

# CME292: Advanced MATLAB for Scientific Computing

## Homework #2 NLA & Opt

Due: Tuesday, April 28, 2015

### Instructions

This problem set will be combined with Homework #3. For this combined problem set, 2 out of the 6 problems are required. You are free to choose the problems you complete.

*Before completing problem set, please see HomeworkInstructions on Coursework for general homework instructions, grading policy, and number of required problems per problem set.*

### Problem 1

A matrix  $\mathbf{A} \in \mathbb{R}^{m \times n}$  can be compressed using the low-rank approximation property of the SVD. The approximation algorithm is given in Algorithm 1.

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**Algorithm 1** Low-Rank Approximation using SVD

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**Input:**  $\mathbf{A} \in \mathbb{R}^{m \times n}$  of rank  $r$  and approximation rank  $k$  ( $k \leq r$ )

**Output:**  $\mathbf{A}_k \in \mathbb{R}^{m \times n}$ , optimal rank  $k$  approximation to  $\mathbf{A}$

- 1: Compute (thin) SVD of  $\mathbf{A} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^T$ , where  $\mathbf{U} \in \mathbb{R}^{m \times r}$ ,  $\mathbf{\Sigma} \in \mathbb{R}^{r \times r}$ ,  $\mathbf{V} \in \mathbb{R}^{n \times r}$
  - 2:  $\mathbf{A}_k = \mathbf{U}(:, 1:k)\mathbf{\Sigma}(1:k, 1:k)\mathbf{V}(:, 1:k)^T$
- 

Notice the low-rank approximation,  $\mathbf{A}_k$ , and the original matrix,  $\mathbf{A}$ , are the same size. To actually achieve compression, the truncated singular factors,  $\mathbf{U}(:, 1:k) \in \mathbb{R}^{m \times k}$ ,  $\mathbf{\Sigma}(1:k, 1:k) \in \mathbb{R}^{k \times k}$ ,  $\mathbf{V}(:, 1:k) \in \mathbb{R}^{n \times k}$ , should be stored. As  $\mathbf{\Sigma}$  is *diagonal*, the required storage is  $(m+n+1)k$  doubles. The original matrix requires storing  $mn$  doubles. Therefore, the compression is useful only if  $k < \frac{mn}{m+n+1}$ .

Recall from lecture that the SVD is among the most expensive matrix factorizations in numerical linear algebra. Many SVD approximations have been developed; a particularly interesting one from [1] is given in Algorithm 2. This algorithm computes a rank  $p$  approximation to the SVD of a matrix. This approximation rank is *different* than the compression rank from Algorithm 1.

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**Algorithm 2** Low-Rank Probabilistic SVD Approximation

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**Input:**  $\mathbf{A} \in \mathbb{R}^{m \times n}$  (usually  $n \ll m$ ), approximation rank  $p$ , and number of power iterations  $q$

**Output:** Approximate SVD of  $\mathbf{A} \approx \mathbf{U}\mathbf{\Sigma}\mathbf{V}^T$

- 1: Generate  $n \times p$  Gaussian test matrix  $\mathbf{\Omega}$
  - 2: Form  $\mathbf{Y} = (\mathbf{A}\mathbf{A}^T)^q\mathbf{A}\mathbf{\Omega}$
  - 3: Compute QR factorization of  $\mathbf{Y}$ :  $\mathbf{Y} = \mathbf{Q}\mathbf{R}$
  - 4: Form  $\mathbf{B} = \mathbf{Q}^T\mathbf{A}$
  - 5: Compute SVD of  $\mathbf{B} = \tilde{\mathbf{U}}\mathbf{\Sigma}\tilde{\mathbf{V}}^T$
  - 6: Set  $\mathbf{U} = \mathbf{Q}\tilde{\mathbf{U}}$
-

In this problem, you will use the Singular Value Decomposition (SVD) to compress an image of your choosing. You will be given the function `get_rgb.m` that accepts the filename of an image and returns the RGB matrices defining the image, stacked vertically to form a single skinny matrix. Also, the function `plot_image_rgb.m` will take stacked RGB matrix and (optionally) a handle to an axes graphics object (default will use `gca`) and plot the corresponding image (the input to `plot_image_rgb.m` is the output of `get_rgb.m`). To “compress” the image, compress the stacked RGB matrix and pass to `plot_image_rgb.m`. Another possibility involves compressing the RGB matrices individually. For this assignment, use the stacked version for the compression.

```

1 function [RGB] = get_rgb(fname)
2
3 % Parse filename to determine extension
4 if isempty(find(fname=='.',1))
5     error('Filename must include extension.');
```

```

6 else
7     ext = fname(find(fname=='.',1,'last')+1:end);
8 end
9
10 A = double(imread(fname,ext));
11 RGB = [A(:, :, 1);A(:, :, 2);A(:, :, 3)];
12
13 end
```

```

1 function [] = plot_image_rgb(RGB,axHan)
2
3 if nargin < 4, axHan=gca; end
4
5 m = size(RGB,1)/3; n = size(RGB,2);
6 A = zeros(m,n,3);
7 A(:, :, 1)=RGB(1:m, :);
8 A(:, :, 2)=RGB(m+1:2*m, :);
9 A(:, :, 3)=RGB(2*m+1:end, :);
10 image(uint8(A), 'Parent', axHan);
11
12 end
```

- Consider an image of  $m \times n$  pixels. The  $R$ ,  $G$ ,  $B$  matrices will be of dimension  $m \times n$ .
- Compress to ranks  $[5, 10, 15, 20, 25, 50, 100, 200, \min(3m, n)]$  and visualize resulting images. Your result should be similar to Figure 1 for the image of Palm Drive.
  - Use `svd` for the SVD step in Algorithm 1
  - (extra credit) Implement the probabilistic SVD algorithm from Algorithm 2 and use this for the SVD step in Algorithm 1. Good values for the approximation rank will depend on the image you choose to compress. Notice that a rank  $p$  approximation to the SVD in Algorithm 2 only admits low-rank approximations of up to rank  $p$  (cannot go up to  $\min(3m, n)$ ). In this case, the ranks  $[5, 10, 15, 20, 25, 50, 100, 200, \min(3m, n)]$  cannot all be used. Choose any 9 interesting ranks to present.
- Plot  $\nu(k)$  for  $k \in \{1, 2, \dots, \min(3m, n)\}$  for the stacked RGB matrix, where

$$\nu(k) = 1 - \frac{\sum_{i=1}^k \sigma_i^2}{\sum_{j=1}^{\min(3m, n)} \sigma_j^2}. \quad (1)$$

and  $\sigma_i$  is the  $i$ th singular value of the matrix. The result should be similar to Figure 2a for the image of Palm Drive. This gives an indication of the compressibility of the image. A fast singular value decay implies the image can be compressed to a small rank with minimal loss in quality.

- Use the singular values generated by `svd` to complete this part.
- (extra credit) Repeat using the singular values generated by Algorithm 2.



Figure 1: Classic SVD, Varying degrees of compression

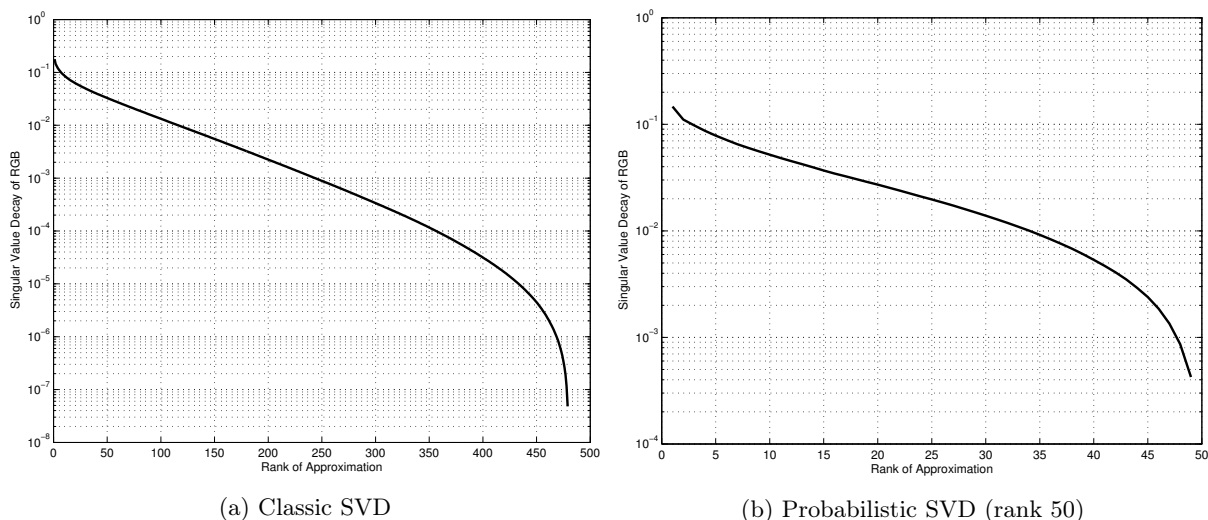


Figure 2: Singular value decay

## Problem 2

In this problem, you will gain experience with sparse matrices, matrix decompositions, and linear system solvers. You will be given a medium-scale sparse matrix in triplet format (`row`, `col`, `val`) as well as a right-hand side vector, `b` (`matrix_medium.mat`). For this matrix, complete the following tasks.

- (a) Convert sparse triplet (`row`, `col`, `val`) into a matrix `A`
- (b) Determine number of nonzeros in `A`
- (c) Compute approximate minimum degree permutation of `A`: `p = amd(A)`
- (d) Plot sparsity structure of matrix `A`
- (e) Compute LU decomposition of  $L*U = A(p, p)$  and plot sparsity structure of both matrix factors
- (f) Determine whether matrix is symmetric positive definite (don't use `eig` as it may be very slow). If matrix is SPD,
  - compute Cholesky factorization of  $A(p, p) = R' * R$  and plot sparsity structure of factor `R`
  - compute incomplete Cholesky factorization of  $A \rightarrow R_{inc}$  with no fill-in
    - `ichol` or `cholinc` depending on your version of MATLAB
  - Plot the sparsity structure of `R_{inc}`. Briefly comment.
- (g) Solve linear system of equations  $A*x = b$ . Define  $M = R_{inc}' * R_{inc}$ .
  - Using `mldivide` (backslash); this will be considered the exact solution of  $A*x = b$
  - Using the following iterative schemes
    - Conjugate Gradient without preconditioning
    - Preconditioned Conjugate Gradient, preconditioned with `M`
    - MINRES without preconditioning
    - MINRES, preconditioned with `M`

- Create a table containing the CPU time and number of iterations required for convergence of each algorithm and the relative error in converged solution, where  $(\text{error} = \text{norm}(\mathbf{x\_backslash} - \dots \mathbf{x\_iterative}) / \text{norm}(\mathbf{x\_backslash}))$ .
- Create a convergence plot of residual vector norm for each algorithm.

Additionally, you will also be given a large-scale sparse matrix in triplet format (`matrix_large.mat`). Repeat parts (a), (b), (c), (f), (g), (h) for the large matrix. Generate a spy plot for only the matrix itself (not the factors as some may be dense). *Warning: The large matrix will be fairly CPU intensive. I recommend running on corn.*

### Problem 3

In this problem, you will write a nonlinear equation solver using Newton-Raphson method. The Newton-Raphson method solves a nonlinear system iteratively by linearizing about the current iterate and finding the root of the linear system. To be explicit, consider the nonlinear function  $\mathbf{R} : \mathbb{R}^n \rightarrow \mathbb{R}^n$ . The goal is to solve  $\mathbf{R}(\mathbf{x}) = \mathbf{0}$ , given some starting value  $\mathbf{x}_0$ . The truncated Taylor series of  $\mathbf{R}$  about point  $\bar{\mathbf{x}}$  is

$$\mathbf{R}(\mathbf{x}) = \mathbf{R}(\bar{\mathbf{x}}) + \frac{\partial \mathbf{R}}{\partial \mathbf{x}}(\bar{\mathbf{x}})(\mathbf{x} - \bar{\mathbf{x}}) + \mathcal{O}(\|\mathbf{x} - \bar{\mathbf{x}}\|^2). \quad (2)$$

Substituting the truncated Taylor series into the equation  $\mathbf{R}(\mathbf{x}) = 0$  we have

$$\mathbf{x} = \bar{\mathbf{x}} - \left[ \frac{\partial \mathbf{R}}{\partial \mathbf{x}}(\bar{\mathbf{x}}) \right]^{-1} \mathbf{R}(\bar{\mathbf{x}}) \quad (3)$$

Applying (2) iteratively, starting from  $\mathbf{x}_0$ , until some convergence criteria is met is the Newton-Raphson method for solving nonlinear equations. Convergence is only guaranteed if  $\mathbf{x}_0$  is *close* to the actual solution.

- The convergence criteria for the nonlinear iterations is an important consideration for algorithmic speed and reliability. Most convergence criteria are based on the norm of the residual  $\|\mathbf{R}(\mathbf{x})\|$ . In this problem, we consider the nonlinear iterations to be converged if *either* of the following are satisfied:

$$\begin{aligned} & - \|\mathbf{R}(\mathbf{x}_n)\| \leq \epsilon_{\text{res}}^{\text{rel}} \|\mathbf{R}(\mathbf{x}_0)\| \\ & - \|\mathbf{R}(\mathbf{x}_n)\| \leq \epsilon_{\text{res}}^{\text{abs}} \quad \text{AND} \quad \|\mathbf{x}_n - \mathbf{x}_{n-1}\| \leq \epsilon_{\text{inc}}^{\text{abs}} \end{aligned}$$

where  $\epsilon_{\text{res}}^{\text{rel}}$ ,  $\epsilon_{\text{res}}^{\text{abs}}$ ,  $\epsilon_{\text{inc}}^{\text{abs}}$  are convergence tolerances. Use the 2-norm for this assignment (`norm(x)`).

- For many situations in computational science, the computation of the Jacobian matrix  $\frac{\partial \mathbf{R}}{\partial \mathbf{x}}$  is very expensive and it is desirable to reduce the number of times it must be computed. Therefore variants of Newton's method have considered *freezing* the Jacobian matrix at some iteration and reusing it until an iteration fails to reduce the residual. Namely, the Jacobian matrix is only recomputed when

$$- \|\mathbf{R}(\mathbf{x}_k)\| > \epsilon \|\mathbf{R}(\mathbf{x}_{k-1})\| \text{ for } \epsilon \in [0, 1]$$

- As Newton's method is *not* guaranteed to converge for an arbitrary starting point, it is good practice to specify a maximum number of nonlinear iterations. If the nonlinear iterations do not converge, a warning message should be issued. It is up to you to decide whether the final iterate  $\mathbf{x}_{N_{\text{max}}}$  is an acceptable solution.

Newton-Raphson's method is summarized in Algorithm 3 (without the "frozen" Jacobian feature). It should be emphasized that Algorithm 3 is just pseudo-code, not an optimized algorithm.

In this problem, your tasks are

**Algorithm 3** Newton-Raphson Method**Input:**  $\mathbf{x}_0 \in \mathbb{R}^n$ ,  $\epsilon_{\text{res}}^{\text{rel}}$ ,  $\epsilon_{\text{res}}^{\text{abs}}$ ,  $\epsilon_{\text{inc}}^{\text{abs}}$ ,  $N_{\text{max}}$ **Output:**  $\mathbf{x}^* \in \mathbb{R}^n$  such that  $\mathbf{R}(\mathbf{x}^*) \approx \mathbf{0}$ 

```

1: for  $k = 0, 1, 2, \dots, N_{\text{max}}$  do
2:    $\Delta \mathbf{x}_k = -\frac{\partial \mathbf{R}}{\partial \mathbf{x}}(\mathbf{x}_k)^{-1} \mathbf{R}(\mathbf{x}_k)$ 
3:    $\mathbf{x}_{k+1} = \mathbf{x}_k + \Delta \mathbf{x}_k$ 
4:   if  $\|\mathbf{R}(\mathbf{x}_{k+1})\| \leq \epsilon_{\text{res}}^{\text{rel}} \|\mathbf{R}(\mathbf{x}_0)\|$  or  $(\|\mathbf{R}(\mathbf{x}_{k+1})\| \leq \epsilon_{\text{res}}^{\text{abs}} \text{ AND } \|\mathbf{x}_{k+1} - \mathbf{x}_k\| \leq \epsilon_{\text{inc}}^{\text{abs}})$  then
5:      $\mathbf{x}^* = \mathbf{x}$ 
6:     return
7:   end if
8:   Issue convergence warning
9:    $\mathbf{x}^* = \mathbf{x}$ 
10: end for

```

- (1) Implement Newton-Raphson's method (Algorithm 3). The input for your Newton-Raphson implementation should be similar to that of `fsolve`, namely, a function handle, a starting point  $\mathbf{x}_0$ , and various tolerances. The function handle should accept a vector and return two outputs, the nonlinear function and Jacobian evaluated at the input vector. Starter code (`newton_raphson`) has been provided in the code distribution for this assignment. Feel free to use this as a starting point or start from scratch.
- (2) Implement the modified Newton-Raphson method that allows for
  - freezing the Jacobian matrix until  $\|\mathbf{R}(\mathbf{x}_k)\| > 0.1 \|\mathbf{R}(\mathbf{x}_{k-1})\|$
- (3) Make sure your algorithm returns the following output information (I recommend storing it in a structure, but it is not required)
  - the number of nonlinear iterations required for convergence,  $k$
  - the residual norm at convergence,  $\|\mathbf{R}(\mathbf{x}_k)\|$
  - the number of Jacobian computations required
  - the residual norm convergence history, i.e.  $\|\mathbf{R}(\mathbf{x}_j)\|$  for  $j \in \{0, \dots, k\}$
- (4) The piece of code below will setup a nonlinear system of equations of dimension 122 and define a function handle whose *first* argument is the nonlinear function and the *second* is the Jacobian. The solution of this nonlinear equation will solve an instance of the nonlinear truss problem defined in Problem 4 of this assignment. *To run this code, the folder `nltruss` must be on your MATLAB search path. Either copy all files in `nltruss` to your current folder or preferably use `addpath('/path/to/nltruss')` to add `nltruss` to your search path.* Starter code provided in `nonlinear_study.m`.

```

1 mesh = 'mesh3';
2 nel = eval([mesh, '()']);
3 [msh,bcs,mat] = setup_problem(0.1*ones(nel,1),mesh);
4 func = @(W) staticResidual(W,msh,mat,bcs);

```

Using the function handle defined above and your Newton-Raphson solver, conduct the following study:

- Solve the nonlinear system using `fsolve`. Turn the Jacobian option on for a fair comparison with your Newton-Raphson implementation. Make sure the tolerances, `TolX` and `TolFun`, and maximum iterations, `MaxIter`, correspond to those you use below. Record the number of iterations. Assume there is one Jacobian evaluation per iteration.



- Plot the mesh and its deformation (the solution of the nonlinear system) using `visualize_truss_loads(msh,bcs,mat,x,1)` where  $\mathbf{x}$  is the solution of the nonlinear system of equations (output of `fsolve`)
  - Solve the nonlinear system using your Newton-Raphson implementation with the following variations. For each, record the residual norm history, number of iterations required for convergence, and number of Jacobian evaluations.
    - Recompute the Jacobian at every iteration and use backslash to solve the linear system
    - Repeat the above three tasks, but reuse the Jacobian until  $\|\mathbf{R}(\mathbf{x}_k)\| > 0.1\|\mathbf{R}(\mathbf{x}_{k-1})\|$
  - I recommend all tolerances be set to  $10^{-8}$  and  $N_{\max} = 1000$ .
- (5) Create a table that summarizes the results of the above study, comparing number of nonlinear iterations, residual at convergence, and number of Jacobian evaluations required for each algorithm (3 in total - `fsolve` and your Newton-Raphson solver with and without recycling the Jacobian).
- (6) Create a plot of residual norm versus nonlinear iteration for both Newton-Raphson algorithms (not necessary for `fsolve`).
- (7) Briefly comment on the results.

## Problem 4

In this problem, you will gain experience with MATLAB's optimization functions, in particular `fmincon`. You will be given a general, *nonlinear* 2D truss code that will be used to define the optimization problems. Prior knowledge of truss analysis or mechanics is not necessary. A brief exposition of the truss analysis framework is provided below.

First, we begin with some terminology and nomenclature. A truss structure will be composed of  $n_v$  nodes or vertices connected by  $n_{el}$  truss elements. Each element  $e$  has an initial length  $L_e$ , cross-sectional area  $A_e$ , elastic modulus  $E_e$ , density  $\rho_e$ , and deformed length (i.e. after loads applied)  $\ell_e$ . A truss element can only carry *normal* forces, not shear or moments. This means the direction of the force within an element must align with the element. By definition, the intersection of two truss elements will be a *pinned* connection (otherwise the members would carry shear/moments). The force carried by element  $e$  will be denoted  $\mathbf{N}_e$ . Each vertex will be subject to forces that are applied externally or transferred from the truss elements. Let  $\mathbf{f}_i^{\text{int}} \in \mathbb{R}^2$  denote the force on vertex  $i$  due to the truss elements, usually called the *internal* force. Also, let  $\mathbf{f}_i^{\text{ext}} \in \mathbb{R}^2$  denote the force on vertex  $i$  due to external loads. Finally,  $\mathbf{u}_i \in \mathbb{R}^2$  denotes the displacement of vertex  $i$  induced by the loads.

Boundary conditions for truss elements can either be *displacement* or *force* boundary conditions. Force boundary conditions correspond to external loads, while displacement boundary conditions indicate prescribed displacements. Every node must have *exactly one* boundary condition for *each* degree of freedom (i.e. the  $x$ - and  $y$ -directions).

Static equilibrium at the nodes leads to the governing equation

$$\mathbf{f}^{\text{int}}(\mathbf{u}) = \mathbf{f}^{\text{ext}}. \quad (4)$$

There are several important quantities that will be used in this assignment as objective or constraints.

- Stress

$$\sigma_e = E_e \frac{\ell_e}{L_e} \quad (5)$$

- Weight

$$W = \sum_{e=1}^{n_{el}} \rho_e A_e \ell_e \quad (6)$$

- Compliance

$$C = \sum_{i=1}^{n_v} \mathbf{f}_i^T \mathbf{u}_i \quad (7)$$

In the truss program you are given, there will be three relevant data structures.

- msh - contains all mesh information
- bcs - contains all boundary condition information
- mat - contains all material information (including area)

The below piece of code demonstrates how to call use the nonlinear truss code. Pay particular attention to the commands up to `solve_truss` as this describes how to solve the nonlinear truss problem given a new vector of elemental areas. Additional starter code (`optimize_truss.m`) has been provided for you in the code distribution.

```
% Bridge mesh
mesh = 'mesh2';

% Determine number of elements in mesh and define vector of areas
nel = eval([mesh, '()']);
A = ones(nel,1);

% Setup mesh, BCs, and materials
[msh,bcs,mat] = setup_problem(A,mesh);

% Solve nonlinear truss equations (and sensitivity equations)
[U,dUdA,F,dFdA] = solve_truss(msh,mat,bcs);

% Plot mesh and deformation
% Vary last argument to scale deformation
visualize_truss_loads(msh,bcs,mat,U,10);

% Compute deformed lengths and sensitivities
[l,dldA] = compute_element_lengths(U,dUdA,msh);

% Compute weight and sensitivities
[W,dWdA] = compute_weight(U,dUdA,msh,mat);

% Compute compliance and sensitivities
[C,dCdA] = compute_compliance(U,dUdA,F,dFdA);

% Compute element force and sensitivities
[N,dNdA] = compute_internal_force(U,dUdA,msh,mat);

% Compute stress and sensitivities
[s,dsdA] = compute_stress(U,dUdA,msh,mat);
```

(1) Solve the following minimization problem

$$\begin{aligned} & \underset{\mathbf{A} \in \mathbb{R}^{n_{el}}}{\text{minimize}} && W(\mathbf{A}) \\ & \text{subject to} && |\sigma_e(\mathbf{A})| \leq \sigma_{\text{fail}} \\ & && 0.01 \leq \mathbf{A} \leq 1, \end{aligned} \quad (8)$$



for the mesh in Figure 4. Use an initial guess of  $\mathbf{A} = 0.1$ . This is accomplished using `mesh = ... 'mesh2'` in the code provided above. I recommend starting (and debugging) with `mesh = 'mesh1'` as the corresponding mesh has fewer elements and is faster to solve/optimize (Figure 3). Also, feel free to play around with `nseg` in `mesh2.m`, which controls the number of spans in the bridge.

- Use `fmincon` - try active set, interior point, and SQP algorithms
- Use both user-defined gradients (provided in `nltruss`) and gradients automatically computed with finite differences (`'GradObj'`, `'GradConstr'`).
- Comment on differences in optimal solution and timings.
- Plot the deformed truss with optimal areas (`visualize_truss_loads.m`)

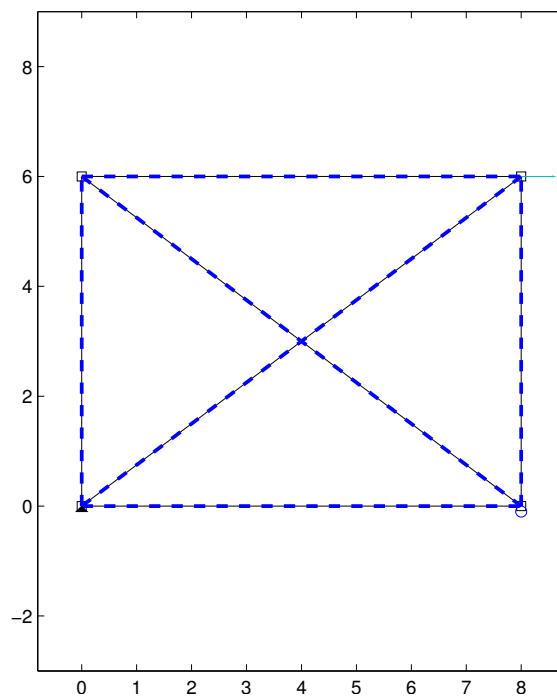


Figure 3: Simple Mesh (`mesh1`)

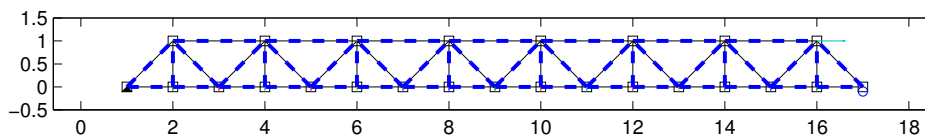


Figure 4: Truss Bridge Mesh (`mesh2`)

(2) Solve the following optimization problem

$$\begin{aligned}
 & \underset{\mathbf{A} \in \mathbb{R}^{n_{el}}}{\text{minimize}} && C(\mathbf{A}) \\
 & \text{subject to} && W(\mathbf{A}) \leq W(\mathbf{A}_0) \\
 & && |\sigma_e(\mathbf{A})| \leq \sigma_{\text{fail}} \\
 & && 0.01 \leq \mathbf{A} \leq 1,
 \end{aligned} \tag{9}$$

where  $\mathbf{A}_0 = 0.1$  is the initial guess.

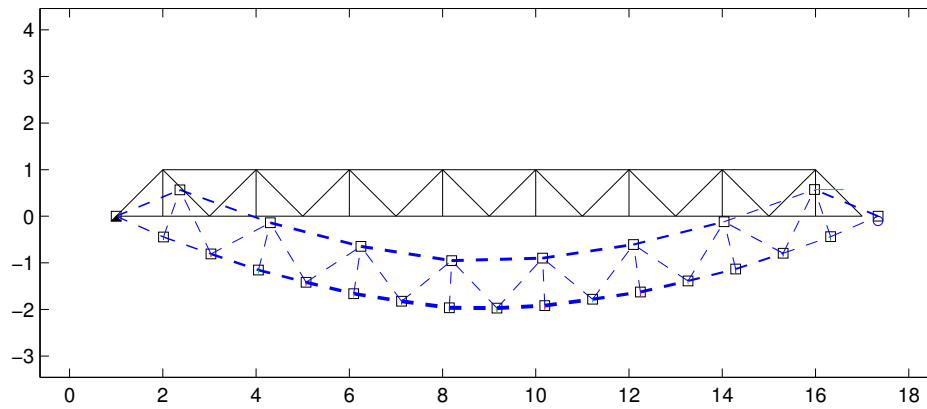


Figure 5: Truss Bridge - Optimized (Min Weight), Deformed (mesh2)

- Use `fmincon` - any optimization algorithm you choose
- Use user-defined gradients (provided in `nltruss`) ('`GradObj`', '`GradConstr`').
- Plot the deformed truss with optimal areas (`visualize_truss_loads.m`)

## References

- [1] N. Halko, P.-G. Martinsson, and J. A. Tropp, "Finding structure with randomness: Probabilistic algorithms for constructing approximate matrix decompositions," *SIAM review*, vol. 53, no. 2, pp. 217–288, 2011.