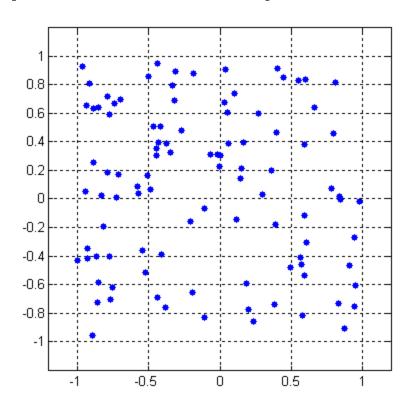
Principal Component Analysis

CSE 6367 – Computer Vision Vassilis Athitsos University of Texas at Arlington

Vector Spaces

- For our purposes, a vector is a tuple of d numbers: X = (x1, x2, ..., xd).
- An example vector space: the 2D plane.
 - Every point in the plot is a vector.
 - Specified by two numbers.



Images are Vectors

 An M-row x N-column image is a vector of dimensionality ...

Images are Vectors

- An M-row x N-column image is a vector of dimensionality.
 - M*N if the image is grayscale.
 - M*N*3 if the image is in color.

Images are Vectors

Consider a 4x3 grayscale image:

```
- A = [A11 A12 A13
A21 A22 A23
A31 A32 A33
A41 A42 A43];
```

- The (Matlab) vector representation Av of A is:
 Av = [A11 A21 A31 A41 A12 A22 A32 A42 A13 A23 A33 A43];
- Mathematically, order does not matter, IF IT IS THE SAME FOR ALL IMAGES.
- In Matlab, to vectorize an image:
 - -Av = A(:);
 - NOTE: The above returns a COLUMN vector.

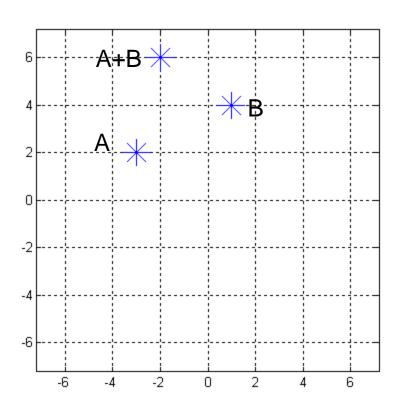
Vector Operations: Addition

- (x1, ..., xd) + (y1, ..., yd) = (x1+y1, ..., xd+yd)
- Example: in 2D:

$$-A = (-3, 2)$$

$$-B = (1, 4)$$

$$-A+B = (-2, 6).$$



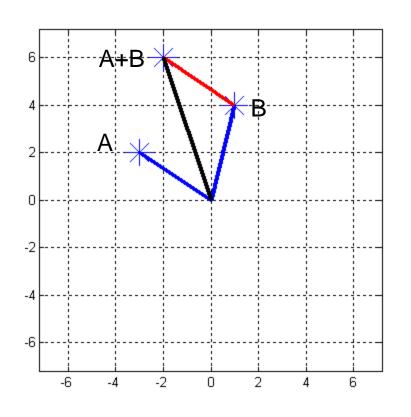
Vector Operations: Addition

- (x1, ..., xd) + (y1, ..., yd) = (x1+y1, ..., xd+yd)
- Example: in 2D:

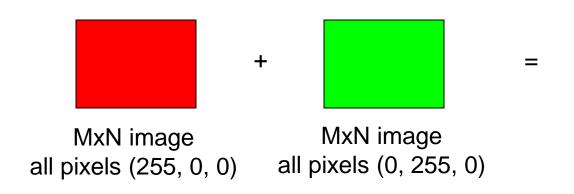
$$-A = (-3, 2)$$

$$-B = (1, 4)$$

$$-A+B = (-2, 6).$$

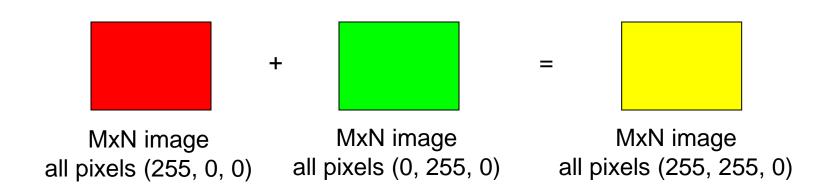


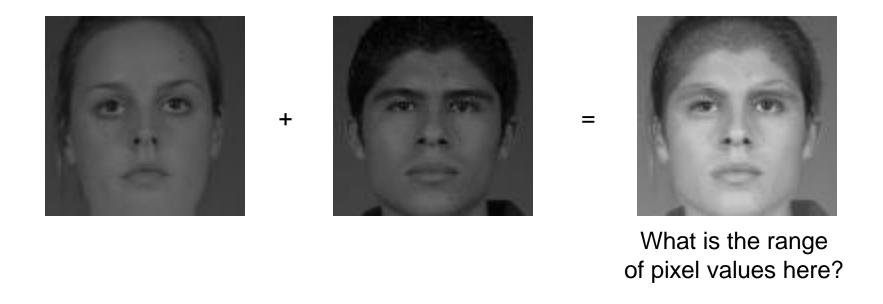
Addition in Image Space



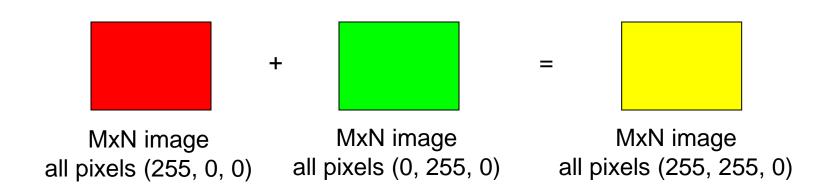


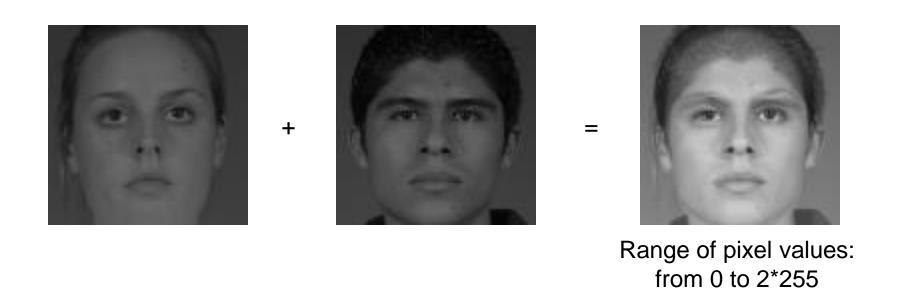
Addition in Image Space





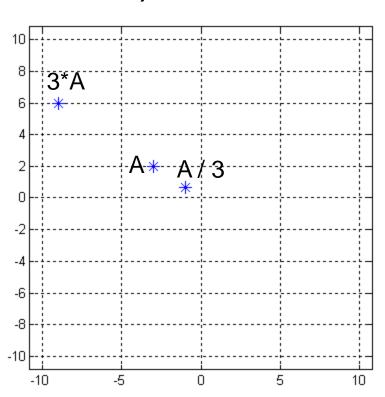
Addition in Image Space





Vector Operations: Scalar Multiplication

- c * (x1, ..., xd) = (c* x1, ..., c * xd)
- (x1, ..., xd) / c = 1/c * (x1, ..., xd)= (x1/c, ..., xd/c)
- Example: in 2D:
 - -A = (-3, 2)
 - -3 * A = (-9, 6)
 - -A/3 = (-1, 0.67).



Multiplication in Image Space

image









image * 0.5

Operations in Image Space

- Note: with addition and multiplication we can easily generate images with values outside the [0, 255] range.
 - These operations are perfectly legal.
 - However, we cannot visualize the results directly.
 - Must convert back to [0 255] range to visualize.

Linear Combinations

- Example: c1 * v1 + c2 * v2 + c3 * v3
 - c1, c2, c3: real numbers.
 - v1, v2, v3: vectors
 - result:

Linear Combinations

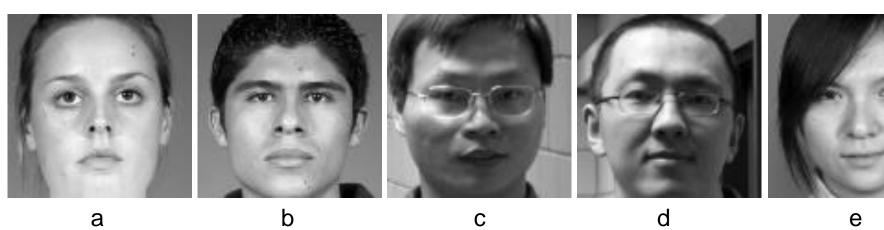
- Example: c1 * v1 + c2 * v2 + c3 * v3
 - c1, c2, c3: real numbers.
 - v1, v2, v3: vectors
 - result: vector.

Vectors Must Be of Same Size

- We cannot add vectors that are not of the same size.
 - -(1,2) + (0,1,0) is NOT DEFINED.
- To add images A and B:
 - IMPORTANT: Most of the time, it only makes sense to add A and B <u>ONLY</u> if they have the same number of rows and the same number of columns.
 - WARNING: Matlab will happily do the following:

```
a = rand(4,3);
b = rand(6,2);
c = a(:) + b(:);
```

Example Linear Combination

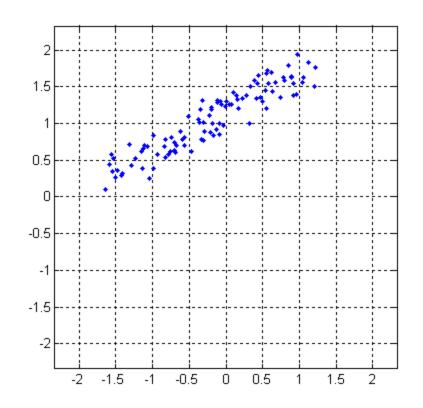




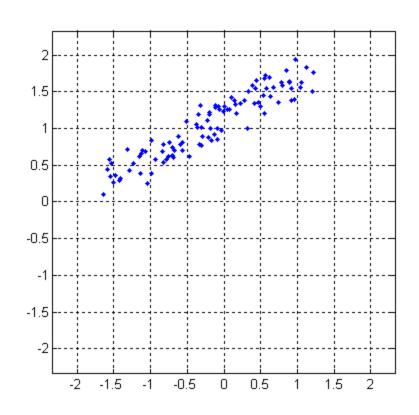
avg

```
a = read_gray('4309d111.bmp');
b = read_gray('4286d201.bmp');
c = read_gray('4846d101.bmp');
d = read_gray('4848d101.bmp');
e = read_gray('4853d101.bmp');
avg = 0.2*a + 0.2*b + 0.2*c + 0.2*d + 0.2*e;
% or, equivalently:
avg = (a+b+c+d+e) / 5;
```

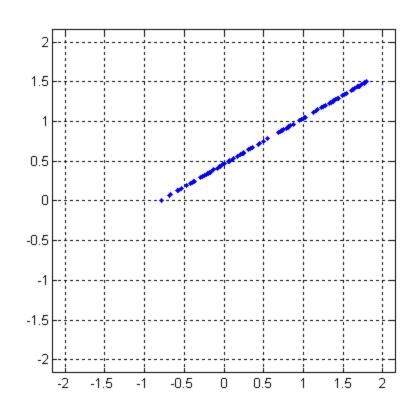
- Consider a set of vectors in a ddimensional space.
- How many numbers do we need to represent each vector?



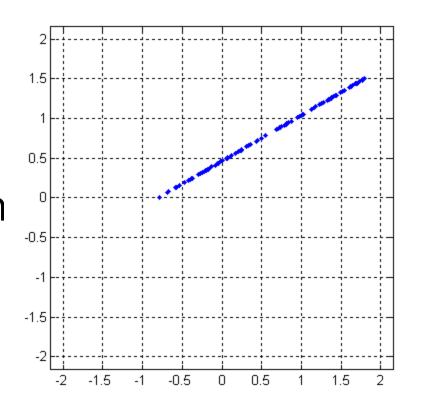
- Consider a set of vectors in a ddimensional space.
- How many numbers do we need to represent each vector?
 - At most d: the same as the number of dimensions.
- Can we use fewer?



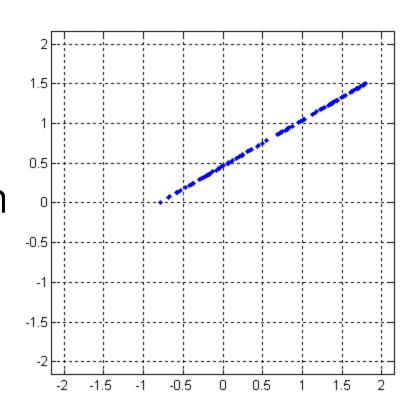
- Consider a set of vectors in a ddimensional space.
- How many numbers do we need to represent each vector?
 - At most d: the same as the number of dimensions.
- Can we use fewer?



- In this example, every point (x, y) is on a line
 -y = ax + b;
- If we have 100 points on this plot, how many numbers do we need to specify them?

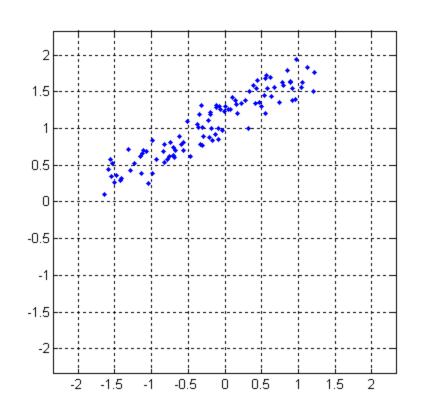


- In this example, every point (x, y) is on a line
 -y = ax + b;
- If we have 100 points on this plot, how many numbers do we need to specify them?
 - 102: a, b, and the x coordinate of each point.
 - Asymptotically: one number per point.



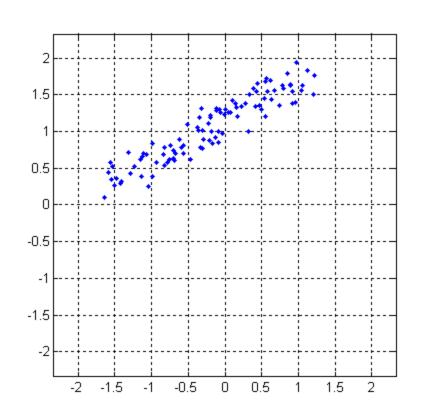
Lossy Dimensionality Reduction

- Suppose we want to project all points to a single line.
- This will be lossy.
- What would be the best line?

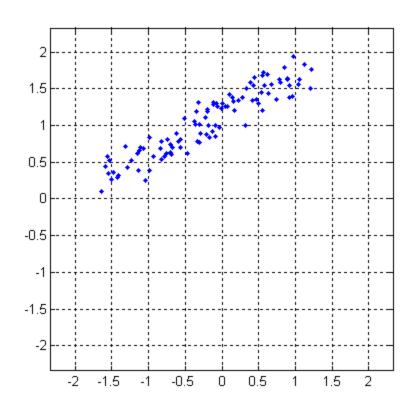


Lossy Dimensionality Reduction

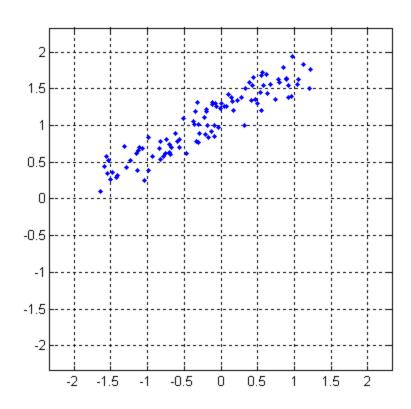
- Suppose we want to project all points to a single line.
- This will be lossy.
- What would be the best line?
- Optimization problem.
 - Infinite answers.
 - We must define how to evaluate each answer.



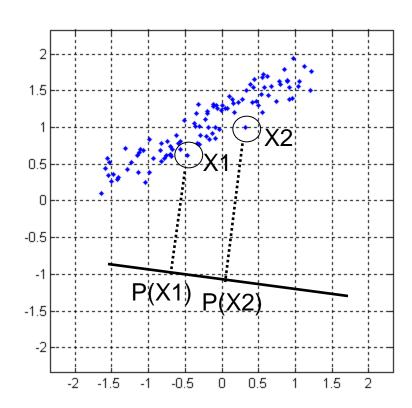
- We want to measure how good a projection P is, GIVEN A SET OF POINTS.
 - If we don't have a specific set of points in mind, what would be the best projection?



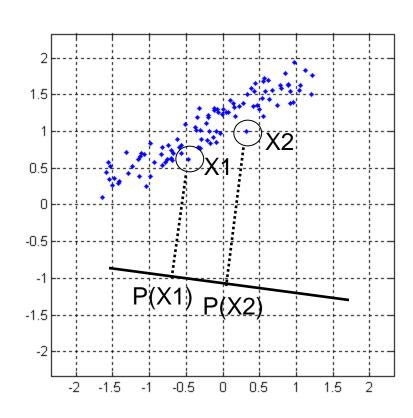
- We want to measure how good a projection P is, GIVEN A SET OF POINTS.
 - If we don't have a specific set of points in mind, what would be the best projection?
 - NONE: all are equally good/bad.



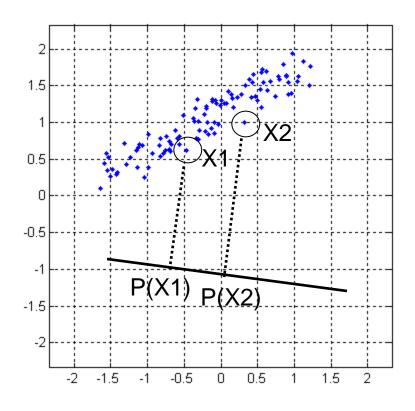
- Consider a pair of points:
 X1, X2.
- D1 = squared distance from X1 to X2.
 - -sum((X1-X2).*(X1-X2))
- D2 = squared distance from P(X1) to P(X2).
- Error(X1, X2) = D1 D2.
 - Will it ever be negative?



- Consider a pair of points:
 X1, X2.
- D1 = squared distance from X1 to X2.
 - -sum((X1-X2).*(X1-X2))
- D2 = squared distance from P(X1) to P(X2).
- Error(X1, X2) = D1 D2.
 - Will it ever be negative?
 - NO: D1 >= D2 always.



- Now, consider the entire set of points:
 - X1, X2, ..., Xn.
- Error(P) = sum(Error(Xi, Xj) | i, j = 1, ..., n, i != j).

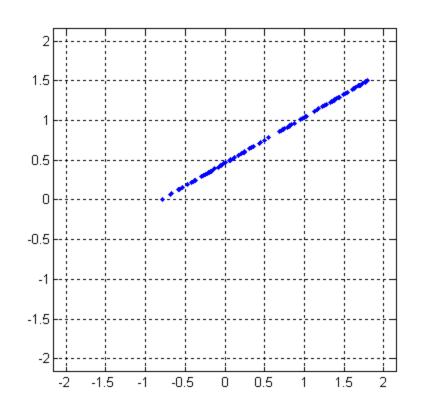


Interpretation:

- We measure how well P preserves distances.
- If P preserves distances,Error(P) = 0.

Example: Perfect Projection Exists

 In this case, projecting to a line oriented at 30 degrees would give zero error.



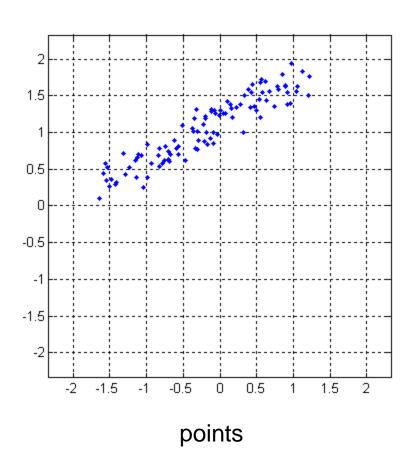
First step: center the data.

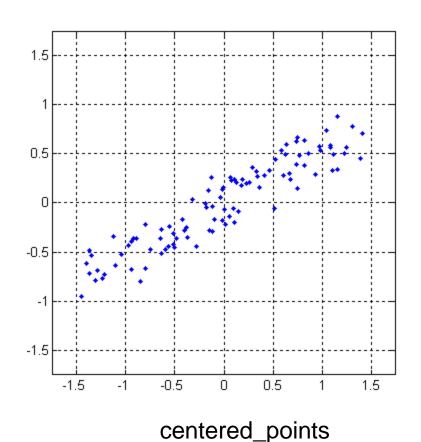
```
number = size(points, 2);
% note that we are transposing twice
average = [mean(points')]';
centered_points = zeros(size(points));

for index = 1:number
          centered_points(:, index) = points(:, index) - average;
end

plot_points(centered_points, 2);
```

First step: center the data.





Second step: compute the covariance matrix.

```
covariance matrix = centered points * centered points';
```

 In the above line we assume that each column is a vector.

Second step: compute the covariance matrix.

```
covariance_matrix = centered_points * centered_points';
```

- In the above line we assume that each column is a vector.
- Third step: compute the eigenvectors and eigenvalues of the covariance matrix.

```
[eigenvectors eigenvalues] = eig(covariance_matrix);
```

Eigenvectors and Eigenvalues

```
eigenvectors =

0.4837 -0.8753
-0.8753 -0.4837

eigenvalues =

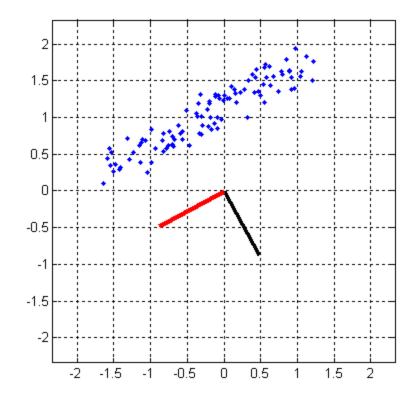
2.0217 0

0 77.2183
```

- Each eigenvector v is a column, that specifies a line going through the origin.
- The importance of the i-th eigenvector is reflected by the i-th eigenvalue.
 - second eigenvalue = 77, first eigenvalue = 2, =>
 second eigenvector is far more important.

Visualizing the Eigenvectors

black: v1 (eigenvalue = 2.02) red: v2 (eigenvalue = 77.2)



```
plot_points(points, 1);
p1 = eigenvectors(:, 1);
p2 = eigenvectors(:, 2);
plot([0, p1(1)], [0, p1(2)], 'k-', 'linewidth', 3);
hold on;
plot([0, p2(1)], [0, p2(2)], 'r-', 'linewidth', 3);
```

PCA Code

```
function [average, eigenvectors, eigenvalues] = ...
                                     compute pca(vectors)
number = size(vectors, 2);
% note that we are transposing twice
average = [mean(vectors')]';
centered vectors = zeros(size(vectors));
for index = 1:number
    centered vectors(:, index) = vectors(:, index) - average;
end
covariance matrix = centered vectors * centered vectors';
[eigenvectors eigenvalues] = eig( covariance matrix);
% eigenvalues is a matrix, but only the diagonal
% matters, so we throw away the rest
eigenvalues = diag(eigenvalues);
[eigenvalues, indices] = sort(eigenvalues, 'descend');
eigenvectors = eigenvectors(:, indices);
```

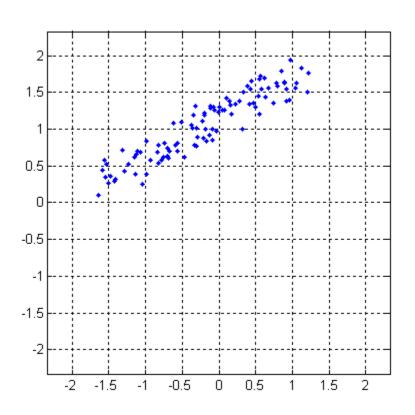
Reducing Dimensions From 2 to 1

- Precompute:
 - the eigenvector P1 with the largest eigenvalue.
 - the mean avg of all the data.

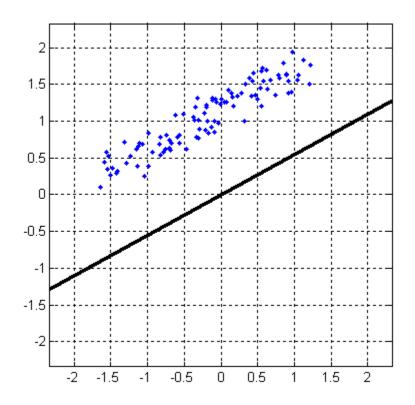
Reducing Dimensions From 2 to 1

- Two key operations:
 - projection to lower-dimensional space.
 - backprojection to the original space.
- Projection:
 - $-P(V) = \langle V avg, P1 \rangle = P1' * (V avg)$
 - Dot product between (V-avg) and P1.
 - NOTE: The eigenvectors that Matlab returns have unit norm.
 - Backprojection:
 - B(P(V)) = P1 * P(V) + avg.

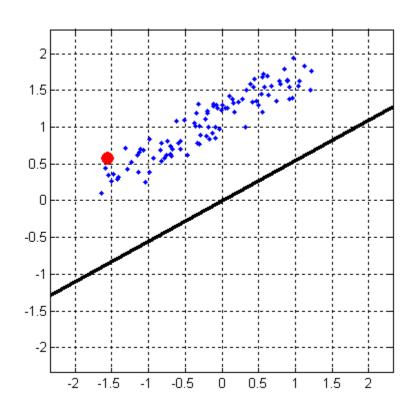
A set of points.



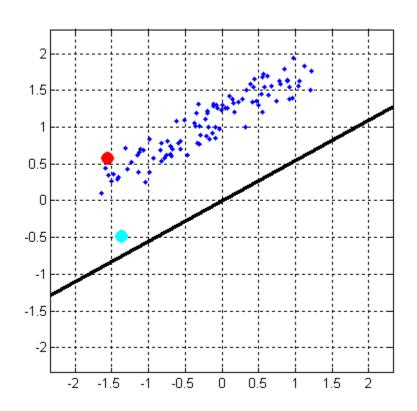
- Do PCA, identify p1 = the first eigenvector.
- We plot the line of direction of p1.



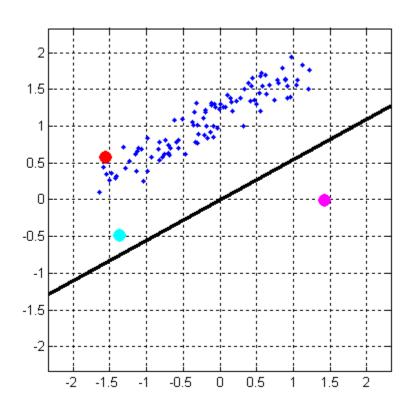
- Choose a point v4 = [-1.556, 0.576]'.
 - Shown in red.



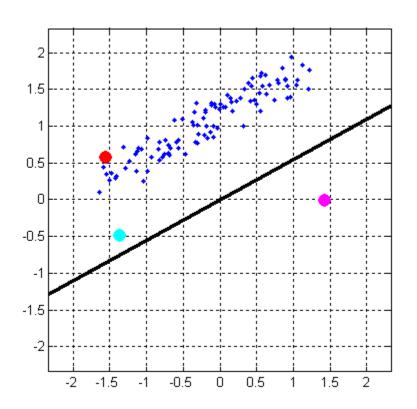
- centered_v4 = v4 average.
 - Shown in cyan.



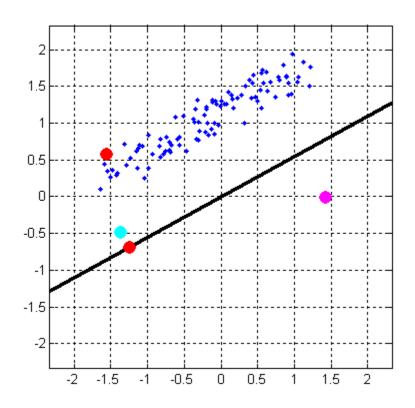
- projection = p1' * (centered_v4);
- result: projection = 1.43 (shown in magenta)



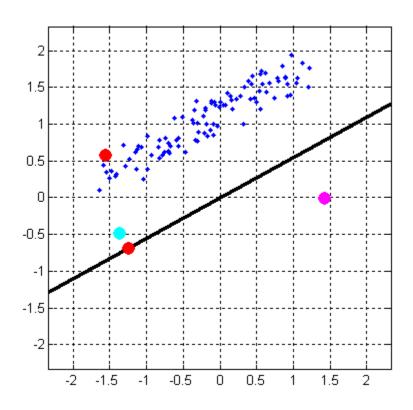
 Note: projection is a single number. To plot it, we actually treat that number as the x value, and use a y value of 0.



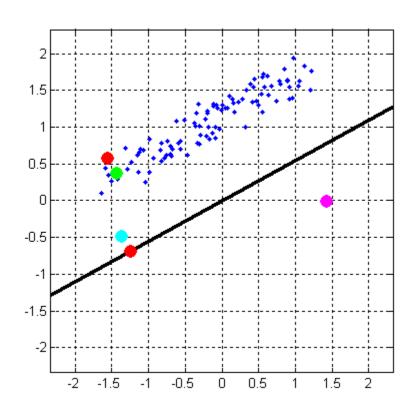
- b1 = p1 * projection;
 - shown in red, on top of black line.
- How are b1 and projection related?



- b1 = p1 * projection;
 - shown in red, on top of black line.
- projection = +-distance of b1 from the origin.



- reconstructed v4 = b1 + average;
 - shown in green.



PCA on Faces

- Motivation: If a face is a 31x25 window, we need 775 numbers to describe the face.
- With PCA, we can approximate face images with much fewer numbers.
- Two benefits:
 - Faster computations.
 - The amount of loss can be used for face detection.

PCA vs Template Matching

- If we use template matching to detect faces, what is the perfect face (easiest to be detected)?
- How about PCA?

PCA vs Template Matching

- Template matching:
 - The average face is the only perfect face (after we normalize for brightness and contrast).
 - As a model, it is not rich enough.
- PCA:
 - There is a space of perfect faces.

Preparing a Dataset

- Get a set of face images.
 - They must be of same size, and aligned, so that eye locations match each other, nose locations match each other, and so on.
 - Make the mean and std of all faces equal.
 - Discards variations in intensity and contrast.

Code for Preprocessing Faces

```
clear;
load faces1032;
number = size(faces, 2);
dimensions = size(faces, 1);
% set mean of all faces to 0, and std to 1.
for index = 1: number
    face = faces(:, index);
    face = (face - mean(face)) / std(face);
    faces(:, index) = face;
end
% do pca
[mean face, eigenvectors, eigenvalues] = compute_pca(faces);
```

Visualizing Eigenfaces



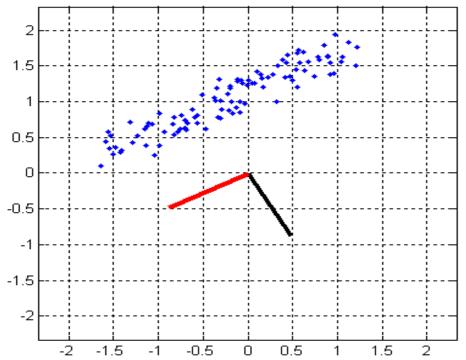
Approximating a Face With 0 Numbers

 What is the best approximation we can get for a face image, if we know nothing about the face image (except that it is a face)?



Equivalent Question in 2D

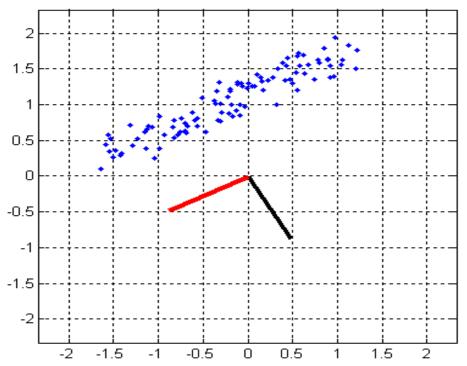
 What is our best guess for a 2D point from this point cloud, if we know nothing about that 2D point (except that it belongs to the cloud)?



Equivalent Question in 2D

 What is our best guess for a 2D point from this point cloud, if we know nothing about that 2D point (except that it belongs to the cloud)?

Answer: the average of all points in the cloud.



Approximating a Face With 0 Numbers

 What is the best approximation we can get for a face image, if we know nothing about the face image (except that it is a face)?



Approximating a Face With 0 Numbers

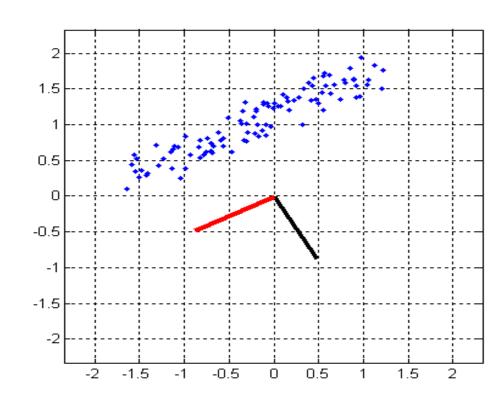
- What is the best approximation we can get for a face image, if we know nothing about the face image (except that it is a face)?
 - The average face.





Guessing a 2D Point Given 1 Number

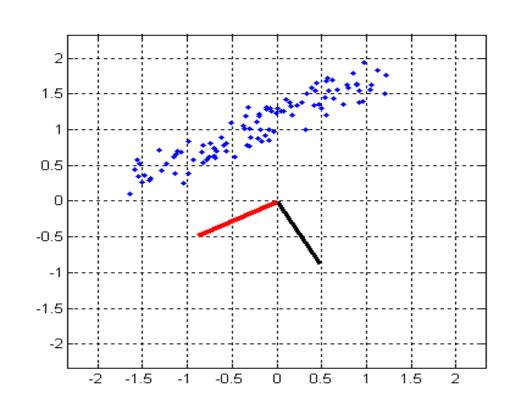
- What is our best guess for a 2D point from this point cloud, if we know nothing about that 2D point, except a single number?
 - What should that number be?



Guessing a 2D Point Given 1 Number

- What is our best guess for a 2D point from this point cloud, if we know nothing about that 2D point, except a single number?
 - What should that number be?

Answer: the projection on the first eigenvector.



Approximating a Face With 1 Numbers

 What is the best approximation we can get for a face image, if we can represent the face with a single number?





Approximating a Face With 1 Numbers

- With 0 numbers, we get the average face.
- 1 number: PCA projection to 1D.
- Reconstruction:
 - average face + <face, eigenvector1> * eigenvector1





Approximating a Face With 1 Number

```
v1 = faces(:, 13);
f1 = reshape(v1, size(mean_face));
k = 0;
%%
x1 = v1' * eigenvectors(:, 1);
term1 = x1 * eigenvectors(:, 1);
term1 = reshape(term1, size(mean_face));
reconstructed1 = mean_face + term1;
```





mean_face

Approximating a Face With 1 Number

```
v1 = faces(:, 13);
f1 = reshape(v1, size(mean_face));
k = 0;
%%
x1 = v1' * eigenvectors(:, 1);
term1 = x1 * eigenvectors(:, 1);
term1 = reshape(term1, size(mean_face));
reconstructed1 = mean_face + term1;
```





Approximating a Face With 2 Numbers

```
x2 = v1' * eigenvectors(:, 2);
term2 = x2 * eigenvectors(:, 2);
term2 = reshape(term1, size(mean_face));
reconstructed2 = reconstructed1 + term2;
```





Approximating a Face With 3 Numbers

```
x = v1' * eigenvectors(:, 3);
term = x * eigenvectors(:, 3);
term = reshape(term1, size(mean_face));
reconstructed3 = reconstructed2 + term;
```





Approximating a Face With 4 Numbers





Approximating a Face With 5 Numbers





Approximating a Face With 6 Numbers





Approximating a Face With 7 Numbers





Approximating a Face With 10 Numbers

```
f1 = faces(:, 13);
f1 = reshape(f1, size(mean_face));
ev = eigenvectors(:, 1:10);
p = pca_projection(f1, mean_face, ev);
b = pca_backprojection(p, mean_face, ev);
reconstructed10 = reshape(b, size(mean_face));
```





PCA Projection Code

```
function result = normalize_face(image_window, mean_face)
% function result = normalize_face(vector)
%
% normalizes the vector so that the size matches that of \
% mean face, the mean is 0 and the std is 1.

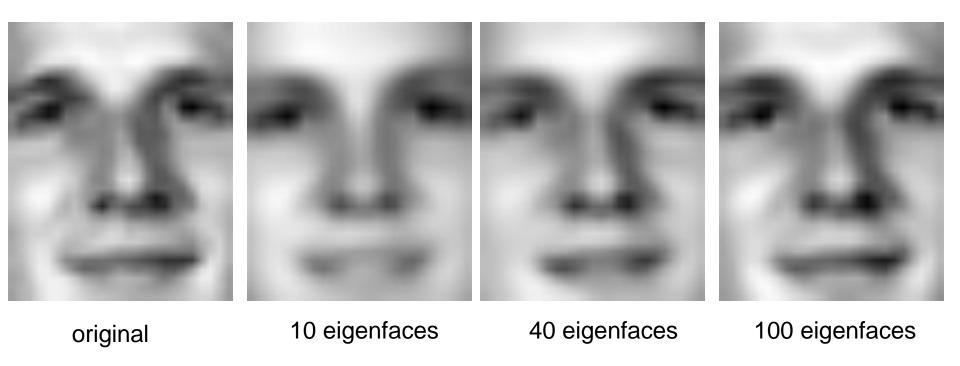
result = imresize(image_window, size(mean_face), 'bilinear');
result = result(:);
result = result - mean(result(:));
result = result / std(result(:));
result(isnan(result)) = 0;
```

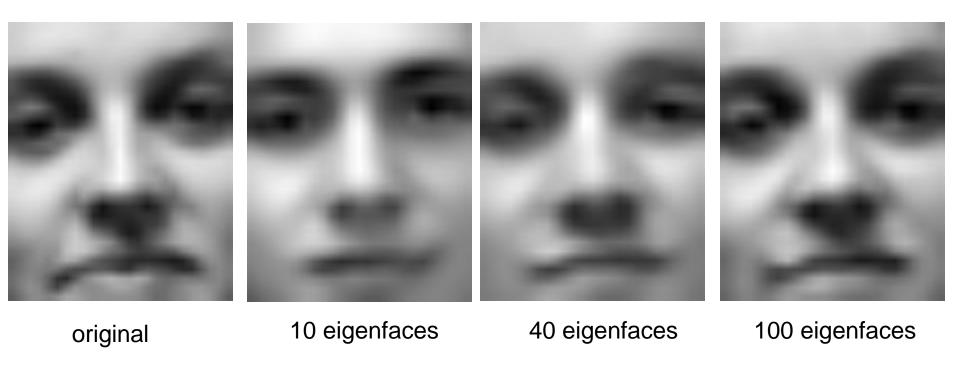
PCA Projection Code

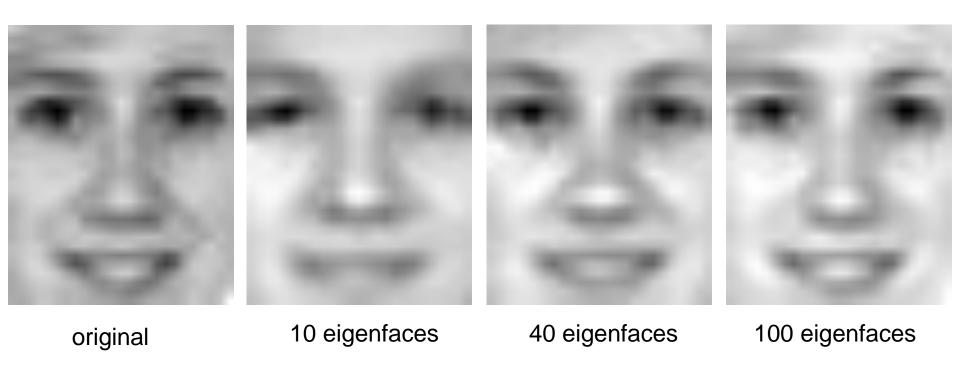
```
function result = eigenface projection(image window, ...
                                       mean face, eigenvectors)
 function result = eigenface projection(image window, ...
응
                                          average, eigenvectors)
응
% the vector is converted to a column vector. Each column in
% eigenvectors should be an eigenvector. The mean is converted
% to a column vector
normalized = normalize face(image window, mean face);
% subtract mean from vector
centered = normalized(:) - mean face(:);
% convert vector to a column vector.
result = eigenvectors' * centered;
```

PCA Backprojection Code

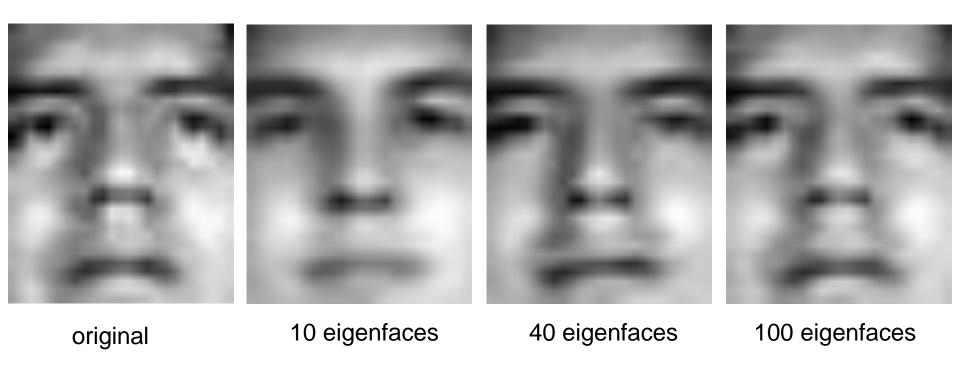
- The PCA projection gives us a few numbers to represent a face.
- The backprojection uses those few numbers to generate a face image.



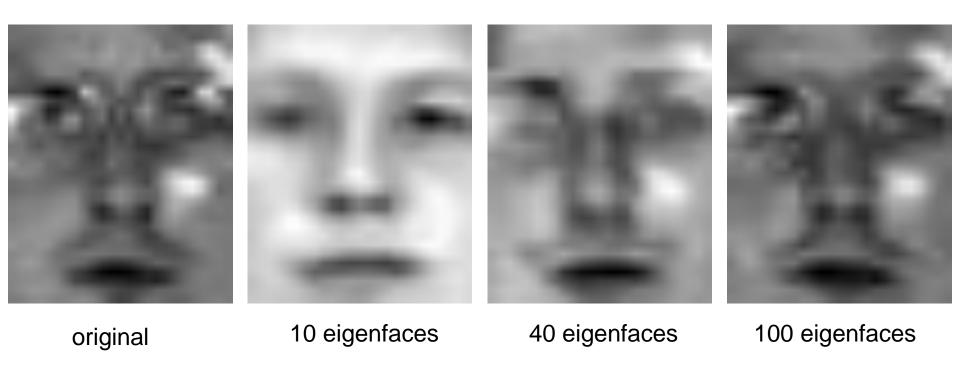




Note: teeth not visible using 10 eigenfaces



Note: using 10 eigenfaces, gaze direction is towards camera

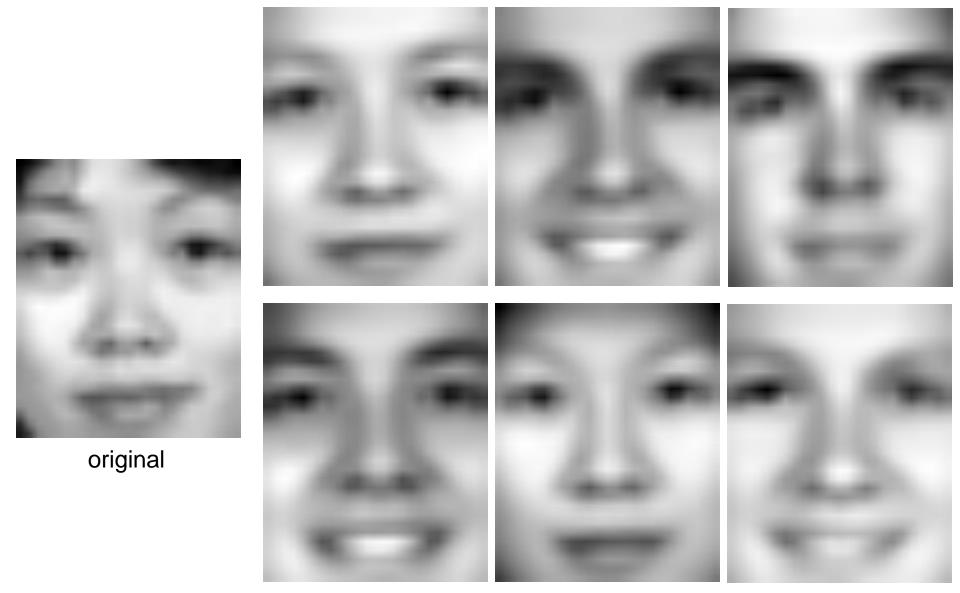


Note: using 10 eigenfaces, glasses are removed

Projecting a Non-Face



How Much Can 10 Numbers Tell?



eigenfaces

Using Eigenfaces for Detection

- Intuition: Why are eigenfaces good for representing faces?
 - Because projection OF FACES to the eigenspace loses little information.
 - However, projection of NON-FACES to the eigenspace of faces loses more information.
- Criterion for face detection: how much is the reconstruction error?
- Quantitative:
 - Define f = face, b = backprojection(projection(f)).
 - Error: norm of (f b).

Evaluating Error for a Window

```
function result = pca score(window, mean face, ...
                            eigenvectors, eigenface number)
% normalize the window
window = imresize(window, size(mean face), 'bilinear');
window = window(:);
window = window - mean(window);
window = window / std(window);
window(isnan(window)) = 0;
top eigenvectors = eigenvectors(:, 1:eigenface number);
projection = pca projection(window, mean face, ...
                            top eigenvectors);
reconstructed = pca backprojection (projection, mean face, ...
                                   top eigenvectors);
diff = reconstructed - window(:);
result = sum(diff .* diff);
```

```
function [max responses, max scales, max rotations] = ...
    template search (image, template, scales, rotations, result number)
% function [result, max scales, max rotations] = ...
     template search (image, template, scales, rotations, result number)
용
읒
% for each pixel, search over the specified scales and rotations,
% and record:
% - in result, the max normalized correlation score for that pixel
% over all scales
% - in max scales, the scale that gave the max score
% - in max rotations, the rotation that gave the max score
% clockwise rotations are positive, counterclockwise rotations are
% negative.
% rotations are specified in degrees
```

```
function [max responses, max scales, max rotations] = ...
    template search (image, template, scales, rotations, result number)
max responses = ones(size(image)) * -10;
max scales = zeros(size(image));
max rotations = zeros(size(image));
for rotation = rotations
    rotated = imrotate(image, -rotation, 'bilinear', 'crop');
    [responses, temp max scales] = ...
        multiscale correlation(rotated, template, scales);
    responses = imrotate(responses, rotation, 'nearest', 'crop');
    temp max scales = imrotate(temp max scales, rotation, ...
                               'nearest', 'crop');
    higher maxes = (responses > max responses);
    max responses(higher maxes) = responses(higher maxes);
    max scales(higher maxes) = temp max scales(higher maxes);
    max rotations(higher maxes) = rotation;
end
```

```
function result = find template(image, template, scales, ...
                                rotations, result number)
% function result = find template(image, template, scales)
응
% returns the bounding boxes of the best matches for the template in
% the image, after searching all specified scales and rotations.
% result number specifies the number of results (bounding boxes).
[max responses, max scales, max rotations] = ...
    template search (image, template, scales, ...
                    rotations, result number);
result = detection boxes(image, template, max responses, ...
                         max scales, result number);
```

```
function [result, boxes] = template detector demo(image, template, ...
                                      scales, rotations, result number)
 function [result, boxes] =
응
              template detector demo(image, template, ...
응
                                      scales, rotations, result number)
% returns an image that is a copy of the input image, with
% the bounding boxes drawn for each of the best matches for
% the template in the image, after searching all specified
% scales and rotations.
boxes = find template(image, template, scales, ...
                      rotations, result number);
result = image;
for number = 1:result number
    result = draw rectangle1(result, boxes(number, 1), ...
                             boxes (number, 2), ...
                             boxes (number, 3), boxes (number, 4));
end
```

Including Information From PCA

- One approach: call pca_score on every possible window.
 - Drawback: slow:
- How can we combine a slow but more accurate approach and a fast but less accurate approach?

Filter-and-refine Approach

- How can we combine a slow but more accurate approach and a fast but less accurate approach?
- Filter step: use the fast approach to get a shortlist of candidates.
- Refine step: use the slow approach to rerank the shortlist and choose the final results.

Filter-and-refine Face Detection

- Filter step: use correlation to identify a number of candidates.
 - number of candidates is 50 in the code.
- Refine step: use pca_score to evaluate those candidates.

```
function result = find faces(image, scales, result number, ...
                             eigenface number)
load face filter;
load final eigens;
% start filter step
preliminary number = 50;
preliminary results = find template(image, face filter, ...
                                    scales, 0, preliminary number);
% start refine step
for number = 1:preliminary number
    top = preliminary results(number, 1);
    bottom = preliminary_results(number, 2);
    left = preliminary results(number, 3);
    right = preliminary results(number, 4);
    window = image(top:bottom, left:right);
    preliminary results(number, 6) = ...
        pca score (window, mean face, eigenvectors, eigenface number);
end
[values, indices] = sort(preliminary results(:, 6), 'ascend');
top indices = indices(1: result number);
result = preliminary results(top indices, :);
```

Recap on PCA for Face Detection

What model for faces are we using?

Recap on PCA for Face Detection

- What model for faces are we using?
 - Faces are reconstructed well using the top K eigenvectors.
 - NOTE: each K defines a different model.
- What is a perfect face under this model?

Recap on PCA for Face Detection

- What model for faces are we using?
 - Faces are reconstructed well using the top K eigenvectors.
 - NOTE: each K defines a different model.
- What is a perfect face under this model?
 - A face spanned using the top K eigenvectors.
 - A face F such that:
 - pca_backprojection(pca_projection(F)) = F.
 - A face on which pca_score returns 0.

A Generalized View of Classifiers

- We have studied two face detection methods:
 - Normalized correlation.
 - PCA.
- Each approach exhaustively evaluates all image suwbwindows.
- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.

Features and Classifiers

- Our goal, in the next slides, is to get a better understanding of:
 - What is a feature?
 - What is a classifier?

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is a feature here?

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is a feature here?
- Two possible answers:
 - Each pixel value is a feature.
 - The feature is the result of normalized correlation with the template.

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is the score of each subwindow?

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is the score of each subwindow?
 - The result of normalized correlation with the template.
 - Arguably, the score is the feature itself.

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- How does the decision depend on the score?

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- How does the decision depend on the score?
 - In find_template, faces are the top N scores.
 - N is an argument to the find_template function.
 - An alternative, is to check if score > threshold.
 - Then we must choose a threshold instead of N.

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- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is a feature here?
 - The result of the pca_projection function.
 - In other words, the K numbers that describe the projection of the subwindow on the space defined by the top K eigenfaces.

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is the score of each subwindow?

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
- What is the score of each subwindow?
 - The result of the pca_score function.
 - The sum of squared differences between:
 - the original subwindow W, and
 - pca_backprojection(pca_projection(W)).

- Each subwindow is evaluated in three steps:
 - First step: extract features.
 - Feature: a piece of information extracted from a pattern.
 - Compute a score based on the features.
 - Make a decision based on the score.
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- Each subwindow is evaluated in three steps:
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Defining a Classifier

- Choose features.
- Choose a scoring function.
- Choose a decision process.
 - The last part is usually straightforward.
 - We pick the top N scores, or we apply a threshold.
- Therefore, the two crucial components are:
 - Choosing features.
 - Choosing a scoring function.

What is a Feature?

- Any information extracted from an image.
- The number of possible features we can define is enormous.
- Any function F we can define that takes in an image as an argument and produces one or more numbers as output defines a feature.
 - Correlation with a template defines a feature.
 - Projecting to PCA space defines features.
 - More examples?

What is a Feature?

- Any information extracted from an image.
- The number of possible features we can define is enormous.
- Any function F we can define that takes in an image as an argument and produces one or more numbers as output defines a feature.
 - Correlation with a template defines a feature.
 - Projecting to PCA space defines features.
 - Average intensity.
 - Std of values in window.