

# Dimensionality Reduction

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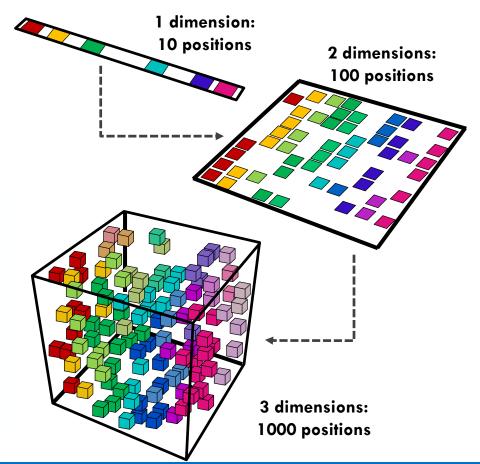
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### **Curse of Dimensionality**

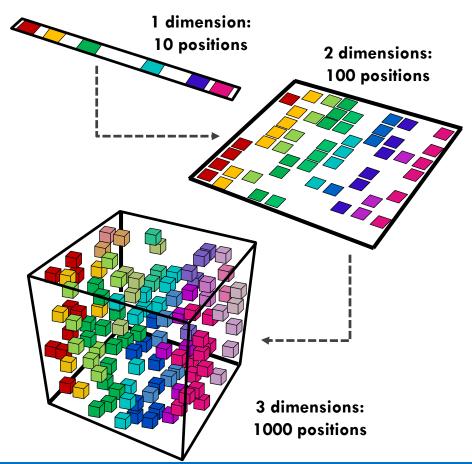
 Theoretically, increasing features should improve performance





### **Curse of Dimensionality**

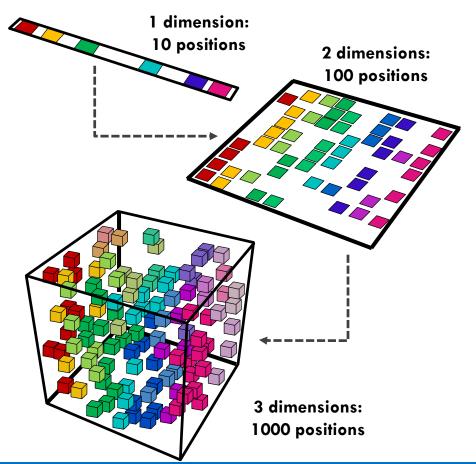
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- In practice, too many features leads to worse performance





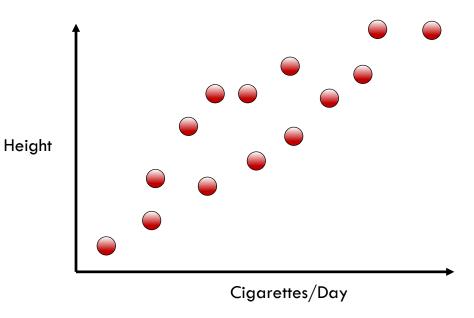
### Curse of Dimensionality

- Theoretically, increasing features should improve performance
- In practice, too many features leads to worse performance
- Number of training examples required increases exponentially with dimensionality





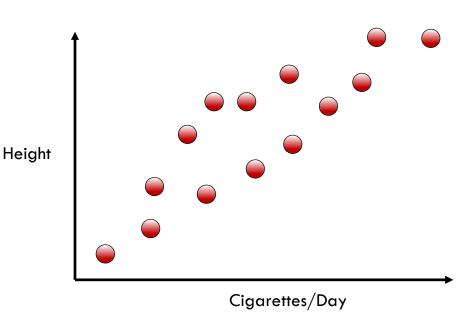
- Data can be represented by fewer dimensions (features)
- Reduce dimensionality by





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- Reduce dimensionality by selecting subset (feature elimination)

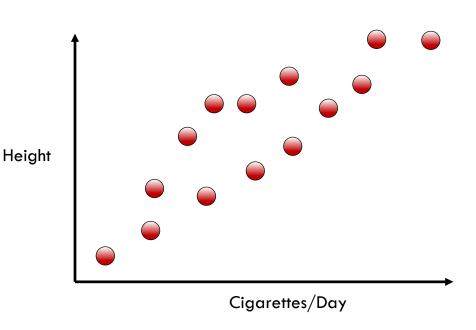
Combine with linear and non-





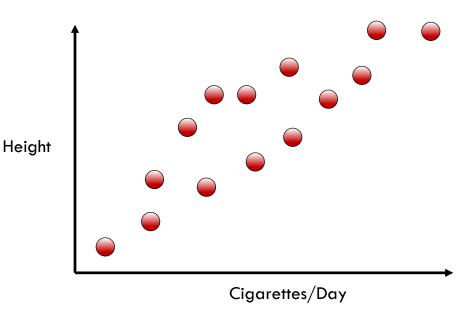
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- Reduce dimensionality by selecting subset (feature elimination)

 Combine with linear and nonlinear transformations



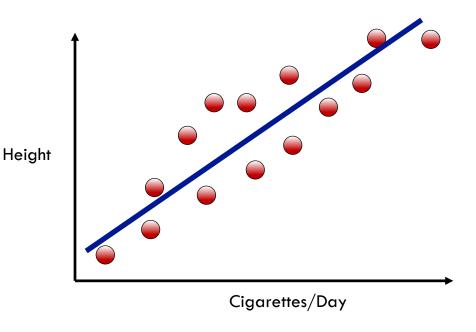


- Two features: height and cigarettes per day
- Both features increase together (correlated)
- Can we reduce number of features to one?



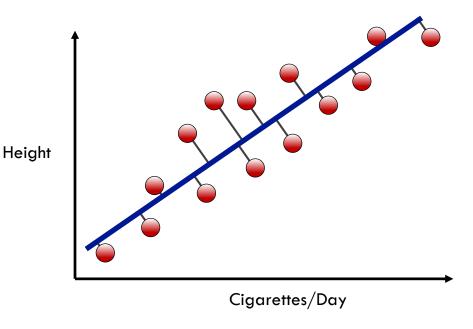


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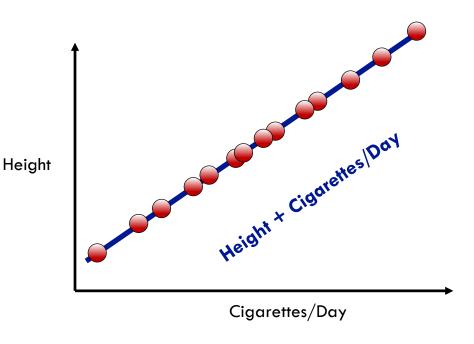


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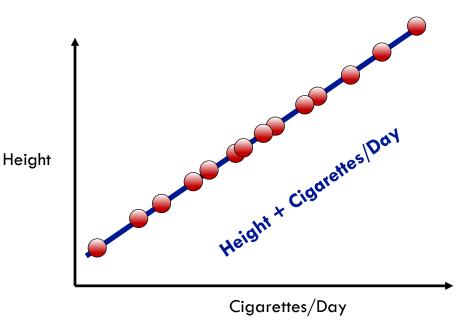


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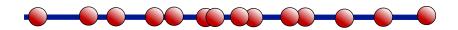


- Create single feature that is combination of height and cigarettes
- This is Principal Component Analysis (PCA)





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Height + Cigarettes/Day

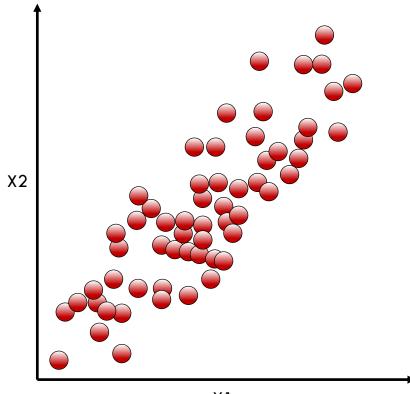


### **Dimensionality Reduction**

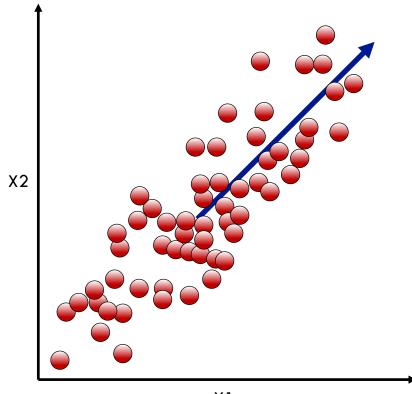
Given an N-dimensional data set (x), find a  $N \times K$  matrix (U):

 $y = U^T x$ , where y has K dimensions and K < N

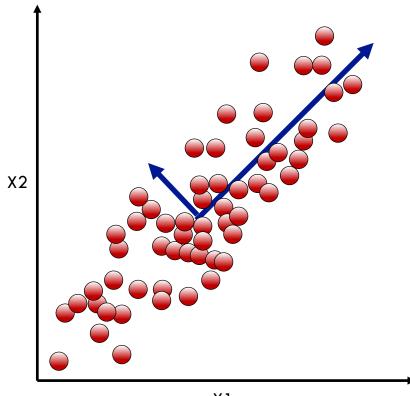
$$x = \begin{bmatrix} x_1 \\ x_2 \\ \dots \\ x_n \end{bmatrix} \xrightarrow{U^T} y = \begin{bmatrix} y_1 \\ y_2 \\ \dots \\ y_k \end{bmatrix} (K < N)$$

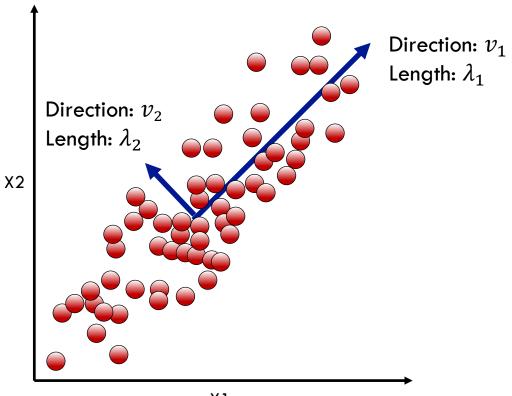












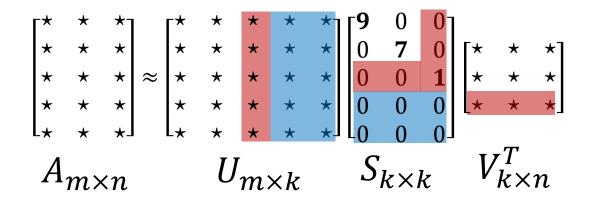
### Single Value Decomposition (SVD)

- SVD is a matrix factorization method normally used for PCA
- Does not require a square data set
- SVD is used by Scikit-learn for PCA



### Truncated Single Value Decomposition

- How can SVD be used for dimensionality reduction?
- Principal components are calculated from US
- "Truncated SVD" used for dimensionality reduction  $(n \rightarrow k)$

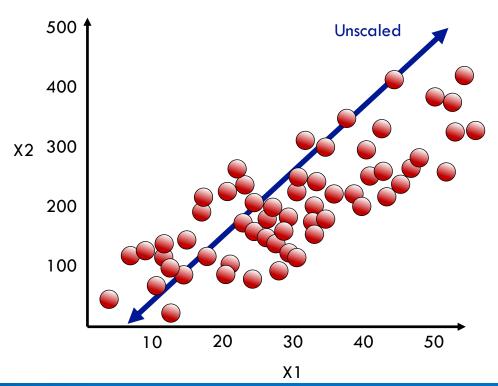




### Importance of Feature Scaling

 PCA and SVD seek to find the vectors that capture the most variance

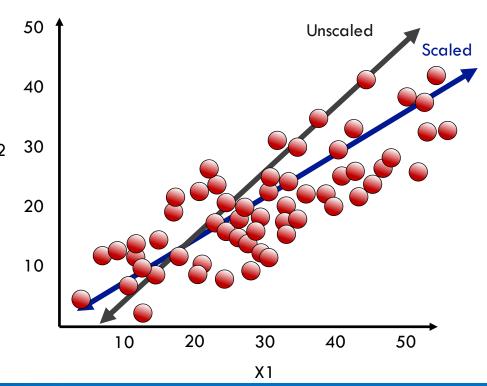
 Variance is sensitive to axis scale





### Importance of Feature Scaling

- PCA and SVD seek to find the vectors that capture the most variance
- Variance is sensitive to axis scale
- Must scale data!





Import the class containing the dimensionality reduction method

from sklearn.decomposition import PCA



#### Import the class containing the dimensionality reduction method

from sklearn.decomposition import PCA

#### Create an instance of the class

PCAinst = PCA(n\_components=3, whiten=True)

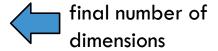


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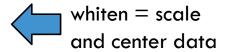


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X\_trans = PCAinst.fit\_transform(X\_train)



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Does not work with sparse matrices



### Truncated SVD: The Syntax

#### Import the class containing the dimensionality reduction method

from sklearn.decomposition import TruncatedSVD

#### Create an instance of the class

SVD = TruncatedSVD(n\_components=3)

#### Fit the instance on the data and then transform the data

X\_trans = SVD.fit\_transform(X\_sparse)

Works with sparse matrices—used with text data for Latent Semantic Analysis (LSA)

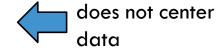


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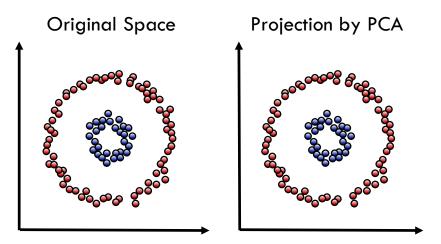
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### Moving Beyond Linearity

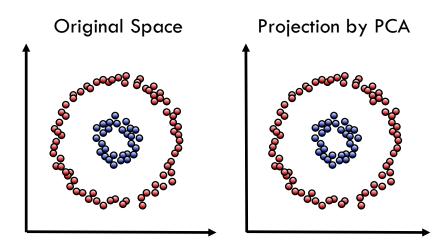
- Transformations calculated with PCA/SVD are linear
- Data can have non-linear features
- · This can cause dimensionality





### Moving Beyond Linearity

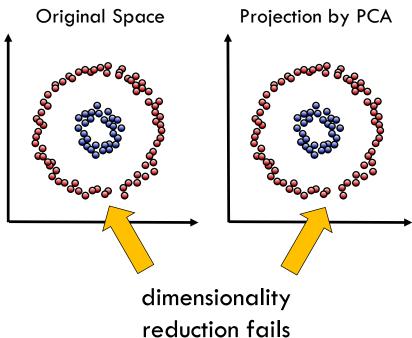
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- Transformations calculated with PCA/SVD are linear
- Data can have non-linear features
- This can cause dimensionality reduction to fail

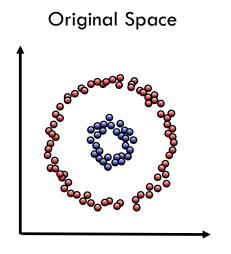


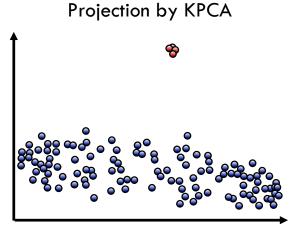


### Kernel PCA

 Solution: kernels can be used to perform non-linear PCA

1.1 1 1.1



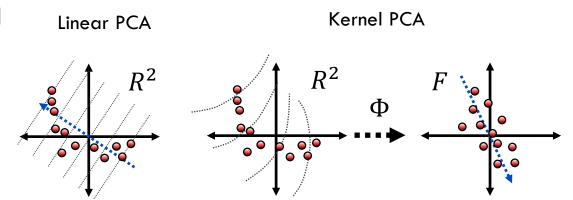




### Kernel PCA

 Solution: kernels can be used to perform non-linear PCA

 Like the kernel trick introduced for SVMs





### Kernel PCA: The Syntax

#### Import the class containing the dimensionality reduction method

from sklearn.decomposition import KernelPCA

#### Create an instance of the class

kPCA = KernelPCA(n\_components=3, kernel='rbf', gamma=1.0)

#### Fit the instance on the data and then transform the data

X\_trans = kPCA.fit\_transform(X\_train)

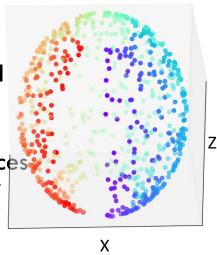


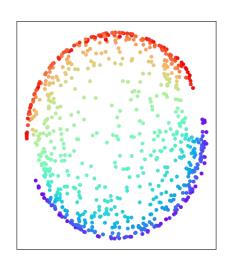
## Multi-Dimensional Scaling (MDS)

Non-linear transformation

 Doesn't focus on maintaining overall variance

Instead, maintains geometric distances between points







## MDS: The Syntax

#### Import the class containing the dimensionality reduction method

from sklearn.manifold import MDS

#### Create an instance of the class

mdsMod = MDS(n\_components=2)

#### Fit the instance on the data and then transform the data

X\_trans = mdsMod.fit\_transform(X\_sparse)

Many other manifold dimensionality methods exist: Isomap, TSNE.

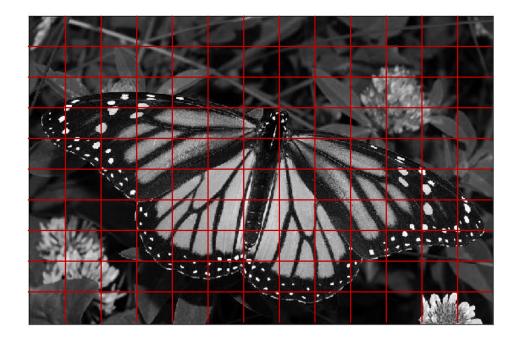


- Frequently used for high dimensionality data
- Natural language processing (NLP)—many word combinations
- Image-based data sets—pixels are features



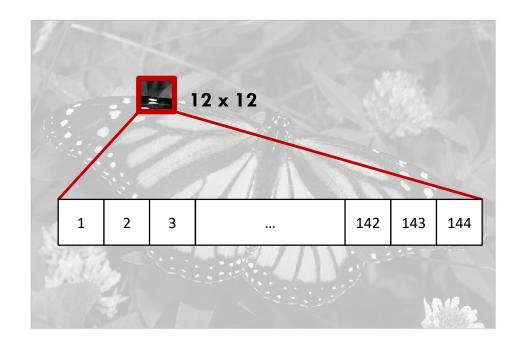


Divide image into 12 x 12 pixel sections





- Divide image into 12 x 12 pixel sections
- Flatten section to create row of data with 144 features





- Divide image into 12 x 12 pixel sections
- Flatten section to create row of data with 144 features
- Perform PCA on all data points

7/	1/100			The second second		80	
ğ	1	2	3	<b>:</b>	142	143	144
	1	2	3		142	143	144
	1	2	3		142	143	144
	1	2	3		142	143	144
	1	2	3		142	143	144
	1	2	3		142	143	144
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## PCA Compression: $144 \rightarrow 60$ Dimensions







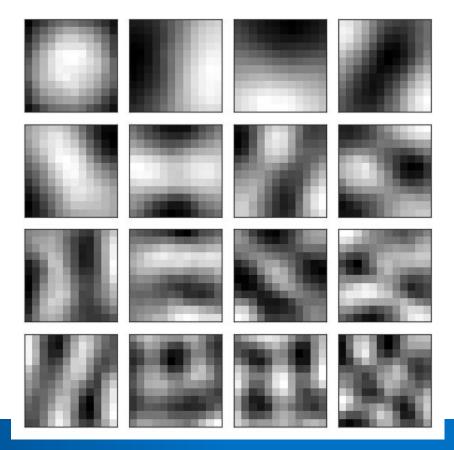
## PCA Compression: $144 \rightarrow 16$ Dimensions







# Sixteen Most Important Eigenvectors





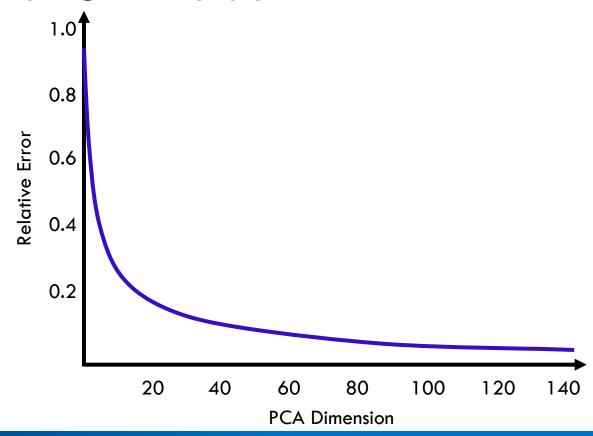
## PCA Compression: $144 \rightarrow 4$ Dimensions





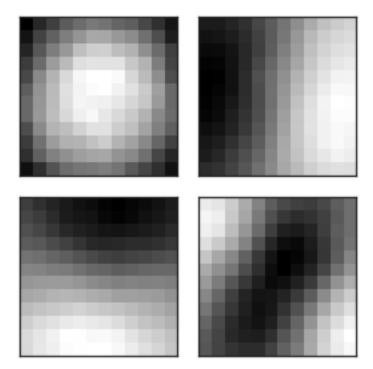


#### L2 Error and PCA Dimension



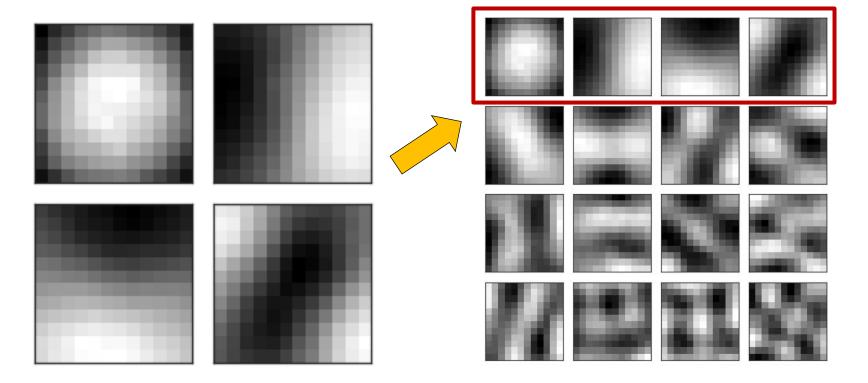


## Four Most Important Eigenvectors





# Four Most Important Eigenvectors





## PCA Compression: $144 \rightarrow 1$ Dimension



