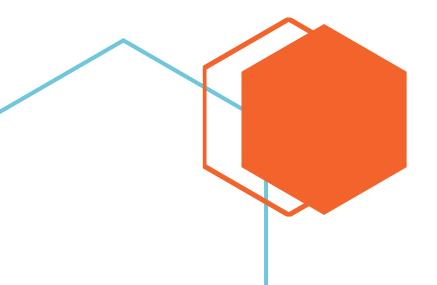
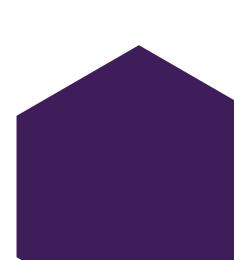


ADMM Project Report

This report is regarding the observations, calculations and scripting of ADMM algorithm used in various daily life applications. The algorithm shows how simple optimization can be even when extended to a million variables.





ADMM Project Report

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Acknowledgement

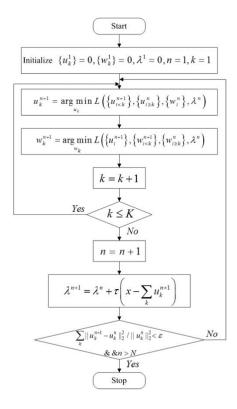
I would like to thank Mr. K.P Soman to grant me an opportunity to make this project report analysis and helping throughout the course. I would also like to thank Mr. Premjith for guiding and helping me throughout course.

What is ADMM?

The alternating direction method of multipliers (ADMM) is an algorithm that solves convex optimization problems by breaking them into smaller pieces, each of which are then easier to handle. It has recently found wide application in a number of areas.

Lagrangian Multiplier

In mathematical optimization, the method of Lagrange multipliers is a strategy for finding the local maxima and minima of a function subject to equality constraints. The basic idea is to convert a constrained problem into a form such that the derivative test of an unconstrained problem can still be applied.



Topics Chosen

- 1. Linear Programming
- 2. Huber fitting
- 3.SVM
- 4. Intersection of Polyhedra
- 5. Quadratic Programming
- 6. Lasso

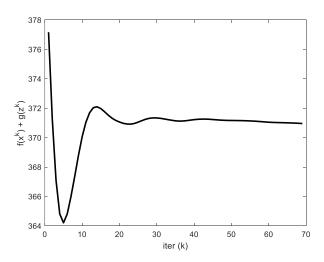
Linear Programming

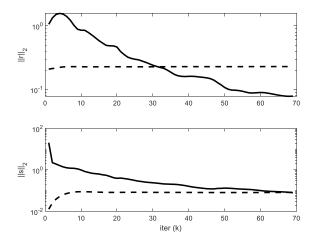
```
function [z, history] = linprog(c, A, b, rho, alpha)
% linprog Solve standard form LP via ADMM
% [x, history] = linprog(c, A, b, rho, alpha);
% Solves the following problem via ADMM:
% minimize
             C'*X
% subject to Ax = b, x >= 0
% The solution is returned in the vector x.
% history is a structure that contains the objective value, the primal and
% dual residual norms, and the tolerances for the primal and dual residual
% norms at each iteration.
% rho is the augmented Lagrangian parameter.
% alpha is the over-relaxation parameter (typical values for alpha are
% between 1.0 and 1.8).
% More information can be found in the paper linked at:
% http://www.stanford.edu/~boyd/papers/distr_opt_stat_learning_admm.html
t start = tic;
QUIET = 0;
MAX ITER = 1000;
ABSTOL = 1e-4;
RELTOL = 1e-2;
[m n] = size(A);
x = zeros(n,1);
z = zeros(n,1);
u = zeros(n,1);
if ~QUIET
  fprintf('%3s\t%10s\t%10s\t%10s\t%10s\n', 'iter', ...
   'r norm', 'eps pri', 's norm', 'eps dual', 'objective');
end
for k = 1:MAX_ITER
  % x-update
  tmp = [rho*eye(n), A'; A, zeros(m)] \setminus [rho*(z - u) - c; b];
  x = tmp(1:n);
  % z-update with relaxation
  zold = z;
  x_hat = alpha*x + (1 - alpha)*zold;
  z = pos(x_hat + u);
  u = u + (x_hat - z);
  % diagnostics, reporting, termination checks
```

```
history.r_norm(k) = norm(x - z);
  history.s norm(k) = norm(-rho*(z - zold));
  history.eps_pri(k) = sqrt(n)*ABSTOL + RELTOL*max(norm(x), norm(-z));
  history.eps_dual(k) = sqrt(n)*ABSTOL + RELTOL*norm(rho*u);
  if ~QUIET
    fprintf('%3d\t%10.4f\t%10.4f\t%10.4f\t%10.2f\n', k, ...
       history.r_norm(k), history.eps_pri(k), ...
       history.s_norm(k), history.eps_dual(k), history.objval(k));
  end
  if (history.r_norm(k) < history.eps_pri(k) && ...
    history.s_norm(k) < history.eps_dual(k))
     break;
  end
end
if ~QUIET
  toc(t_start);
end
end
function obj = objective(c, x)
  obj = c'*x;
end
Example
randn('state', 0);
rand('state', 0);
n = 500; % dimension of x
m = 400; % number of equality constraints
c = rand(n,1) + 0.5; % create nonnegative price vector with mean 1
x0 = abs(randn(n,1)); % create random solution vector
A = abs(randn(m,n)); % create random, nonnegative matrix A
b = A*x0;
[x history] = linprog(c, A, b, 1.0, 1.0);
K = length(history.objval);
h = figure;
```

history.objval(k) = objective(c, x);

```
plot(1:K, history.objval, 'k', 'MarkerSize', 10, 'LineWidth', 2); ylabel('f(x^k) + g(z^k)'); xlabel('iter (k)'); g = figure; subplot(2,1,1); semilogy(1:K, max(1e-8, history.r_norm), 'k', ... 1:K, history.eps_pri, 'k--', 'LineWidth', 2); ylabel('||r||_2'); <math display="block">subplot(2,1,2); semilogy(1:K, max(1e-8, history.s_norm), 'k', ... 1:K, history.eps_dual, 'k--', 'LineWidth', 2); ylabel('||s||_2'); xlabel('iter (k)');
```



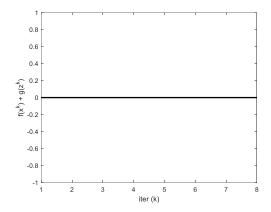


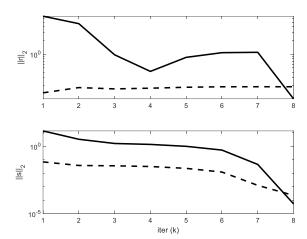
Intersection of Polyhedra

```
function [z, history] = polyhedra_intersection(A1, b1, A2, b2, rho, alpha)
t start = tic;
QUIET = 0;
MAX_{ITER} = 1000;
ABSTOL = 1e-4;
RELTOL = 1e-2;
n = size(A1,2);
x = zeros(n,1);
z = zeros(n,1);
u = zeros(n,1);
if ~QUIET
  fprintf('%3s\t%10s\t%10s\t%10s\t%10s\t%10s\n', 'iter', ...
   'r norm', 'eps pri', 's norm', 'eps dual', 'objective');
end
for k = 1:MAX_ITER
  % x-update
  % use cvx to find point in first polyhedra
  cvx_begin quiet
    variable x(n)
    minimize (sum_square(x - (z - u)))
    subject to
       A1*x <= b1
  cvx end
  % z-update with relaxation
  zold = z;
  x_hat = alpha*x + (1 - alpha)*zold;
  % use cvx to find point in second polyhedra
  cvx_begin quiet
    variable z(n)
    minimize (sum_square(x_hat - (z - u)))
    subject to
       A2*z \le b2
  cvx_end
  u = u + (x_hat - z);
  % diagnostics, reporting, termination checks
```

```
history.objval(k) = 0;
  history.r_norm(k) = norm(x - z);
  history.s_norm(k) = norm(-rho*(z - zold));
  history.eps pri(k) = sqrt(n)*ABSTOL + RELTOL*max(norm(x), norm(-z));
  history.eps_dual(k) = sqrt(n)*ABSTOL + RELTOL*norm(rho*u);
  if ~QUIET
    fprintf('%3d\t%10.4f\t%10.4f\t%10.4f\t%10.2f\n', k, ...
      history.r_norm(k), history.eps_pri(k), ...
       history.s_norm(k), history.eps_dual(k), history.objval(k));
  end
  if (history.r_norm(k) < history.eps_pri(k) && ...
    history.s_norm(k) < history.eps_dual(k))
     break;
  end
end
if ~QUIET
  toc(t_start);
end
end
Example:
randn('state', 0);
rand('state', 0);
         % dimension of variable
n = 5:
m1 = 10; % number of faces for polyhedra 1
m2 = 12; % number of faces for polyhedra 2
c1 = 10*randn(n,1);
                        % center of polyhedra 1
c2 = -10*randn(n,1);
                        % center of polyhedra 2
% consider the following picture:
%
%
     a1
% c ----> x
% from the center "c", we travel along vector "a1" (not necessarily a unit
% vector) until we reach x. at "x", a1 'x = b. a point y is to the left of x
% if a1'y \leq b.
%
% pick m1 random directions with different magnitudes
A1 = diag(1 + rand(m1,1))*randn(m1,n);
% the value of b is found by traveling from the center along the normal
```

```
% vectors in A1 and taking its inner product with A1.
b1 = diag(A1*(c1*ones(1,m1) + A1'));
% pick m2 random directions with different magnitudes
A2 = diag(1 + rand(m2,1))*randn(m2,n);
% the value of b is found by traveling from the center along the normal
% vectors in A1 and taking its inner product with A1.
b2 = diag(A2*(c2*ones(1,m2) + A2'));
% find the distance between the two polyhedra--make sure they overlap by
% checking if the distance is 0
cvx_begin quiet
  variables x(n) y(n)
  minimize sum_square(x - y)
  subject to
    A1*x <= b1
    A2*y \le b2
cvx_end
% if the distance is not 0, expand A1 and A2 by a little more than half the
% distance
if norm(x-y) > 1e-4,
  A1 = (1 + 0.5*norm(x-y))*A1;
  A2 = (1 + 0.5*norm(x-y))*A2;
  % recompute b's as appropriate
  b1 = diag(A1*(c1*ones(1,m1) + A1'));
  b2 = diag(A2*(c2*ones(1,m2) + A2'));
end
[x history] = polyhedra_intersection(A1, b1, A2, b2, 1.0, 1.0);
K = length(history.objval);
h = figure;
plot(1:K, history.objval, 'k', 'MarkerSize', 10, 'LineWidth', 2);
ylabel('f(x^k) + g(z^k)'); xlabel('iter (k)');
g = figure;
subplot(2,1,1);
semilogy(1:K, max(1e-8, history.r_norm), 'k', ...
  1:K, history.eps_pri, 'k--', 'LineWidth', 2);
ylabel('| |r| |_2');
subplot(2,1,2);
semilogy(1:K, max(1e-8, history.s_norm), 'k', ...
  1:K, history.eps_dual, 'k--', 'LineWidth', 2);
ylabel('| |s| |_2'); xlabel('iter (k)');
```





Support Vector Machine

```
function [xave, history] = linear svm(A, lambda, p, rho, alpha)
% linear_svm Solve linear support vector machine (SVM) via ADMM
% [x, history] = linear_svm(A, lambda, p, rho, alpha)
% Solves the following problem via ADMM:
  minimize (1/2) \mid |w| \mid 2^2 + \lambda s
% where A is a matrix given by [-y_j*x_j -y_j], lambda is a
% regularization parameter, and p is a partition of the observations in to
% different subsystems.
% The function h_i(w, b) is a hinge loss on the variables w and b.
% It corresponds to h_j(w,b) = (Ax + 1)_+, where x = (w,b).
% This function implements a *distributed* SVM that runs its updates
% serially.
% The solution is returned in the vector x = (w,b).
% history is a structure that contains the objective value, the primal and
% dual residual norms, and the tolerances for the primal and dual residual
% norms at each iteration.
% rho is the augmented Lagrangian parameter.
% alpha is the over-relaxation parameter (typical values for alpha are
% between 1.0 and 1.8).
% More information can be found in the paper linked at:
% http://www.stanford.edu/~boyd/papers/distr opt stat learning admm.html
t_start = tic;
QUIET = 0;
MAX ITER = 1000;
ABSTOL = 1e-4;
RELTOL = 1e-2;
[m, n] = size(A);
N = max(p);
% group samples together
for i = 1:N,
  tmp{i} = A(p==i,:);
end
A = tmp;
x = zeros(n,N);
z = zeros(n,N);
u = zeros(n,N);
```

```
if ~QUIET
  fprintf('%3s\t%10s\t%10s\t%10s\t%10s\n', 'iter', ...
   'r norm', 'eps pri', 's norm', 'eps dual', 'objective');
end
for k = 1:MAX_ITER
       % x-update
  for i = 1:N,
    cvx_begin quiet
       variable x_var(n)
       minimize (sum(pos(A{i}*x_var + 1)) + rho/2*sum_square(x_var - z(:,i) + u(:,i)))
    cvx_end
    x(:,i) = x_var;
  end
  xave = mean(x,2);
  % z-update with relaxation
  zold = z;
  x_hat = alpha*x + (1-alpha)*zold;
  z = N*rho/(1/lambda + N*rho)*mean(x hat + u, 2);
  z = z*ones(1,N);
  % u-update
  u = u + (x_hat - z);
  % diagnostics, reporting, termination checks
  history.objval(k) = objective(A, lambda, p, x, z);
  history.r norm(k) = norm(x - z);
  history.s_norm(k) = norm(-rho*(z - zold));
  history.eps_pri(k) = sqrt(n)*ABSTOL + RELTOL*max(norm(x), norm(-z));
  history.eps_dual(k) = sqrt(n)*ABSTOL + RELTOL*norm(rho*u);
  if ~QUIET
    fprintf('%3d\t%10.4f\t%10.4f\t%10.4f\t%10.2f\n', k, ...
       history.r_norm(k), history.eps_pri(k), ...
       history.s_norm(k), history.eps_dual(k), history.objval(k));
  end
  if (history.r_norm(k) < history.eps_pri(k) && ...
    history.s norm(k) < history.eps dual(k))
     break:
```

```
end
end
if ~QUIET
  toc(t_start);
end
end
function obj = objective(A, lambda, p, x, z)
  obj = hinge_loss(A,x) + 1/(2*lambda)*sum_square(z(:,1));
end
function val = hinge_loss(A,x)
  val = 0;
  for i = 1:length(A)
    val = val + sum(pos(A{i}*x(:,i) + 1));
  end
end
Example:
rand('seed', 0);
randn('seed', 0);
n = 2;
m = 200;
N = m/2;
M = m/2;
% positive examples
Y = [1.5+0.9*randn(1,0.6*N), 1.5+0.7*randn(1,0.4*N);
2*(randn(1,0.6*N)+1), 2*(randn(1,0.4*N)-1)];
% negative examples
X = [-1.5+0.9*randn(1,0.6*M), -1.5+0.7*randn(1,0.4*M);
2*(randn(1,0.6*M)-1), 2*(randn(1,0.4*M)+1)];
x = [X Y];
y = [ones(1,N) - ones(1,M)];
A = [-((ones(n,1)*y).*x)'-y'];
xdat = x';
lambda = 1.0;
% partition the examples up in the worst possible way
% (subsystems only have positive or negative examples)
p = zeros(1,m);
p(y == 1) = sort(randi([1 10], sum(y==1), 1));
p(y == -1) = sort(randi([11 20], sum(y == -1), 1));
```

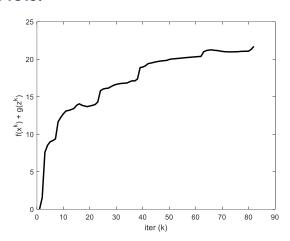
```
[x history] = linear_svm(A, lambda, p, 1.0, 1.0);

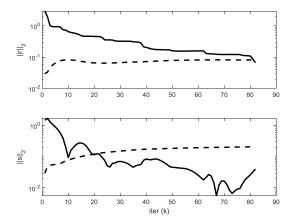
K = length(history.objval);

h = figure;
plot(1:K, history.objval, 'k', 'MarkerSize', 10, 'LineWidth', 2);
ylabel('f(x^k) + g(z^k)'); xlabel('iter (k)');

g = figure;
subplot(2,1,1);
semilogy(1:K, max(1e-8, history.r_norm), 'k', ...
1:K, history.eps_pri, 'k--', 'LineWidth', 2);
ylabel('||r||_2');

subplot(2,1,2);
semilogy(1:K, max(1e-8, history.s_norm), 'k', ...
1:K, history.eps_dual, 'k--', 'LineWidth', 2);
ylabel('||s||_2'); xlabel('iter (k)');
```



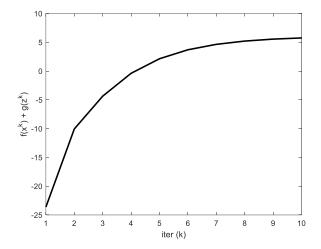


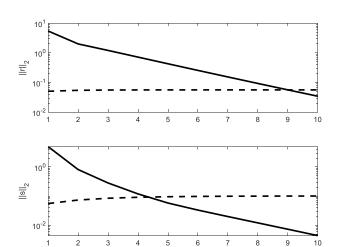
Quadratic Programming

```
function [z, history] = quadprog(P, q, r, lb, ub, rho, alpha)
% quadprog Solve standard form box-constrained QP via ADMM
% [x, history] = quadprog(P, q, r, lb, ub, rho, alpha)
% Solves the following problem via ADMM:
% minimize
               (1/2)*x'*P*x + q'*x + r
  subject to lb <= x <= ub
% The solution is returned in the vector x.
% history is a structure that contains the objective value, the primal and
% dual residual norms, and the tolerances for the primal and dual residual
% norms at each iteration.
%
% rho is the augmented Lagrangian parameter.
% alpha is the over-relaxation parameter (typical values for alpha are
% between 1.0 and 1.8).
%
% More information can be found in the paper linked at:
% http://www.stanford.edu/~boyd/papers/distr_opt_stat_learning_admm.html
%
t start = tic;
QUIET = 0;
MAX ITER = 1000;
ABSTOL = 1e-4;
RELTOL = 1e-2;
n = size(P,1);
x = zeros(n,1);
z = zeros(n,1);
u = zeros(n, 1);
if ~QUIET
  fprintf('%3s\t%10s\t%10s\t%10s\t%10s\n', 'iter', ...
   'r norm', 'eps pri', 's norm', 'eps dual', 'objective');
end
for k = 1:MAX_{ITER}
  if k > 1
```

```
x = R \setminus (R' \setminus (rho^*(z - u) - q));
  else
     R = \text{chol}(P + \text{rho*eye(n)});
     x = R \setminus (R' \setminus (rho^*(z - u) - q));
  end
  % z-update with relaxation
  zold = z;
  x hat = alpha*x + (1-alpha)*zold;
  z = min(ub, max(lb, x_hat + u));
  % u-update
  u = u + (x_hat - z);
  % diagnostics, reporting, termination checks
  history.objval(k) = objective(P, q, r, x);
  history.r_norm(k) = norm(x - z);
  history.s_norm(k) = norm(-rho*(z - zold));
  history.eps_pri(k) = sqrt(n)*ABSTOL + RELTOL*max(norm(x), norm(-z));
  history.eps_dual(k) = sqrt(n)*ABSTOL + RELTOL*norm(rho*u);
  if ~QUIET
     fprintf('%3d\t%10.4f\t%10.4f\t%10.4f\t%10.4f\t%10.2f\n', k, ...
       history.r_norm(k), history.eps_pri(k), ...
       history.s_norm(k), history.eps_dual(k), history.objval(k));
  end
  if (history.r_norm(k) < history.eps_pri(k) && ...
    history.s_norm(k) < history.eps_dual(k))
     break;
  end
end
if ~QUIET
  toc(t_start);
end
end
function obj = objective(P, q, r, x)
  obj = 0.5*x'*P*x + q'*x + r;
end
```

```
Example:
randn('state', 0);
rand('state', 0);
n = 100;
% generate a well-conditioned positive definite matrix
% (for faster convergence)
P = rand(n);
P = P + P';
[V D] = eig(P);
P = V*diag(1+rand(n,1))*V';
q = randn(n,1);
r = randn(1);
I = randn(n, 1);
u = randn(n,1);
lb = min(l, u);
ub = max(l,u);
[x history] = quadprog(P, q, r, lb, ub, 1.0, 1.0);
K = length(history.objval);
h = figure;
plot(1:K, history.objval, 'k', 'MarkerSize', 10, 'LineWidth', 2);
ylabel('f(x^k) + g(z^k)'); xlabel('iter (k)');
g = figure;
subplot(2,1,1);
semilogy(1:K, max(1e-8, history.r_norm), 'k', ...
  1:K, history.eps_pri, 'k--', 'LineWidth', 2);
ylabel('||r||_2');
subplot(2,1,2);
semilogy(1:K, max(1e-8, history.s_norm), 'k', ...
  1:K, history.eps_dual, 'k--', 'LineWidth', 2);
ylabel('| |s| |_2'); xlabel('iter (k)');
```





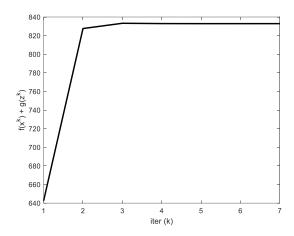
Huber fitting

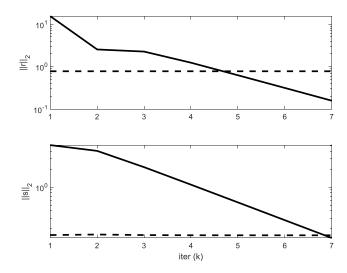
```
function [x, history] = huber_fit(A, b, rho, alpha)
% huber_fit Solves a robust fitting problem
% [z, history] = huber_fit(A, b, rho, alpha);
% solves the following problem via ADMM:
  minimize 1/2*sum(huber(A*x - b))
% with variable x.
% The solution is returned in the vector x.
% history is a structure that contains the objective value, the primal and
% dual residual norms, and the tolerances for the primal and dual residual
% norms at each iteration.
% rho is the augmented Lagrangian parameter.
% alpha is the over-relaxation parameter (typical values for alpha are
% between 1.0 and 1.8).
%
% More information can be found in the paper linked at:
% http://www.stanford.edu/~boyd/papers/distr_opt_stat_learning_admm.html
t_start = tic;
QUIET = 0;
MAX_{ITER} = 1000;
ABSTOL = 1e-4;
RELTOL = 1e-2;
[m, n] = size(A);
% save a matrix-vector multiply
Atb = A'*b;
x = zeros(n,1);
z = zeros(m,1);
u = zeros(m, 1);
% cache factorization
[L U] = factor(A);
```

```
if ~QUIET
  fprintf('%3s\t%10s\t%10s\t%10s\t%10s\n', 'iter', ...
   'r norm', 'eps pri', 's norm', 'eps dual', 'objective');
end
for k = 1:MAX_ITER
  % x-update
  q = Atb + A'*(z - u);
  x = U \setminus (L \setminus q);
  % z-update with relaxation
  zold = z;
  Ax_hat = alpha*A*x + (1-alpha)*(zold + b);
  tmp = Ax hat - b + u;
  z = rho/(1 + rho)*tmp + 1/(1 + rho)*shrinkage(tmp, 1 + 1/rho);
  u = u + (Ax_hat - z - b);
  % diagnostics, reporting, termination checks
  history.objval(k) = objective(z);
  history.r_norm(k) = norm(A*x - z - b);
  history.s_norm(k) = norm(-rho*A'*(z - zold));
  history.eps_pri(k) = sqrt(n)*ABSTOL + RELTOL*max([norm(A*x), norm(-z), norm(b)]);
  history.eps_dual(k) = sqrt(n)*ABSTOL + RELTOL*norm(rho*u);
  if ~QUIET
    fprintf('%3d\t%10.4f\t%10.4f\t%10.4f\t%10.4f\t%10.2f\n', k, ...
       history.r_norm(k), history.eps_pri(k), ...
       history.s_norm(k), history.eps_dual(k), history.objval(k));
  end
  if history.r norm(k) < history.eps pri(k) && ...
    history.s_norm(k) < history.eps_dual(k);
    break
  end
end
if ~QUIET
  toc(t_start);
end
end
```

```
function p = objective(z)
  p = (1/2*sum(huber(z)));
end
function z = shrinkage(x, kappa)
  z = pos(1 - kappa./abs(x)).*x;
end
function [L U] = factor(A)
  [m, n] = size(A);
  if (m \ge n) % if skinny
    L = chol(A'*A, 'lower');
  end
  % force matlab to recognize the upper / lower triangular structure
  L = sparse(L);
  U = sparse(L');
end
Example:
randn('seed', 0);
rand('seed',0);
              % number of examples
m = 5000;
n = 200:
            % number of features
x0 = randn(n,1);
A = randn(m,n);
A = A*spdiags(1./norms(A)',0,n,n); % normalize columns
b = A*x0 + sqrt(0.01)*randn(m,1);
b = b + 10*sprand(m,1,200/m); % add sparse, large noise
[x history] = huber_fit(A, b, 1.0, 1.0);
K = length(history.objval);
h = figure;
plot(1:K, history.objval, 'k', 'MarkerSize', 10, 'LineWidth', 2);
ylabel('f(x^k) + g(z^k)'); xlabel('iter (k)');
g = figure;
subplot(2,1,1);
semilogy(1:K, max(1e-8, history.r_norm), 'k', ...
  1:K, history.eps_pri, 'k--', 'LineWidth', 2);
ylabel('||r||_2');
subplot(2,1,2);
```

semilogy(1:K, max(1e-8, history.s_norm), 'k', ... 1:K, history.eps_dual, 'k--', 'LineWidth', 2); ylabel('||s||_2'); xlabel('iter (k)');





Lasso

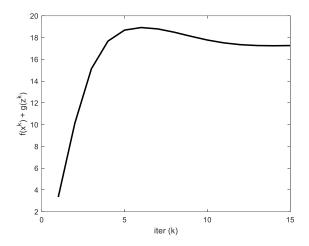
```
function [z, history] = lasso(A, b, lambda, rho, alpha)
% lasso Solve lasso problem via ADMM
% [z, history] = lasso(A, b, lambda, rho, alpha);
% Solves the following problem via ADMM:
  minimize 1/2* | | Ax - b | |_2^2 + \lambda | | x | |_1
% The solution is returned in the vector x.
% history is a structure that contains the objective value, the primal and
% dual residual norms, and the tolerances for the primal and dual residual
% norms at each iteration.
% rho is the augmented Lagrangian parameter.
% alpha is the over-relaxation parameter (typical values for alpha are
% between 1.0 and 1.8).
%
% More information can be found in the paper linked at:
% http://www.stanford.edu/~boyd/papers/distr_opt_stat_learning_admm.html
%
t_start = tic;
QUIET = 0;
MAX ITER = 1000;
ABSTOL = 1e-4;
RELTOL = 1e-2;
[m, n] = size(A);
% save a matrix-vector multiply
Atb = A'*b:
x = zeros(n,1);
z = zeros(n,1);
u = zeros(n,1);
% cache the factorization
[L U] = factor(A, rho);
if ~QUIET
  fprintf('%3s\t%10s\t%10s\t%10s\t%10s\n', 'iter', ...
   'r norm', 'eps pri', 's norm', 'eps dual', 'objective');
```

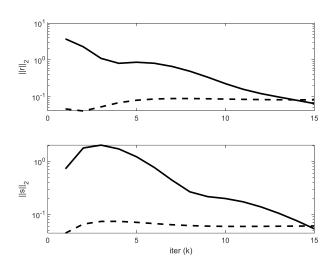
```
end
```

```
for k = 1:MAX_ITER
  % x-update
  q = Atb + rho^*(z - u); % temporary value
  if (m \ge n) % if skinny
    x = U \setminus (L \setminus q);
              % if fat
    x = q/rho - (A'*(U \setminus (L \setminus (A*q))))/rho^2;
  end
  % z-update with relaxation
  zold = z;
  x_hat = alpha*x + (1 - alpha)*zold;
  z = shrinkage(x_hat + u, lambda/rho);
  % u-update
  u = u + (x_hat - z);
  % diagnostics, reporting, termination checks
  history.objval(k) = objective(A, b, lambda, x, z);
  history.r_norm(k) = norm(x - z);
  history.s_norm(k) = norm(-rho*(z - zold));
  history.eps_pri(k) = sqrt(n)*ABSTOL + RELTOL*max(norm(x), norm(-z));
  history.eps_dual(k) = sqrt(n)*ABSTOL + RELTOL*norm(rho*u);
  if ~QUIET
    fprintf('%3d\t%10.4f\t%10.4f\t%10.4f\t%10.4f\t%10.2f\n', k, ...
       history.r_norm(k), history.eps_pri(k), ...
       history.s_norm(k), history.eps_dual(k), history.objval(k));
  end
  if (history.r_norm(k) < history.eps_pri(k) && ...
    history.s_norm(k) < history.eps_dual(k))
     break;
  end
end
if ~QUIET
  toc(t_start);
end
end
function p = objective(A, b, lambda, x, z)
```

```
p = (1/2*sum((A*x - b).^2) + lambda*norm(z,1));
end
function z = shrinkage(x, kappa)
  z = max(0, x - kappa) - max(0, -x - kappa);
end
function [L U] = factor(A, rho)
  [m, n] = size(A);
  if (m \ge n) % if skinny
    L = \text{chol}(A'*A + \text{rho*speye(n)}, 'lower');
              % if fat
    L = \text{chol}(\text{speye(m)} + 1/\text{rho}^*(A^*A'), 'lower');
  end
  % force matlab to recognize the upper / lower triangular structure
  L = sparse(L);
  U = sparse(L');
end
Example:
randn('seed', 0);
rand('seed',0);
m = 1500;
              % number of examples
n = 5000;
             % number of features
p = 100/n;
             % sparsity density
x0 = sprandn(n,1,p);
A = randn(m,n);
A = A*spdiags(1./sqrt(sum(A.^2))',0,n,n); % normalize columns
b = A*x0 + sqrt(0.001)*randn(m,1);
lambda_max = norm(A'*b, 'inf');
lambda = 0.1*lambda_max;
[x history] = lasso(A, b, lambda, 1.0, 1.0);
K = length(history.objval);
h = figure;
plot(1:K, history.objval, 'k', 'MarkerSize', 10, 'LineWidth', 2);
ylabel('f(x^k) + g(z^k)'); xlabel('iter (k)');
g = figure;
subplot(2,1,1);
semilogy(1:K, max(1e-8, history.r_norm), 'k', ...
```

```
1:K, history.eps_pri, 'k--', 'LineWidth', 2); ylabel('||r||_2'); subplot(2,1,2); semilogy(1:K, max(1e-8, history.s_norm), 'k', ... 1:K, history.eps_dual, 'k--', 'LineWidth', 2); ylabel('||s||_2'); xlabel('iter (k)');
```





References

https://web.stanford.edu/~boyd/papers/admm/