

Introduction to Computer Vision

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Probability, PCA, covariance and
classification

Reading

Szeliski

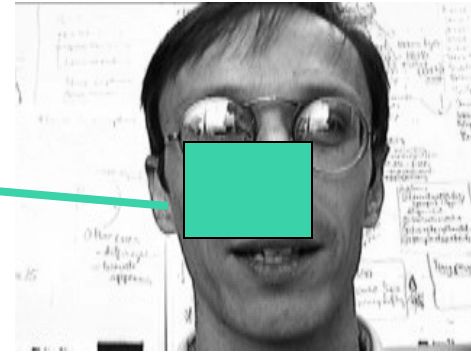
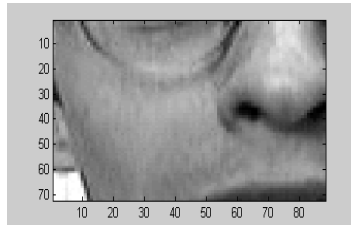
14.1, Face Recognition (including PCA)

A1.1 and 1.2, SVD and PCA

Goals

- Finish probability and classification
 - Everything you need assignment 2
- Wed/Fri: Motion and prep for assign 3

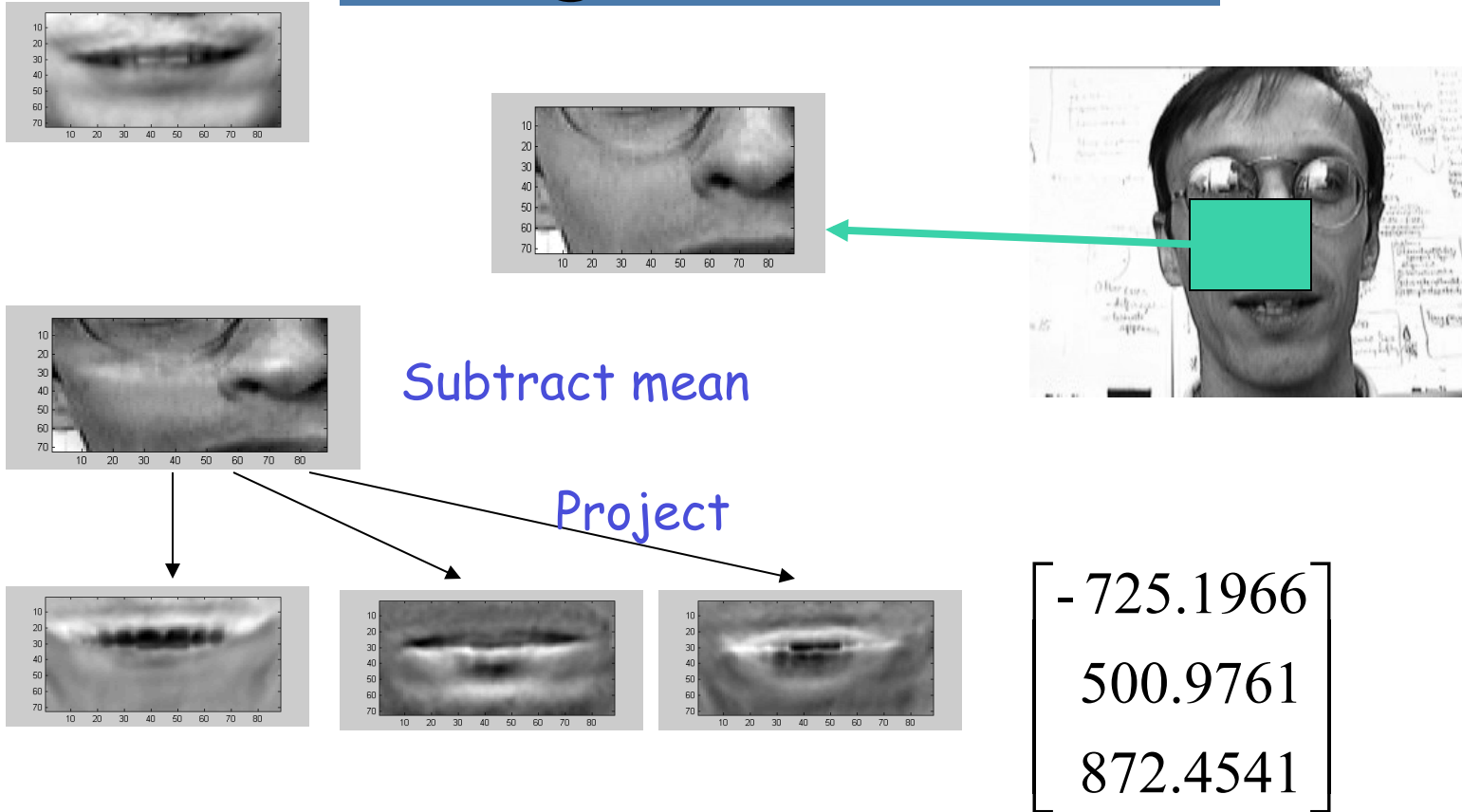
Images as Vectors



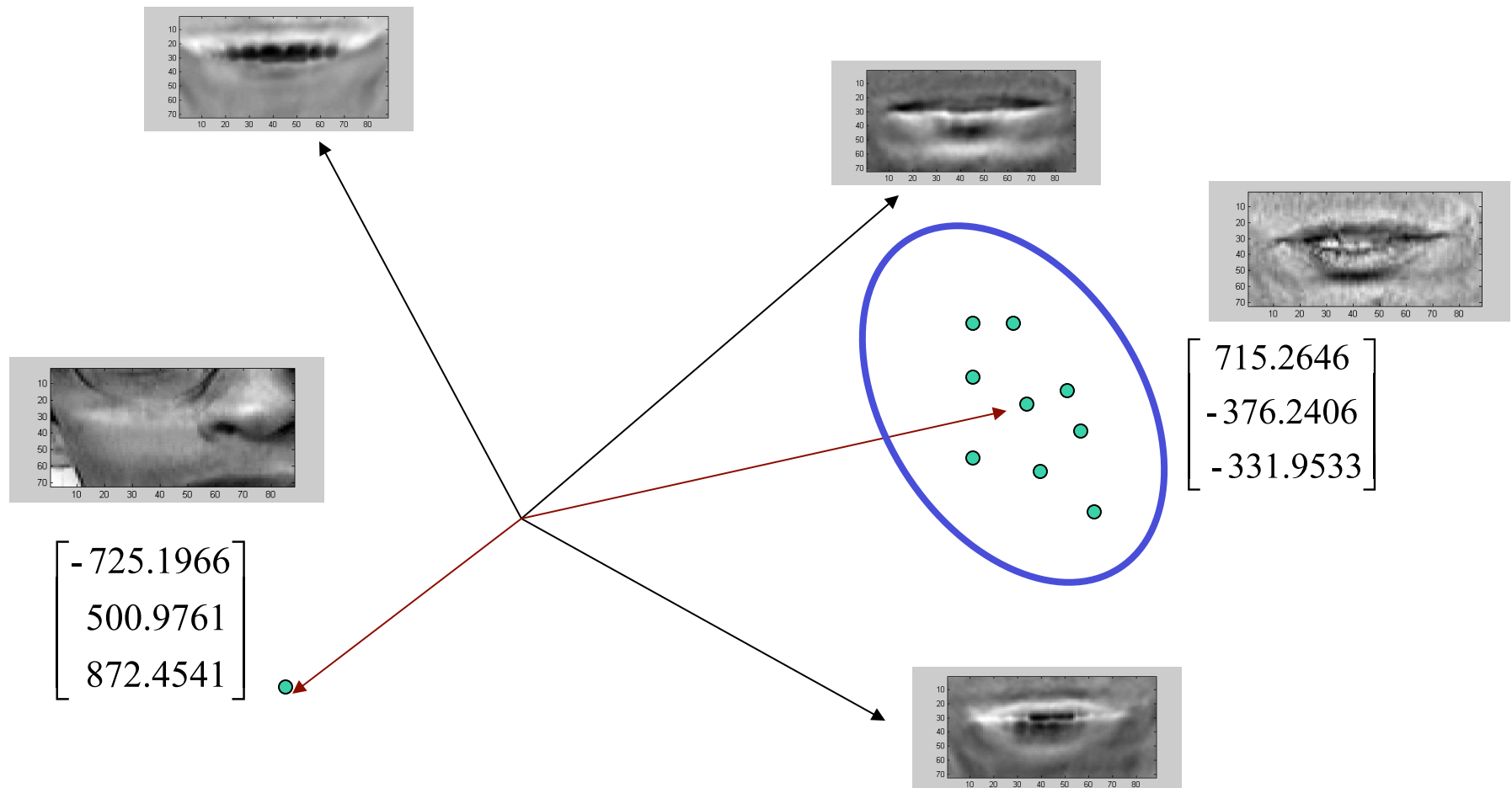
$$= \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_{n \times m} \end{bmatrix}$$

Is it a mouth?

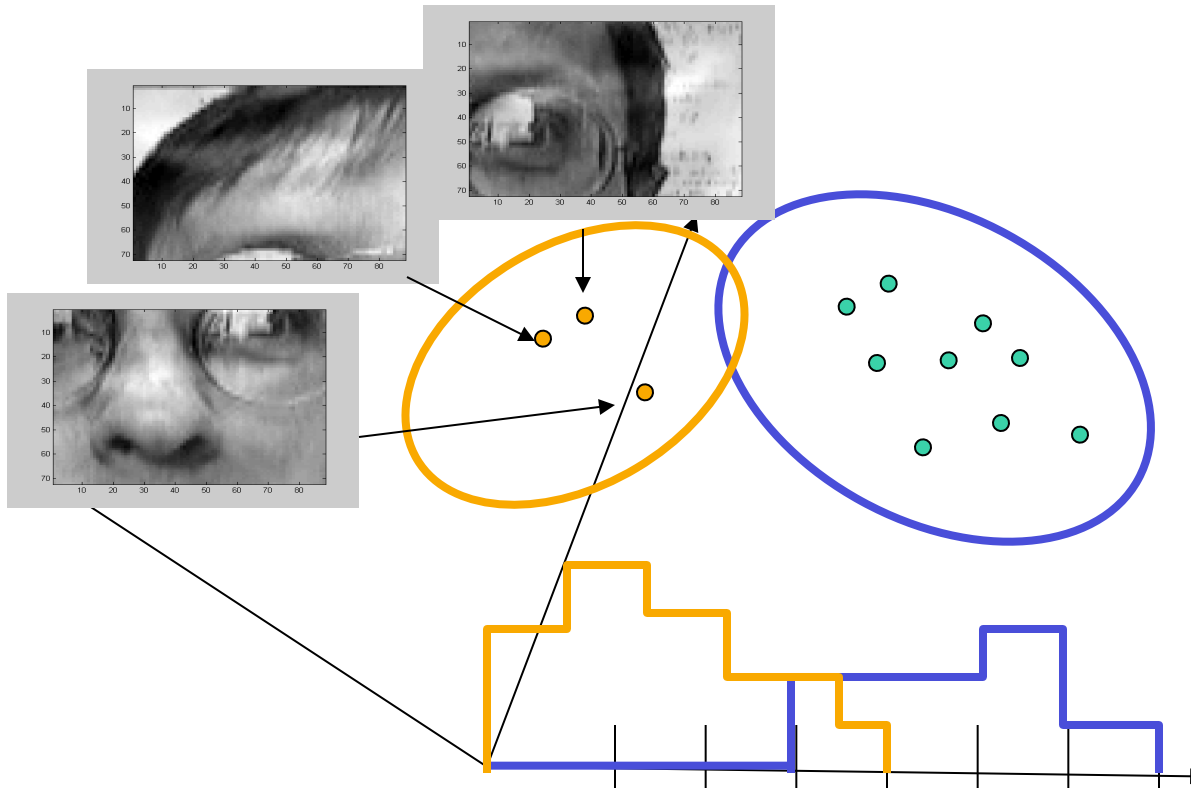
Images as Vectors



Mouth Space

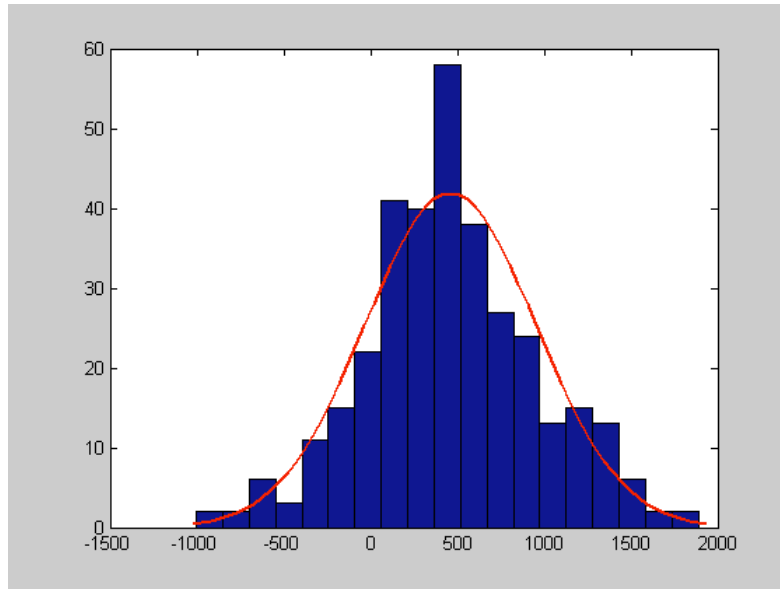


Classification

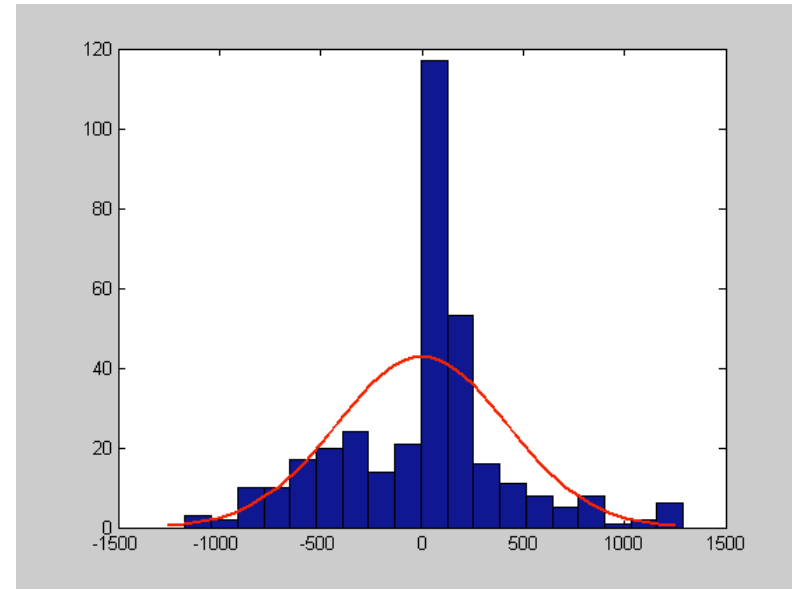


Classification

Imagine we just consider one dimension (one linear coefficient).

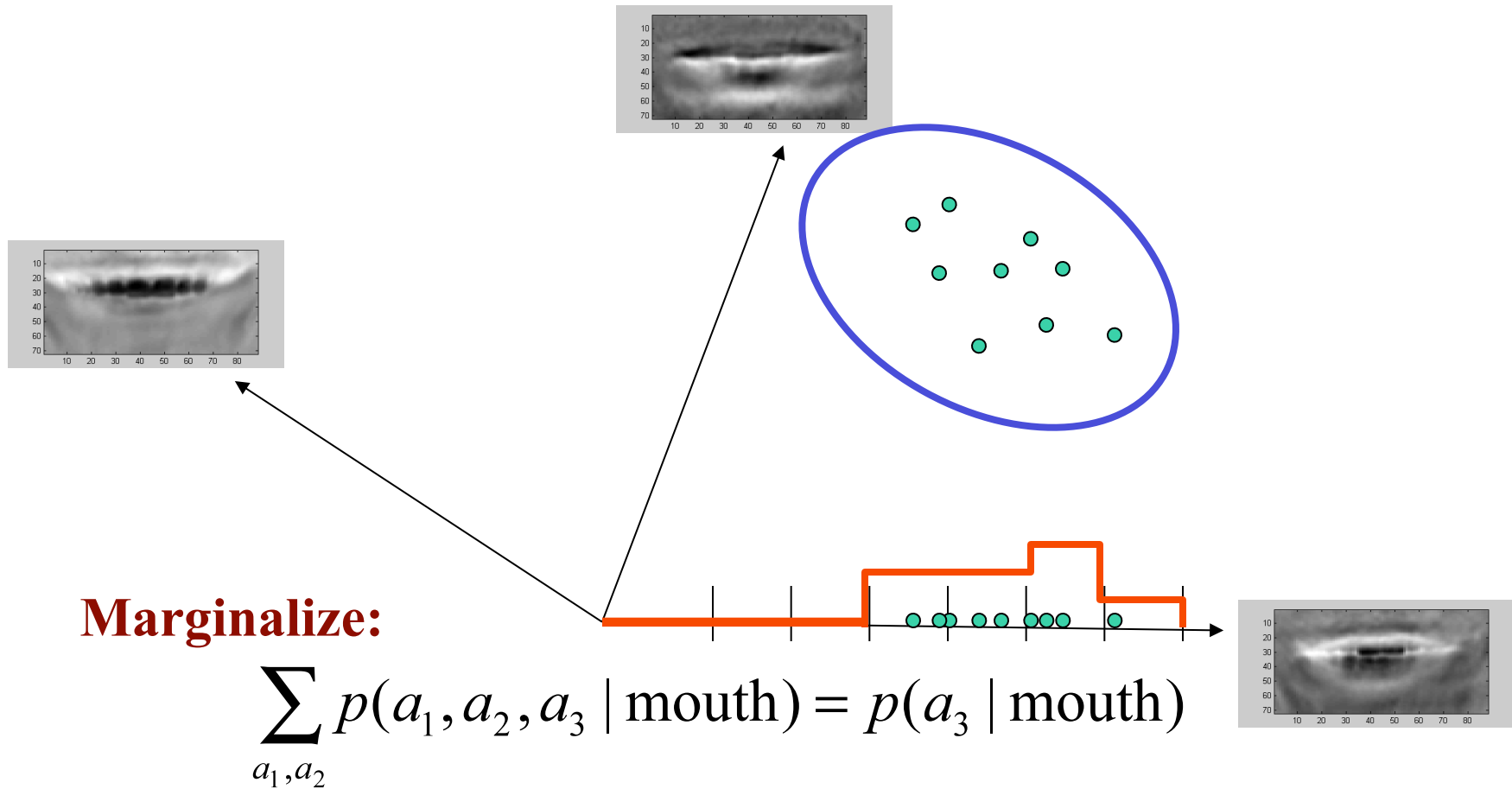


$$p(a_3 \mid \neg\text{mouth})$$



$$p(a_3 \mid \text{mouth})$$

Probabilistic Model



Marginalization

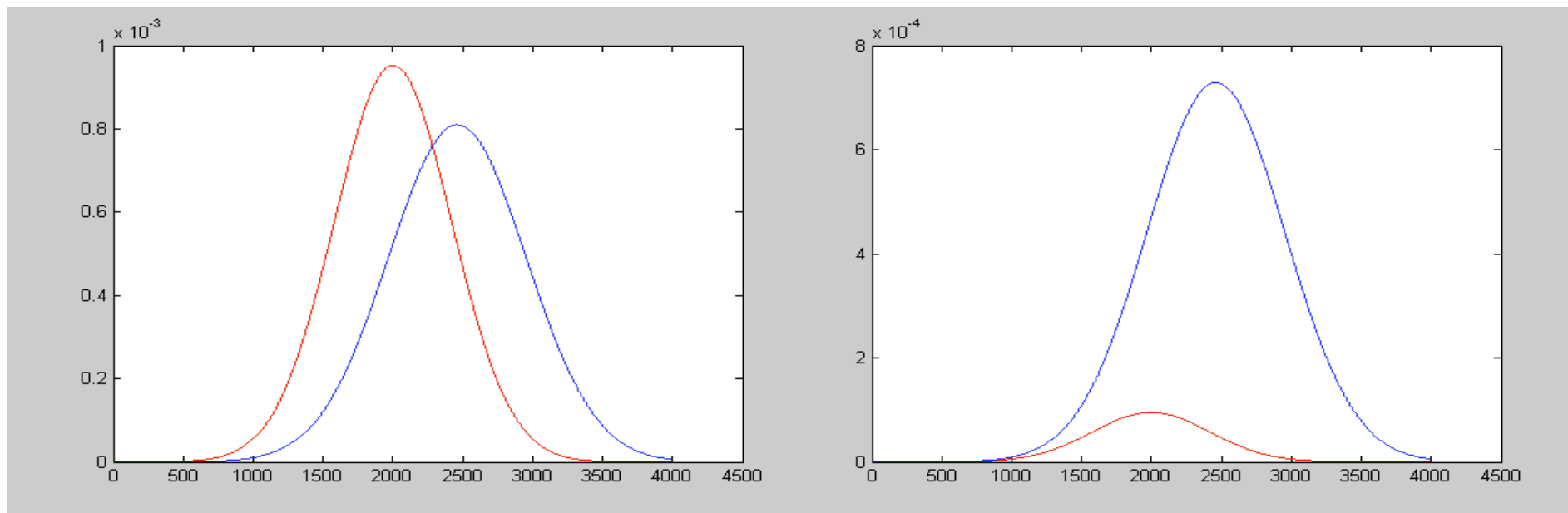
$$p(a,b) = p(a | b)p(b)$$

$$p(a) = \sum_b p(a | b)p(b) = \sum_b p(a,b)$$

Posterior Probability

$$p(\text{mouth} | a_3) = \frac{\overset{\text{likelihood}}{p(a_3 | \text{mouth})} \overset{\text{prior}}{p(\text{mouth})}}{\underset{\text{normalization constant (independent of mouth)}}{p(a_3)}}$$

Maximum A Posteriori Classification

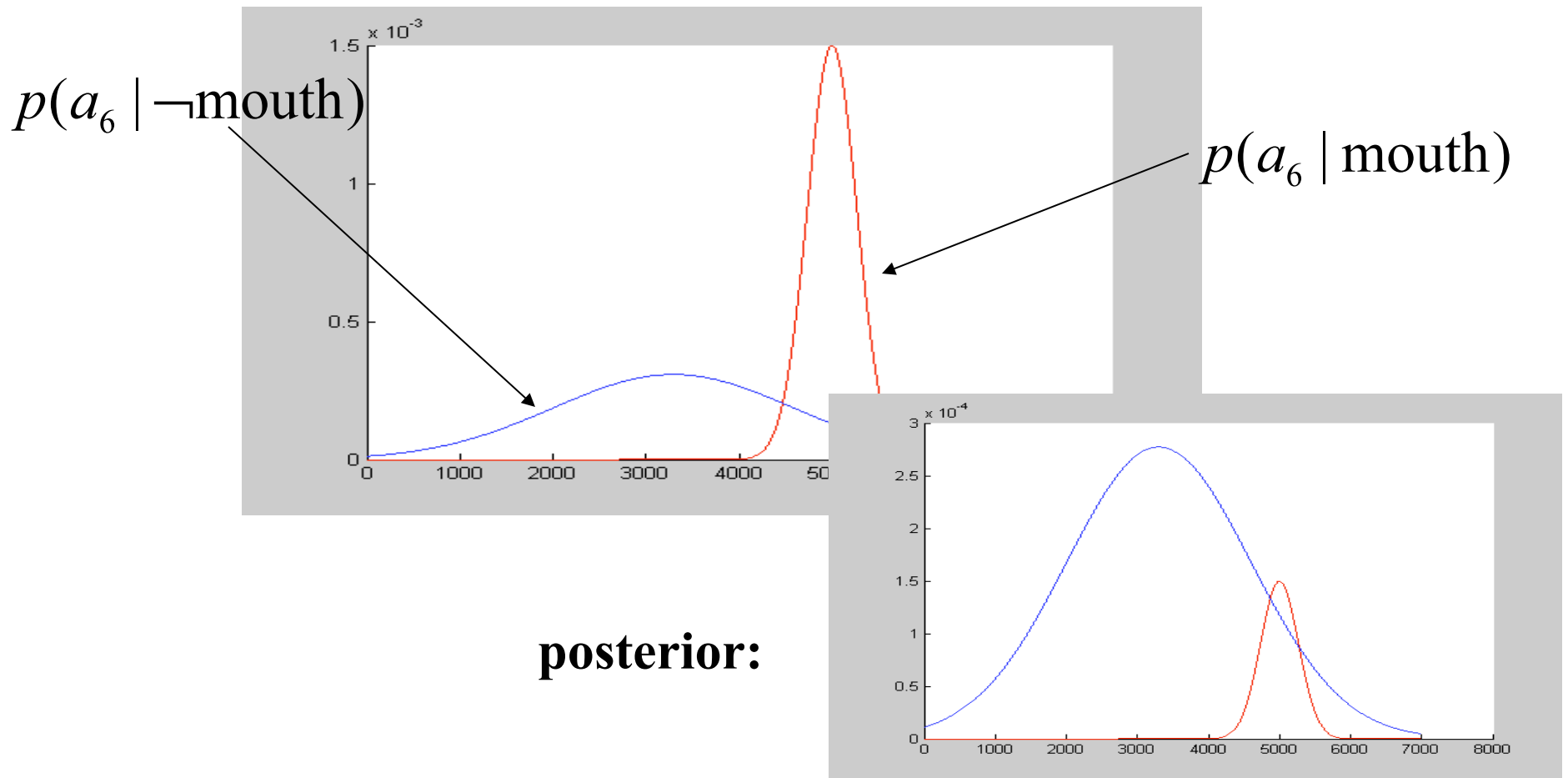


likelihood

posterior

From a_3 alone, it looks like MAP classification will always prefer the not-mouth interpretation.

What about the other coefficients?



Conditional Independence

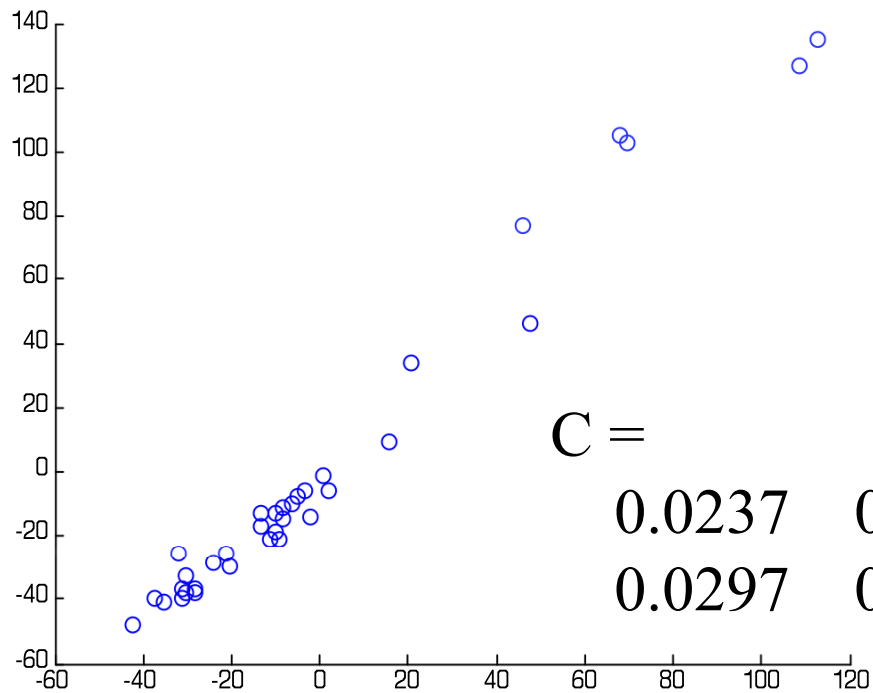
$$p(a_1, a_2, \dots, a_M \mid \text{mouth}) = \prod_{i=1}^M p(a_i \mid \text{mouth})$$

Conditional Independence

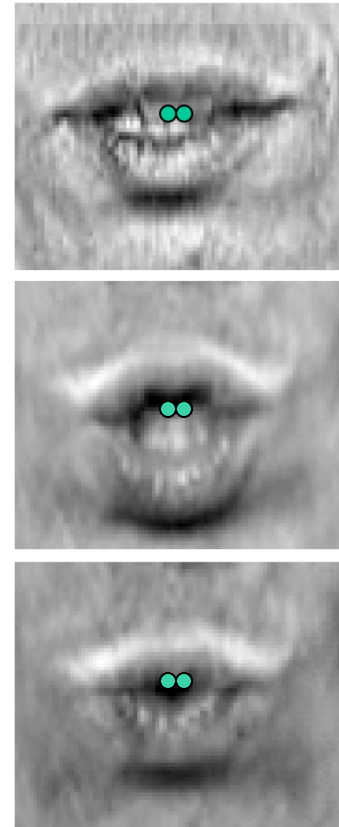
$$p(a_1, a_2, \dots, a_M \mid \text{mouth}) = \prod_{i=1}^M p(a_i \mid \text{mouth})$$

Where does this break?

Example: Covariance

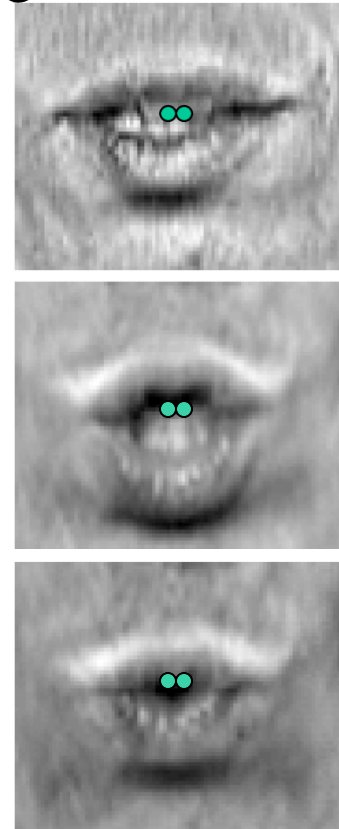
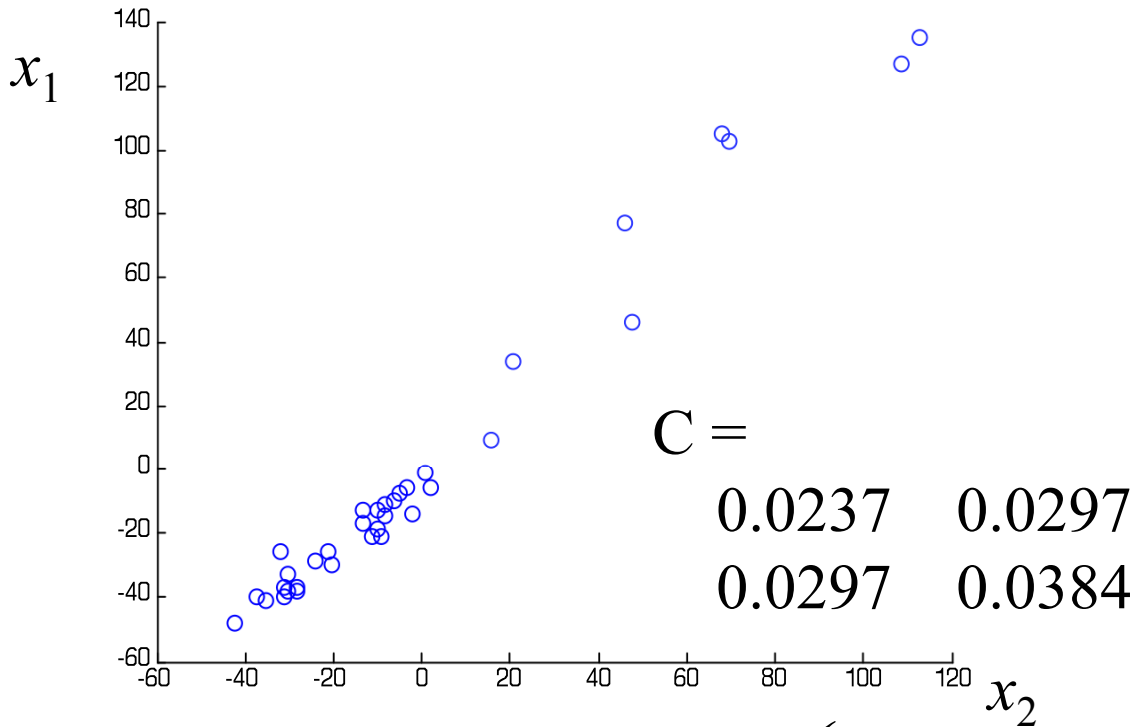


$$C = \begin{bmatrix} 0.0237 & 0.0297 \\ 0.0297 & 0.0384 \end{bmatrix}$$



$$C = \text{cov}(A(:,30*88+46)/255, A(:,30*88+47)/255)$$

Example: Covariance



$$p(\bar{x}) = \frac{1}{(2\pi)^{D/2} |C|^{1/2}} \exp\left(-\frac{1}{2}(\bar{x} - \bar{\mu})^T C^{-1}(\bar{x} - \bar{\mu})\right)$$

Example: Covariance

```
mu=[0 0];
```

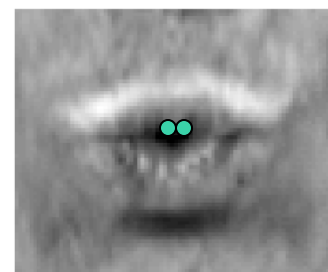
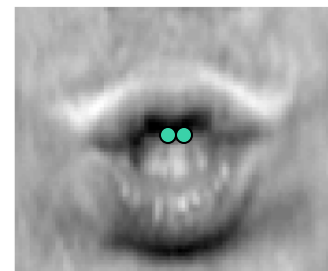
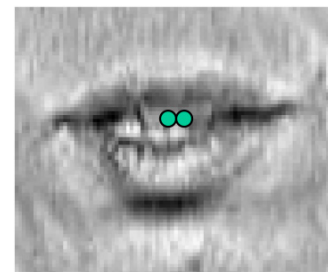
```
% draw 500 samples from a multivariate
```

```
% Gaussian
```

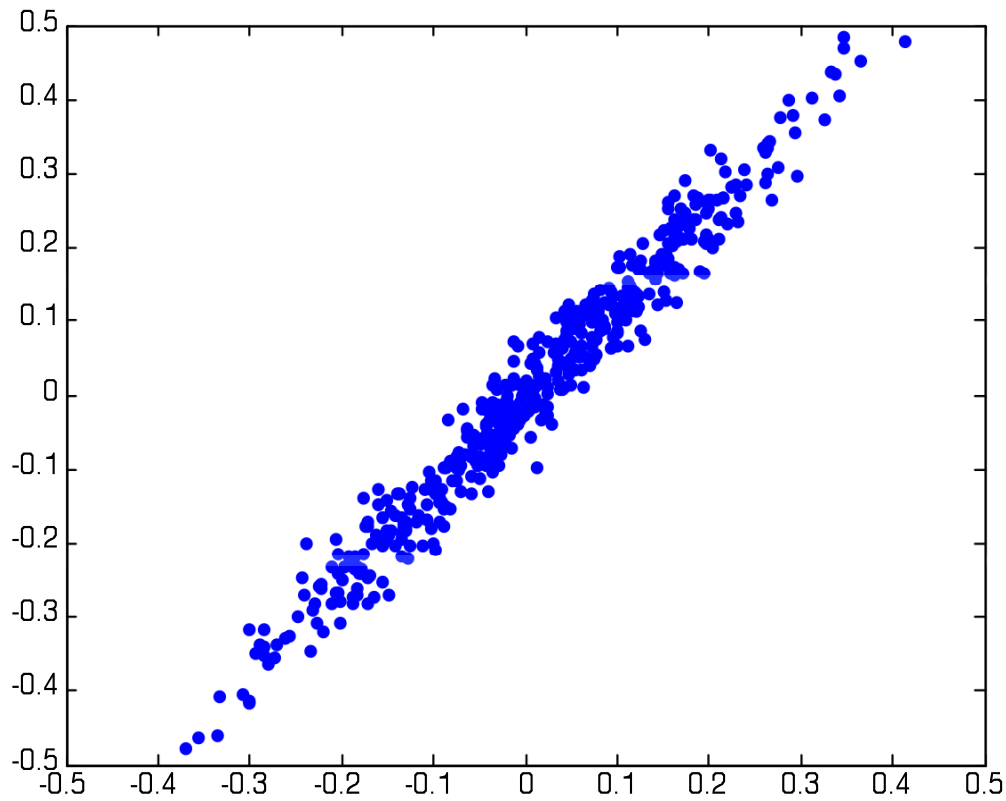
```
r = mvnrnd(mu, C, 500);
```

```
plot(r(:,1), r(:,2), '.');
```

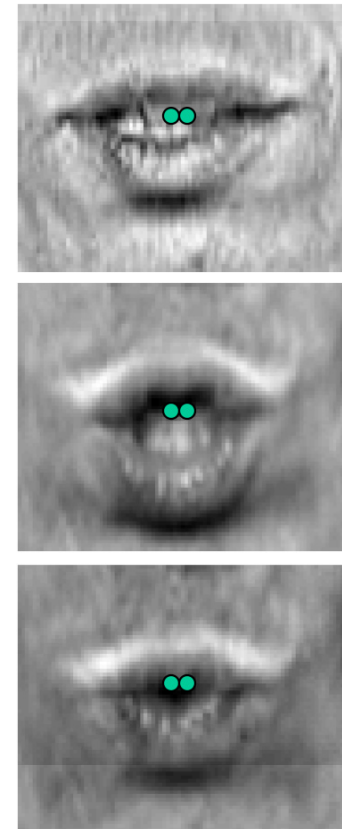
```
axis([-0.5 0.5 -0.5 0.5])
```



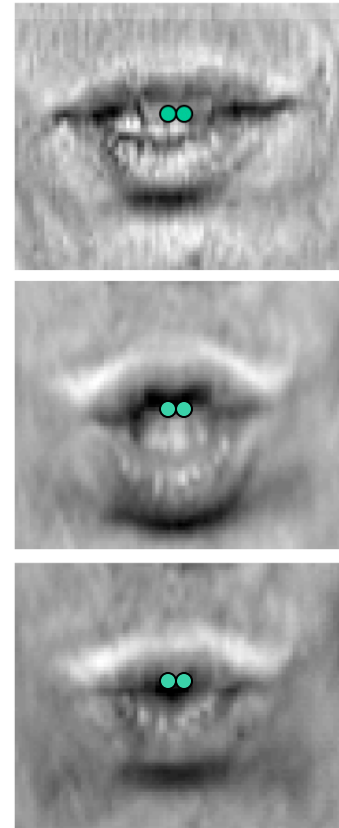
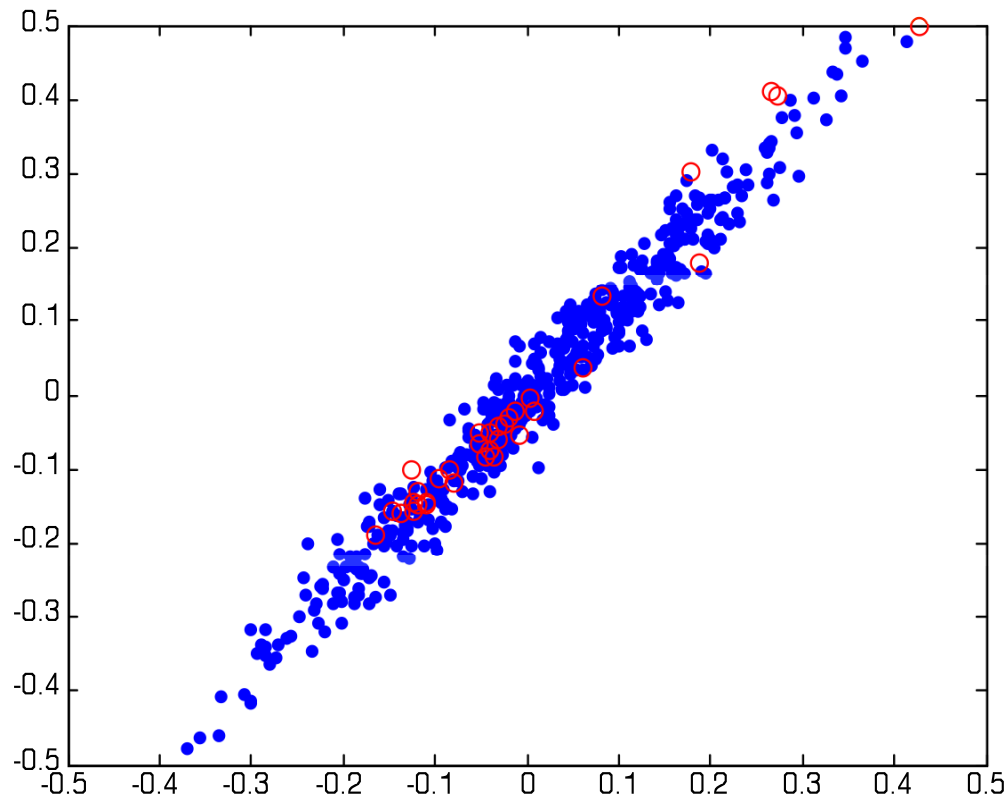
Example: Covariance



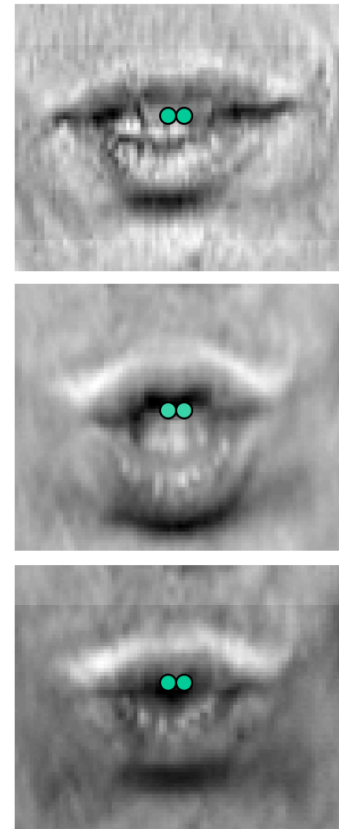
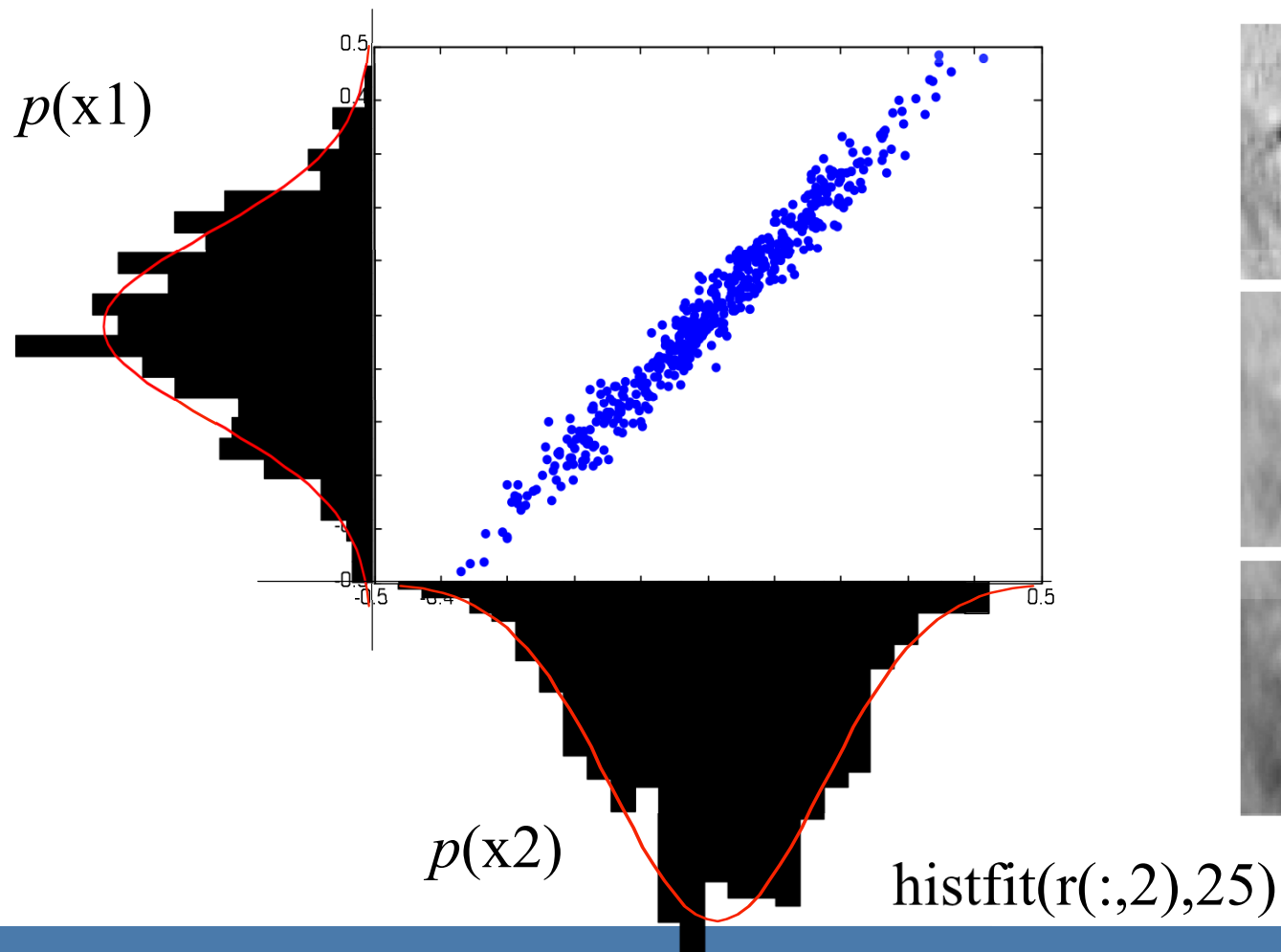
Samples from $p(x_1, x_2)$



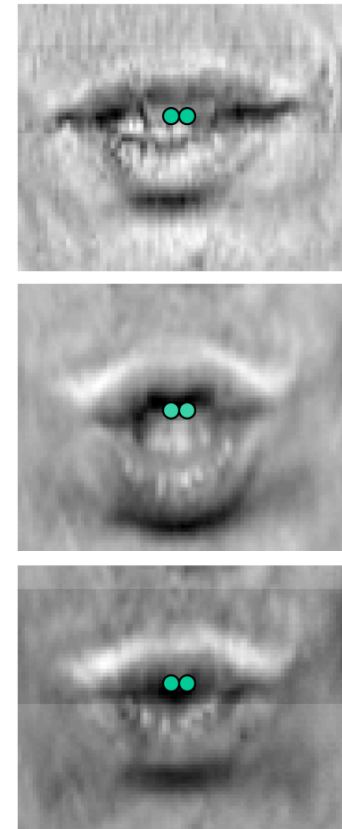
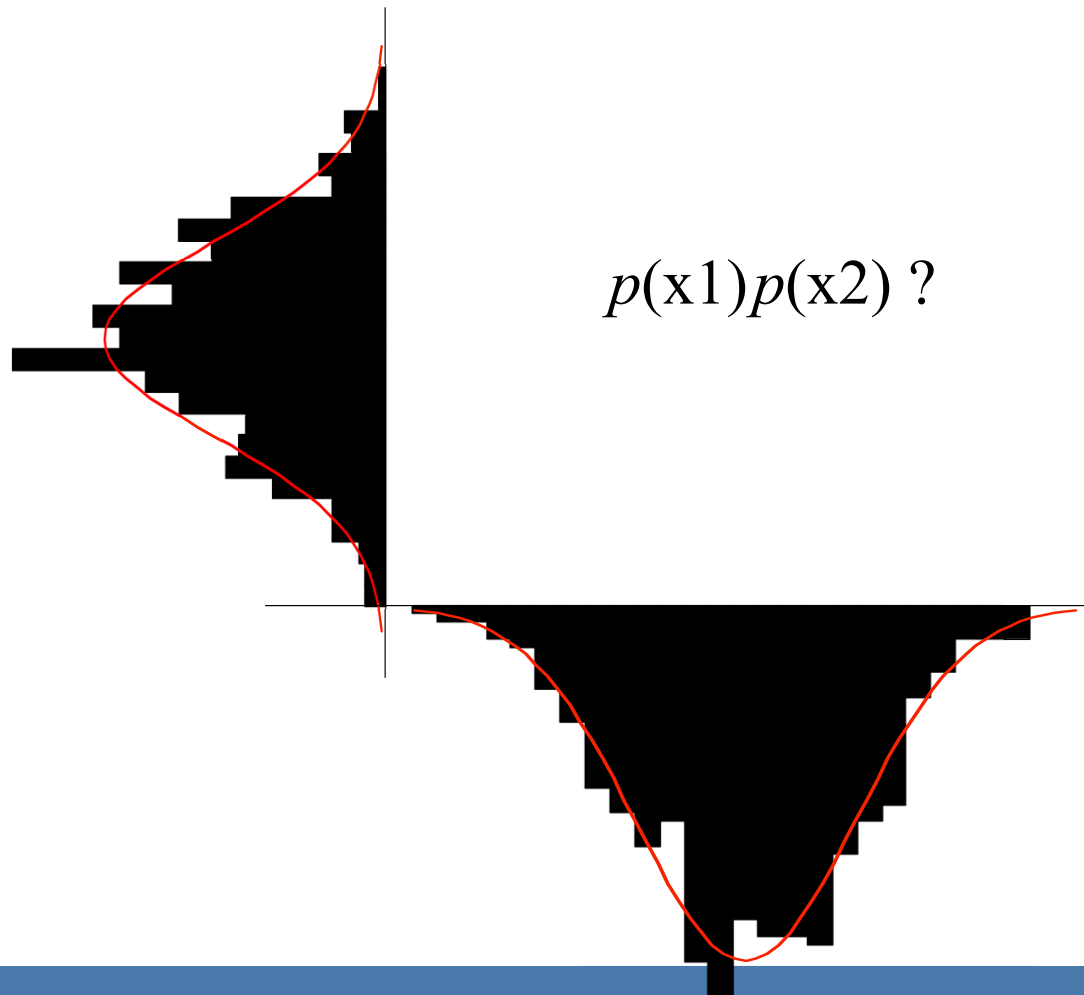
Example: Covariance



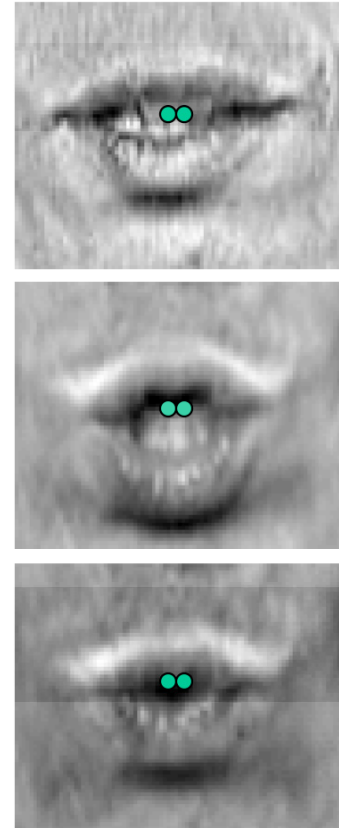
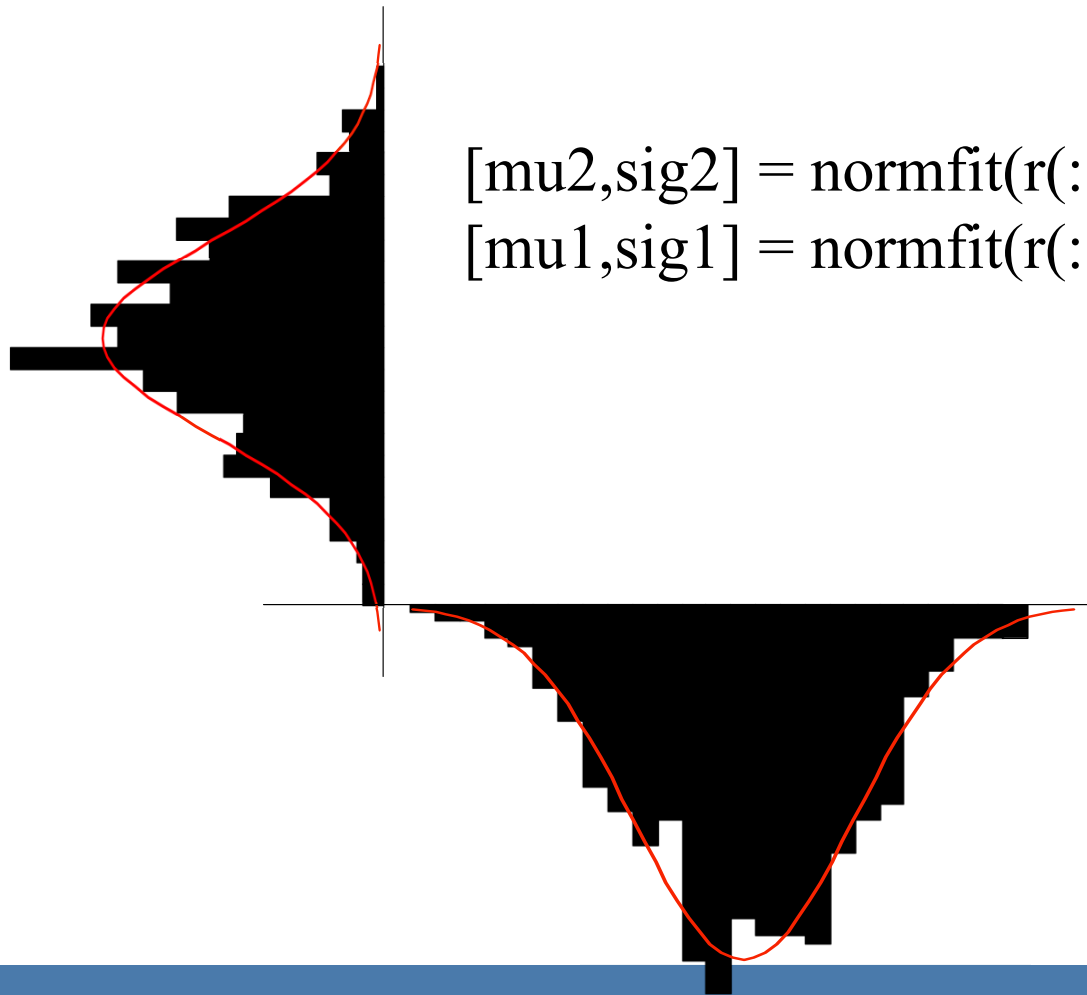
Marginals



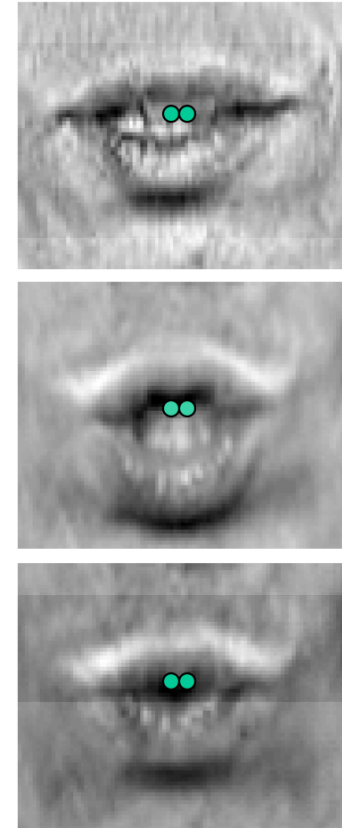
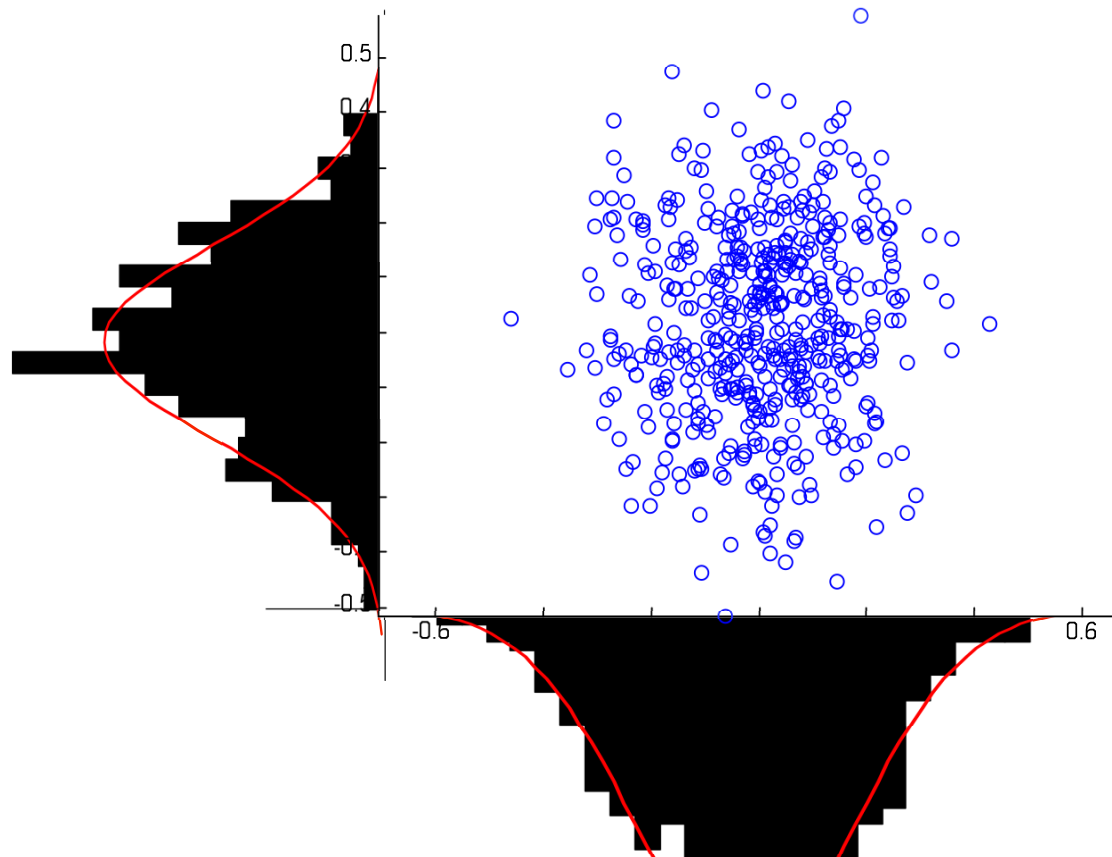
Independence



Independence

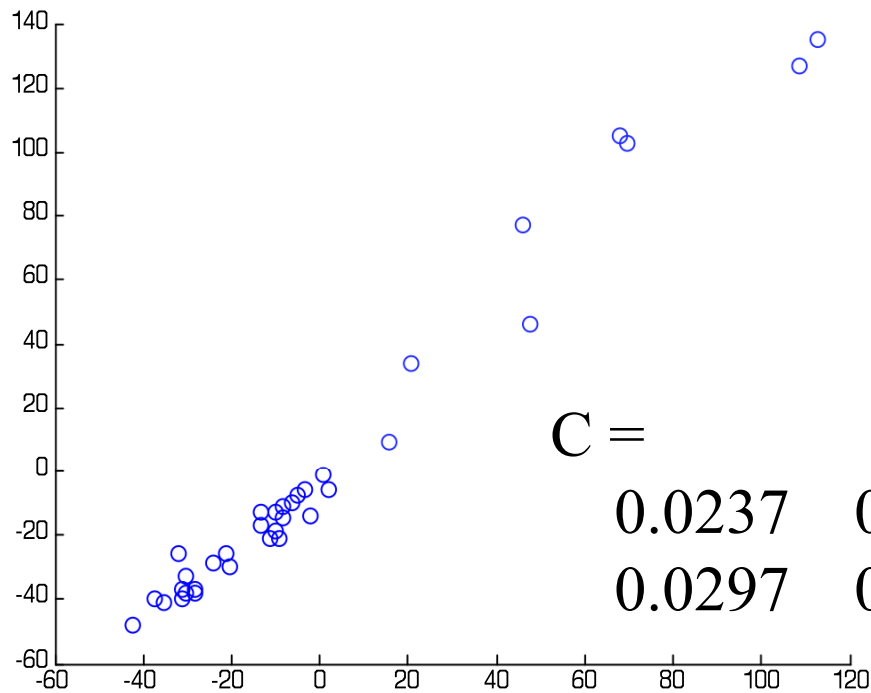


Independence



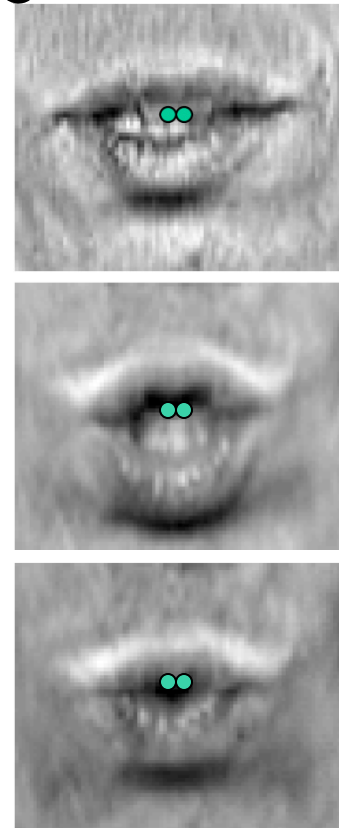
```
scatter(normrnd(mu1,sig1,500,1),normrnd(mu2,sig2,500,1))
```


Example: Covariance



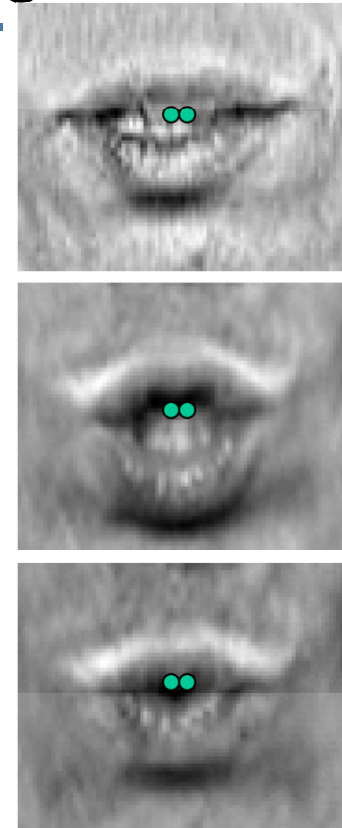
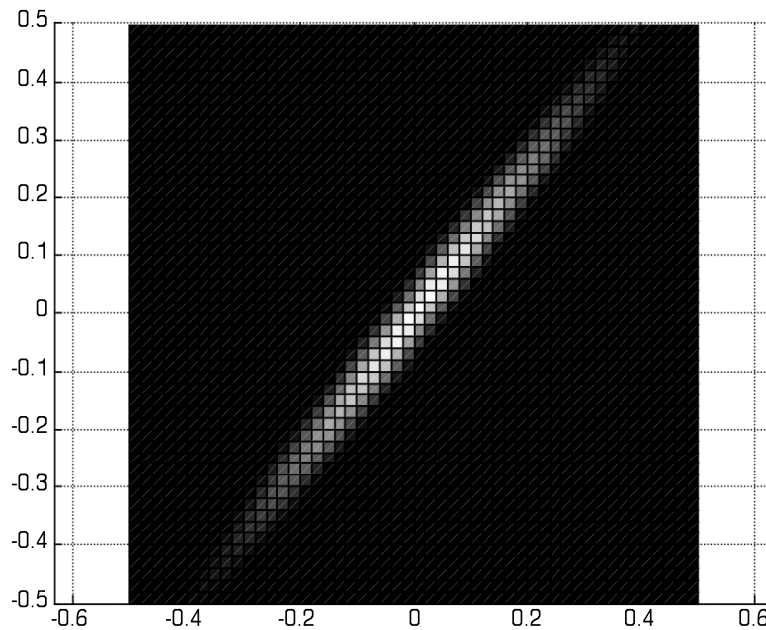
C =

$$\begin{matrix} 0.0237 & 0.0297 \\ 0.0297 & 0.0384 \end{matrix}$$



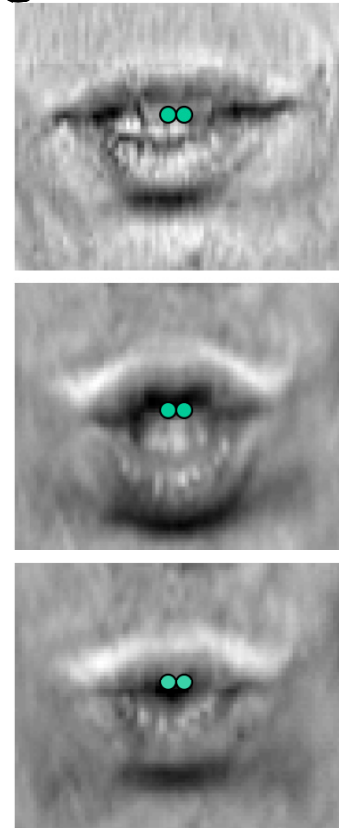
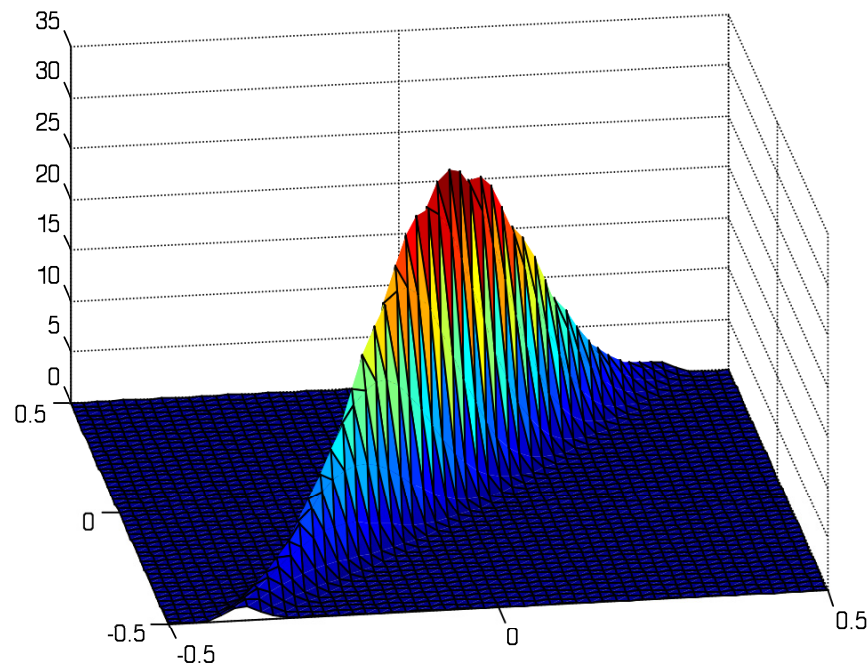
$$p(\bar{x}) = \frac{1}{(2\pi)^{D/2} |C|^{1/2}} \exp\left(-\frac{1}{2}(\bar{x} - \bar{\mu})^T C^{-1}(\bar{x} - \bar{\mu})\right)$$

Example: Covariance



```
[X1, X2]=meshgrid(-0.5:0.02:0.5, -0.5:0.02:0.5);  
X = [X1(:) X2(:)];  
p=mvnpdf(X, mu, C);  
surf(X1, X2, reshape(p,size(X1,1), size(X1,2)));
```

Example: Covariance



```
[X1, X2]=meshgrid(-0.5:0.02:0.5, -0.5:0.02:0.5);  
X = [X1(:) X2(:)];  
p=mvnpdf(X, mu, C);  
surf(X1, X2, reshape(p,size(X1,1), size(X1,2))); colormap default;
```

Whitening

```
[U,D]=eig(C)
```

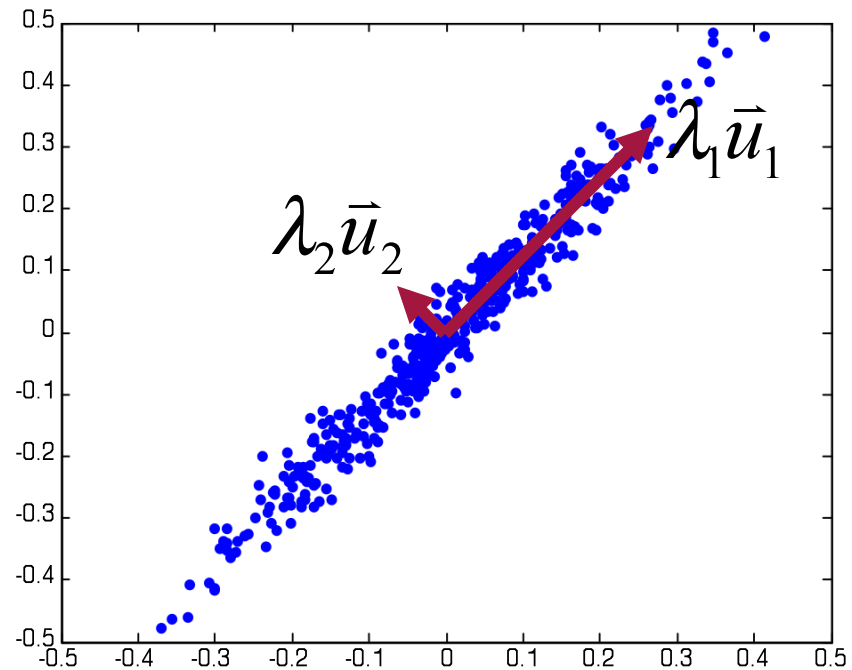
```
U =
```

```
-0.7876  0.6162  
 0.6162  0.7876
```

```
D =
```

```
0.0004    0  
 0    0.0617
```

```
plot(r(:,1), r(:,2), '.');
```



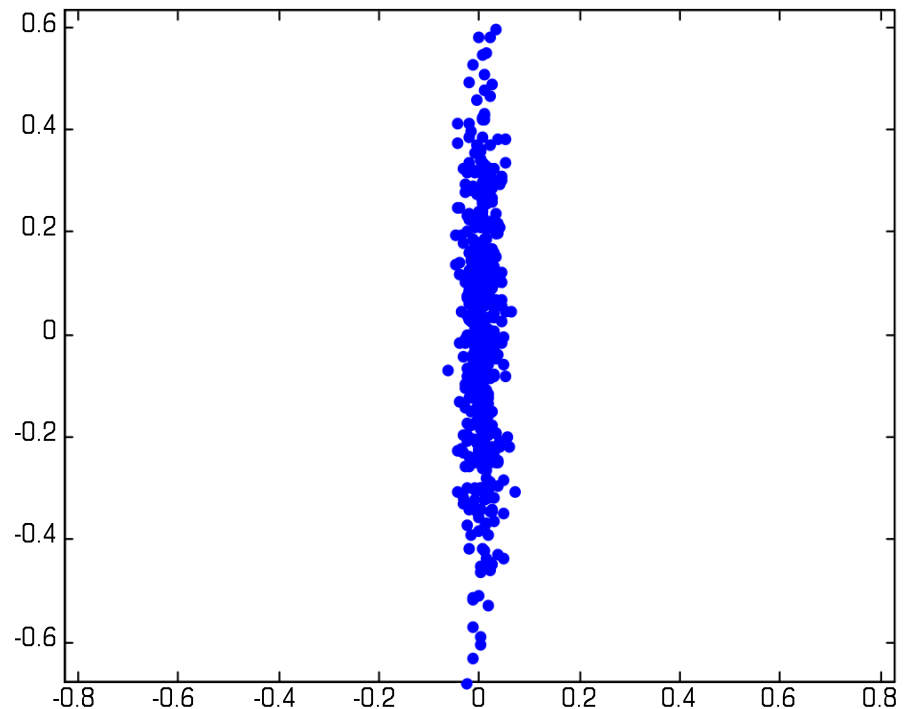
Whitening

```
[U, D]=eig(C)
% project points onto basis
coeffs = r*U;
plot(coeffs(:,1), coeffs(:,2), '!');
axis true
```

$$\tilde{x}^n = \bar{x}^n U$$

\bar{x}^n zero mean

$$\text{cov}(\tilde{x}) = D$$

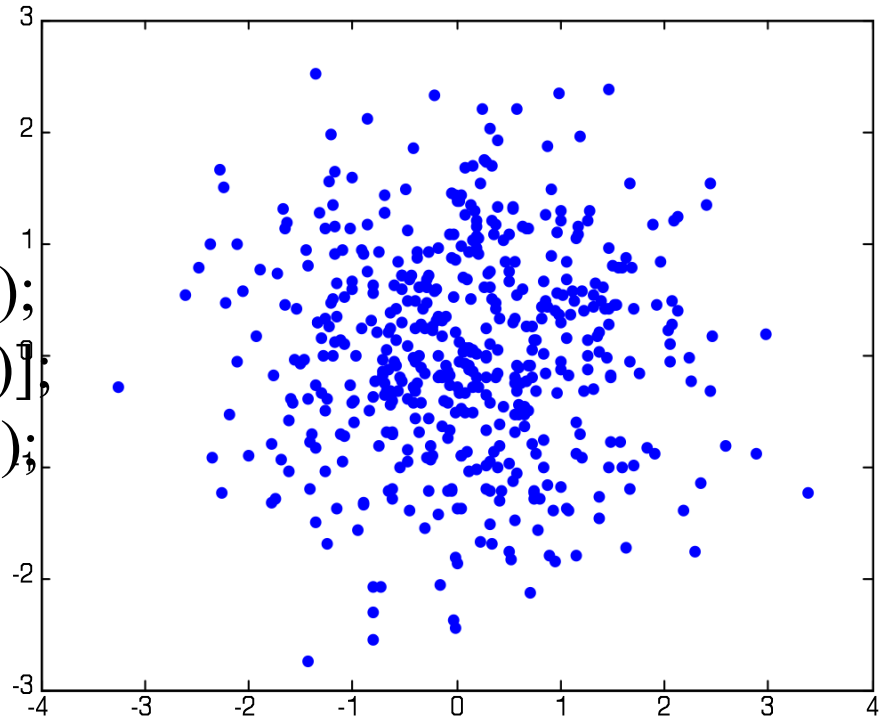


Whitening

```
[U, D]=eig(C)
% project points onto basis
coeffs = r*U;
coeffs2=[coeffs(1,+)/sqrt(D(1,1));
         coeffs(2,+)/sqrt(D(2,2))];
plot(coeffs2(:,1), coeffs2(:,2) ,'.');
axis true
```

$$\tilde{\mathbf{x}}^n = \bar{\mathbf{x}}^n \mathbf{U} \mathbf{D}^{-1/2}$$

$$\bar{\mathbf{x}}^n \text{ zero mean} \quad \text{cov}(\tilde{\mathbf{x}}) = \mathbf{I}$$



Diagonal Covariance

$$p(\bar{x}) = \frac{1}{(2\pi)^{D/2} |C|^{1/2}} \exp\left(-\frac{1}{2}(\bar{x} - \bar{\mu})^T C^{-1}(\bar{x} - \bar{\mu})\right)$$

Determinant is just the product of the diagonals (ie variances).

$$p(\bar{x}) = \prod_{i=1:D} p(x_i) = \prod_i \frac{1}{\sqrt{2\pi\sigma_i}} \exp\left(-\frac{1}{2}(\bar{x}_i - \bar{\mu}_i)^2 / \sigma_i^2\right)$$

Some Facts

$$C = E[(\bar{x} - \bar{\mu})(\bar{x} - \bar{\mu})^T] = \begin{bmatrix} \sigma_{xx} & \sigma_{xy} \\ \sigma_{yx} & \sigma_{yy} \end{bmatrix}$$

If x and y are statistically independent then $s_{xy}=0$.

If $s_{xy}=0$, then x and y are uncorrelated.

Uncorrelated does not imply statistically independent.

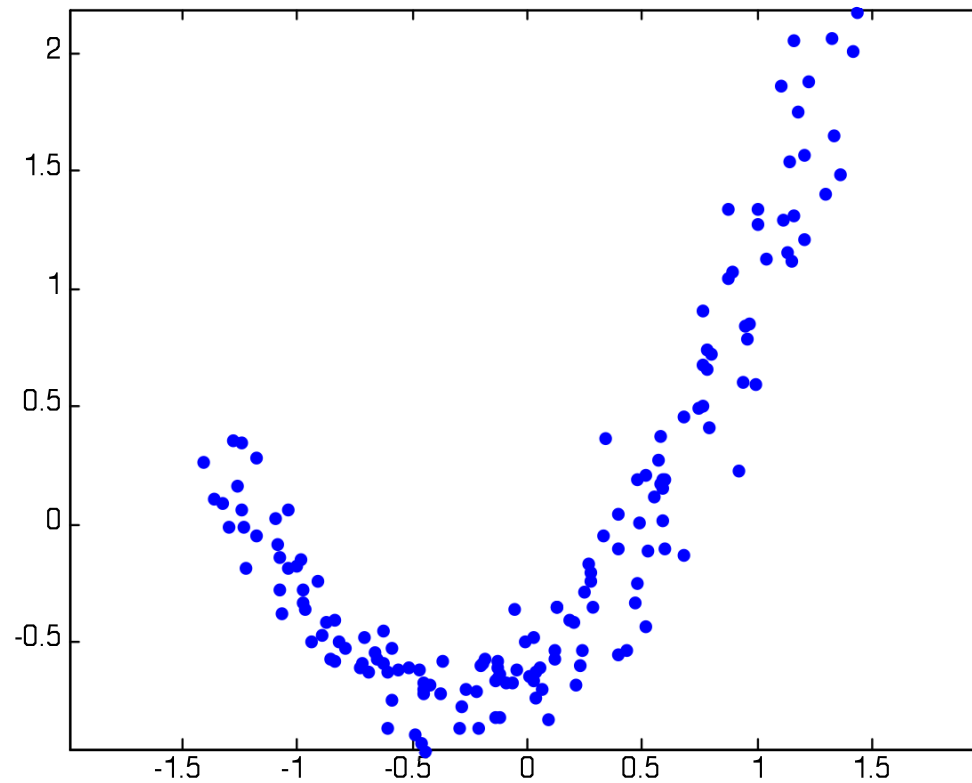
Uncorrelated and Gaussian does.

PCA de-correlates the directions but unless the data is Gaussian, the coefficients are not statistically independent.

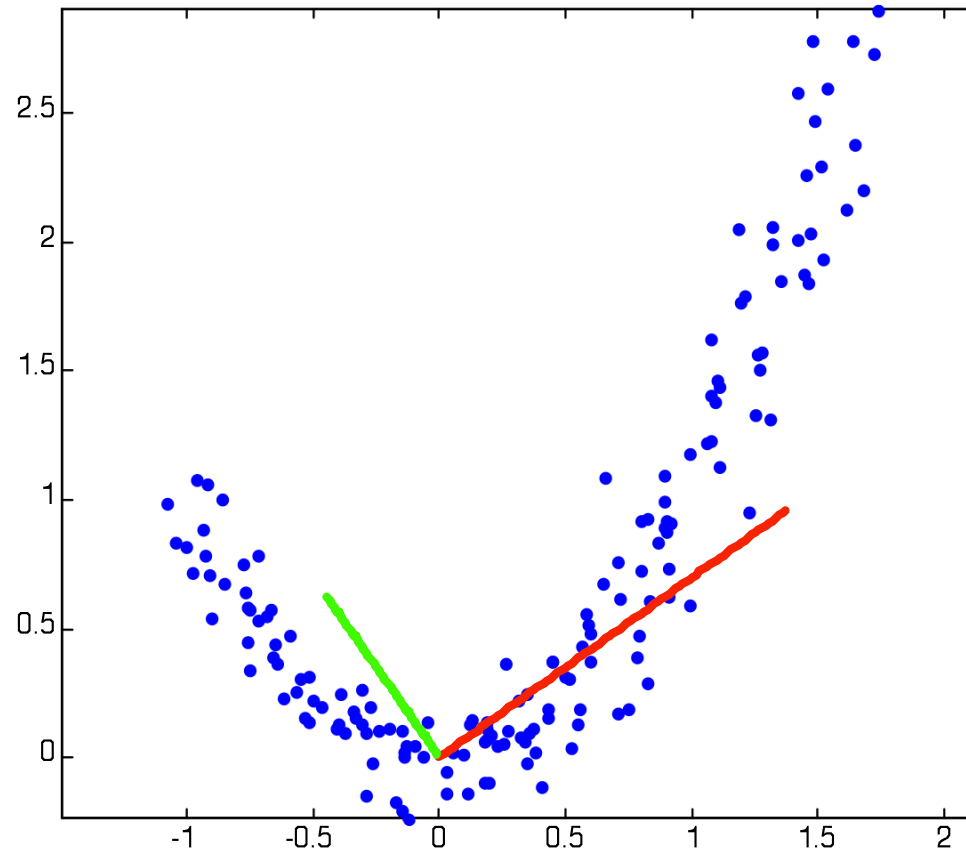
Why does decorrelated not imply statistically independent?

- PCA takes into account the second-order statistics in the data (in the covariance matrix).
- The covariance matrix captures correlation.
- PCA decorrelates the data.
- But covariance is only a second order statistic.
- Gaussians are fully described by their first and second order statistics (mean and covariance) –decorrelating then results in statistical independence
- But if the data has non-zero higher order statistics, decorrelating will not make the dimensions statistically independent.

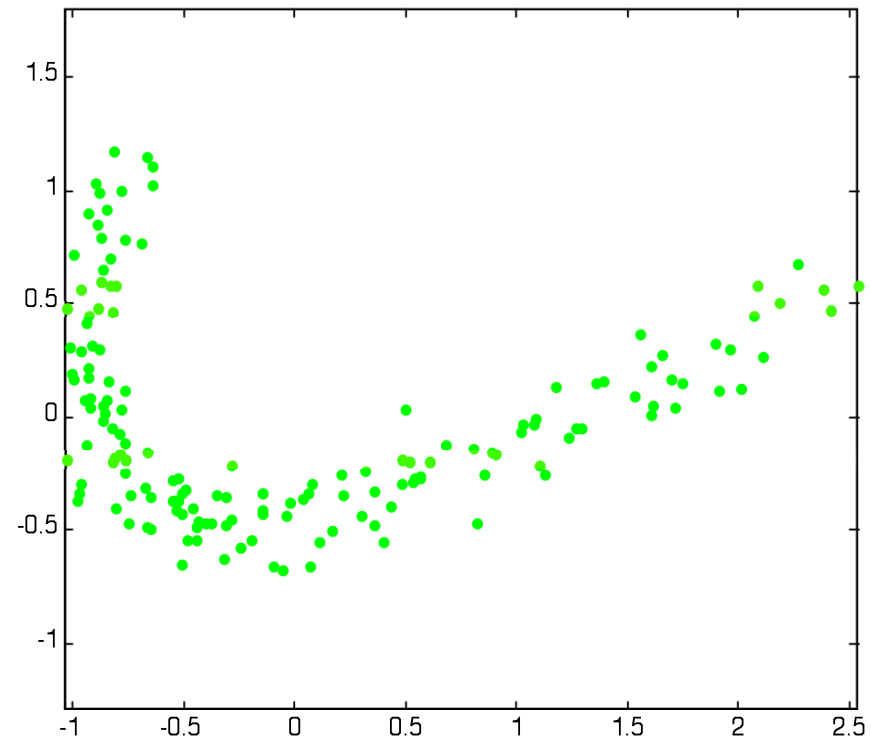
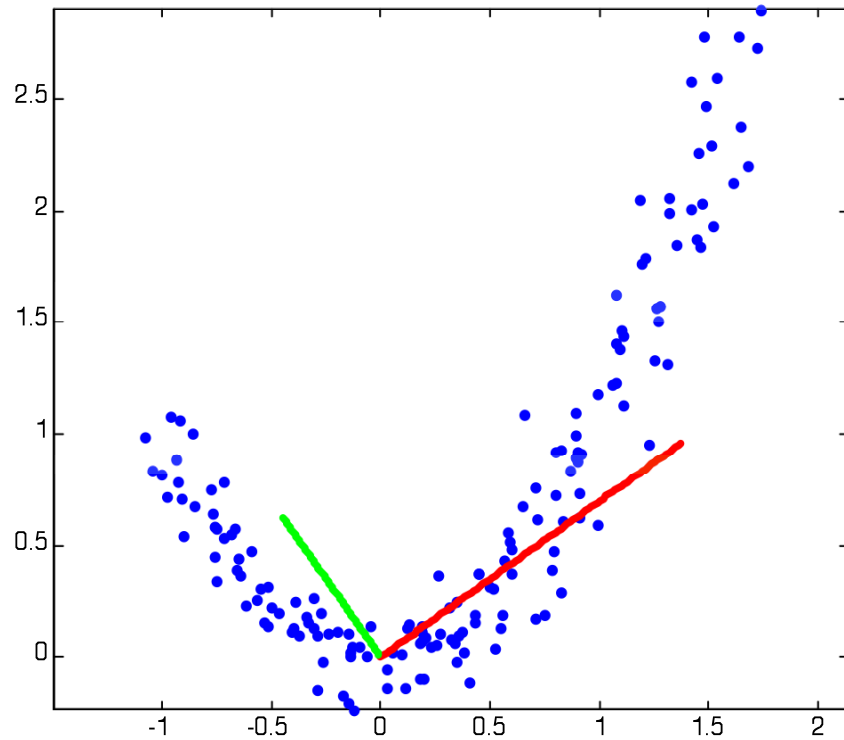
PCA and non-Gaussian data



PCA and non-Gaussian data



PCA and non-Gaussian data

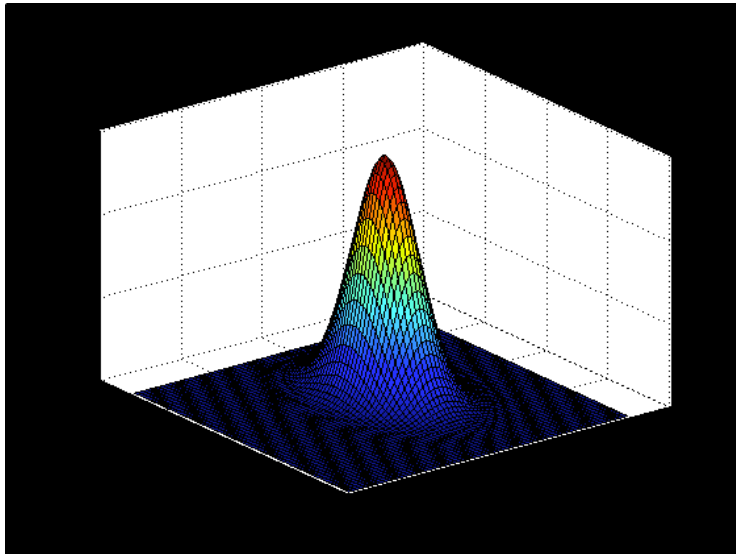


Decorrelated: $\text{corr}(X'u_1, X'u_2)$
ans = $-5.8981e-017$

$\text{cov}(X'u_1, X'u_2)$
0.9834 -0.0000
-0.0000 0.2010

Not statistically independent.

PCA and Covariance



`surf(m36)`

$A_n = A - \text{mean}(A)$

$U = \text{eigenvectors}$

$U_{36} = \text{matrix of linear coeffs}$

Let's look at how a_3 and a_6 co-vary.

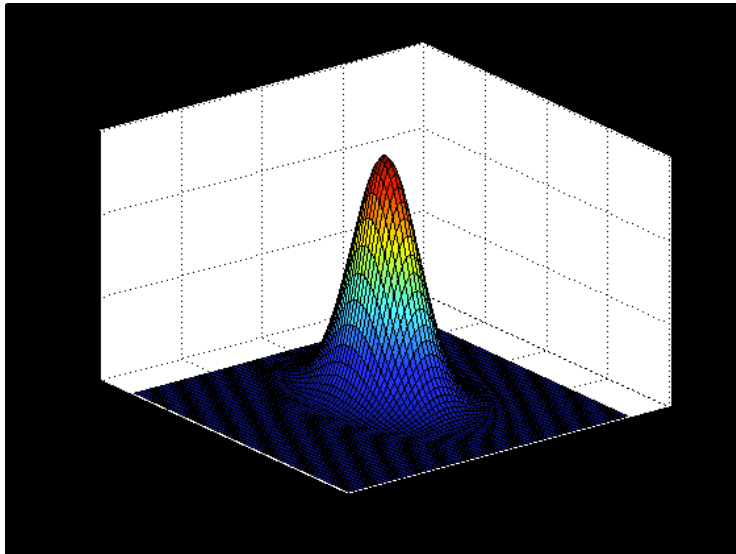
```
U36=[An*U(:,3) An*U(:,6)];
```

```
C36=cov(U36)
```

```
mu36=[mean(An*U(:,3)) mean(An*U(:,6))]
```

```
m36=mvnpdf(X, mu36, C36);
```

Covariance

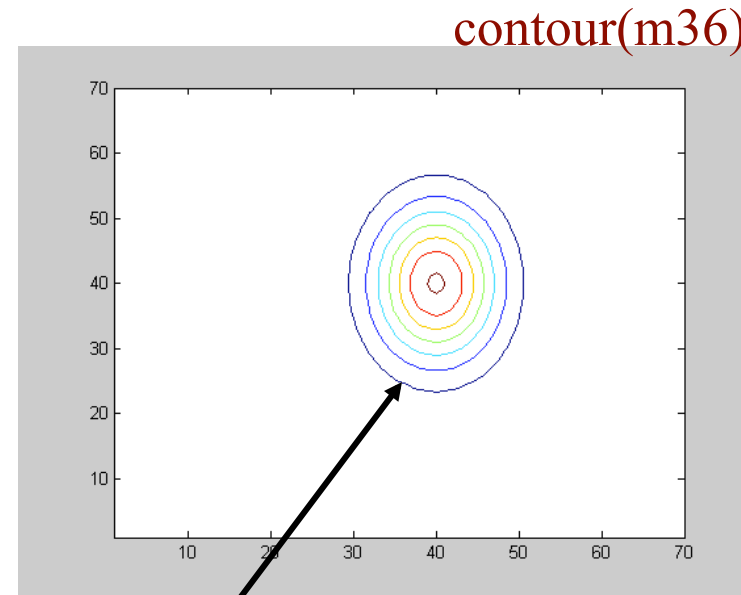
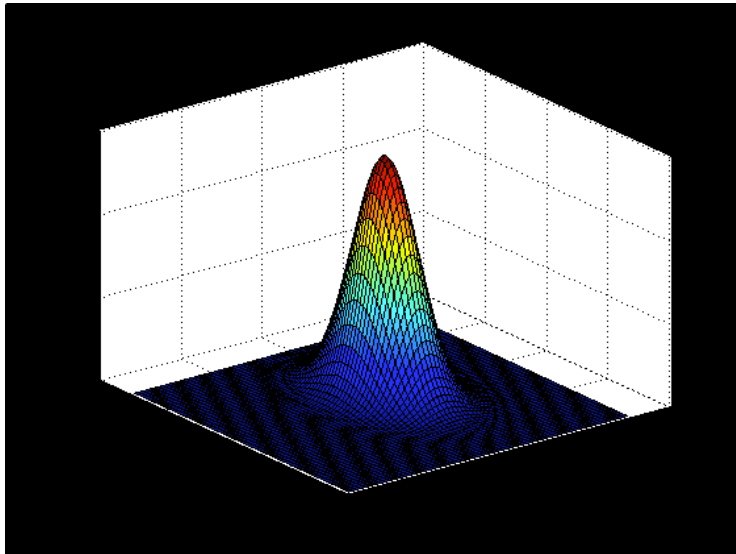


Multivariate Gaussian
(Normal)

Mahalanobis distance Δ^2

$$p(\bar{x}) = \frac{1}{(2\pi)^{D/2} |C|^{1/2}} \exp\left(-\frac{1}{2} \overbrace{(\bar{x} - \bar{\mu})^T C^{-1} (\bar{x} - \bar{\mu})}^{\Delta^2}\right)$$

Covariance Ellipse



hyperellipsoids of constant Mahalanobis distance Δ^2

Note the ellipse is axis-aligned. Why?

Mahalanobis distance

$$p(\bar{x}) = \frac{1}{(2\pi)^{D/2} |C|^{1/2}} \exp\left(-\frac{1}{2}(\bar{x} - \bar{\mu})^T C^{-1}(\bar{x} - \bar{\mu})\right)$$

$$\tilde{x} = \bar{x} - \bar{\mu}$$

$$d(\tilde{x}) = (\tilde{x}^T C^{-1} \tilde{x})$$

$$C = USU^T$$

Mahalanobis Distance

$$\begin{aligned}d(\tilde{x}) &= (\tilde{x}^T C^{-1} \tilde{x}) \\&= \tilde{x}^T (USU^T)^{-1} \tilde{x} \\&= \tilde{x}^T US^{-1}U^T \tilde{x} \\&= y^T S^{-1} y \\&= \sum_{i=1}^D \frac{y_i^2}{\lambda_i} \\&\approx \sum_{i=1}^M \frac{y_i^2}{\lambda_i}\end{aligned}$$

Linear coefficients

$$y = U^T \tilde{x}$$

Error in approximation?

- Above measures “distance in feature space”.
- Residual error is “distance from feature space”. This can be approximated.
- See Moghaddam & Pentland paper on website.
- Taking approximate error into account improves detection for problem 3.