

Introduction to structural and physicochemical features calculated by propy (2012)

1. AACComposition module

1.1. Amino acid composition

1.2. Dipeptide composition

1.3. Tripeptide composition

2. Autocorrelation module

2.1. Normalized Moreau-Broto autocorrelation descriptors

2.2. Moran autocorrelation descriptors

2.3. Geary autocorrelation Descriptors

3. CTD module

3.1. Composition descriptors

3.2. Transition descriptors

3.3. Distribution descriptors

4. QuasiSequenceOrder module

4.1. Sequence-order-coupling numbers

4.2. Quasi-sequence-order (QSO) descriptors

5. PseudoAAC module

5.1. Type I pseudo amino acid composition descriptors

5.2. Type II pseudo amino acid composition descriptors