Coursework Four Results: Jamal Khan - jzk09

The mean vector for Hepatitis C data set:

1.164 3.916 0.448 5.679 4.865 3.778 4.127 3.604 0.217

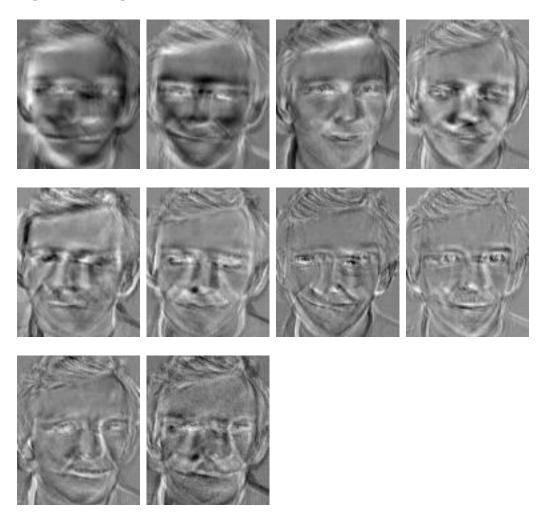
The covariance matrix for Hepatitis C data set:

 $5.880 \ 0.245 \ -0.004 \ -0.284 \ -0.346 \ -0.417 \ -0.481 \ -0.267 \ -0.067 \ 0.245 \ 7.572 \ 0.073 \ -0.358 \ -0.346 \ 0.257 \ 2.273 \ 0.946 \ -0.032 \ -0.004 \ 0.073 \ 0.247 \ -0.116 \ -0.039 \ -0.003 \ 0.100 \ 0.066 \ -0.005 \ -0.284 \ -0.358 \ -0.116 \ 16.960 \ 7.969 \ 4.443 \ -1.036 \ -0.306 \ 0.047 \ -0.346 \ -0.346 \ -0.039 \ 7.969 \ 8.361 \ 4.921 \ -0.763 \ -0.752 \ -0.001 \ -0.417 \ 0.257 \ -0.003 \ 4.443 \ 4.921 \ 6.683 \ -0.087 \ -0.376 \ -0.141 \ -0.481 \ 2.273 \ 0.100 \ -1.036 \ -0.763 \ -0.087 \ 20.583 \ 1.122 \ 0.014 \ -0.267 \ 0.946 \ 0.066 \ -0.306 \ -0.752 \ -0.376 \ 1.122 \ 4.317 \ 0.119 \ -0.067 \ -0.032 \ -0.005 \ 0.047 \ -0.001 \ -0.141 \ 0.014 \ 0.119 \ 0.170$

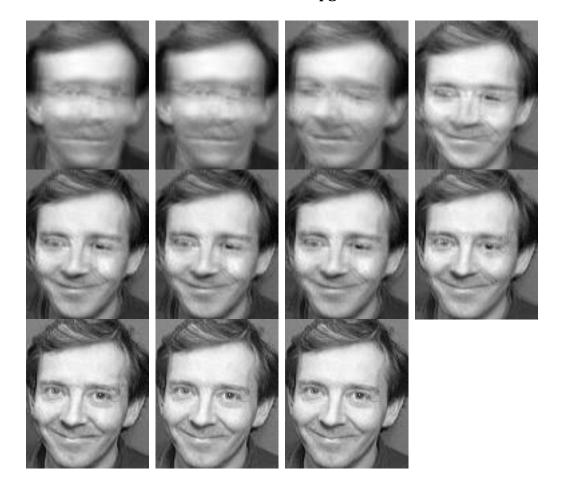
The component magnitudes for image 'c.pgm':

1110,709 1537,080 410,837 -1887,013 941,699 -892,087

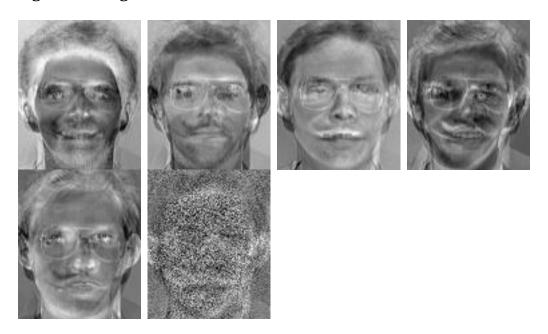
 ${\bf Eigenface\ images\ for\ task\ 4.3\ are\ below:}$



Partial reconstructions for task 4.4 of 'c.pgm 'are below:



Eigenface images for new basis are below:



Partial reconstructions for 'c.pgm', using new basis:

