Table 2. Numerical results from global fitting of two-state motional-narrowing model to single-molecule probability distributions

Variable[1]	Substrate-free AK	AK + AMP – PNP/AMP
k_{open}	120 ± 40 s^{-1}	160 ± 40 s^{-1}
k_{close}	220 ± 70 s^{-1}	440 ± 110 s^{-1}
a	0.45 ± 0.02	0.42 ± 0.04
b	0.76 ± 0.01	0.73 ± 0.01
$\sigma_{0.15}$	0.24 ± 0.10	0.13 ± 0.02
$\sigma_{0.14}$	0.25 ± 0.03	0.14 ± 0.01
$\sigma_{0.13}$	0.25 ± 0.03	0.18 ± 0.01
$\sigma_{0.12}$	0.26 ± 0.03	0.20 ± 0.01
$\sigma_{0.11}$	0.28 ± 0.03	0.22 ± 0.01
$\sigma_{0.10}$	0.28 ± 0.02	0.23 ± 0.01
$\sigma_{0.09}$	0.27 ± 0.03	0.24 ± 0.01
$\sigma_{0.08}$	0.29 ± 0.04	0.25 ± 0.02
$\sigma_{0.07}$	0.29 ± 0.04	0.26 ± 0.02

[1] ?/Au: Please confirm “Variable” as an appropriate heading or provide another.