**Figure S3. LMI matrices describing correlated motions. Average LMI matrices achieved with 5ns time-windows.** Examples with simulations of different lengths (i.e 100 ns (A-C) and one μs(D)) and of different replicates of the same system (i.e. r1 (A), r5 (B) and r6 (C)) for DriFREE simulations. The LMI matrices are overall very similar, showing a robust description of the correlated motions upon averaging over five ns in the target proteins.



D

C

B

A