## Supporting Information PCAViz: An Open-Source Python/JavaScript Toolkit for Visualizing Molecular Dynamics Simulations in the Web Browser

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Table S1. PCAViz accuracy when applied to a LARP1 simulation. Using various cumulative-variance and rounding-precision parameters, we compressed a 100-frame simulation of LARP1 (1,326 non-hydrogen atoms) with the PCAViz Compressor. We then decompressed each simulation to recover the atomic Cartesian coordinates. To judge PCAViz accuracy, we calculated the average RMSD (plus or minus the standard deviation) between each frame of the original trajectory and the corresponding frame of the PCAViz-processed trajectory. We also include the file size of each output JSON file so users can judge the extent of compression. The number of top components that PCAViz retained in order to account for the

user-specified cumulative-variance cutoff is given in parentheses.

Cumulative-Variance Cutoff (Number of Components)	Precision:	Precision:	Precision:
	Tenth	Hundredth	Thousandth
20% (1)	1.55 Å ± 0.18	1.47 Å ± 0.18	1.46 Å ± 0.18
	48K	56K	65K
25% (1)	1.55 Å ± 0.18	1.47 Å ± 0.18	1.46 Å ± 0.18
	48K	56K	65K
30% (2)	1.54 Å ± 0.17	1.38 Å ± 0.14	1.37 Å ± 0.14
	56K	66K	78K
40% (3)	1.52 Å ± 0.17	1.32 Å ± 0.12	1.30 Å ± 0.12
	64K	75K	89K
50% (5)	1.49 Å ± 0.16	1.23 Å ± 0.10	1.20 Å ± 0.10
	81K	94K	111K
60% (9)	1.45 Å ± 0.16	1.10 Å ± 0.080	1.08 Å ± 0.080
	113K	131K	157K
70% (15)	1.42 Å ± 0.16	0.99 Å ± 0.054	0.95 Å ± 0.056
	162K	188K	227K
75% (20)	1.41 Å ± 0.16	0.91 Å ± 0.049	0.87 Å ± 0.052
	203K	235K	285K
80% (27)	1.39 Å ± 0.16	0.82 Å ± 0.040	0.77 Å ± 0.044
	260K	300K	365K
90% (48)	1.36 Å ± 0.16	0.62 Å ± 0.038	0.55 Å ± 0.046
	431K	497K	608K
100% (99)	1.34 Å ± 0.16	0.31 Å ± 0.033	0.032 Å ± 0.0035
	845K	978K	1.2 M

**Table S2. PCAViz accuracy when applied to a TEM-1 simulation.** Using various cumulative-variance and rounding-precision parameters, we compressed a 1,000-frame simulation of TEM-1 β-lactamase (2,030 non-hydrogen atoms) with the PCAViz Compressor. We then decompressed each simulation to recover the atomic Cartesian coordinates. To judge PCAViz accuracy, we calculated the average RMSD (plus or minus the standard deviation) between each frame of the original trajectory and the corresponding frame of the PCAViz-processed trajectory. We also include the file size of each output JSON file so users can judge the extent of compression. The number of top components that PCAViz retained in order to

account for the user-specified cumulative-variance cutoff is given in parentheses.

Cumulative-Variance Cutoff (Number of Components)	Precision:	Precision:	Precision:
	Tenth	Hundredth	Thousandth
20% (2)	0.92 Å ± 0.068	0.85 Å ± 0.056	0.85 Å ± 0.056
	94K	111K	131K
25% (4)	0.92 Å ± 0.068	0.82 Å ± 0.053	0.81 Å ± 0.053
	125K	147K	174K
30% (5)	0.91 Å ± 0.069	0.80 Å ± 0.052	0.79 Å ± 0.052
	141K	165K	196K
40% (10)	0.90 Å ± 0.070	0.75 Å ± 0.047	0.74 Å ± 0.048
	217K	254K	303K
50% (19)	0.89 Å ± 0.070	0.69 Å ± 0.037	0.67 Å ± 0.038
	354K	412K	497K
60% (34)	0.88 Å ± 0.071	0.62 Å ± 0.031	0.60 Å ± 0.031
	581K	676K	818K
70% (61)	0.86 Å ± 0.072	0.55 Å ± 0.023	0.52 Å ± 0.023
	988K	1.2M	1.4M
75% (84)	0.86 Å ± 0.072	0.51 Å ± 0.020	0.48 Å ± 0.019
	1.4M	1.6M	1.9M
80% (115)	0.86 Å ± 0.073	0.47 Å ± 0.017	0.43 Å ± 0.016
	1.8M	2.1M	2.5M
90% (240)	0.85 Å ± 0.073	0.36 Å ± 0.013	0.30 Å ± 0.0089
	3.6M	4.2M	5.2M
100% (1000)	0.84 Å ± 0.074	0.21 Å ± 0.016	0.021 Å ± 0.0017
	15M	17M	21M