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Mon, 07/30/2018 - 11:16 #1

humayun

new package af	ter 2D classification
Hi all,	
	cation, when I try to make a new package of refined tting this following error message.
-bash-4.2\$ /prog Aborted	grams/share/capsules/lib/job.sh: line 113: 2192 (core dumped) "\$SB_EXECFILE" "\$@
Any idea how to	solve it? thank you!

Humayun

timgrant

Hi Humayun,

Hi Humayun,

Making a new package should not run other programs, and so you shouldn't get an error from a script - unless you are running cisTEM from inside a script?

If running in a script, perhaps you are redirecting the output? In which case, do you see any output from the actualy cisTEM program?

Thanks,

Tim

Mon, 07/30/2018 - 16:49 (Reply to #2)

humayun

Hi Tim,
thanks for your reply. I am not running cisTEM in a script, I am using the GUI.
Still stuck with the same error, any other comments or suggestions?
Thanks,
Humayun

timgrant

Your error is coming from the

Your error is coming from the script "/programs/share/capsules/lib/job.sh" what is line 113 of that script?

Tim

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