

```
% Import data
data_HW3 = readtable("Top 100 Genes.xlsx","ReadRowNames",true);
```

```
% Inspect table
head(data_HW3,5)
```

```
ans = 5×101 table
```

...

	A_23_P342744	A_24_P246891	A_23_P24535	A_23_P75362	A_32_P76399
1 GENE_SYMBOL	'LIX1L'	'NEU4'	'TTC12'	'IFITM10'	'EIF3L'
2 GENE_NAME	'Lix1 homolog...	'sialidase 4'	'tetratrico...	'interferon i...	'eukaryotic t...
3 GSM1912920	'-0.0999'	'-0.96104'	'0.28358'	'-0.31814'	'-0.26628'
4 GSM1912921	'1.0914'	'1.6402'	'-1.4203'	'-0.54017'	'-0.089449'
5 GSM1912922	'-0.37438'	'1.7162'	'0.17951'	'0.75482'	'0.4971'

```
test_patients = readcell("Patient IDs - Test Set.xlsx");
train_patients = readcell("Patient IDs - Training Set.xlsx");
```

```
% Using readmatrix correctly loads data as class "double"
%data_HW3_2 = readmatrix("Top 100 Genes & Rand 15 Patients.xlsx");
```

```
% Create new X and Y matrices (have to convert class if using readtable)
Xtrain = str2double(data_HW3{train_patients, 1:100});
Xtest = str2double(data_HW3{test_patients, 1:100});

Ytrain = table2array(data_HW3(train_patients, end));
Ytest = table2array(data_HW3(test_patients, end));
```

Multi-linear Regression using Hold-out Validation

Normal Regression

```
% Find correlation of all 100 genes and then extract data for top 15 genes
r_100 = corr(Xtrain,Ytrain);
[r_15,index_15] = maxk(abs(r_100),15);
```

```
% Create new training and test predictor data sets with top 15 genes
Xtrain_15 = Xtrain(:,index_15);
Xtest_15 = Xtest(:,index_15);
```

```
mdl = fitlm(Xtrain_15,Ytrain)
```

```
mdl =
Linear regression model:
y ~ 1 + x1 + x2 + x3 + x4 + x5 + x6 + x7 + x8 + x9 + x10 + x11 + x12 + x13 + x14 + x15
```

Estimated Coefficients:

	Estimate	SE	tStat	pValue
(Intercept)	627.65	54.797	11.454	2.7843e-12
x1	163.8	175.67	0.93244	0.35881
x2	-31.076	115.96	-0.26799	0.79061
x3	-1.6132	93.205	-0.017308	0.98631
x4	-4.73	140.2	-0.033738	0.97332
x5	252.87	152.73	1.6557	0.10857
x6	134.88	119.03	1.1332	0.26641
x7	-56.536	128.64	-0.4395	0.66356
x8	107.58	63.946	1.6824	0.10322
x9	145.38	70.929	2.0497	0.049532
x10	141.14	51.1	2.762	0.0098648
x11	177.62	122.02	1.4556	0.15623
x12	-276.22	212.1	-1.3023	0.20306
x13	-19.133	92.052	-0.20785	0.8368
x14	-257.19	123.27	-2.0864	0.045844
x15	48.972	119.86	0.40858	0.68585

Number of observations: 45, Error degrees of freedom: 29

Root Mean Squared Error: 212

R-squared: 0.718, Adjusted R-Squared: 0.572

F-statistic vs. constant model: 4.92, p-value = 0.00012

```
Ypred_norm = predict(mdl,Xtest_15);
r_norm = corr(Ytest,Ypred_norm)
```

```
r_norm = 0.0816
```

```
r2_norm = r_norm^2
```

```
r2_norm = 0.0067
```

```
RMSE = sqrt(mean((Ypred_norm-Ytest).^2))
```

```
RMSE = 359.6592
```

```
avg_error = mean(abs(Ypred_norm-Ytest))
```

```
avg_error = 296.3538
```

Lasso Regression

```
[B1, Fit] = lasso(Xtrain_15,Ytrain,'CV',5);
B1_coeff = B1(:,Fit.Index1SE)
```

```
B1_coeff = 15x1
    143.2955
         0
    15.2578
         0
    28.6057
     3.0056
         0
    34.1621
    38.1426
    30.7815
```

:

```
B1_intercept = Fit.Intercept(Fit.Index1SE)
```

```
B1_intercept = 599.6518
```

```
Ypred_lasso = Xtest_15 * B1_coeff + B1_intercept;
```

```
r_lasso = corr(Ypred_lasso,Ytest)
```

```
r_lasso = 0.1174
```

```
r2_lasso = r_lasso^2
```

```
r2_lasso = 0.0138
```

```
RMSE_lasso = sqrt(mean((Ypred_lasso-Ytest).^2))
```

```
RMSE_lasso = 314.1729
```

```
avg_error_lasso = mean(abs(Ypred_lasso-Ytest))
```

```
avg_error_lasso = 257.2454
```

Stepwise Regression

```
[B2,~,~,~,stats] = stepwisefit(Xtrain_15,Ytrain);
```

```
Initial columns included: none
```

```
Step 1, added column 1, p=3.57549e-06
```

```
Step 2, added column 10, p=0.000962066
```

```
Step 3, added column 15, p=0.0485883
```

```
Final columns included: 1 10 15
```

'Coeff'	'Std.Err.'	'Status'	'P'
[393.7172]	[90.9898]	'In'	[9.4619e-05]
[-0.3560]	[99.5721]	'Out'	[0.9972]
[64.4157]	[69.1039]	'Out'	[0.3568]
[52.2210]	[64.1681]	'Out'	[0.4206]
[43.3180]	[90.0316]	'Out'	[0.6330]
[42.2050]	[67.8657]	'Out'	[0.5375]
[29.5171]	[95.3956]	'Out'	[0.7586]
[74.9424]	[55.4607]	'Out'	[0.1842]
[100.5944]	[53.7852]	'Out'	[0.0688]
[65.2065]	[31.8982]	'In'	[0.0474]
[131.6148]	[87.8463]	'Out'	[0.1419]
[5.7396]	[88.2548]	'Out'	[0.9485]
[88.5513]	[70.0971]	'Out'	[0.2138]
[-38.5347]	[82.3811]	'Out'	[0.6425]
[149.7515]	[73.6695]	'In'	[0.0486]

```
Ypred_step = Xtest_15(:,[1,10,15])*B2([1,10,15]) + stats.intercept;
```

```
r_stepwise = corr(Ypred_step,Ytest)
```

```
r_stepwise = 0.1894
```

```
r2_stepwise = r_stepwise^2
```

```
r2_stepwise = 0.0359
```

```
RMSE_stepwise = sqrt(mean((Ypred_step-Ytest).^2))
```

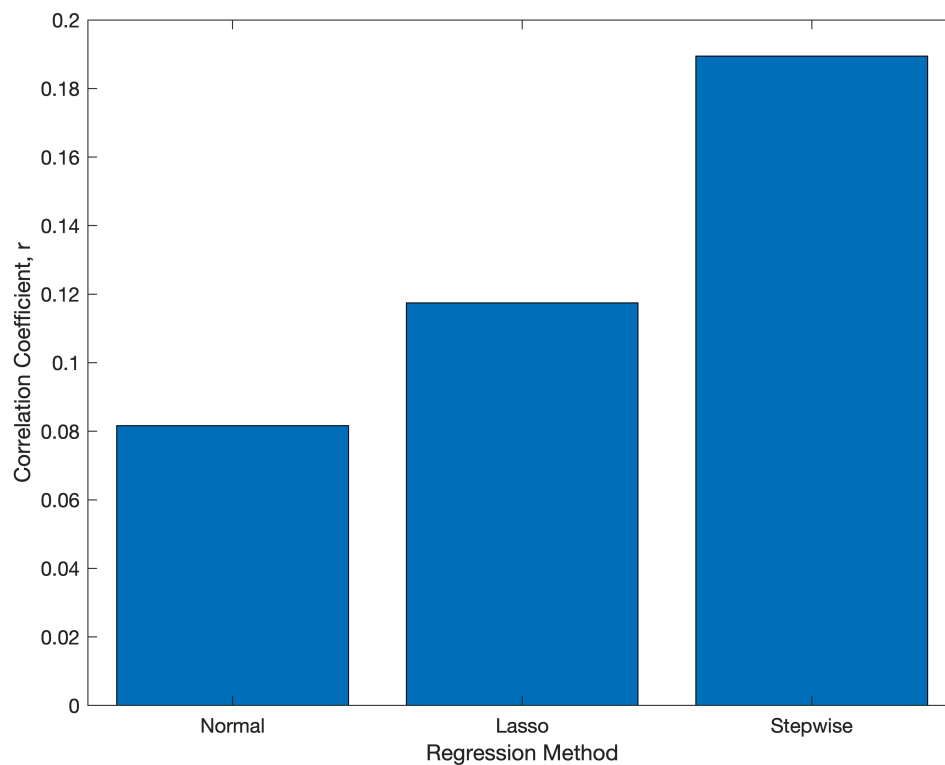
```
RMSE_stepwise = 347.2721
```

```
avg_error_stepwise = mean(abs(Ypred_step-Ytest))
```

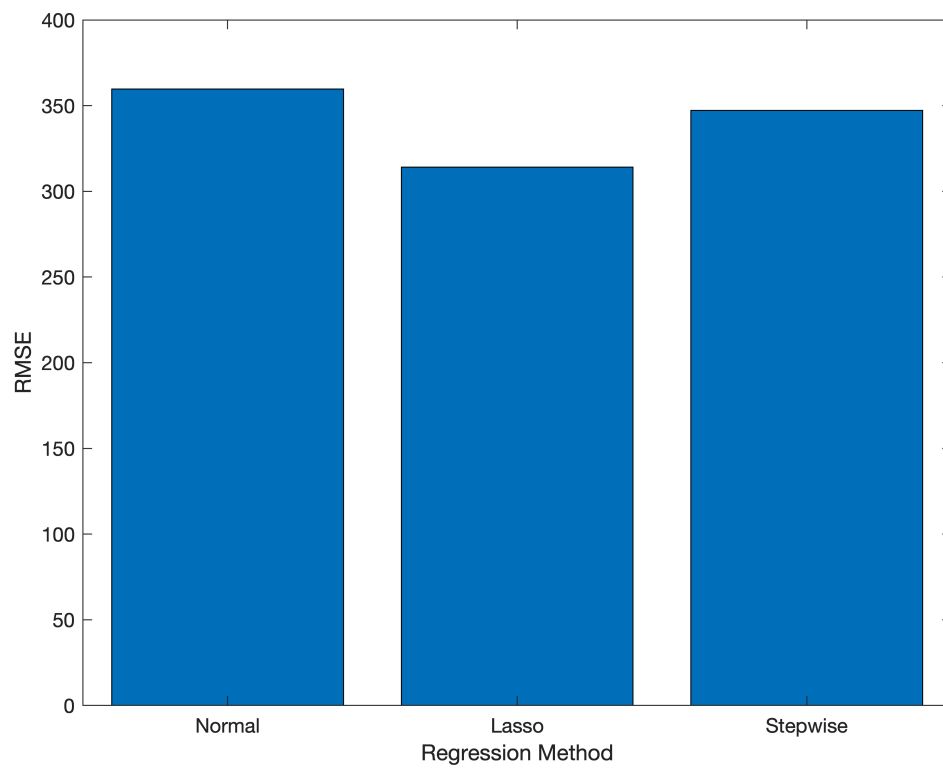
```
avg_error_stepwise = 281.9262
```

Compare Results of 3 Methods

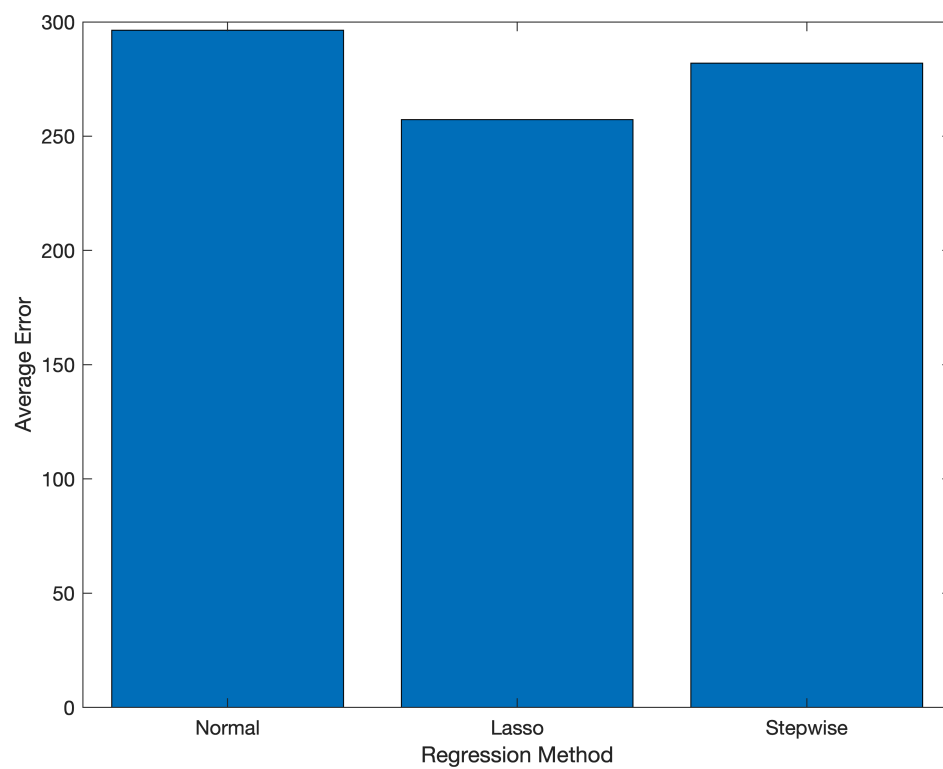
```
% Create labels for bar graphs  
x = categorical({'Normal','Lasso','Stepwise'});  
x = reordercats(x,{'Normal','Lasso','Stepwise'});  
  
% Correlation bar graph  
bar(x,[r_norm,r_lasso,r_stepwise])  
xlabel("Regression Method")  
ylabel("Correlation Coefficient, r")
```



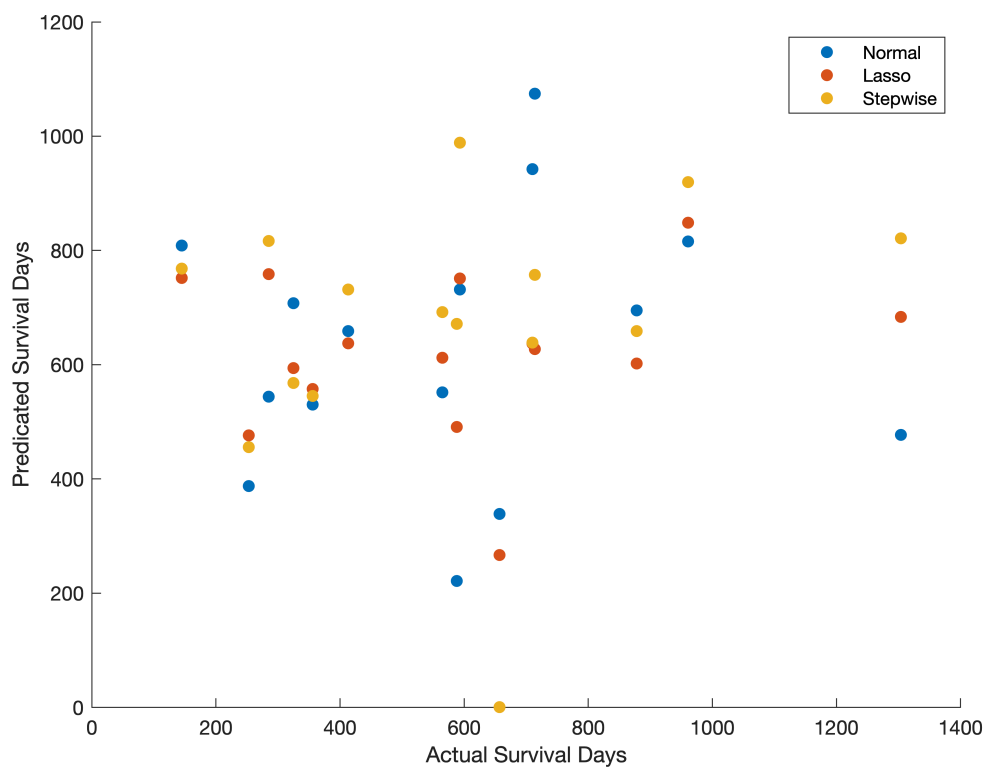
```
% RMSE bar graph  
bar(x,[RMSE,RMSE_lasso,RMSE_stepwise])  
xlabel("Regression Method")  
ylabel("RMSE")
```



```
% Avg Error bar graph  
bar(x,[avg_error,avg_error_lasso,avg_error_stepwise])  
xlabel("Regression Method")  
ylabel("Average Error")
```



```
scatter(Ytest,Ypred_norm,"filled")
xlabel("Actual Survival Days")
ylabel("Predicated Survival Days")
hold on
scatter(Ytest,Ypred_lasso,"filled")
hold on
scatter(Ytest,Ypred_step,"filled")
hold off
legend("Normal","Lasso","Stepwise")
```



```
boxplot(data_HW3.SurvivalDays)
```

