Supplementary table. Sedimentation velocity parameters for complexes of C.Esp1396I with 35 bp DNA fragments containing Pesp1396ICR and Pesp1396IM binding sites

| Species | Sa | S _{20,w} ^b | M _r ^c | $M_r^{\ d}$ | f/f ₀ ^e |
|-----------------|------------------|--------------------------------|-----------------------------|-------------|-------------------------------|
| | $(x 10^{-13} s)$ | $(x 10^{-13} s)$ | (kDa) | (kDa) | |
| CR-promoter | 2.69 | 2.81 | 23.7 | 22.2 | 1.95 |
| 4:1 protein:DNA | 3.94 | 4.12 | 60.0 | 60.2 | 1.64 |
| M-promoter | 2.59 | 2.71 | 23.5 | 22.2 | 2.00 |
| 2:1 protein:DNA | 3.29 | 3.39 | 40.3 | 41.2 | 1.71 |

^a Experimental sedimentation coefficient obtained from SEDFIT

^b Corrected sedimentation coefficient

^c Experimental M_r from c(M) plot

^d Theoretical M_r for DNA and DNA-protein complexes

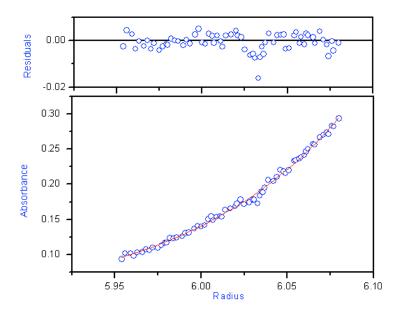
^e Frictional ratio obtained from SEDFIT

Supplementary Figure 1. Sedimentation equilibrium analysis of the dimerisation of C.Esp1396I

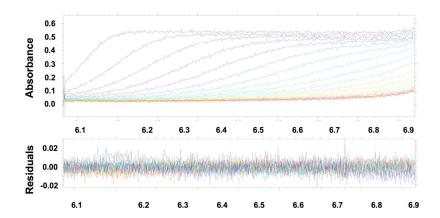
Data were collected at 15, 18 and 21 hours, monitoring absorption at a wavelength of 275 nm. A representative absorption profile is shown after 15 hours, with a rotor speed of 28,000 rpm and protein concentration $64 \mu M$.

Supplementary Figure 2. Sedimentation velocity analysis of C.Esp1396I-DNA complexes.

Data from a single velocity experiment (Pesp1396IM DNA duplex), together with the residuals from the fit. For clarity, every fifth scan is shown.



Supplementary Figure 1.



Supplementary Fig. 2