## **Text S6. Relation between power of a signal and atomic MSF**

In the current study, atomic position, dihedral angle, and atomic distance trajectory signals are processed, so a more convenient measure than energy may be mean square fluctuations (MSF). When the signal is mean-centered, i.e. deviation of the trajectory from its mean value, MSF is equivalent to the power of the signal:

$$
MSF = \frac{1}{N} \sum_{n=0}^{N-1} |x_n|^2 = \frac{E}{N} = P
$$
\n(S1)

$$
\text{MSF} = P = \frac{E}{N} = \frac{1}{N^2} \sum_{k=0}^{N-1} \left| X_k \right|^2 = \sum_{k=0}^{N-1} \left| \frac{X_k}{N} \right|^2 \tag{S2}
$$

By this way, contribution of  $k^{\text{th}}$  sinuoidal component to the MSF of the trajectory is found to be equal to 2 *N*  $\left|\frac{X_k}{X}\right|^2$ . DFT may be applied to each of the three Cartesian coordinates (*r* = 1,2,3)

of an atom.

$$
X_n^r = \sum_{n=0}^{N-1} x_n^r e^{-\frac{i2\pi kn}{N}}
$$
 (S3)

In this case, MSF of a single atom, using Parseval's relation, can be obtained by

$$
MSF = \sum_{r=1}^{3} \frac{1}{N} \sum_{n=0}^{N-1} \left| x_n^r \right|^2 = \sum_{r=1}^{3} \sum_{k=0}^{N-1} \left| \frac{X_k^r}{N} \right|^2 = \sum_{k=0}^{N-1} \sum_{r=1}^{3} \left| \frac{X_k^r}{N} \right|^2 \tag{S4}
$$

Here, the contribution of the  $k^{\text{th}}$  sinusoidal component to the MSF of an atom (power at the  $k^{\text{th}}$ 

frequency component) is equal to  $\sum_{r=1}^{3}$ 1 2 *r r k N*  $\frac{X_k^r}{X_k}$ . Generalization of this equation to *R* number of C<sub> $\alpha$ </sub>

atoms corresponding to a specific region in the protein, contribution of the  $k<sup>th</sup>$  sinusoidal

component to the residue-averaged MSF can be computed by  $\frac{1}{R} \sum_{r=1}^{3R}$ *r r k N X R* 3 1  $\frac{1}{2} \sum_{k=1}^{3R} \left| \frac{X_k^r}{\cdot} \right|^2$ .