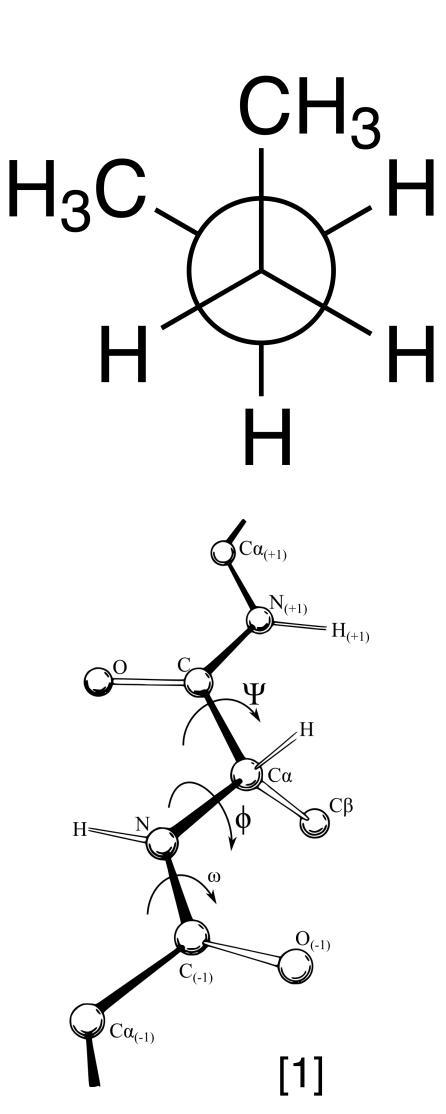
3.2.2 Dimensionality reduction

- This module will consist of a mini-lecture and exercises on dimensionality reduction
- At the end of this module, you should be able to answer the following questions:
 - What are the benefits of dimensionality reduction?
 - In principal components analysis, what are the meanings of the eigenvectors and eigenvalues?
 - What does it mean to project a configuration onto an eigenvector?

Dimensionality reduction with principal component analysis

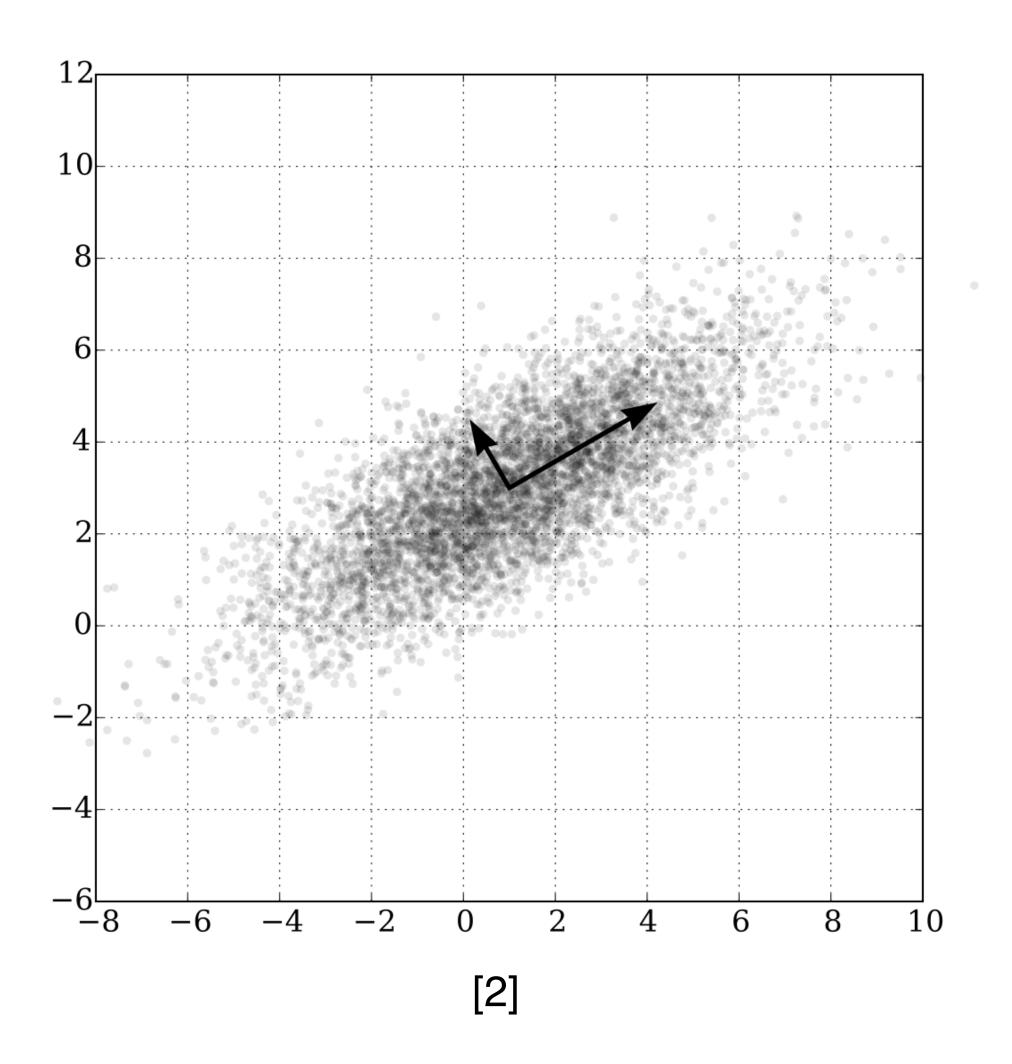
Dimensionality reduction

- Biomolecular simulations have 3N dimensions, but in practice, functionally relevant macromolecular motion can be described with a lot fewer.
- Cartesian coordinates (x, y, and z) of each atom are the most common description but molecular systems can be described by other coordinate systems
- You have probably already seen
 - the free energy of butane as a function of the dihedral angle
 - the Ramachandran diagram, where protein backbones are described by the φ and ψ angles



Principal component analysis (PCA)

- An automated way to do dimensionality reduction
- A linear transformation of coordinates in decreasing order of variance
 - First principal component has the largest variance
 - Second principal component has second largest variance
 - And so forth
- Dimensions can be reduced by keeping the highest-variance dimensions
- See https://en.wikipedia.org/wiki/
 Principal component analysis



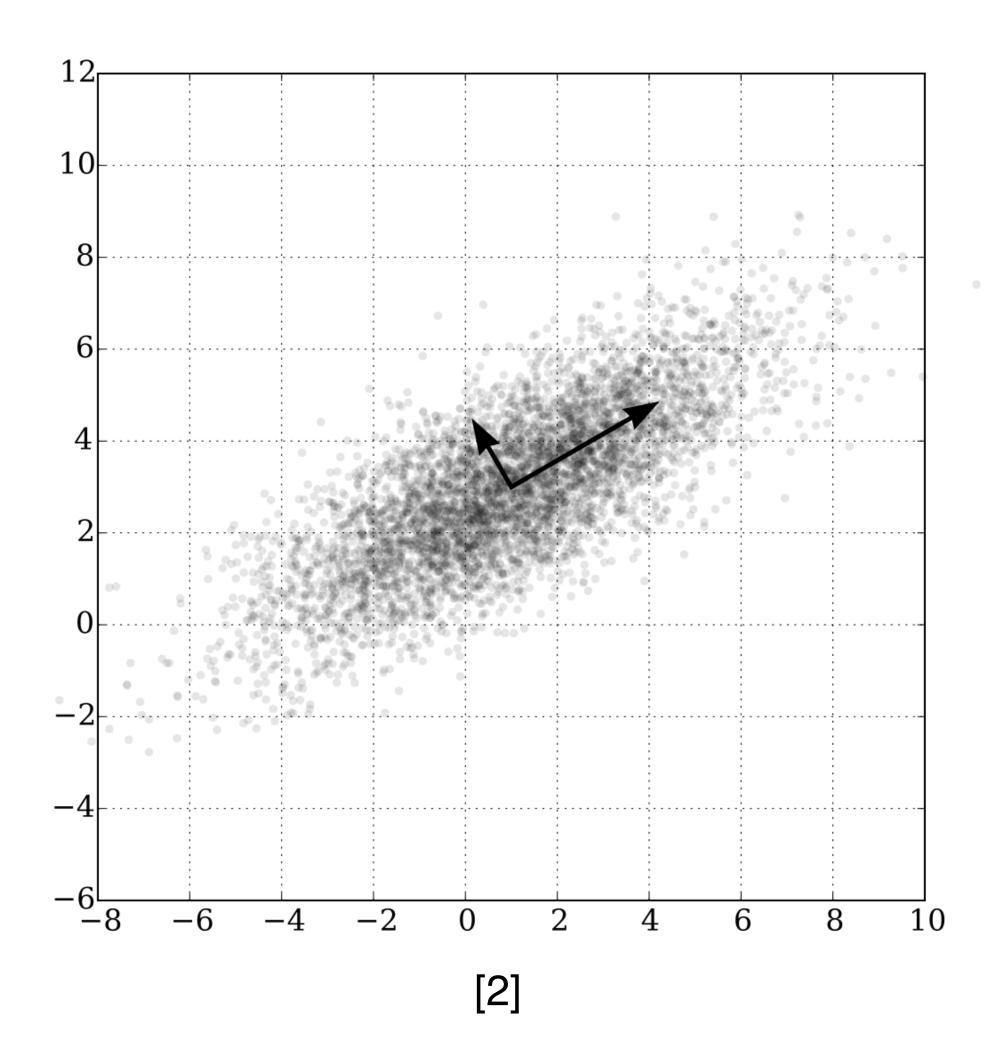
The matrices of PCA

- **CV** = **PV**
- C: covariance matrix
 - C_{kl} is the covariance between dimensions k and l.
 - Usually empirically estimated from data.
- V: matrix of column eigenvectors
 - V_{kI} is the
 - importance of the original coordinate k
 - in the transformed coordinate I.

- known as the principal components
- the columns are orthonormal vectors
- P: diagonal matrix of eigenvalues
 - $P_{kl} = \lambda_{kl}$ if k = 1
 - $P_{kl} = 0$ otherwise
 - variances in transformed coordinate system
 - scaling of the eigenvectors
- All three matrices have the same size

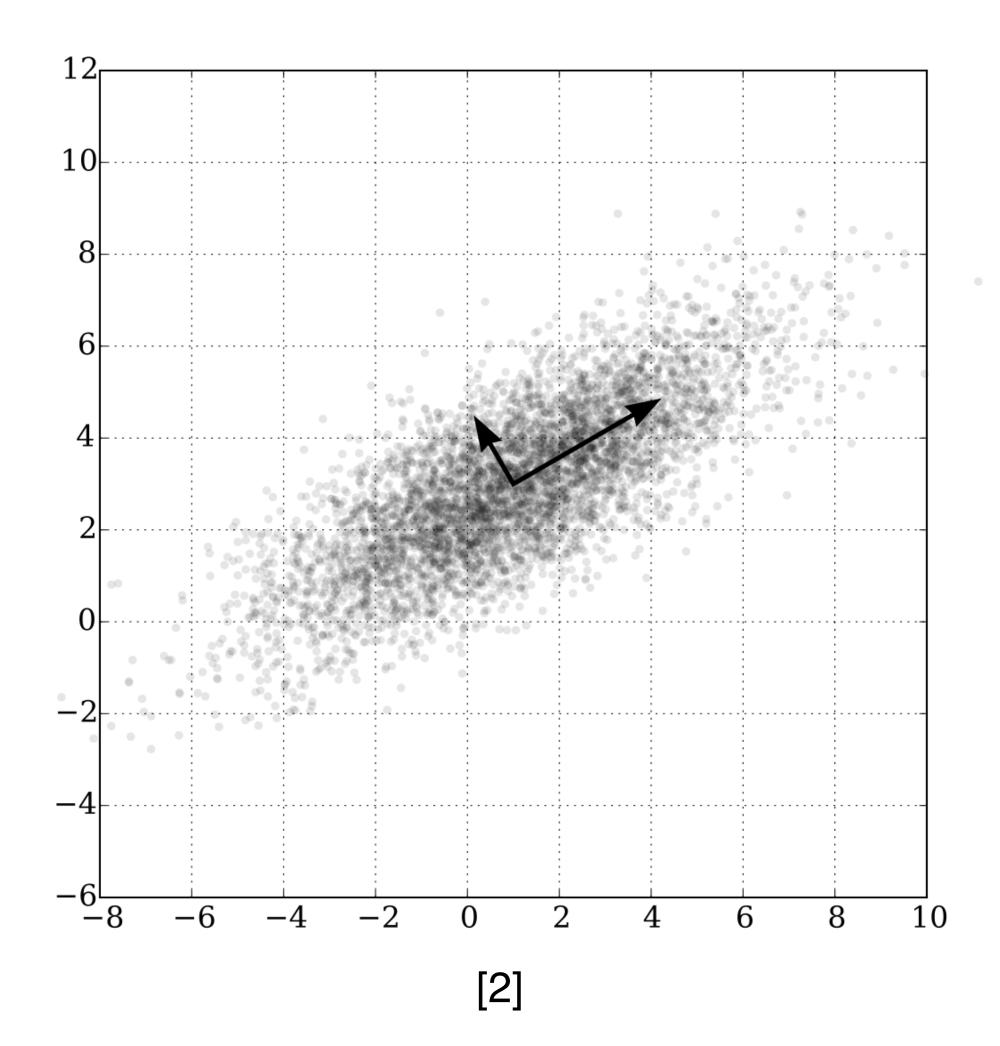
Question

- Approximately, what is the first principal component/eigenvector?
- The vector is ~ (3.5, 2). The magnitude is ~
 4. The normalized eigenvector is ~ (0.875, 0.5).



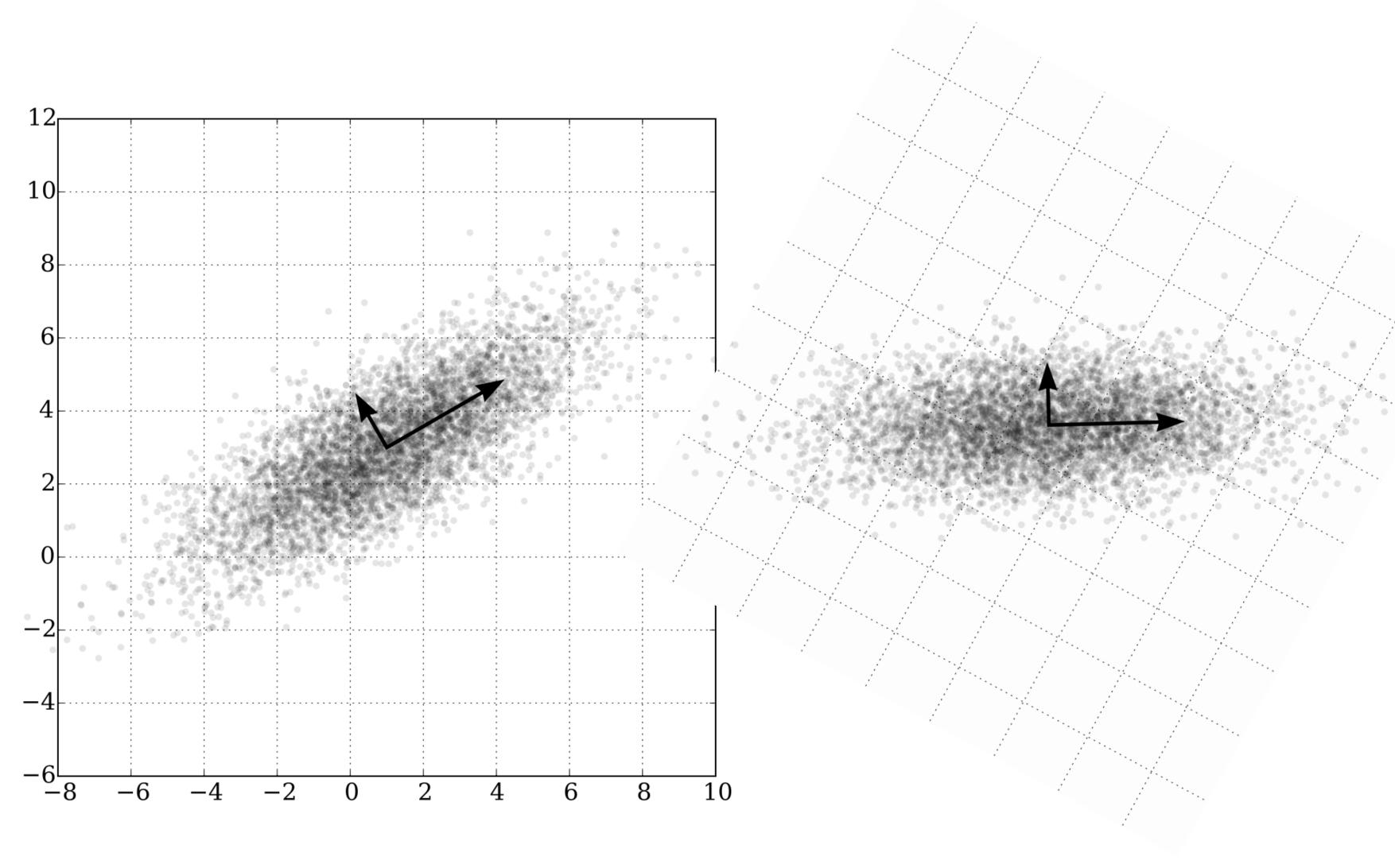
Question

- How do the first and second eigenvalue compare?
- The first eigenvalue is approximately three times larger than the second



Question

What would this
 plot look like if the x
 axis were
 projections onto
 the first PC and the
 y axis were
 projections on the
 second PC?



PCA analysis of molecular simulation

- Load and run the jupyter notebook https://github.com/CCBatIIT/
 modelingworkshop/exercises/mpro/RIKEN_trajectory/analysis/DimerPCA.ipynb, which demonstrates PCA analysis for a simulation of mpro
- What part of the enzyme has the strongest weight in the first three principal components?
- What is the dot product between a principal component and itself? What is the dot product between a principal component and another principal component?
- What fraction of the variance is accounted for in the first three principal components?
- At approximately what point in the trajectory does the largest conformational shift occur?

Time-lagged independent component analysis (TICA)

- Like PCA, based on eigenvector and eigenvalues of a matrix
 - autocorrelation instead of covariance matrix
 - isolates slow motions into coordinates
- Often used in Markov state models (MSMs)
- Slow is often but not always functionally relevant

Review

- What are the benefits of dimensionality reduction?
- In principal components analysis, what are the meanings of the eigenvectors and eigenvalues?
- What does it mean to project a configuration onto an eigenvector?