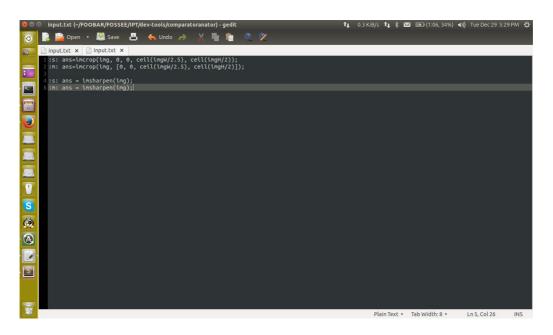
First of all, a 3d matrix in matlab and a list in scilab will not match. Thus, for the tool to work, your functions must return a matrix for an image. To do this, use the migrator tool in dev-tools/migrator.

Input.txt

The input file to the comparison tool will be 'input.txt', whose format is as follows.

Each comparison has a Scilab function, preceded by ':s:' and it's Matlab equivalent, preceded by ':m:', as shown below.



Since the functions are tested for multiple images, input.txt will use the variables img, imgH (image height) and imgW (image width) to generate specific commands.

[Matlab and Scilab have different rounding rules. So please use 'ceil()' while using imgH or imgW].

After manually writing input.txt, do the following steps:

Generating output.txt

- 1. Run matlab using the command 'matlab -nojvm', from within the comparator directory.
- 2. Run, 'dfa_ttd_maker', and give 'input.txt' as the input file.
- 3. fd = fopen('output.txt', 'w');
- 4. Next, to include colored images in the comparison, run 'dfa(ttd, fd, 'sample_images/color/')', and for greyscale images, run 'dfa(ttd, fd, 'sample images/greyscale/')'.
- 5. Exit Matlab using 'exit'.

```
utput.txt (~/FOOBAR/FOSSEE/IPT/dev-tools/comparatoranator) - ge
       📄 ๊ Open 🔻 💆 Save 🖺 🤚 Undo 🧀 🐰 🖺
                :s: img = imread('sample_images/color/10r.bmp'); imgH = size(img, 1); imgH = size(img, 2);
:m: img = imread('sample_images/color/10r.bmp'); imgH = size(img, 1); imgH = size(img, 2);
                     ans=imcrop(img, 0, 0, ceil(imgW/2.5), ceil(imgH/2));
ans=imcrop(img, [0, 0, ceil(imgW/2.5), ceil(imgH/2)]);
                :s: ans = imsharpen(img);
:m: ans = imsharpen(img);
                :s: img = imread('sample_images/color/10r.jpg'); imgH = size(img, 1); imgW = size(img, 2);
:m: img = imread('sample images/color/10r.jpg'); imgH = size(img, 1); imgW = size(img, 2);
                :s: ans=imcrop(img, 0, 0, ceil(imgW/2.5), ceil(imgH/2));
:m: ans=imcrop(img, [0, 0, ceil(imgW/2.5), ceil(imgH/2)]);
                :s: ans = imsharpen(img);
:m: ans = imsharpen(img);
9
S
               :s: img = imread('sample_images/color/10r.png'); imgH = size(img, 1); imgW = size(img, 2);
:m: img = imread('sample_images/color/10r.png'); imgH = size(img, 1); imgW = size(img, 2);
Ö
                :s: ans=imcrop(img, 0, 0, ceil(imgW/2.5), ceil(imgH/2));
:m: ans=imcrop(img, [0, 0, ceil(imgW/2.5), ceil(imgH/2)]);
:s: ans = imsharpen(img);
:m: ans = imsharpen(img);
                :s: img = imread('sample_images/color/10r.tiff'); imgH = size(img, 1); imgW = size(img, 2);
:m: img = imread('sample_images/color/10r.tiff'); imgH = size(img, 1); imgW = size(img, 2);
                     ans=imcrop(img, 0, 0, ceil(imgW/2.5), ceil(imgH/2));
ans=imcrop(img, [0, 0, ceil(imgW/2.5), ceil(imgH/2)]);
                                                                                                                                                                                       Plain Text ▼ Tab Width: 8 ▼ Ln 49, Col 54 INS
```

Running the comparison(s)

Now, open a terminal session inside dev-tools/comparinatoranator and type 'make'.

Give 'output.txt' as the input file. Next, specify whether you want to build the toolbox or not.

```
Copyright 1984-2014 The MathWorks, Inc.
                   R2014b (8.4.0.150421) 64-bit (glnxa64)
                             September 15, 2014
For online documentation, see http://www.mathworks.com/support
For product information, visit www.mathworks.com.
Enter input file (Enter for stdin): output.txt
 ---- Do you want me to run the builder? (Runs GUI) -----
     1) Yes
     2) No
Select a menu number: 2
Comparing...
scilab command: img = imread('sample_images/color/10r.bmp'); imgH = size(img,
); imgW = size(img, 2);
natlab command: img = imread('sample images/color/10r.bmp'); imgH = size(img,
); imgW = size(img, 2);
sending goScilab
```

When the terminal says 'sending goScilab', open another terminal session the same directory, and run:

"scilab-cli -f sciScript.sce &> scilogs.txt".

After the Matlab script stops running, all the comparison results are logged in 'logs.txt', as shown below.

```
Nothing to compare for scilab img = imread('sample_images/color/10r.bmp'); imgH = size(img, 1); imgW = size(img, 2); and matlab img = imread('sample_images/color/10r.bmp'); imgH = size(img, 1); imgW = size(img, 2);.

Match for scilab ans=imcrop(img, 0, 0, ceil(imgW/2.5), ceil(imgH/2)); and matlab ans=imcrop(img, [0, 0, ceil(imgW/2.5), ceil(imgH/2)]);

Mismatch for scilab ans = imsharpen(img); and matlab ans = imsharpen(img);

Nothing to compare for scilab img = imread('sample_images/color/10r.jpg'); imgH = size(img, 1); imgW = size(img, 2); and matlab img = imread('sample_images/color/10r.jpg'); imgH = size(img, 1); imgW = size(img, 2);.

Match for scilab ans=imcrop(img, 0, 0, ceil(imgW/2.5), ceil(imgH/2)); and matlab ans=imcrop(img, [0, 0, ceil(imgW/2.5), ceil(imgH/2)]);

Mismatch for scilab ans = imsharpen(img); and matlab ans = imsharpen(img);

Nothing to compare for scilab img = imread('sample_images/color/10r.png'); imgH = size(img, 1); imgW = size(img, 2); :

''Answer of the structure of the size of the s
```

Whenever a mismatch is encountered, the results are logged in 'sciLogs.txt'.

[Note: The above tool was coded and tested with Matlab 2014b and Scilab 5.5.0.]