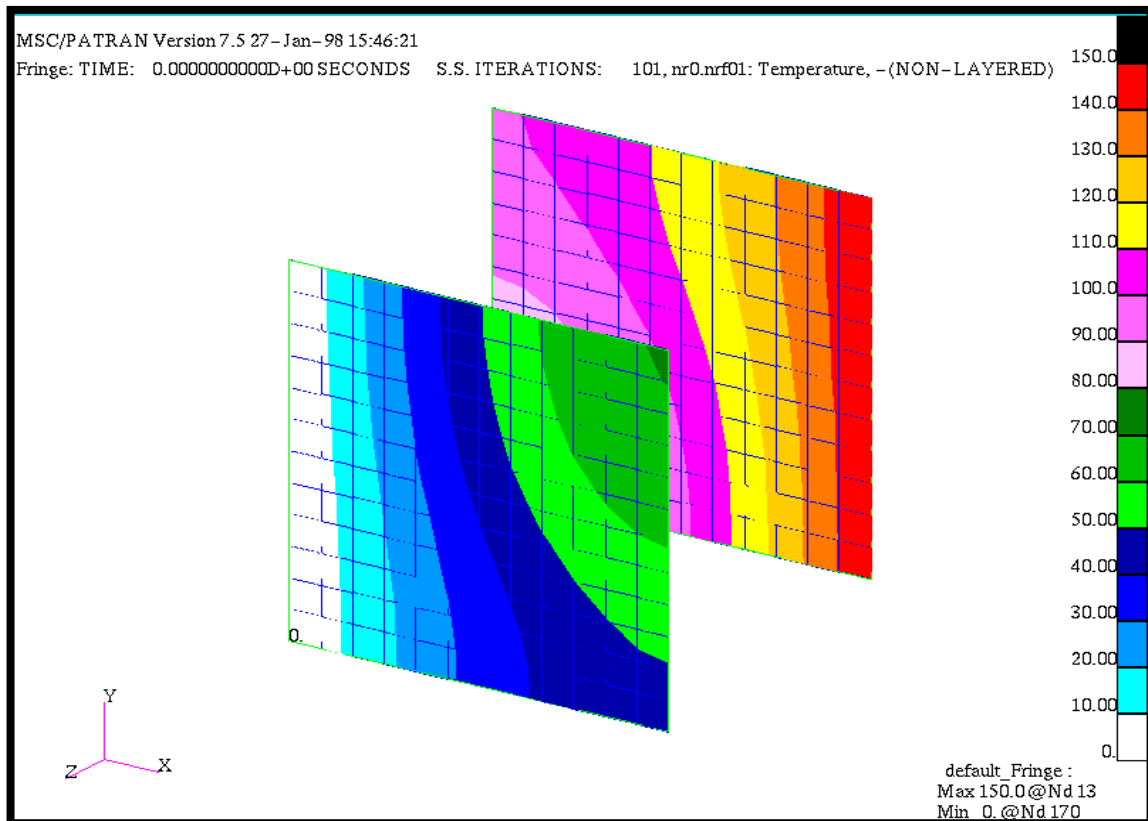


## Exercise 21

# *Optimizing Performance of Radiation Interchange Analysis*



### Objective:

- Modify the database of exercise\_14 and the template.dat.apnd file in order to increase analysis speed and reduce file size
- Rerun and monitor the analysis and compare CPU time of the run and file size to those of Exercise 14



## Model Description:

In this Exercise we will reopen the database created in Exercise 14 and modify some LBC's and the template.dat.apnd file. These modifications will significantly reduce the execution time of both the radiation interchange calculations as well as the thermal analysis network run. Also, the size of several of the files will be significantly reduced.

Any analyst who uses the radiation interchange capability of MSC/Thermal should become practiced in using the available flags and settings which will increase execution speed and reduce storage demands.

## Exercise Overview:

- Open the existing database named **exercise\_14.db**.
- Use **Load/BCs/Modify/Radiation** to modify the existing radiation boundary conditions.
- Create a new radiation Load/BC for Surface 2.
- Change the Job Name in the Analysis form to **exercise\_21**.
- Modify the **template.dat.apnd** file to include a collapse flag.
- Submit the model for analysis and use the commands described to monitor its progress.
- Debug, if necessary and resubmit after deleting all the files in the jobnamed subdirectory.
- Read in results file and plot results.
- Compare CPU times and File sizes.
- **Quit** MSC/PATRAN.

## Exercise Procedure:

### Open an existing database

1. Open the existing database named **exercise\_14.db**.

Within your window environment change directories to the microcircuit.db working directory. Run MSC/PATRAN by typing **p3** in your xterm window.

Next, select **File** from the *Menu Bar* and open the existing microcircuit database.

<b>File</b>	
<b>Open</b>	
Database List	<b>exercise_14.db</b>
<b>OK</b>	

MSC/PATRAN will open a Viewport and change various *Control Panel* selections from a ghosted appearance to a bold format.

### Use Load/BC Modify

2. Use **Load/BCs/Modify/Radiation** to modify the existing radiation boundary conditions.

In order to give a different Convex Surface ID flag to each surface it is necessary to modify the Input Data Form of the existing Load/BC for radiation.

◆ <b>Load/BC</b>	
<b>Modify/Radiation</b>	
Select Set to Modify	<your set name>
<b>Modify Data...</b>	
Convex Surface ID	1
Obstr Flag (0=Obst, 1=No-Obst)	1
<b>OK</b>	
<b>Modify Application Region...</b>	
Application Region	<b>Surface 1 (delete Surface 2)</b>
<b>OK</b>	
<b>Apply</b>	

3. Create a new radiation Load/BC for Surface 2.

**Create a new Load/BC**

By limiting the application region of the previous Load/BC to Surface 1, it is necessary to create a second Load/BC for Surface 2. Obviously the Application Region will be Surface 2. The Input Data Form will be the same as the last Load/BC but will have a different Convex Surface ID flag.

<b>◆ Load/BC</b>	
<b>Create/Radiation/Element Uniform</b>	
New Set Name	Rad2
Target Element Type	2D
<b>Input Data...</b>	
Vfac Template ID	200
Convex Surface ID	2
Obstu Flagg (0= Obst, 1= non-Obst)	1
Enclosure ID	1
<b>OK</b>	
<b>Select Application Region</b>	<select Surface 2>
<b>ADD</b>	
<b>OK</b>	
<b>Apply</b>	

4. Change the Job Name in the Analysis form to **exercise\_21**.

**Change Job Name**

On the Analysis Form change the *Job Name* to **exercise\_21**. This will create a new subdirectory of files for this analysis which will facilitate comparing data between the two runs, **exercise\_14** and **exercise\_21**.

<b>◆ Analysis</b>	
Job Name	exercise_21

## Modify template

5. Modify the **template.dat.apnd** file to include a collapse flag.

Open a Unix shell and edit the **template.dat.apnd** file associated with **exercise\_14**. If necessary make any existing **template.dat.apnd** file out of the way and reinstate the **template.dat.apnd** file associated with **exercise\_14**. Edit the **tempalte.dat.apnd** file to include a collapse flag.

```
*=====
VFAC 100
0.1 1.0 0 0 0.0 0.0 0 100
VFAC 200
0.1 1.0 0 0 0.0 0.0 0 200
*=====
```

The main advantage of using COLLAPSE to collapse radiosity nodes is that this will result in a much smaller number of radiation resistors in the model. A smaller number of resistors usually means that the thermal analysis will proceed faster. In the best cases, the number of radiation resistors may be reduced by about a factor of four for 2D Cartesian or axisymmetric models and by about a factor of 16 for 3D models.

## Submit the model

6. Submit the model for analysis and use the commands described to monitor its progress.

Return to the open [Analysis Form](#) and check **Apply**. After the [Command Line History Window](#) stops scrolling, change focus to the UNIX window and affect the **cd exercise\_21** command with a carriage return. Repeated execution of **ls** within the jobname subdirectory will show you the progress of your analysis: Once the file **vf.msg.01** appears, type:

```
$ tail -f vf.msg.01 <CR>
```

This will provide a continuous status of the viewfactor run.

When viewfactor is complete it will end the status with a message, Successful Execution Completed.

Use the **<Ctrl>c** key combination to terminate the tail function.

Again input a sequence of **ls** commands until a **stat.bin** file appears in the directory list. Once you see the **stat.bin** file type:

```
$ qstat c
```

to monitor the progress of the network analysis. This command will self terminate after 20 repetitions or upon job completion. Monitor the data from the **qstat** command to determine the numerical status of the analysis.

Check for the existence of an nr0.nrf.01 results file. If it exists the numerical analysis is complete and successful.

7. Debug, if necessary, and resubmit after deleting all the files in the jobnamed subdirectory.

If Step 5 does not yield a results file then determine what went wrong.

Is there a patqb.log file?

If so, then is there a patq.msg file? If there is no patqb.log file then look in the MSC/PATRAN Command Line History Window or in the PATRAN interface for any error messages.

If there is a patqb.log file and no patq.msg file then look for error messages in patqb.log.

If there is a patq.msg file then look for error messages in it.

If there are no error messages in the patq.msg file but this analysis requests that a viewfactor run be made then is there a vf.msg file?

If there is a vf.msg file then look for error messages in it.

For this analysis answering the above questions should provide a clue to the problem.

Once the error is found and resolved Repeat Steps 4 and 5. Remember that now many of the files will have an extension index which has been incremented by 1, e.g., vf.msg.01 to vf.msg.02. If it is convenient you may delete all the files from the exercise\_21 Job Named subdirectory prior to resubmitting the analysis.

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**Debug**

## Read and plot results

### 8. Read in results file and plot results.

From within MCS/PATRAN the only indication that the analysis has successfully finished is the existence of an nrX.nrf.01 results file in a subdirectory one level below your working directory.

Recall that p3 was initiated from a working directory which contained the microcircuit.db database file. The analysis, initiated from within MSC/PATRAN, created a new subdirectory with the same name as the *Job Name*; it should be named exercise\_21/. By using **Read Result** in the Analysis form and **Selecting Results File...** you can filter down to the *Job Name* subdirectory and check for the existence of the results file.

◆ Analysis	
Read Results/Result Entities	
Select Results File...	
Directories	<path>/exercise_21
Filter	
Available Files	nr0.nrf.01
OK	
Select Rslt Template File...	
Files	pthermal_1_nodal.res_tmpl
OK	
Apply	

To plot the results to posted FEM use the **Results Application radio button**.

◆ Results	
Create/Quick Plot	
Select Result Cases	<select the second> TIME: 0.000000000D+00 S...
Select Fringe Result	Temperature,
Apply	



Select the *Fringe Attributes* icon.



Display:	Element Edges
Label Style...	
Label Format:	Fixed
Significant figures	4 <use slider bar>
OK	
Apply	

9. Compare CPU times and File sizes.

Use the `qstat` command in each of the Job Name subdirectories to find the **CPU Time** data and record it in the following table.

Use the `ls -al v*` command in each Job Name subdirectory to record the size of the `vfnode.dat`, `vfraw.dat`, and `vfres.dat` files in the following table.

Subdirectories	Exercise_14	Exercise_21
CPU Time (sec.)		
Vfnode.dat (bytes)		
Vfraw.dat (bytes)		
Vfres.dat (bytes)		

The size and speed improvement are significant.

10. **Quit** MSC/PATRAN.

To stop MSC/PATRAN select **F**ile on the *Menu Bar* and select **Q**uit from the drop-down menu.

**Compare files**

**Quit MSC/Patran**

