the on-line help for this tool for a detailed account of all the utilities of this tool.

Enter a band number in the "band" box, click on the scan number you wish to see, and click "disp_spd" to display a subset of the data. Note that to examine a different scan, you must click on the previous scan number again to "unset" it; otherwise there will be two scans selected and the tool will complain (multiple scans can be selected for producing an AAR within mask_inter, but only one scan can be displayed at a time). If you click the "plotmode" button, a background window will pop up in which you can set the plot scale; we recommend that you choose a fixed y-scale, as this is a good way of finding detectors for which dark subtraction didn't work properly, or for finding noisy detectors or other strange features. Each time you change the plot scale, click "disp_spd" again to display the data with the new plot values.

To mask out a specific section of a detector, click "select" and the detector of interest, which will cause the data from that detector to show in the bottom left window. Then click "mask" and use the left mouse button to select the region to be masked out (e.g., in the case of a memory effect). If you see the same glitch in all 12 detectors of a band, you can use the "do to all" button after masking out the glitch in one of the detectors. If you judge the input from an entire detector to be bad, you can click on the detector number in the list to the right-hand side of the display, and then click "faulty" to set it all bad.

IMPORTANT NOTE: Using the "faulty" button to set a detector "bad" is NOT reversible at the current time, and sets the specified detector bad for all scans. Setting the detector bad by masking all points out using the "mask" button IS reversible (using "unmask"), and only sets the detector bad for the specified scan.

Saving the masked SPD should be considered if a significant amount of masking has been done.

10) *spd_U*=**aas_updowncal**(*spd_DM*) (help: IA AUTO ANALYSIS)

Currently no interactive tool to correct for differences between SWS06 'up' and 'down' scans is available. Therefore the pipeline routine is used, which, due the lack of an proper automatic up-down calibration procedure, essentially does nothing to the data.

A patch is available for SWS02-like, short range scans with SWS06, of e.g. spectral lines, which were too broad for SWS02 line scans. Here an additional pseudo scan table sctbl_pseudo_S02.txt is created, which pretends the SWS06 DARK–UP–DOWN–DARK sequences to consist of SWS02 DARK–UPDOWN–DARK scans. Additionally a procedure file sctbl_6to2_patch.pro is created, which e.g. masks the data points inbetween the SWS06 UP–DOWN scans. A very careful inspection of the pseudo scan table and the procedure file is strongly recommended before use!

10') *sctbl_pseudo_S02*=**make_sctbl**(*spd*,**undoc=2**) (help: IA UTILITIES)
print_sctbl,*sctbl_pseudo_S02*,**file=**'*sctbl_pseudo_S02.txt*'
@sctbl_pseudo_S02
spd_U=**updown_inter**(*spd_DM*, *sctbl_pseudo_S02*) (help: IA UTILITIES)
save,*spd_U*,**file=**'*spd_U.save*',**/xdr**

We refer the reader to the document 'Reducing AOT SWS02 Data in IA' for a description of the updown_inter routine.

11) **plot_phot**,*spd_U*,*sctbl*,[1],13,**/all** (help: IA UTILITIES)
plot_phot,*spd_U*,*sctbl*,[1],25,**/all**
plot_phot,*spd_U*,*sctbl*,[1],37,**/all**

The routine plot_phot can be used to look at photometric check data. The command lines given above request IA to plot the first photometric check for all 12 detectors of bands 2,

3, and 4 (which start with detector numbers 13, 25, and 37); the band 1 photometric check is not used. If you have more than one photometric check, you can replace the [1] above with a range of values (e.g. [1,3]) and compare them on the same plot. By specifying a particular detector and removing the /all flag, you can look at the photometric checks for that detector.

Data points already flagged as bad by the system will be marked (see help file for color code). If these are the only bad points and the photometric checks look sensible, the standard routines will handle the bad points appropriately. Note that the default is to always use the first photometric check, so if it is corrupted it should be replaced with another. If you suspect your photometric data has problems which are severe enough to affect the ultimate calibration, such as a corrupted first photometric check or large spikes not already flagged, consult an IA guru.

12) *spd_U*=**respcal**(*spd_U*) (help: IA AUTO ANALYSIS)
spd_U=**fluxcon**(*spd_U*)
spd_U=**velcor**(*spd_U*)

The routines respcal and fluxcon perform responsivity calibration and flux conversion and apply various calibration files; respcal uses CAL files 13, 19, and 25; fluxcon uses CAL file 13. The output of fluxcon looks like the following:

```
Warning(SPCP): some phot. check data might be doubtful       Det:26 Uv: 2
Warning(SPCP): some phot. check data might be doubtful       Det:39 Uv: 2
Warning(SPCP): some phot. check data might be doubtful       Det:46 Uv: 3
Warning(SPCP): some phot. check data might be doubtful       Det:48 Uv: 2
```

If the number following "Uv:" (i.e. the unvalid data points) is greater than 3, the photometric check data for that detector indicated may be suspect.

The task velcor performs a correction to the heliocentric frame based on the velocity of the satellite. The velocity is taken from the SPD FITS header. 0.0 km/s are applied, if the SPD header is not properly filled (cf. step 4).

13) *spd_F*=**flat_inter**(*spd_U,sctbl*) (help: IA UTILITIES)
save,*spd_F*,**file**='*spd_F.save*',/**xdr**

The first command starts up the flatfield correction tool flat_inter. This task corrects the flatfield of each detector in the band with respect to the others. As with "updown_inter", the correction can be a "gain" or "offset" correction, and is made on the basis of smoothed data. It is generally recommended to use the "median" filter, and to exclude poor points. As with updown_inter, you should make sure that the fit you have chosen does not introduce spurious features, and vary the smoothing or correction parameters accordingly. The default values are Res1=500 and Res2=100; try lower values, especially for band 4 (LW2:) for very faint sources, and larger values may be useful for very bright sources. In general, it is a good idea to keep the relation Res1$\sim$5 Res2. As before, the 'gain' option should only be used with strong signal. NOTE: be sure to press "return" after changing each entry; otherwise, strange effects may result if new values are not registered by the tool.

Faulty detectors might be specified with the faulty=[...] keyword, which contains the numbers of the bad detectors as a vector. These detectors are flatfielded anyhow, but not used for the definition of the flatfield itself.

14) *spd_FM*=**mask_inter**(*spd_FM,sctbl*) (help: IA UTILITIES)
save,*spd_FM*,**file**='*spd_FM.save*',/**xdr**

Now that all calibrations have been applied, you can go back into the interactive masking tool used in step 9 to mask out noise spikes or outliers; for instance, a signal which is clearly spurious, such as a strong spike detected only in the up or down scan of all 12

detectors, will bias the resulting spectrum. This masking step may be best done after following the rest of the steps described below and looking at an unmasked version of the output spectrum to see where lines have been detected, and returning to this step to produce a refined dataset.

Click the "w-mode" button to plot in wavelength units, and specify the wavelength range of interest in the "plotmode" pop-up window. You can use the "fluxclip" button to mask out outlying data by hand. When you click this button, the data from all detectors is plotted in the lower left hand window, and you can define a series of boxes with the left mouse button; the points inside these boxes are then clipped. Clicking the right mouse button returns you to the main mask_inter tool. If you want to remove an unwanted glitch (glitches are marked with yellow asterisks), click "select" to specify the detector, and if needed click "zoom" and use the cursor to select the section of the scan holding the offending point (both up and down scans will be shown, with wavelength units along the upper x axis of the plot). Then click "unglitch", and click the mouse cursor on either side of the glitched point. The unglitched point will show up as a white cross, like the rest of the good data.

You can see the effects of your masking by using the "extract_aar" and "aar_f" buttons, which will produce a filtered AAR (with the resolution specified in the "resol" window) of the specific section of spectrum that you have selected with band, scan, and x_range. It is recommended to use kappa=2.5-3 and resol>2000, especially for faint sources. Further, you can output your final, masked AAR result to a file by clicking on the "save_aar" button. This action will pop up a window where you can specify the name of the file you wish to write, and what format: XDR, IDL SAVE, or FITS. The first two options both write IDL save files which can be restored in an IA session (the first using the XDR format described in step 4). Repeat for each line.

15) *spd_FM*=**killwait**(*spd_FM*) (help: IA UTILITIES)

aar=**extract_aar**(*spd_FM*,**/all**) (help: IA AUTO ANALYSIS)

save,*aar*,**file='***aar.save*','**/xdr**

status=sap_wfits('*aar.fits*','*aar*)** (help: ISAP)

range=**aarfilt(***aar*,**[6.92,7.32],1200.,kappa=3.0,/noglitch)**

The first task flags "wait" points as "bad"; it is useful on SWS01, SWS02, and SWS06 data. The next task extracts an AAR from the reduced SPD. The keyword /all, which has to be used due to current software problems, makes sure that all scans, including the reference scans are passed to the AAR (cf. step 6). The following two commands save the AAR as an IDL save set and a FITS file. Note that all the masked and glitched points are still in the AAR, but are ignored in aarfilt on request by keyword /noglitch.

The last task produces a final, filtered AAR for a particular SWS06 scan; you should repeat this step for each individual range. In this example, the wavelength range is 6.92-7.32 microns, the resolution is 1200, and kappa is set to 3 (3σ). Save the individual filtered files as the whole AAR was saved above. If you have faulty detectors that were not masked out before, you can do so with this command as well–see the help file.

After this step, the basic reduction of SWS06 data is complete, but there is likely significant fringing, especially for lines lying in bands 2 and 3. These can be good to deal with for SWS06 data, if they cover a long baseline containing many fringes. There are different tools you can use to defringe, either by trying to remove the fringes or fitting a line shape; these include **linefit** (in ISAP), **aarfringe**, or **multifit**. Consult your IA guru for strategies for recovering as much information as possible.

Finishing up:

Type "help" to look at the list of IDL structures you have generated in this data reduction section. Make sure to save all files which were produced from the most time-consuming steps (i.e., any of the interactive tools) as IDL save sets and FITS files before exiting from IA. Then you can restore the SPD or AAR at any stage and redo some of the reduction if you find it necessary at a later date.

Reducing AOT SWS07 Data in IA[31]

E. Sturm

and

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and

H. Feuchtgruber

and

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NOTE: This document is subject to change due to ongoing IA/ISAP software development.

Before you begin:

This guide is intended to give a basic description of how to use IA tasks to reduce AOT7 data: it assumes that the FITS files containing the raw data have already been retrieved from the archive, and gives only a representative use of each command. This document provides background information for the accompanying IA script. For a full description of input parameters and alternate uses of a particular command, consult the IA help tool. This help tool is accessed by typing a "?" on the command line after starting an IA session. There are many possible help topics, so for each command described below, we have included, to the right hand side, the help topic where it can be found.

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All IDL command syntax in an IA data reduction session is shown in **boldface** below, with variable and file names indicated in *italics*. All variable names are meant to be representative, but of course can be chosen by the user. These instructions suggest saving output after particularly time-consuming steps; the user should weigh the value of additional saves against the constraints of disk space. We also recommend that you work in a separate directory for each of your datasets, to prevent confusion.

Before starting IA, you should close all applications which use color (e.g., Netscape, Mosaic, or additional IDL sessions), as they will interfere with the IA color table. After IA has been started, you may restart these applications without affecting the IA session. In addition, for some tools a better performance is achieved if your window manager is set such that a window is active when the pointer is moved into it, but it is not automatically brought to the foreground. In general, it is advisable to keep the text window from which you started IA accessible, as some error messages from the interactive tools will be displayed in this window (note that some error messages will appear in pop-up windows of their own).

To begin: Start your IA session by typing **ia** at the system prompt. Then follow the instructions below.

1) **cal_select** (help: IA UTILITIES)

This tool is used to select calibration files. See Appendix B.4 of the IA User's Manual for descriptions of each calibration file. The cal_select command opens a window which lists all current calibration files, and there are two possible settings for each CAL file: 'BASE' or 'TEST'. The choice will depend on the current status of the calibration files, so ask an IA guru.

It is possible to load a cal_select.dat file (a binary file saved out of a previous run of cal_select) by clicking the 'load' button. NOTE: if using the 'load' option, the 'path' box must be filled in or cal_select will crash.

After either setting the appropriate CAL files manually or loading your calibration file, click 'OK' to exit the cal_select tool.

2) *erd*=**read_ferd**('xxxYYYzz') (help: IA INPUT OUTPUT)

This command reads in the ERD from the raw data files. The 8 digits in parentheses refer to the revolution number (xxx), target number (YYY), and observation sequence number (zz) from the archive listing. You may also use the command "read_fspd" to read in the SPD directly and skip to step 5; however, the pipeline reduction is ever-changing, and starting with the ERD allows you to take advantage of the most recent version of the reduction software.

3) *spd*=**dspd**(*erd*) (help: IA DERIVE SPD)

This command converts the ERD to an SPD, using the following CAL files: (1; 2; 2a,b; 3; 4; 5; 6; 12; 16a,b,c,e; 18; 23; 24).

4) *spd_orig*=**read_fspd**('*SWSPxxxYYYzz.FITS*') (help: IA INPUT OUTPUT)
spd.header=*spd_orig.header*
save,*spd*,**file=**'*spd.save*'**,/xdr**

Unfortunately, the full header is not properly saved by dspd; the two commands in this step correct this problem by reading in the SPD (spd_orig) directly and copying its header into your recently derived SPD (spd). The filename given to read_fspd is in the format of files coming directly from the archive, where 'xxxYYYzz' are the same digits entered in step 2 above. The third command saves the SPD into an IDL save set, which can be restored in later sessions of IA if necessary. In that case, typing

restore,'*spd.save*' (help: ROUTINES)

will restore the IDL structure *spd* to your IA session.

NOTE: An SPD is rather large, and the use of the option /xdr makes the saving process

significantly faster. The /xdr (for eXternal Data Representation) option is not necessary to save smaller quantities such as the sctbl or aotlr alone (e.g., step 6), but it is needed if you wish to export any save sets between VMS and UNIX machines (see the IDL help file on "save" for more information.)

5) **list_header,***spd* (help: IA UTILITIES)
set_plot,'ps
device,file='*aper_plot.ps***',/land**
aper_plot,*spd*,'5A',60. (help:IA UTILITIES)
aper_plot,*spd*,'5C',60.
aper_plot,*spd*,'6',60.
device,/close
set_plot,'x

These tasks give you more information about your dataset. The list_header command lists the FITS header, and in particular, you should note the position angle of your observations: the real position angle is (roll angle) + 90 degrees. The remaining commands in this step create a postscript file containing plots of the four SWS apertures on the sky (for bands 1 & 2; 3ACD; 3E; and 4). The value of 60. above is the declination scale of the plot in arcseconds. Consult your local system manager for printer options to determine how to print the postscript file locally.

6) *sctbl*=**make_sctbl(***spd***)** (help: IA UTILITIES)
print_sctbl,*sctbl*,**file='***sctbl.txt***',/spawn**
save,*sctbl*,**file='***sctbl.save***'**

These commands create, print out, and save the scan table, which gives such information as the sequence of observations (darks, photometric checks, and science scans) and wavelengths covered by each band. The print_sctbl command prints the scan table to the

screen as well as to the specified file name. The /spawn option is a local MPE command to print the specified text file to the default printer; again, consult your local system manager for printer options.

7) *aar*=**daar(***spd*,**/all)** (help: IA AUTO ANALYSIS)
save,*aar*,**file='***aar.save*',**/xdr**
status=sap_wfits('*aar.fits*',*aar*) (help: ISAP)

This task extracts an AAR from the reduced SPD, and the following two commands save the AAR as an IDL save set and a FITS file.

8) *index*=**where(***aar.data.det* **le 48)**
aar_aot6=**cut_aar(***aar, index*) (help: ISAP)
status=sap_wfits('*aar_aot6.fits*',*aar_aot6*) (help: ISAP)

If your F-P observation is merged with a grating observation (short wavelength section) it is recommended to split the AAR and to cut out the grating part for subsequent processing. Grating observations merged with F-P observations are always of type AOT6. The first command finds the index of all data belonging to detectors 1 to 48, i.e. to the grating part. F-P observations are made with detectors 49 (SW) and 51 (LW). Detectors 50 and 52 are redundant detectors which are not used for observations. The second command extracts a "sub-AAR", containing only the AOT6 part. The F-P part will be extracted in step 9. The last command saves the AOT6 AAR as a FITS file (e.g. for subsequent processing with ISAP).

9) *index*=**get_aotb_index(***aar,'5B',stat*) (help: ISAP)
subind=**where(***aar.data(index).wave* **ge 13 and $**

aar.data(index).wave **le 14)**

aar_fp=**cut_aar**(*aar, index(subind)*)

The order and aperture information in the scan table tells you which of the F-P bands 5A, 5B, 5C, 5D and 6 are present in your observation. See Table 1. If you have several different lines in one AOT band (see the scan table) you should further split the AAR. Subsequent processing is easier if there is a separate AAR for each spectral line. This splitting is done using the sequence get_aotb_index - where - cut_aar. In the above example first the index of all band 5B data and then the index of all data points of band 5B within the wavelength range (13-14 μm) is determined. Again "cut_aar" is used to extract a sub-AAR containing the desired data only.

Table 1: Relation between aperture, order, AOT band and wavelength for SWS07 detectors

Aperture	Order	AOT Band	Wavelength
1	3	5A	11.4 - 12.2
1	2	5B	12.2 - 16.0
2	2	5C	16.0 - 19.0
2	1	5D	19.0 - 26.0
3	1	6	26.0 - 44.5

10) **aar_fp.data.sdir=1**

The scan direction for F-P data is always 1, i.e. "up". Due to a software problem (which will be solved, soon) some of the valid F-P data sometimes are assigned a scan direction -1 ("down"), hence, would not be treated correctly in some of the following processing steps. To make sure that all data are included, simply overwrite the sdir tag of the AAR structure with a 1.

11) *flat_ref*=**sws_rebin(***aar_fp*,**resol=10000,over=6,lines=0, $**
method='mean') (help:IA UTILITIES)

In this step a reference spectrum for the flatfielding in step 12 is created. There are several ways to do this. In the above example the reference is created through a rebinning of the data itself to a new grid of relatively low resolution. An alternative would be to use one of the scans which you consider a good one. In an AOT7 several scans of a line are counted as different lines, so you can select the scan via its associated line number:

good_line=**3**

index=**where(***aar_fp.data.line* **eq** *good_line***)**

flat_ref=**define_aar(data=***aar_fp.data(index)***)** (help:IA UTILITIES)

The "define_aar" command creates a new AAR, in this case containing only the line 3 data. (alternatively "cut_aar" could be used).

12) *aar_flat*=**sws_flatfield(***aar_fp***,2,ref=** *flat_ref***,/scaling)** (help:IA UTILITIES)

plot_aar,*aar_fp***,ps=1**

plot_aar,*aar_flat***,ps=2,/oplot**

Now the data are "flat fielded", i.e., to remove remaining differences in the relative flux calibration, the different scans are shifted to a common level. This common level is defined through the reference spectrum, which was created in the last step. The routine tries to fit polynomials of a given order (in this example it is 2) to the different scans. The correction is then done as shifting in an additive way (default) or as a scaling (used in this example). Scaling should be used only for strong sources where additive offsets due to uncertainties in the dark current subtraction can be neglected. The two plot commands allow you to check the result. First the original data are ploted (as + symbols) and then the flat fielded data are overplotted (as * symbols).

13) *aar_clip*=**sigclip(***aar_flat***,resol=30000,lines=0,stop=0, $**

sigma=2.5,/nodata) (help:IA UTILITIES)

plot_aar,*aar_flat***,ps=1**

plot_aar,*aar_clip***,ps=2,/oplot**

Sigma clipping is recommended to get rid of outlyers and, to a certain extent, glitch tails. For each resolutin bin the median and standard deviation are calculated. Each point off by more than SIGMA standard deviations is flagged as NODATA (but not yet removed). By specifying the option "/nodata" all the corresponding data points are actually removed ¿from the data set. The two plot commands again allow a check of the result like in the previous step.

14) *aar_reb*=**sws_rebin(***aar_clip*,**resol=30000,over=6,lines=0, $**
method='mean')
save,*aar_reb*,**file='***aar_reb.save*','**/xdr**
status=sap_wfits('*aar_reb.fits*','*aar_reb*)

A final rebinning can be used to average all scans into a single, smoother spectrum. The rebinning can be performed in various ways (mean, median, flux conserving, weighted or not, etc.). Please consult the on-line help for a detailed description of this routine. For the high F-P resolution there is not much difference between "mean" and "flux conserving".

Finishing up:

Type "help" to look at the list of IDL structures you have generated in this data reduction section. Make sure to save all files which were produced from the most time-consuming steps (i.e., any of the interactive tools) as IDL save sets and FITS files before exiting from IA. Then you can restore the SPD or AAR at any stage and redo some of the reduction if you find it necessary at a later date.