

## 1. Running a JOB – One Page Instructions

2. At the colon sign type **LOG IN [10, xxx]**, and hit enter (xxx-your account)
3. Type **MODE**, enter the chuck size you need for your substrate
4. Edit your job, make sure all info is correct
5. Select proper chuck for your substrate size and thickness (follow instructions for target thickness). If needed add the shim on the back side.
6. Load the sample on chuck, turn on the chuck vacuum
7. Load chuck with substrate on the stage, turn on the stage vacuum
8. Load the reticle, manually align it to the system (follow instructions for aligning the reticle to the system)
9. For **Global alignment** use command: **EX** or **EXEC**
10. **EX job name\pass name** and hit enter
11. For **Local alignment** use command: **MAP**
12. **MAP job name\mapping pass, pass name** and hit enter
13. Press **RESET** on the small keyboard
14. Press **MAN** and **S/C** simultaneously
15. Wait about 5-10 seconds
16. Press **MAN** again
17. Stage with substrate loaded on the chuck will move to alignment position
18. Proceed to the wafer alignment to the system, make sure alignment looks good
19. After alignment, press **EXP** on a small keyboard and the job will be executed
20. Unload your sample
21. Unload the reticle
22. Type **LOG OUT** to leave the system
23. Write in the log book info about the job you executed