IRAF TUTORIAL SESSIONS

Exercise 1.

This is a short exercise that should help acquaint you with the basics of IRAF.

Starting up IRAF can be a bit complicated (you have to be in the right directory, using the right kind of terminal window, etc.) but on our local machines here there is a script to take care of all of this for you - just type "xiraf" and it should start up iraf in a new window, as well as opening an image display program called ds9. (If this is the first time you've run iraf, it will ask you about creating a login.cl [which is the main iraf user preferences file] - just hit return to accept the default action.)

In this, as in the other exercises, the commands after the prompt (either % or cl>) are meant to be typed by the user; the #-sign indicates a comment.

The exercise uses two images, with file names im010.fits and im011.fits, which you may already find in your ~/iraf directory (which is the directory you're already in by default if you started IRAF as indicated above). If they aren't there, you can copy them to your current directory with

cl> cp /data/astro121/iraf_intro/im*.fits .

Notice that there is a period at the end of that command, separated from the filename by a space - that looks like punctuation, but in fact it is part of the command, since '.' is linux-speak for 'the current directory'. Also, if you know a little unix/linux, you may recognize that this is just the standard linux cp command - we have IRAF configured locally to pass a common subset of linux commands directly out to the shell to be executed. For those that aren't pre-defined, you can always execute a linux command from within IRAF by preceding it with a !, as in

cl> ! emacs file.log & # Starts an emacs window under linux.

Our local installation of IRAF is configured to work directly with FITS files by default, where "FITS" stands for Flexible Image Transport System, a standard astronomical image format. Older versions of IRAF (V2.10.4 and earlier, or anything earlier than the end of 1997) worked with a different format where each file was split into a separate header and pixel files. If you encounter files with extensions .imh and .pix, these are IRAF formatted image header and pixel files, respectively.

cl> dir # list the files in the current directory # and make sure you have the .fits files. The next step is to familiarize yourself a bit with some of the basic operations of IRAF. Tasks with similar/related functionality are grouped together in packages. Tasks, and sometimes packages, have parameter files that control their operation.

cl>	diskspace	#	show the available diskspace
cl>	path	#	show the current working directory
cl>	dir	#	list the files in your current directory
cl>	dir *.fits	#	list just the fits files in your directory
c1>	dir uparm	#	list the files in the uparm directory - a
			place where parameter files are stored
c1>	dir l+	#	do a long directory listing
c1>	nackage	#	show what packages are currently loaded
c1>	package	# #	baln for the nackage "images"
CL/	nelp images	# #	help for tack "imboder" using a more
([>	pherp imneader	#	sophisticated pager - it lets you:
	<space bar=""></space>	#	go to next page of help
	b	#	go back one page
	?	#	view options
	a	#	quit the help for this task
cl>	lpar imh	#	this doesn't work (trv it!) - task names
-	1		and parameters need enough characters
			specified to be identifiable
c1>	lpar imheader	#	note query and hidden parameters - query
	ipur imieudei		narameters are listed first with hidden
			parameters following in parentheses
c1>	uploarn imboad	#	unloarn parameters go back to defaults
c1>	imbood im010	# #	chart hander listing what does it tall you?
CL/	imbood	# #	by default imbader will list all images with
C[>	Imneau	#	by default immedder will list all images will
<u>_</u> 1	impaad im010 l	щ	Line extensions listed in the parameter intist
CL>	Timeau Timere L+	#	Long version of the output - shows all
			lieduer lietus
cl>	epar imhead	#	modify the task so it always types the long
	·		header listing - choose yes for the parameter
			entry for "longheader" followed by a return
	: q	#	<pre># exit eparam mode and "save" the new parameters</pre>
	•		[exiting with crtl-C does NOT update the
			parameter file] You can also exit and
			save with ctrl-D.
cl>	lpar imhead	#	was the edited parameter saved?
cl>	imhead im011 pag	ge	·
c1>	dir uparm	#	what is the uparm directory used for? Can you
			find the parameter file for "imheader"? The
			name is a combination of the package (imutil)
			and task name i e imlimhear par
c1>	unlearn imhead	#	go back to the default parameter setting
$c_{1>}$	dir unarm	#	what just hannened?
c1>	imhead im010 1+ 1	π n ⁻	and just happened:
CL/		н #	note the keyword "evotime"
c1>	unlearn bealact	#	HOLE LIE REYNOLD EXPERIME
c1>	lnar healact		
CL/	ipai iiselett		

What have you learned about parameters? What is the difference between query and hidden parameters? What is the uparm directory? Do you understand the different syntaxes that we have been using in task executions? If you have questions see the Beginner's Guide mentioned below in the References section.

A Short Exercise

There are 2 images in your directory, im010 and im011, of the same field, but one frame has been slightly shifted from the other. We want to shift the second frame so it aligns with the first frame, and then average these two frames together.

cl> dir im*.fits	<pre># note the two frames</pre>
<pre>cl> imhead im*.fits</pre>	# check to see if they are the same field
cl> unlearn display	
cl> lpar display	<pre># look at the parameters for the display task</pre>
cl> display im010 1	<pre># load the first image into the image display</pre>

Let us diverge for a moment. This is probably a good time to acquaint yourself with the image display program ds9, if you have not used it before. You should know how to zoom, pan, window, and blink. Most options can be done with the mouse. Notice the pull down menus at the top of the window. Pull each menu down and review the selections use the left mouse button to pull down the menus. At the center of the window are some point and click options - clicking in the top row of buttons changes what buttons are available in the second row for doing various tasks (e.g. clicking the "Zoom" button displays "+", "-", "to fit", etc.). At the top of the window are rough coordinates of the current cursor position as well as an estimate of the pixel value at that cursor position. The two boxes in the upper right corner are "panner" boxes - one shows you what part of the image you are currently looking at (right now we are displaying the full image), and one shows you a zoomed-in view under the current mouse position.

The lookup table can be modified by moving the cursor up and down in the window (contrast) or left and right (brightness) while holding down the right mouse button. Try it. (You can always get back to default settings by clicking "Frame", and then "reset".)

Panning (changing which part of the image you're displaying) is done with the middle mouse button. Move the cursor to an interesting part of the image - click once with the middle mouse button and that part of the image will be centered. You can also drag the blue box in the top panner window to move the region displayed. To zoom in, choose the Zoom button, then click "+" or "-" to zoom in or out; you can also zoom in and out with the scroll wheel on the mouse. Play with this until you understand how it works. Go back to the unzoomed image when you are finished.

We want to blink this field and that of im011. So we will need to load the second image into the second frame buffer - you have 4 frame buffers available for images.

cl> display im011 2

Now try to match the lookup tables of both images. Adjust the contrast and brightness for the second image that you should be looking at now. (Remember, just click and drag with the right mouse button.) Twiddle things until you have black stars on a white sky background. Now look at Frame 1 again (push "Frame", then "next"). Go back to frame 2, and select "Frame -> Match -> Colorbar" from under the "Frame" menu. Now go to frame 1 - they should have the same contrast and brightness now. Then blink between the two frames by pressing the "Blink" button. (You can stop the blinking by clicking "single", or you can blink manually by just repeatedly clicking "Next".) Notice the small shift between the frames? Stop the blink mode by pressing the "single" button.

Play a bit with ds9 until you feel comfortable with it. It has many more features that we will not cover here - see the documentation online, or under the "Help" menu in ds9.

Now back to our original problem of computing and correcting for these shifts.

cl>	display im010 1	#	redisplay im010
cl>	unlearn imexamine		
cl>	lpar imexamine	#	we want to compute the shifts using this task interactively
cl>	imexamine	#	<pre>move the cursor into the ds9 window - notice that it has changed to a blinking circle - imexamine has put us into "interactive image cursor mode"</pre>

- a. put the cursor on some stars and press the "a" key information will be printed in the xgterm window: the x-coord and y-coord values are on the first line with additional information on the second line - see the help page for imexamine for details
- b. with the cursor in the image window type "?" to see the cursor help - exit cursor help with "q" in the xgterm window

c. what we would like to do is select 3 relatively bright stars and measure their positions in both frames and then compute the differences - we will use the average of the differences to shift the second frame so it aligns with the first

mark 3 stars with the "a" key - space them over the image

d. with the cursor in the image window, type "d" - you will be queried in the xgterm window for the next image - enter "im011" and frame "1"

measure the same three stars in this image

e. quit "imexamine" with "q" in the ds9 window

Compute the average shift, i.e. what shift do we want to apply to the second image so that it aligns with the first? The shifts that I computed are -0.53 and -1.68 - do you agree?

cl>	lpar help	
cl>	help imshift sec=description #	look at just one part of the
		help page for the task that
		we want to use
cl>	help imshift sec=example	
cl>	unlearn imshift	
cl>	lpar imshift	
cl>	<pre>imshift im011 shift011 ??? ???</pre>	# shift im011 with the
		appropriate values
cl>	dir	<pre># Note that you have</pre>
		created a new image!

Now, blink im010 and shift011 and see if we did a good job. Do you remember how to do this? Congratulate yourself if things look ok!!

Let's try a little image arithmetic now. We have two frames, im010 and shift011, that we want to average. Let's do it two ways.

cl> unlearn imsum imarith

[Note the concern here about the pixel type, both for the calculation and for the output image.]

c. [Hopefully the results are the same for both operations.]

cl> unlearn imstatistics
cl> lpar imstat
cl> imstat aver*.fits

Notice that when you change hidden parameters on the command line that they are NOT "learn"ed! How do you "learn" parameters?

History -----

IRAF can redirect terminal output to a file, as well as pipe output from one task into the input of another task. There is also a history and recall mechanism.

cl> history	#	prints history tree
cl> ^	#	recall and execute last command, can also
		include number to execute any task in tree
cl> e lpar	#	recall last lpar command - allows you to
		edit command line before executing
		with "return" -
		use the arrow keys to move cursor, delete
		or insert to the left of the cursor
cl> e	#	recall last command - use up/down arrows
		to go up/down history tree
cl> history 100	#	look at last 100 commands
cl> history 100 > hfile	#	redirect output to a file
cl> page hfile	#	page the file
cl> history 100 page	#	alternate method avoiding intermediate
		file

What is the difference between ">" and "|" ?

More on Plotting

Let's explore plotting in IRAF a bit. Some plot tasks are interactive and others are not. You can always replot whatever was in the last plot buffer and play with it.

cl> phelp plot cl> contour shift011	<pre># list help for "plot" package # make a contour plot of your shifted image - if it is not already the frame that you are looking at in the ds9 window, re-display it. Compare the plot with the displayed image</pre>
cl> surface shift011	# draw a surface plot of the same image
cl> implot shift011	<pre># this is an interactive plotting task that is useful for inspecting 2-d images - type "?" - note the cursor</pre>

commands (small letters and : commands) - these will differ from task to task (if they are interactive) - they are NOT global c - column plot, column read from cursor position :l 100 - plot line 100 :c 150 200 - plot average of columns 150-200 :.write meta # save plot in file (Spend some time becoming familiar with "implot" - you will use it often - try the global keys as well.) How can you expand the plot other than with "Z"? look at the "e" key - put both cursors at the lower left corner of a box defining the region you wish to zoom, press "e", then move both cursors to the upper right corner of the box and press "e" again :l 100 - to get the full plot size back # exit q cl> implot dev\$wpix # plot the image that is distributed with the IRAF system :w world # this image has RA and DEC information stored in its header; use it for the x axis. (Why doesn't the y axis display this way as well?) :f %H # convert to hh:mm:ss :.write meta # place the cursor on the middle of the С galaxy and type "c" to get a column plot :f %h # convert to dd:mm:ss :.write meta # exit q

We have saved several plots in a file called meta. Let's look at those plots now.

cl>	unlearn gkimosaic	gkidir
cl>	gkidir meta	<pre># list plot file</pre>
cl>	lpar gkimosaic	
cl>	gkimos meta	<pre># plot the meta file</pre>
	q	<pre># to quit plot mode</pre>

Now let's print out one of the plots. Choose one of them (use the gkidir or gkimosaic commands again to look them over). Note its frame number (as given by gkidir) and then plot it using

cl> gkiextract meta 1 | stdplot

Here I chose frame 1 - use a different one if you wish. In general, you can print from within most plotting tasks by hitting the "=" key (or by typing :.snap), which takes a snapshot of the graphics window and sends it to the printer. Type "help cursor" for more info on this and other graphics cursor commands.

For the next few minutes it may be profitable to spend a bit more time familiarizing yourself with the task IMEXAMINE, since it is such a powerful tool.

cl>	display	dev\$pix	1	#	this is the IRAF distributed
					image, M51. Wpix and pix are
					the same image but wpix has the
					world coordinate information in
					its header

[window the image until it looks good]

cl> imexamine

[put the cursor on the star at	224, 131]
?	<pre># list cursor options - type "q" in</pre>
	the text window to exit list
Z	# print out pixel values around
	cursor
m	# print statistics within box
[move the cursor to a glob on a	a sniral arm
	# surface nlot
·enar	# edit the narameters for the last
	nlot - you must type this from the
	ds9 window although the command
	will appear in the graphics window -
	exit epar mode in the xgterm window
	with :q
1	# plot a line
g	<pre># go to "interactive graphics</pre>
	cursor mode" - the global cursor
	keys can now be used to fuss
4.0	with the graphics plot
:naverage 10	# average 10 lines - since the graphics
	window is now the active window type
i	# go back to "interactive image
	cursor mode"
[try other options]	
q	# quit the task - you can quit
	in either interactive mode

What did the xmag and ymag parameters do?

Print a copy of your image by choosing "Print" from the "File" menu in ds9.

If you are finished with this exercise you may want to clean your directory up a bit with the deletion commands.

cl>	dir		
cl>	delete hfile,meta	#	deletes ordinary files
cl>	imdelete *.fits ver+	#	deletes image files - you may want to keep im*.fits around but we will not be using them again

Why there is a special IMDELETE command for the IRAF images? This is actually a holdover from using IRAF format images which were stored in two separate files, header and pixels. The IMDELETE command would delete both files at once. With FITS files, everything is in one file, so you could also just say

cl> delete *.fits

or

cl> rm *.fits # Actually passes it out to the shell

It's best to use the ver+ option (notice that this is how you "turn on" yes/no parameters without editing the parameter file) when you delete *.anything so that you don't accidentally wipe out a bunch of files! To set this as the default, you can type "epar imdelete", change "verify" to "yes", and save the parameters; the same for the "delete" command. The parameter files are saved in your "uparm" directory for your next IRAF session.

cl> logout

You should always log out of IRAF first before shutting down the window environment with the mouse. Then log off the linux system, if you wish.

% logout

References

A Beginner's Guide to Using IRAF, by Jeannette Barnes, August 1993.

DS9 version 1.9.7.1 documentation

(Note: this exercise was written by Jeannette Barnes of NOAO and modified by Eric Jensen to reflect the use of FITS files as the default image format and DS9 [rather than ximtool] as the image display program.)

Last update Feb 2014.