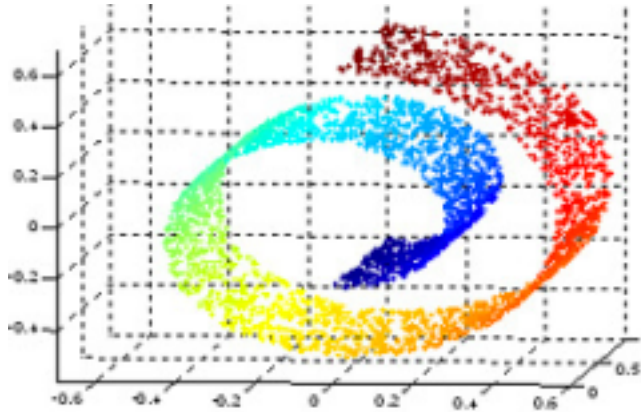


ML4Bio
Lecture #3: Dimensionality
Reduction

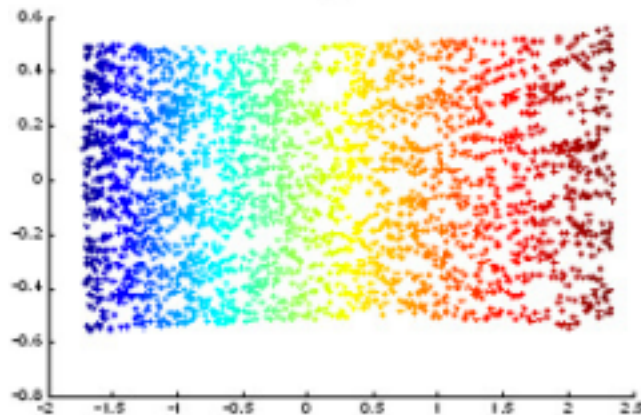
March 9th, 2016

Quaid Morris

Review: Dimensionality reduction



(a)



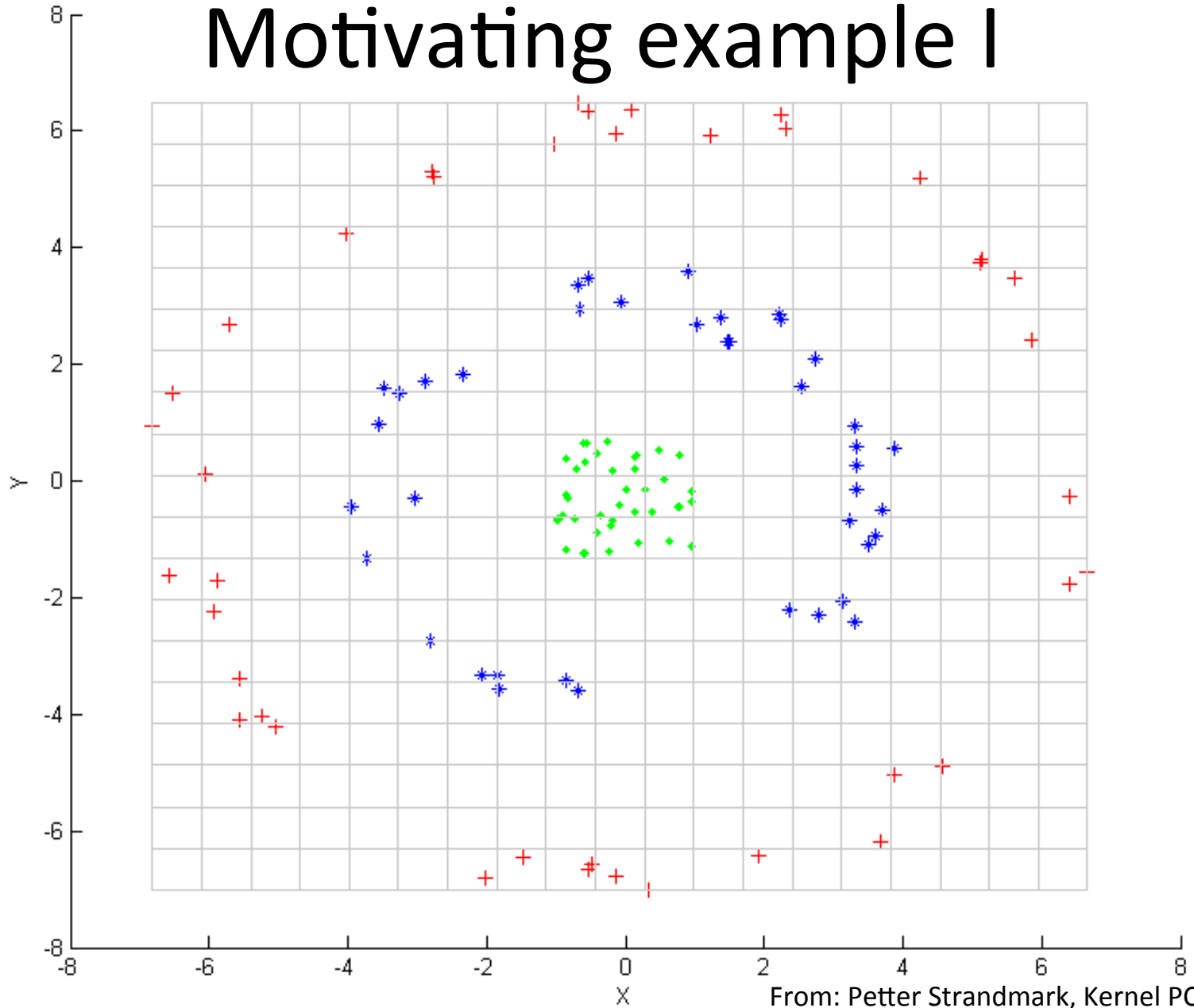
(c)

Do the datapoints lie on a lower dimensional “*manifold*”?

If so, what is the dimensionality?

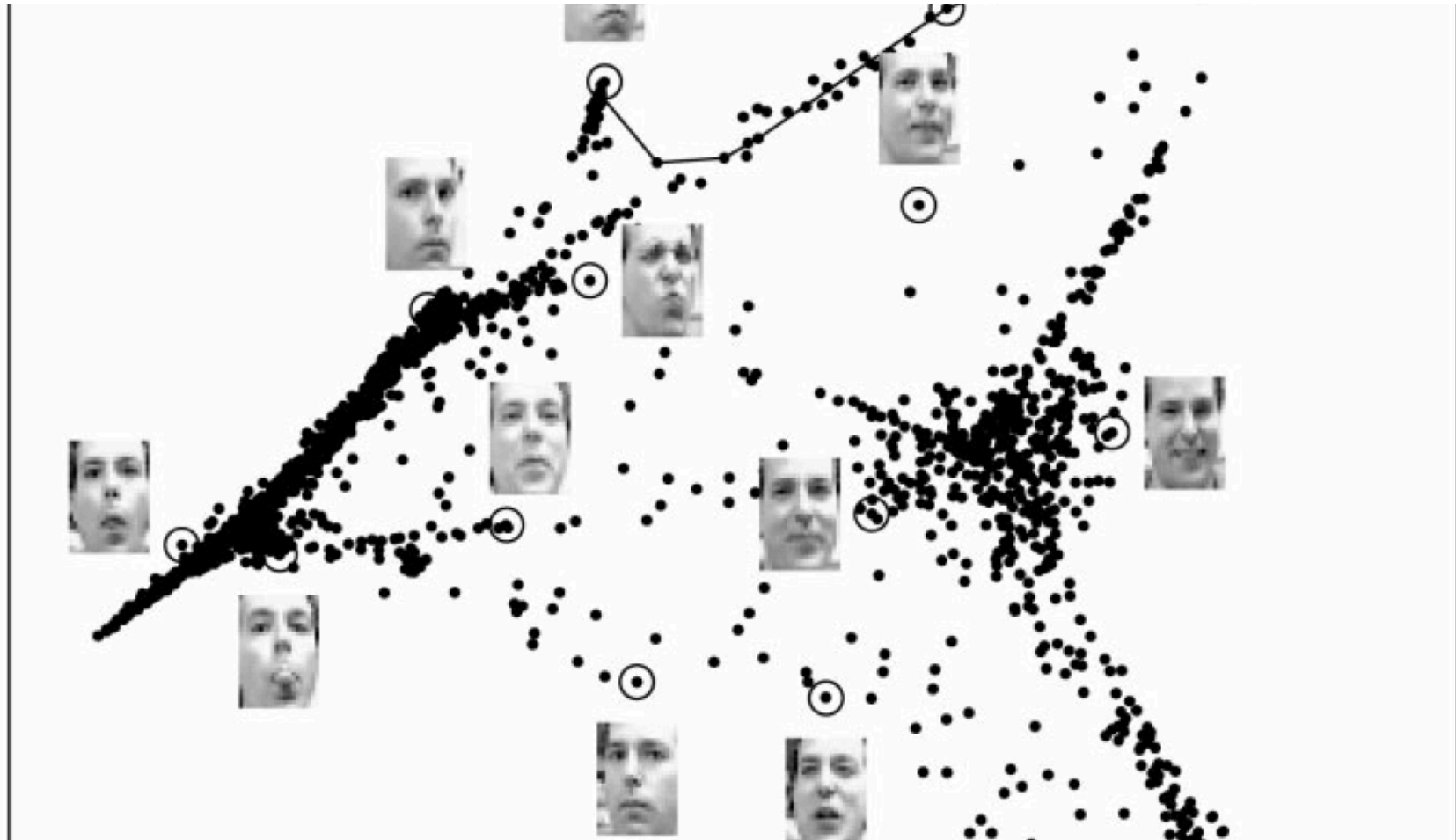
How far apart are two datapoints, if you can only travel on the manifold?

Motivating example I

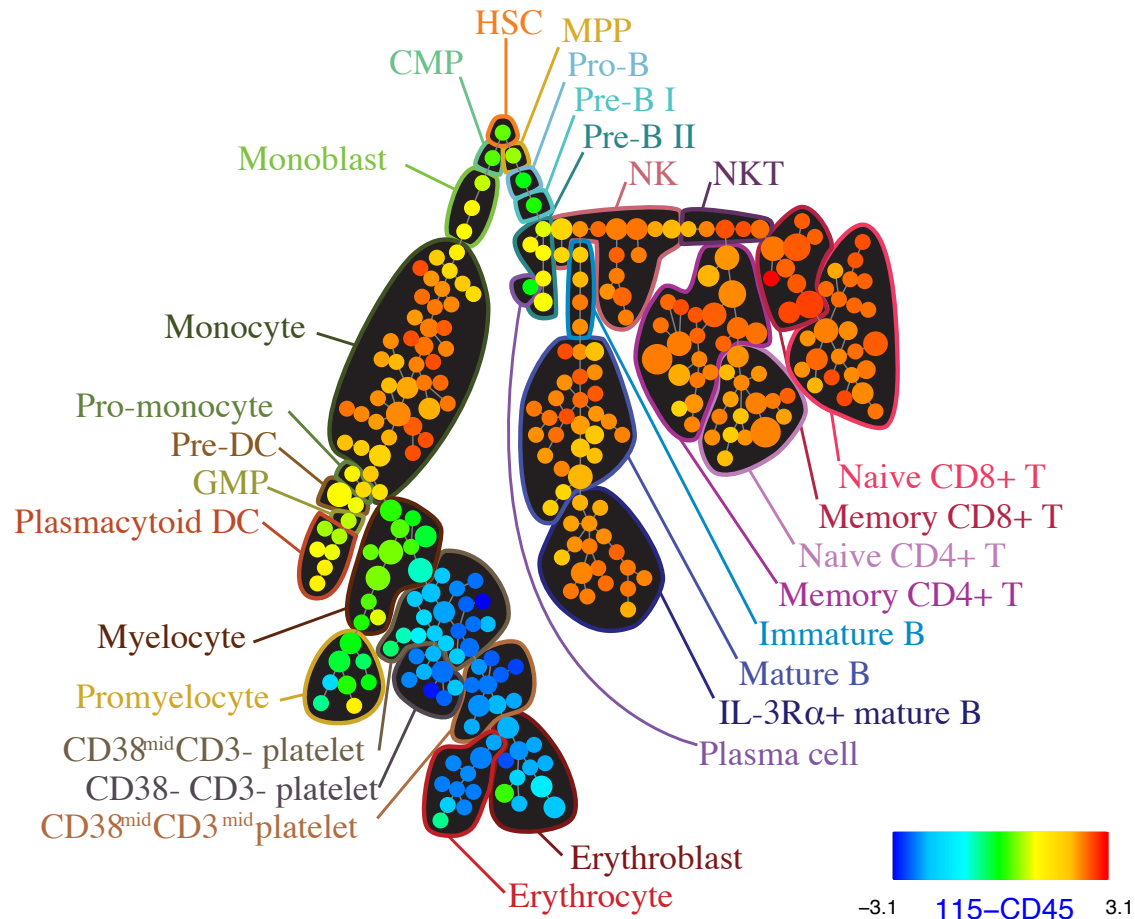


From: Petter Strandmark, Kernel PCA Wikipedia

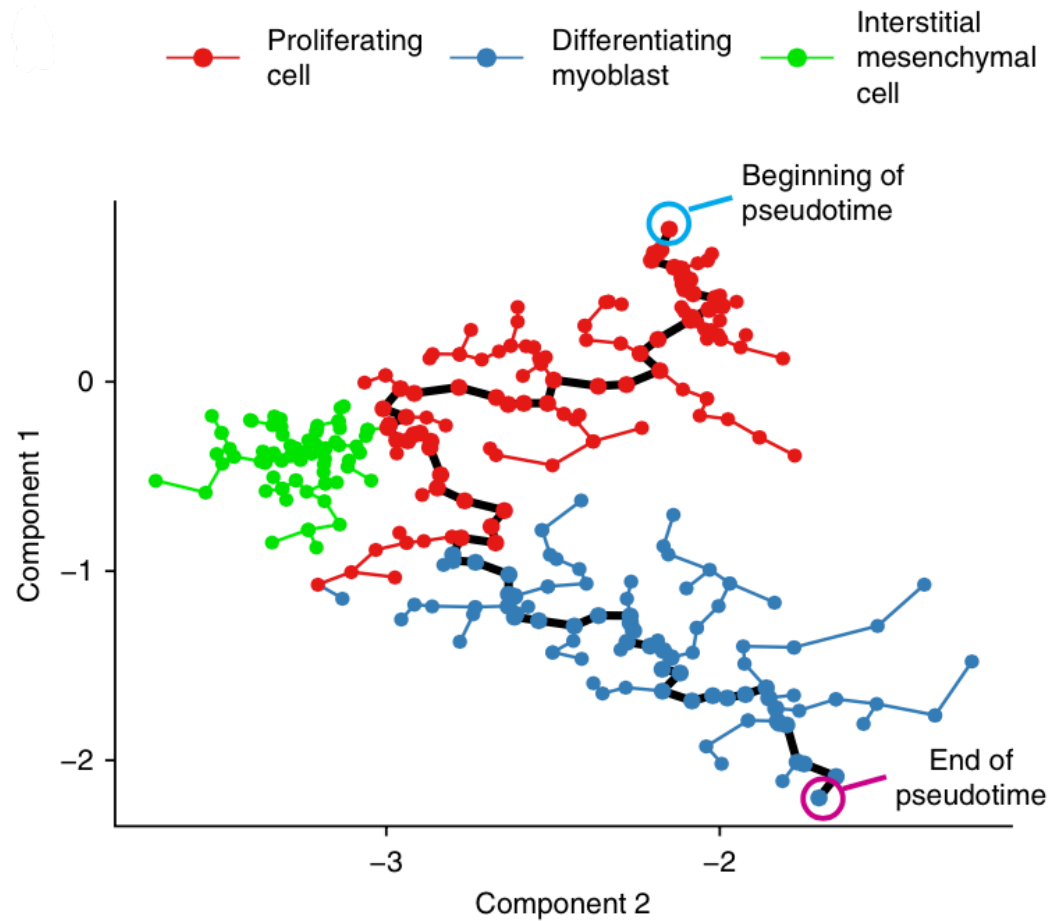
Motivating example 2: Faces



Motivating example 3: SPADE



Motivating example 4: PCA

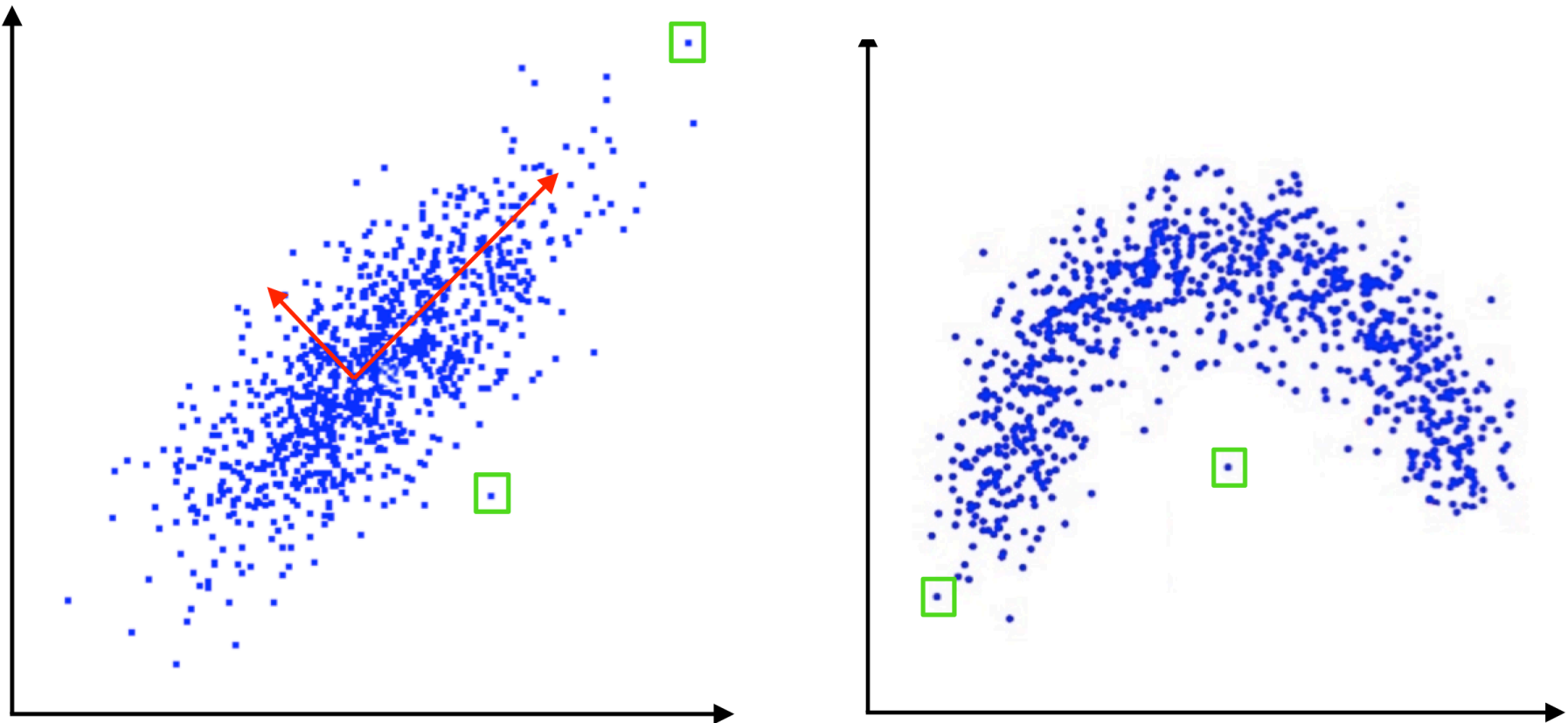


From "monocle": Trapnell et al, NatBio 2014

What is dimensionality reduction?

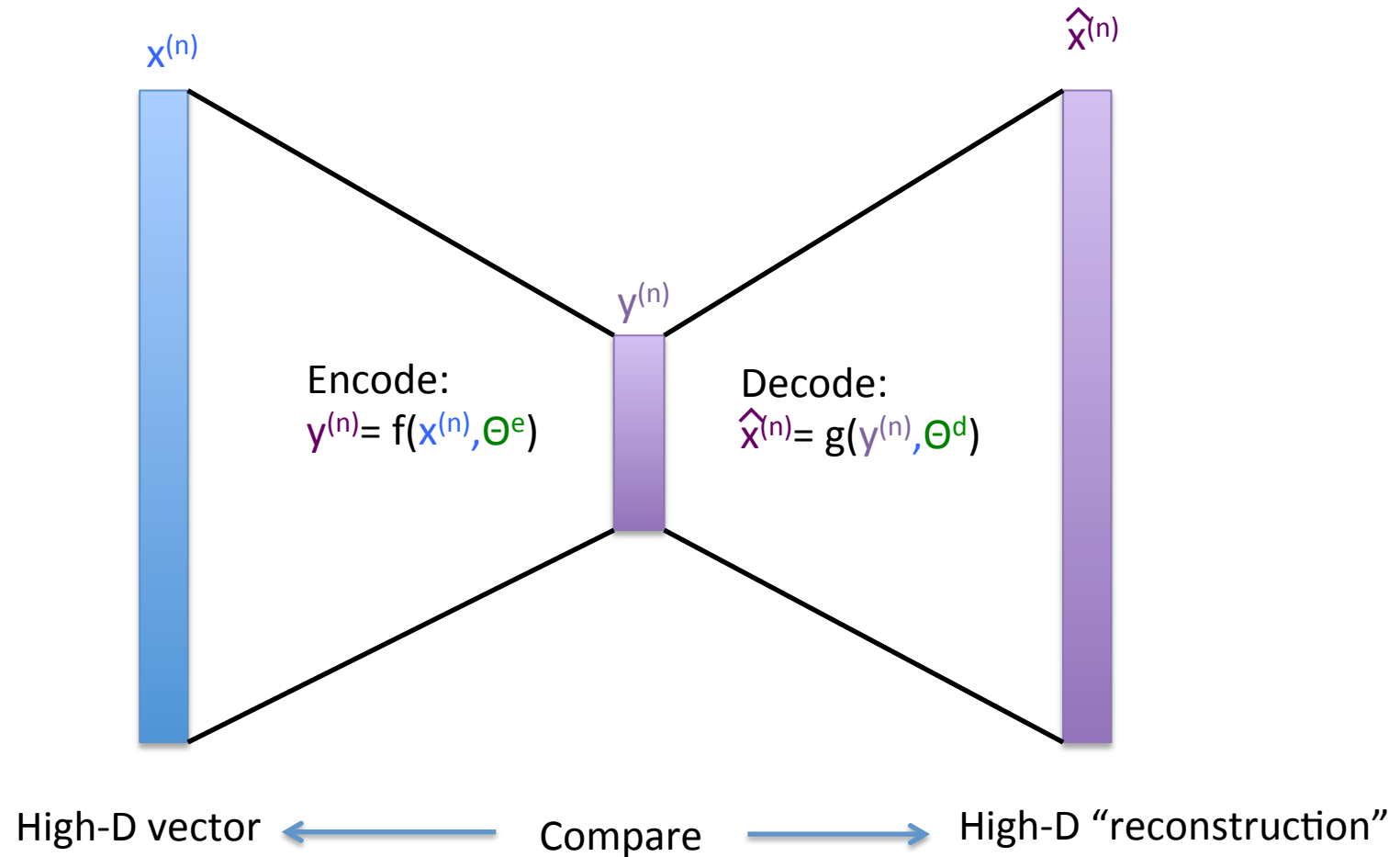
Why do dimensionality reduction?

Why do dimensionality reduction? “outlier detection”



How to do dimensionality reduction?

Strategy #1: “autoencoder”



Four parts of the autoencoder:

1. **Data:** $D = \{x^{(1)}, x^{(2)}, \dots, x^{(N)}\}$
2. **Model** – encoding function $y^{(n)} = f(x^{(n)}, \Theta^e)$ and decoding function $\hat{x}^{(n)} = g(y^{(n)}, \Theta^d)$, each with their own parameters (or shared) parameters
3. **Objective function** –
$$E(\Theta^e, \Theta^d) = \sum_n (x^{(n)} - \hat{x}^{(n)})^T (x^{(n)} - \hat{x}^{(n)})$$
4. **Optimization method** – analytic (PCA) for linear functions, gradient descent otherwise.

Examples: PCA (linear) & deep autoencoders (non-linear)

Objective function example

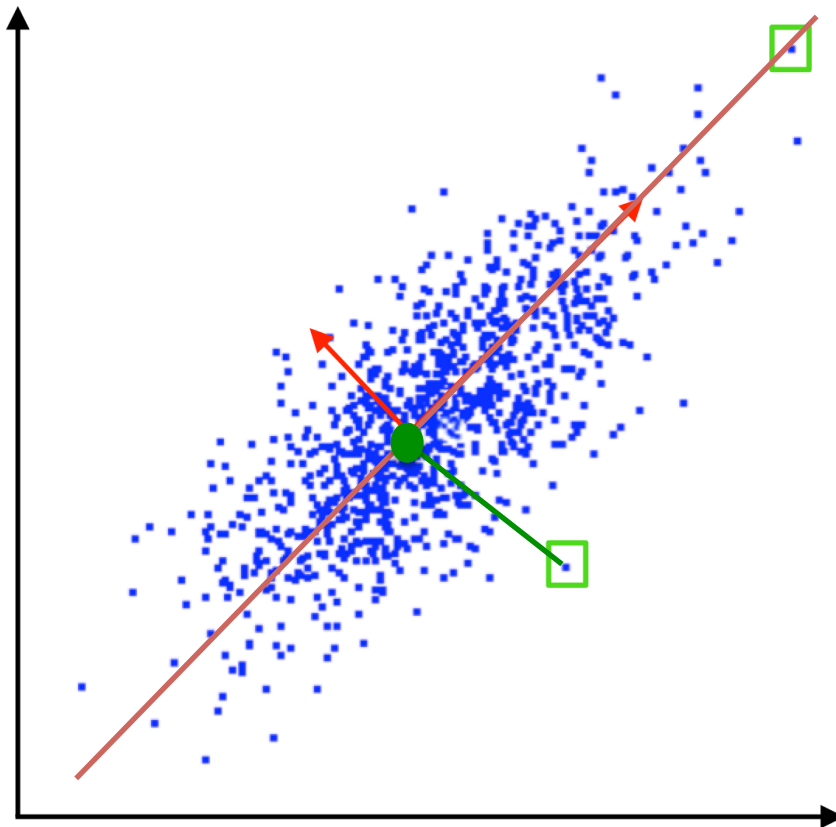
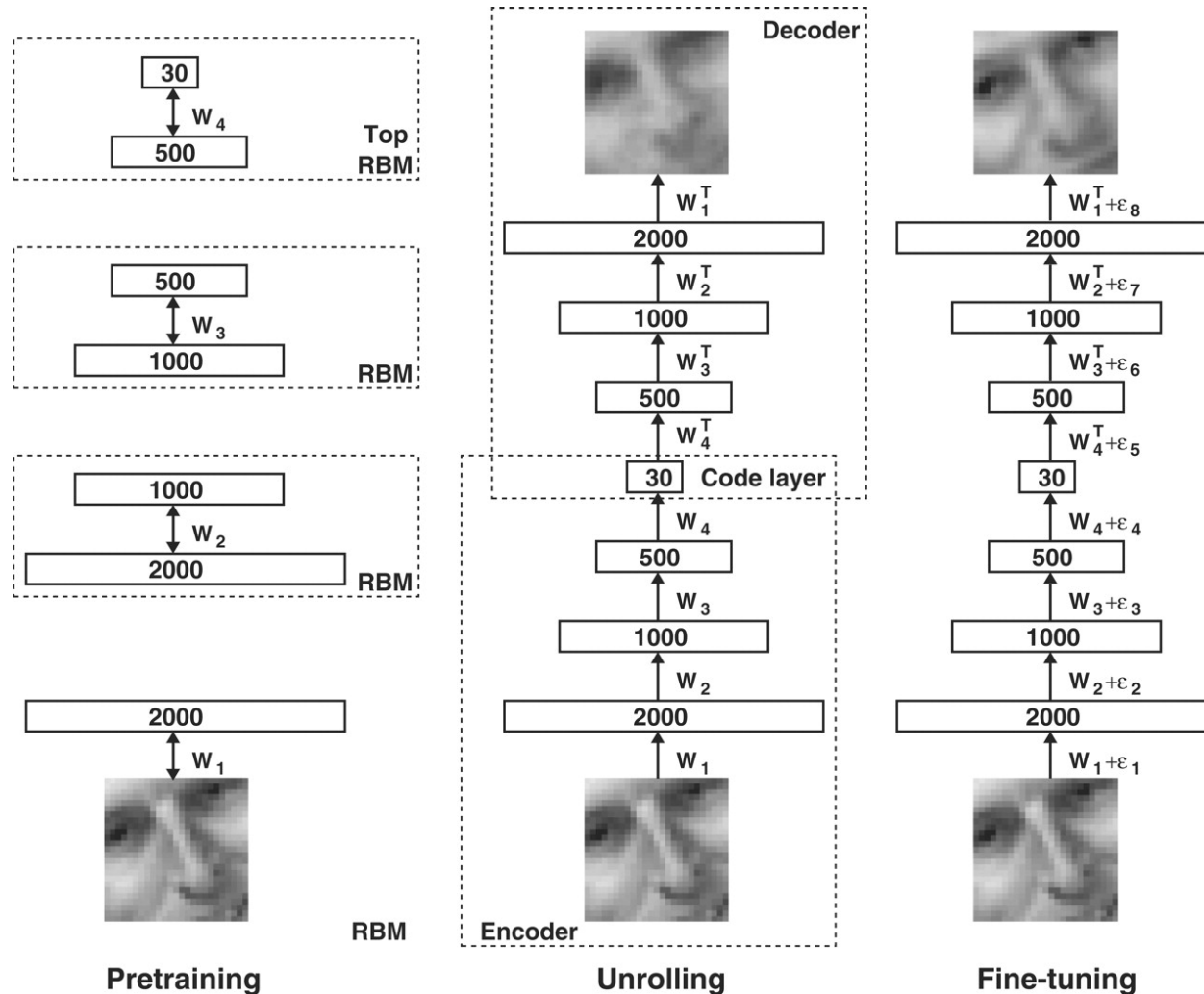


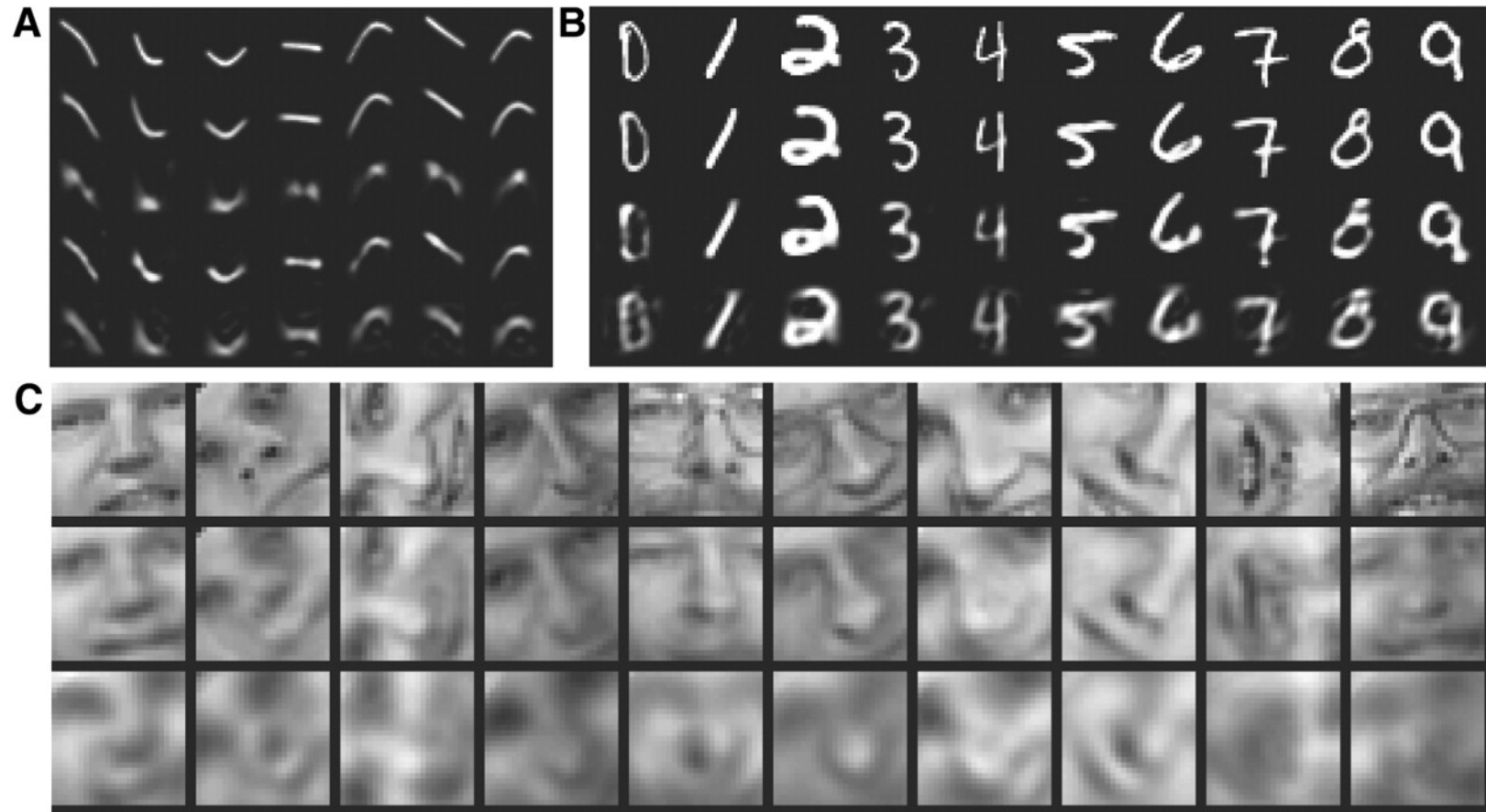
Fig. 1. Pretraining consists of learning a stack of restricted Boltzmann machines (RBMs), each having only one layer of feature detectors.



G. E. Hinton, and R. R. Salakhutdinov Science
2006;313:504-507



Fig. 2. (A) Top to bottom: Random samples of curves from the test data set; reconstructions produced by the six-dimensional deep autoencoder; reconstructions by “logistic PCA” (8) using six components; reconstructions by logistic PCA and standard PCA using 18 components.

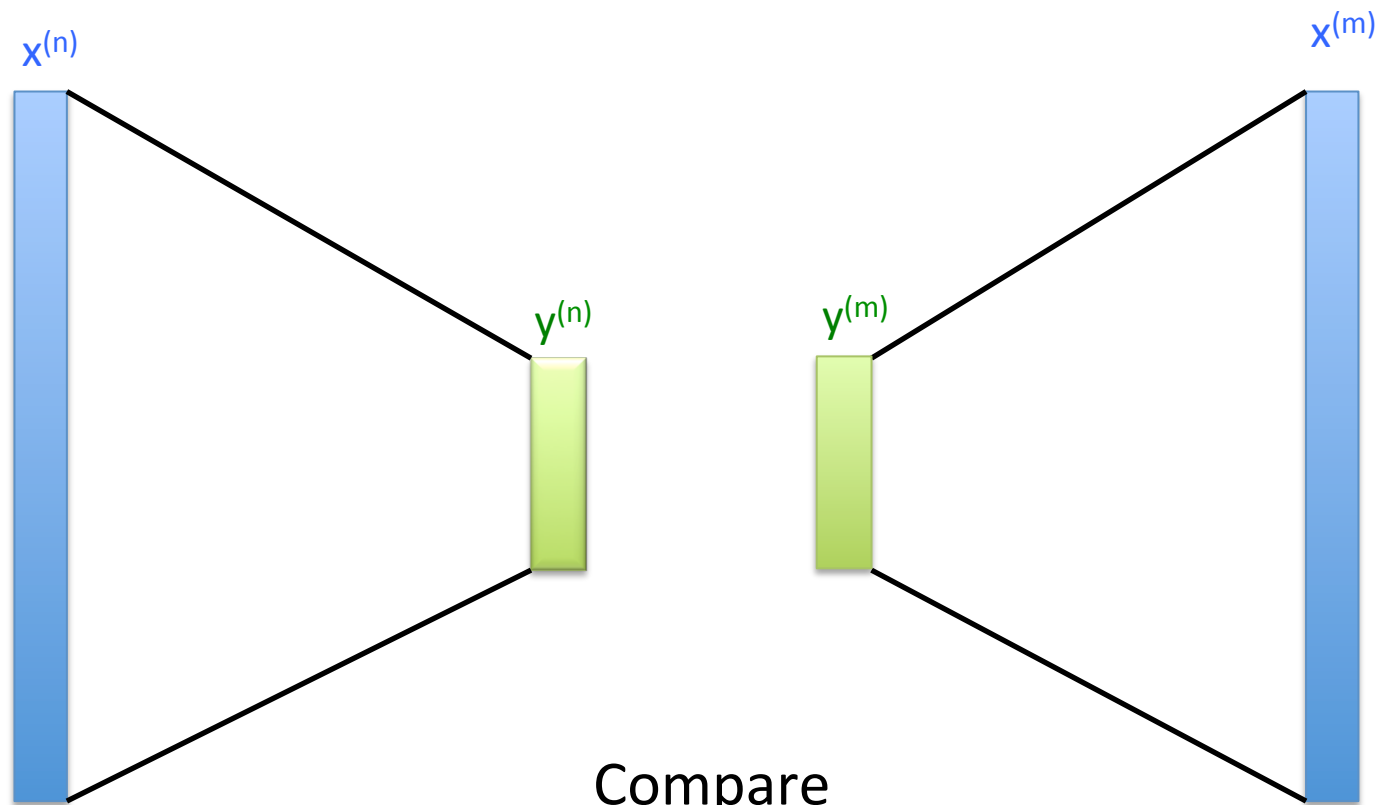


G. E. Hinton, and R. R. Salakhutdinov *Science*
2006;313:504-507



How to do dimensionality reduction?

Strategy #2: “recover distances”
e.g. Multidimensional scaling (MDS)



Compare

$$d(\mathbf{x}^{(n)}, \mathbf{x}^{(m)}) = (\mathbf{x}^{(n)} - \mathbf{x}^{(m)})^T (\mathbf{x}^{(n)} - \mathbf{x}^{(m)})$$

$$\text{to } d(\mathbf{y}^{(n)}, \mathbf{y}^{(m)})$$

Four parts of MDS & friends:

1. **Data:** $D = \{x^{(1)}, x^{(2)}, \dots, x^{(N)}\}$
2. **Model** – parameters are low-d coordinates of each point:

$$\Theta = \{y^{(1)}, y^{(2)}, \dots, y^{(N)}\}$$

1. **Objective function** –

$$E(\Theta) = \sum_{n,m} [d(x^{(n)}, x^{(m)}) - d(y^{(n)}, y^{(m)})]^2$$

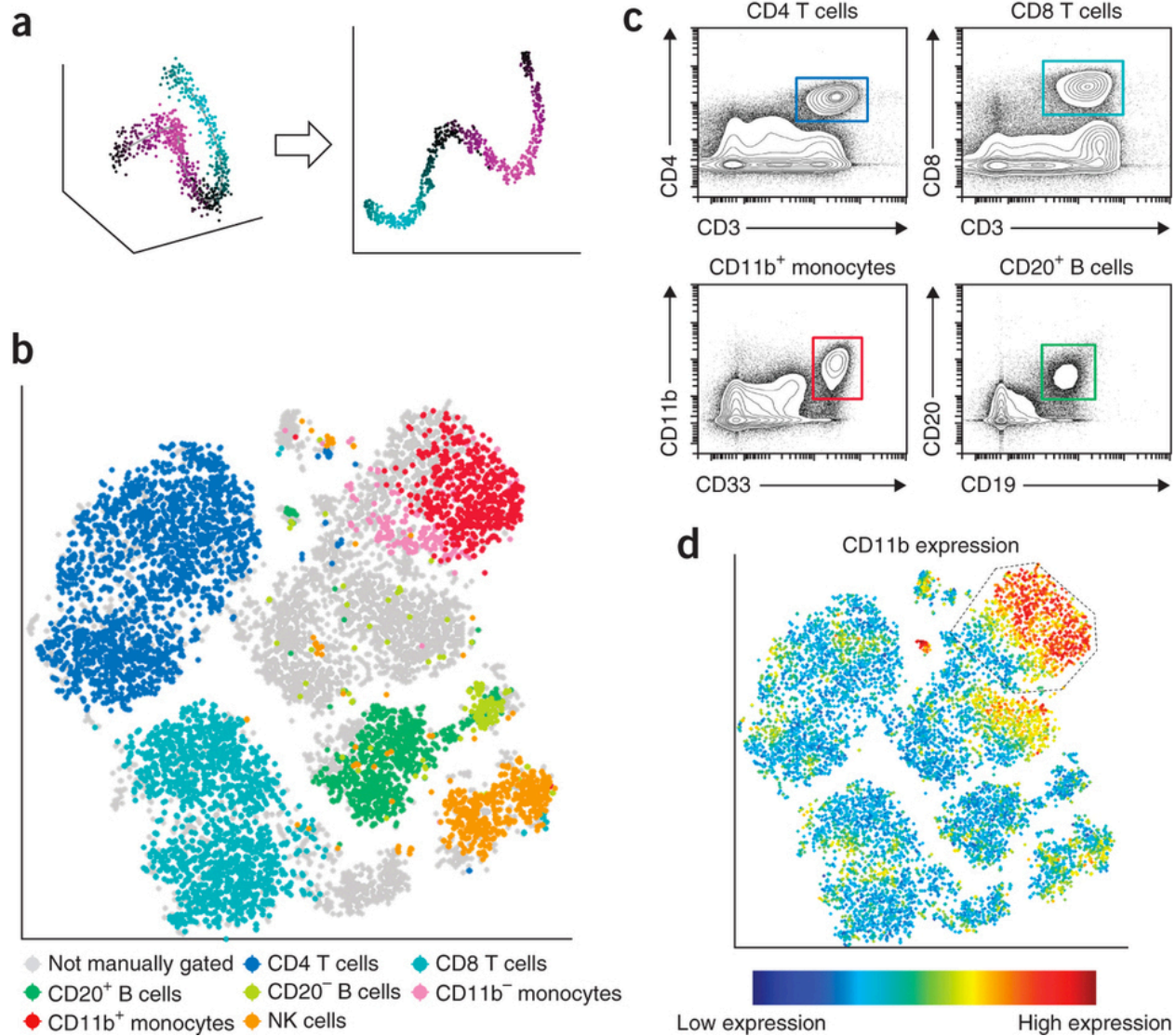
2. **Optimization method** – gradient descent usually.

Examples: MDS, ISOMap, Local linear embedding, SNE, t-SNE

Important variations

1. **ISOMap (Langford, Science 1999)**: use “network distance” rather than Euclidean
2. **SNE (Hinton)**: use “similarity” rather than distance, slightly different objective function
3. **t-SNE (Hinton) / viSNE (Dana Pe'er)**: ignore long distances, use SNE-style objective function
4. **LLE (Roweis, Science 1999)**: use local linear models, ignore long distances

viSNE example



Amir et al,
NatBio 2013

Examples

