# Parallel MATLAB: Single Program Multiple Data

John Burkardt (FSU) & Gene Cliff (AOE/ICAM) 1:30pm - 3:30pm, Thursday, 02 June 2011 Mathematics Commons Room

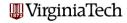
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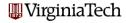
FSU: Florida State University

**AOE**: Department of Aerospace and Ocean Engineering **ICAM**: Interdisciplinary Center for Applied Mathematics



# MATLAB Parallel Computing

- SPMD: Single Program, Multiple Data
- QUAD Example
- Distributed Arrays
- IMAGE Example
- CONTRAST Example
- CONTRAST2: Messages
- Batch Computing
- Conclusion



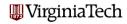
#### SPMD: is not PARFOR

#### Lecture #1: PARFOR - 31 May

The **parfor** command, described in a previous lecture, is easy to use, but it only lets us do parallelism in terms of loops. The only choice we make is whether a loop is to run in parallel.

- We can't determine how the loop iterations are divided up;
- we can't be sure which worker runs which iteration;
- we can't examine the work of any individual worker.

Using **parfor**, the individual workers are *anonymous*, and all the memory is shared (or copied and returned).

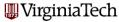


# SPMD: is Single Program, Multiple Data

#### Lecture #2: SPMD - 2 June

The **SPMD** command (today's lecture) is like a very simplified version of **MPI**. There is one client process, supervising workers who cooperate on a single program. Each worker (sometimes also called a "lab") has an identifier, knows how many workers there are total, and can determine its behavior based on that ID.

- each worker runs on a separate core (ideally);
- each worker uses separate workspace;
- a common program is used;
- workers meet at synchronization points;
- the client program can examine or modify data on any worker;
- any two workers can communicate directly via messages.



# SPMD: Getting Workers

An **spmd** program needs workers to cooperate on the program.

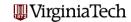
So on a desktop, we issue an interactive matlabpool request:

'ithaca\_2009b', 'matlabpool', 7)

```
matlabpool open local 4
  results = myfunc ( args );

or use batch to run in the background on your desktop:
  job = batch ( 'myscript', 'Configuration', 'local', ...
  'matlabpool', 4 )

or send the batch command to the Ithaca cluster:
  job = batch ( 'myscript', 'Configuration', ...
```



#### SPMD: The SPMD Environment

MATLAB sets up one special worker called the client.

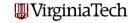
MATLAB sets up the requested number of workers, each with a copy of the program. Each worker "knows" it's a worker, and has access to two special functions:

- numlabs(), the number of workers;
- labindex(), a unique identifier between 1 and numlabs().

The empty parentheses are usually dropped, but remember, these are functions, not variables!

If the client calls these functions, they both return the value 1! That's because when the client is running, the workers are not. The client could determine the number of workers <u>available</u> by

```
n = matlabpool ( 'size' )
```



#### SPMD: The SPMD Command

The client and the workers share a single program in which some commands are delimited within blocks opening with **spmd** and closing with **end**.

The client executes commands up to the first **spmd** block, when it pauses. The workers execute the code in the block. Once they finish, the client resumes execution.

The client and each worker have separate workspaces, but it is possible for them to communicate and trade information.

The value of variables defined in the "client program" can be referenced by the workers, but not changed.

Variables defined by the workers can be referenced or changed by the client, but a special syntax is used to do this.

# SPMD: How SPMD Workspaces Are Handled

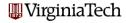
|                            | Client |   |   |     | Worker 1 |   |   |     | Worker     |      |    |
|----------------------------|--------|---|---|-----|----------|---|---|-----|------------|------|----|
|                            |        |   |   |     |          |   |   |     |            |      |    |
|                            | a      | b | е |     | С        | d | f |     | С          | d    | f  |
| 2.                         |        |   |   |     |          |   |   |     |            |      |    |
| a = 3;                     | 3      |   | _ | - ! | _        | _ | _ | - 1 | _          | _    | _  |
| b = 4;                     | 3      | 4 | - |     | -        | - | - |     | -          | -    | -  |
| spmd                       |        |   |   |     |          |   |   |     |            |      |    |
| <pre>c = labindex();</pre> | 3      | 4 | - |     | 1        | - | - |     | 2          | -    | -  |
| d = c + a;                 | 3      | 4 | - |     | 1        | 4 | - |     | 2          | 5    | _  |
| end                        |        |   |   |     |          |   |   |     |            |      |    |
| $e = a + d\{1\};$          | 3      | 4 | 7 |     | 1        | 4 | - |     | 2          | 5    | -  |
| $c{2} = 5;$                | 3      | 4 | 7 | -   | 1        | 4 | - |     | 5          | 6    | _  |
| spmd                       |        |   |   |     |          |   |   |     |            |      |    |
| f = c * b;                 | 3      | 4 | 7 |     | 1        | 4 | 4 |     | 5          | 6    | 20 |
| end                        |        |   |   |     |          |   |   | Į   | <b></b> V: | irgi | ni |

# SPMD: When is Workspace Preserved?

A program can contain several **spmd** blocks. When execution of one block is completed, the workers pause, but they do not disappear and their workspace remains intact. A variable set in one **spmd** block will still have that value if another **spmd** block is encountered.

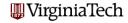
You can imagine the client and workers simply alternate execution.

In MATLAB, variables defined in a function "disappear" once the function is exited. The same thing is true, in the same way, for a MATLAB program that calls a function containing **spmd** blocks. While inside the function, worker data is preserved from one block to another, but when the function is completed, the worker data defined there disappears, just as the regular MATLAB data does.

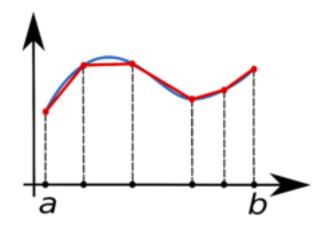


# MATLAB Parallel Computing

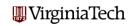
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# QUAD: The Trapezoid Rule



Area of one trapezoid = average height \* base.



# QUAD: The Trapezoid Rule

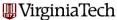
To estimate the area under a curve using one trapezoid, we write

$$\int_{a}^{b} f(x) dx \approx (\frac{1}{2}f(a) + \frac{1}{2}f(b)) * (b - a)$$

We can improve this estimate by using n-1 trapezoids defined by equally spaced points  $x_1$  through  $x_n$ :

$$\int_a^b f(x) dx \approx \left(\frac{1}{2}f(x_1) + f(x_2) + \dots + f(x_{n-1}) + \frac{1}{2}f(x_n)\right) * \frac{b-a}{n-1}$$

If we have several workers available, then each one can get a part of the interval to work on, and compute a trapezoid estimate there. By adding the estimates, we get an approximate to the integral of the function over the whole interval.

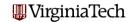


# QUAD: Use the ID to assign work

To simplify things, we'll assume our original interval is [0,1], and we'll let each worker define a and b to mean the ends of its subinterval. If we have 4 workers, then worker number 3 will be assigned  $\left[\frac{1}{2},\frac{3}{4}\right]$ .

To start our program, each worker figures out its interval:

```
fprintf ( 1, ' Set up the integration limits:\n');
spmd
  a = ( labindex - 1 ) / numlabs;
  b = labindex / numlabs;
end
```



#### QUAD: One Name Must Reference Several Values

Each worker is a program with its own workspace. It can "see" the variables on the client, but it usually doesn't know or care what is going on on the other workers.

Each worker defines **a** and **b** but stores *different values* there.

The client can "see" the workspace of all the workers. Since there are multiple values using the same name, the client must specify the index of the worker whose value it is interested in. Thus  $a\{1\}$  is how the client refers to the variable a on worker 1. The client can read or write this value.

MATLAB's name for this kind of variable, indexed using curly brackets, is a composite variable. It is very similar to a cell array.

The workers can "see" the client's variables and inherits a copy of their values, but cannot change the client's data. **WirginiaTech** 

# QUAD: Dealing with Composite Variables

So in QUAD, each worker could print **a** and **b**:

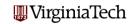
```
spmd
 a = ( labindex - 1 ) / numlabs;
 b = labindex / numlabs;
 fprintf (1, 'A = %f, B = %f\n', a, b);
end
    — or the client could print them all —
spmd
 a = (labindex - 1) / numlabs:
 b = labindex / numlabs;
end
for i = 1 : 4 <-- "numlabs" wouldn't work here!
 fprintf (1, ' A = \%f, B = \%f \setminus n', a\{i\}, b\{i\});
                                            VirginiaTech
end
```

#### QUAD: The Solution in 4 Parts

Each worker can now carry out its trapezoid computation:

#### with result:

- 2 Partial approx 0.874676
- 4 Partial approx 0.567588
- 1 Partial approx 0.979915
- 3 Partial approx 0.719414



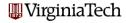
# QUAD: Combining Partial Results

We really want one answer, the sum of all these approximations.

One way to do this is to gather the answers back on the client, and sum them:

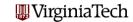
```
quad = sum ( quad_part{1:4} );
fprintf ( 1, ' Approximation %f\n', quad );
with result:
```

Approximation 3.14159265



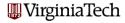
### QUAD: Source Code for QUAD\_FUN

```
function value = quad_fun ( n )
 fprintf ( 1, 'Compute_limits\n' );
 spmd
   a = (labindex - 1) / numlabs;
   b = labindex / numlabs;
    fprintf ( 1. '__Lab_%d_works_on_[%f.%f].\n'. labindex . a. b ):
 end
 fprintf ( 1, 'Each_lab_estimates_part_of_the_integral.\n' );
 spmd
   x = linspace (a, b, n);
   fx = f(x):
   quad_part = (b - a) * (fx(1) + 2 * sum (fx(2:n-1)) + fx(n)) ...
     / 2.0 / (n-1);
    fprintf ( 1. '__Approx_%f\n', quad_part ):
 end
 quad = sum ( quad_part{:} ):
 fprintf ( 1, '__Approximation_=_%f\n', quad )
 return
end
```



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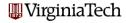


# **DISTRIBUTED:** Conjugate Gradient Setup

It is possible to use what amounts to **SPMD** programming without explicitly using the **spmd** statement. That's because many MATLAB functions and operators are capable of carrying out algorithms that involve the cooperation of multiple workers with separate workspaces.

The user might only see the "client" copy of MATLAB; special commands or options distribute the data to the available workers, who then cooperate to carry out the computation.

Again, this is "really" **SPMD** programming, except that the MathWorks staff had to write the **spmd** blocks, hidden inside MATLAB's functions.



#### **DISTRIBUTED:** The Client Can Distribute

If the client process has a 300x400 array called **a**, and there are 4 SPMD workers, then the simple command

```
ad = distributed ( a );
```

distributes the elements of **a** by columns:

```
Worker 1 Worker 2 Worker 3 Worker 4
Col: 1:100 | 101:200 | 201:300 | 301:400 ]
Row
1 [abcd|efgh|ijkl|mnop]
2 [ABCD|EFGH|IJKL|MNOP]
... [****|****|****|
300 [1234]5678|9012|3456]
```

By default, the last dimension is used for distribution. Wirginia Tech

#### DISTRIBUTED: Workers Can Get Their Part

Once the client has distributed the matrix by the command

```
ad = distributed ( a );
```

[9012]

then each worker can make a local variable containing its part:

```
spmd
  al = getLocalPart ( ad );
  [ ml, nl ] = size ( al );
end
On worker 3, [ml, nl] = (300, 100), and al is
  [ijkl]
  [IJKL]
  [****]
```

Notice that local and global column indices will differ! Wirginia Tech

#### **DISTRIBUTED:** The Client Can Collect Results

The client can access any worker's local part by using curly brackets. Thus it could copy what's on worker 3 by

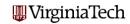
```
worker3_array = al{3};
```

However, it's likely that the client simply wants to collect all the parts and put them back into one normal MATLAB array. If the local arrays are simply column-sections of a 2D array:

```
a2 = [a1{:}]
```

Suppose we had a 3D array whose third dimension was 3, and we had distributed it as 3 2D arrays. To collect it back:

```
a2 = al{1};
a2(:,:,2) = al{2};
a2(:,:,3) = al{3};
```

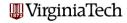


#### **DISTRIBUTED:** Methods to Gather Data

Instead of having an array created on the client and distributed to the workers, it is possible to have a distributed array constructed by having each worker build its piece. The result is still a distributed array, but when building it, we say we are building a **codistributed** array.

Codistributing the creation of an array has several advantages:

- The array might be too large to build entirely on one core (or processor);
- The array is built faster in parallel;
- You skip the communication cost of distributing it.



# **DISTRIBUTED:** Accessing Distributed Arrays

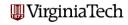
The command al = getLocalPart ( ad ) makes a local copy of the part of the distributed array residing on each worker. Although the workers could reference the distributed array directly, the local part has some uses:

- references to a local array are faster;
- the worker may only need to operate on the local part; then it's easier to write al than to specify ad indexed by the appropriate subranges.

The client can copy a distributed array into a "normal" array stored entirely in its memory space by the command

```
a = gather ( ad );
```

or the client can access and concatenate local parts.



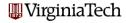
# **DISTRIBUTED:** Conjugate Gradient Setup

Because many MATLAB operators and functions can automatically detect and deal with distributed data, it is possible to write programs that carry out sophisticated algorithms in which the computation never explicitly worries about where the data is!

The only tricky part is distributing the data initially, or gathering the results at the end.

Let us look at a conjugate gradient code which has been modified to deal with distributed data.

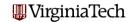
Before this code is executed, we assume the user has requested some number of workers, using the interactive **matlabpool** or indirect **batch** command.



# **DISTRIBUTED**: Conjugate Gradient Setup

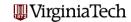
```
% Script to invoke conjugate gradient solution
% for sparse distributed (or not) array
         = 1000:
        = 5000:
    nnz
          = 1/10:
                    % reciprocal condition number
    rc
   A = sprandsym(N, nnz/N^2, rc, 1); % symmetric, positive definite
   A = distributed(A); %A = distributed.sprandsym() is not available
    b = sum(A, 2):
    [x, e\_norm] = cg\_emc(A, b);
    fprintf( 1. 'Error_residual_:___%8.4e_\n'. e_norm):
    np = 8;
    fprintf( 1, 'First_few_x_values:_\n');
    fprintf(1, '-x(-\%02i-)----\%8.4e-\n', [1:np; gather(x(1:np))']);
```

sprandsym sets up a sparse random symmetric array **A**. distributed 'casts' **A** to a distributed array on the workers



# **DISTRIBUTED**: Conjugate Gradient Iteration

```
function [x, resnrm] = cg_emc(A, b, x0, tol, itmax)
% Conjugate gradient iteration for A \times = b,
% with symmetric, positive-definite A
% (from Gill, Murray and Wright, p 147
% Possibly supply missing input parameters ((omitted)
% initialization
     p = zeros(size(x0));
     beta = 0:
     r = A*x0 - b
     rknrm= r'*r:
     x = x0;
     iter = 0:
% CG loop
     while 1
         p = beta*p - r:
         tmp = A*p;
         alpha = rknrm/(p'*tmp):
         x = x + alpha*p;
         r = r + alpha*tmp;
         rkpnrm= r'*r;
         beta = rkpnrm/rknrm:
         rknrm = rkpnrm;
         iter = iter + 1;
         resnrm= norm(A*x - b);
         if iter >= itmax || resnrm <= tol
             break
         end
     end % while 1 > 0
```

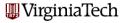


#### **DISTRIBUTED:** Comment

In the conjugate gradient example, we have emphasized how trivial it is to extend a MATLAB algorithm to a distributed memory problem. Essentially, all you have to do is invoke **distributed()**; the operational commands don't change.

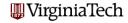
There are two comments worth making, in the interest of honesty:

- Not all MATLAB operators have been extended to work with distributed memory. In particular, (the last time we asked), the backslash or "linear solve" operator x=A\b can't be used yet for sparse distributed arrays.
- Getting "real" data (as opposed to matrices full of random numbers) properly distributed across multiple processors involves more choices and more thought than is suggested by the example we have shown!



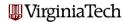
#### **DISTRIBUTED:** 1D Finite Difference Heat

In the next example, we demonstrate a mixed approach wherein the stiffness matrix (K) is initially constructed as a **codistributed** array on the workers. Each worker then modifies its localPart, and also assembles the local contribution to the forcing term (F). The *local* forcing arrays are then used to build a **codistributed** array.



#### **DISTRIBUTED:** 1D Finite Difference Heat code

```
Script to invoke codistributed solve for 1D heat equn by finite difference
 N = 100:
  spmd
          = -2*codistributed.eve(N): % diagonal part
     locP = getLocalPart(codistributed.colon(1, N));
% Loop over columns entering unity above/below the diagonal entry
% Note that columns 1 and N are exceptions
     for jj=locP(1):locP(end)
        if |i| = 1; K(|i|-1, |i|) = 1; end % above the diagonal
        if j = N; K(j + 1, j) = 1; end % below the diagonal
     end
          = (10*labindex())*ones(size(locP'))/N^2;\% funky rhs
     codist = codistributor1d(1, codistributor1d.unsetPartition, [N, 1]);
           = codistributed.build(locf. codist): % distribute the array(s)
  end
   T = K \setminus F; % mldivide is defined for distributed arrays
    plot ((1:N)/N, T, 'ro')
```



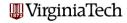
#### DISTRIBUTED: 2D Finite Element Heat Model

Finally, we consider an example that combines SPMD and distributed data to solve a steady state heat equations in 2D, using the finite element method.

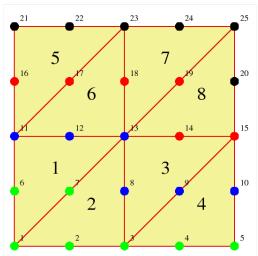
Each worker is assigned a subset of the finite element nodes. That worker is then responsible for constructing the columns of the (sparse) finite element matrix associated with those nodes.

Although the matrix is assembled in a distributed fashion, it has to be gathered back into a standard array before the linear system can be solved, because sparse linear systems can't be solved as a distributed array (yet).

This example is available as in the fem\_2D\_heat folder.



#### **DISTRIBUTED:** The Grid



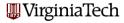
#### **DISTRIBUTED**: Finite Element System matrix

The discretized heat equation results in a linear system of the form

$$Kz = F + b$$

where  $\mathbf{K}$  is the stiffness matrix,  $\mathbf{z}$  is the unknown finite element coefficients,  $\mathbf{F}$  contains source terms and  $\mathbf{b}$  accounts for boundary conditions.

In the parallel implementation, the system matrix  $\mathbf{K}$  and the vectors  $\mathbf{F}$  and  $\mathbf{b}$  are distributed arrays. The default distribution of  $\mathbf{K}$  by columns essentially associates each SPMD worker with a group of finite element nodes.

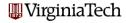


#### **DISTRIBUTED:** Finite Element System Matrix

To assemble the matrix, each worker loops over all elements. If element E contains any node associated with the worker, the worker computes the entire local stiffness matrix  $K_{i,j}$ . Columns of  $\mathbf{K}$  associated with worker nodes are added to the local part of  $K_{i,j}$ . The rest are discarded (which is OK, because they will also be computed and saved by the worker responsible for those nodes ).

When element 5 is handled, the "blue", "red" and "black" processors each compute K. But blue only updates column 11 of  $\mathbf{K}$ , red columns 16 and 17, and black columns 21, 22, and 23.

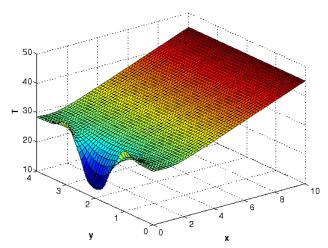
At the cost of some redundant computation, we avoid a lot of communication.

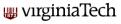


# Assemble Codistributed Arrays - code fragment

```
spmd
   Set up codistributed structure
   Column pointers and such for codistributed arrays.
   Vc = codistributed.colon(1, n_equations);
    IP = getLocalPart(Vc):
    IP_1= IP(1); IP_end = IP(end); %first and last columns of K on this lab
    co_dist_Vc = getCodistributor(Vc); dPM = co_dist_Vc.Partition;
% sparse arrays on each lab
    K_{ab} = sparse(n_{equations}, dPM(labindex));
  Build the finite element matrices — Begin loop over elements
    for n el=1:n elements
      nodes_local = e_conn(n_el ,:); % which nodes are in this element
   % subset of nodes/columns on this lab
      lab_nodes_local = extract( nodes_local . IP_1 . IP_end):
  ... if empty do nothing, else accumulate K_lab, etc end
    end % n_el
% Assemble the 'lab' parts in a codistributed format.
  syntax for version R2009b
    codist_matrix = codistributor1d(2, dPM, [n_equations, n_equations]);
   K = codistributed.build(K_lab, codist_matrix );
                                                                 WirginiaTech
   end % spmd
```

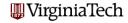
#### DISTRIBUTED: The Results





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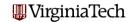


### **IMAGE:** Image Processing in Parallel

Here is a mysterious SPMD program to be run with 3 workers:

```
x = imread ('balloons.tif'):
v = imnoise (x, 'salt_&_pepper', 0.30);
yd = distributed (y);
spmd
  yl = getLocalPart ( yd );
  yl = medfilt2 (yl, [3, 3]);
end
z(1:480,1:640,1) = v(1);
z(1:480.1:640.2) = v(2):
z(1:480.1:640.3) = v(3):
figure;
subplot (1, 3, 1); imshow (x); title (X');
subplot ( 1, 3, 2 ); imshow ( y ); title ( 'Y' );
subplot ( 1, 3, 3 ); imshow ( z ); title ( 'Z' );
```

Without comments, what can you guess about this program?



# IMAGE: Image → Noisy Image → Filtered Image

Original image



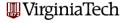
Noisy Image



Median Filtered Image



This filtering operation uses a 3x3 pixel neighborhood. We could blend *all* the noise away with a larger neighborhood.

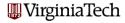


# IMAGE: Image → Noisy Image → Filtered Image

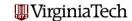
```
Read a color image, stored as 480x640x3 array.
x = imread ('balloons.tif');
 Create an image Y by adding "salt and pepper" noise to X.
y = imnoise (x, 'salt_&pepper', 0.30);
 Make YD, a distributed version of Y.
yd = distributed (y);
 Each worker works on its "local part", YL.
spmd
 vI = getLocalPart (vd);
  yl = medfilt2 (yl, [3, 3]);
end
The client retrieves the data from each worker.
z(1:480,1:640,1) = v(1);
z(1:480,1:640,2) = v(2);
z(1:480.1:640.3) = v(3):
 Display the original, noisy, and filtered versions.
figure;
subplot ( 1, 3, 1 ); imshow ( x ); title ( 'Original_image' );
                                                                   VirginiaTech
subplot ( 1, 3, 2 ); imshow ( y ); title ( 'Noisy_Image' );
                                         'Median_Filtered_Image (**)
subplot ( 1, 3, 3 ): imshow ( z ): title (
```

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```
Get 4 SPMD workers
matlabpool open 4
Read an image
x = imageread( 'surfsup.tif' ):
Since the image is black and white, it will be distributed by columns
xd = distributed(x):
 Each worker enhances the contrast on its portion of the picture
spmd
    xI = getLocalPart(xd);
    xl = nlfilter(xl, [3, 3], @adjustContrast);
    xI = uint8(xI):
end
Concatenate the submatrices to assemble the whole image
xf_spmd = [x|\{:\}];
matlabpool close
```





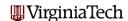




When a filtering operation is done on the client, we get picture 2. The same operation, divided among 4 workers, gives us picture 3. What went wrong?

Each pixel has had its contrast enhanced. That is, we compute the average over a 3x3 neighborhood, and then increase the difference between the center pixel and this average. Doing this for each pixel sharpens the contrast.

```
+----+
| P11 | P12 | P13 |
+----+
| P21 | P22 | P23 |
+----+
| P31 | P32 | P33 |
+----+
| P22 <- C * P22 + ( 1 - C ) * Average
```

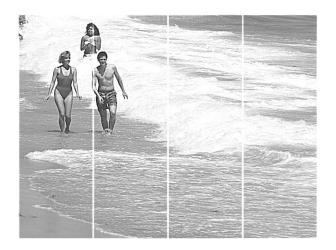


# CONTRAST: Image $\rightarrow$ Contrast Enhancement $\rightarrow$ Image2

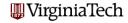
When the image is divided by columns among the workers, artificial internal boundaries are created. The algorithm turns any pixel lying along the boundary to white. (The same thing happened on the client, but we didn't notice!)

| Worker 1        | Worker 2        |     |
|-----------------|-----------------|-----|
| ++              | ++              | +   |
| P11   P12   P13 | P14   P15   P16 | P17 |
| ++              | ++              | +   |
| P21   P22   P23 | P24   P25   P26 | P27 |
| ++              | ++              | +   |
| P31   P32   P33 | P34   P35   P36 | P37 |
| ++              | ++              | +   |
| P41   P42   P43 | P44   P45   P46 | P47 |
| ++              | ++              | +   |

Dividing up the data has created undesirable artifacts! Wirginia Tech

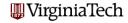


The result is spurious lines on the processed image.



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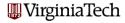
#### CONTRAST2: Workers Need to Communicate

The spurious lines would disappear if each worker could just be allowed to peek at the last column of data from the previous worker, and the first column of data from the next worker.

Just as in MPI, MATLAB includes commands that allow workers to exchange data.

The command we would like to use is **labSendReceive()** which controls the simultaneous transmission of data from all the workers.

```
data_received = labSendReceive ( to, from, data_sent );
```



#### CONTRAST2: Who Do I Want to Communicate With?

```
spmd
 xl = getLocalPart ( xd );
  if (labindex ~= 1)
   previous = labindex - 1;
  else
   previous = numlabs;
  end
  if ( labindex ~= numlabs )
   next = labindex + 1;
  else
   next = 1;
  end
```

### CONTRAST2: First Column Left, Last Column Right

```
column = labSendReceive ( previous, next, xl(:,1) );
if (labindex < numlabs)
 xl = [xl, column];
end
column = labSendReceive ( next, previous, x1(:,end) );
if (1 < labindex)
 xl = [column, xl];
end
```

#### CONTRAST2: Filter, then Discard Extra Columns

```
xl = nlfilter ( xl, [3,3], @enhance_contrast );
if ( labindex < numlabs )</pre>
  xl = xl(:,1:end-1);
end
if (1 < labindex)
  xl = xl(:, 2:end);
end
xl = uint8 (xl);
```

end

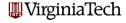
## CONTRAST2: Image $\rightarrow$ Enhancement $\rightarrow$ Image2





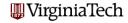


Four SPMD workers operated on columns of this image. Communication was allowed using **labSendReceive**.



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#### **BATCH: Indirect Execution**

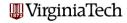
We can run quick, local interactive jobs using the **matlabpool** command to get parallel workers.

The **batch** command is an alternative which allows you to execute a MATLAB script (using either **parfor** or **spmd** statements) in the background on your desktop...or on a remote machine.

The **batch** command includes a **matlabpool** argument that allows you to request a given number of workers.

For remote jobs, the number of cores or processors you are asking for is the matlabpool **plus one**, because of the client.

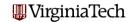
Since Ithaca allocates cores in groups of 8, it makes sense to ask for 7, or 15, or 23 or 31 workers, for instance.



#### BATCH: PRIME\_FUN is the function

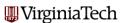
```
function total = prime_fun ( n )
 spmd
    nlo = (n * (labindex - 1)) / numlabs + 1;
    nhi = ( n * labindex ) / numlabs;
    if (nlo = 1)
     n\dot{l}o = 2:
    end
    total_part = 0:
    for i = nlo : nhi
      prime = 1:
      for i = 2 : i - 1
       if ( mod ( i, j ) == 0 )
          prime = 0:
          hreak
       end
      end
      total_part = total_part + prime;
   end
   total_spmd = gplus ( total_part );
  end
  total = total_spmd {1};
  return
```

end



#### BATCH: PRIME\_SCRIPT runs the function

```
%% PRIME_SCRIPT is a script to call PRIME_FUN.
   Discussion:
%
     The BATCH command runs scripts, not functions. So we have to write
     this short script if we want to work with BATCH!
%
%
   Licensing:
%
     This code is distributed under the GNU LGPL license.
   Modified:
%
     27 March 2010
%
%
%
   Author:
     John Burkardt
  n = 10000;
  fprintf ( 1, '\n' );
  fprintf ( 1, 'PRIME_SCRIPT\n' );
  fprintf ( 1, '__Count_prime_numbers_from_1_to_%d\n', n );
  total = prime_fun ( n );
```



# BATCH: Using the BATCH Command

```
job = batch ( 'prime_script', ...
  'configuration', 'local', ... <-- Run it locally.
  'matlabpool', 7)
                     <-- 7 workers, 1 client.
wait ( job ); <-- One way to find out when job is done.
load ( job ); <-- Load the output variables from
                  the job into the MATLAB workspace.
total
              <-- We can examine the value of TOTAL.
destroy ( job ); <-- Clean up
```

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## BATCH: Using the BATCH Command

The wait command pauses your MATLAB session.

Using batch, you can submit multiple jobs:

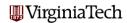
```
job1 = batch ( ... )
job2 = batch ( ... )
```

Using get, you can check on any job's status:

```
get ( job1, 'state' )
```

Using **load**, you can get the whole workspace, or you can examine just a single output variable if you specify the variable name:

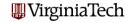
```
total = load ( job2, 'total' )
```



#### **BATCH: The BATCH Command**

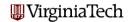
```
job_id = batch (
   'script_to_run', ...
   'configuration', 'local' or 'ithaca_2009b', ...
   'FileDependencies', 'file' or {'file1','file2'}, ...
   'CaptureDiary', 'true', ... (the default)
   'CurrentDirectory', '/home/burkardt/matlab', ...
   'PathDependencies', 'path' or {'path1','path2'}, ...
   'matlabpool', number of workers (can be zero!) )
```

Note that you do not include the file extension when naming the script to run, or the files in the FileDependencies.



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## **CONCLUSION: Summary of Examples**

The QUAD example showed a simple problem that could be done as easily with SPMD as with PARFOR. We just needed to learn about composite variables!

The CONJUGATE GRADIENT example showed that many MATLAB operations work for distributed arrays, a kind of array storage scheme associated with SPMD.

The HEAT example shows how to construct local arrays and assemble these to **distributed** arrays. This enables treatment of very large problems.

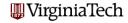
The IMAGE and CONTRAST examples showed us problems which can be broken up into subproblems to be dealt with by SPMD workers. We also saw that sometimes it is necessary for these workers to communicate, using a simple message-passing system.

### Conclusion: Desktop Experiments

Virginia Tech has a limited number of concurrent MATLAB licenses, which include the Parallel Computing Toolbox.

This is one way you can test parallel MATLAB on your desktop machine.

If you don't have a multicore machine, you won't see any speedup, but you may still be able to run some "parallel" programs.



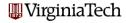
### Conclusion: Ithaca Experiments

If you want to work with parallel MATLAB on Ithaca, you must first get an account, by going to this website:

http://www.arc.vt.edu/index.php

Under the item **Services & Support** select **User Accounts**.

On the new page, under Ithaca Account Requests, select ARC Systems Account Request Form. Fill in the information and submit it. Although you're asked to describe the research you want the account for, you can say that this account is to experiment with Ithaca to see if it is suitable for your work.



### Conclusion: Desktop-to-Ithaca Submission

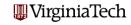
If you want to use parallel MATLAB regularly, you may want to set up a way to submit jobs from your desktop to Ithaca, without logging in directly.

This requires defining a configuration file on your desktop, adding some scripts to your MATLAB directory, and setting up a secure connection to Ithaca.

The steps for doing this are described in the document:

```
https://portal.arc.vt.edu/matlab/...
RemoteMatlabSubmission.pdf
```

We will be available to help you with this process.



#### Conclusion: VT MATLAB LISTSERV

There is a local LISTSERV for people interested in MATLAB on the Virginia Tech campus. We try **not** to post messages here unless we really consider them of importance!

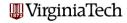
Important messages include information about workshops, special MATLAB events, and other issues affecting MATLAB users.

To subscribe to the mathworks listserver, send email to:

listserv@listserv.vt.edu.

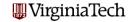
The body of the message should simply be:

subscribe mathworks firstname lastname



#### CONCLUSION: Where is it?

- MATLAB Parallel Computing Toolbox User's Guide 4.3 www.mathworks.com/access/helpdesk/help/pdf\_doc/distcomp/... distcomp.pdf
- Gaurav Sharma, Jos Martin, MATLAB: A Language for Parallel Computing, International Journal of Parallel Programming, Volume 37, Number 1, pages 3-36, February 2009.
- http://scholar.vt.edu/.../spmd\_codes
  - quad
  - conjugate\_gradient
  - fd\_1D\_heat
  - fem\_2D\_heat
  - image
  - contrast and contrast2
  - batch\_prime



#### AFTERWORD: PMODE

PMODE allows interactive parallel execution of MATLAB commands. PMODE achieves this by defining and submitting a parallel job, and it opens a Parallel Command Window connected to the labs running the job. The labs receive commands entered in the Parallel Command Window, process them, and send the command output back to the Parallel Command Window.

pmode start 'local' 2 will initiate pmode; pmode exit will
destroy the parallel job and end the pmode session

This may be a useful way to experiment with computations on the **labs**.

