

Supplementary Information for:

Sequence-dependent DNA hybridisation rates in the absence of secondary structure

Sophie Hertel^{1,^}, Richard E. Spinney^{1,2,^}, Stephanie Y. Xu¹, Thomas E. Ouldridge³, Richard G. Morris^{1,2}, and Lawrence K. Lee^{1,4*}

¹ EMBL Australia Node for Single Molecule Science, School of Medical Sciences, UNSW Sydney, 2052, Australia

² School of Physics, University of New South Wales - Sydney 2052, Australia

³ Department of Bioengineering and Centre for Synthetic Biology, Imperial College London, London, SW7 2AZ, United Kingdom

⁴ ARC Centre of Excellence in Synthetic Biology, University of New South Wales, Sydney, Australia

[^] These authors contributed equally

* To whom correspondence should be addressed. Email: lawrence.lee@unsw.edu.au

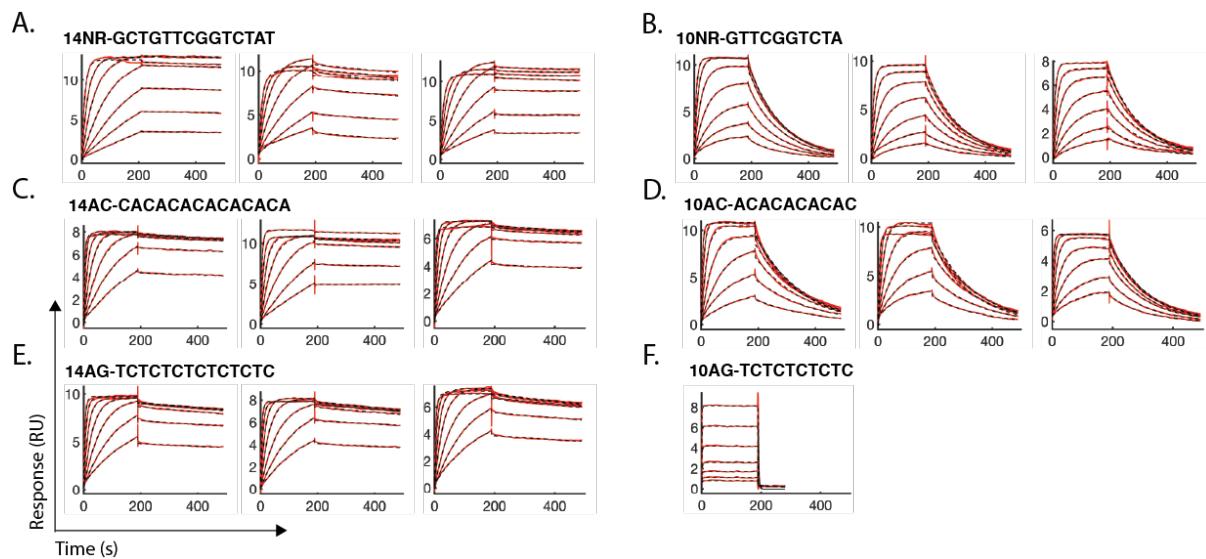


Figure S1. Replicate SPR sensorgrams for repetitive AC and non-repetitive sequences. Raw sensorgrams (red) with mono-exponential fits for association and dissociation (dashed black) are shown for sequences associated with Figure 1 in main text. Curves are serial dilutions from 100 nM to 1.5625 nM (A,B) 50 nM to 0.78125 nM (C-E) and 800 nM to 12.5 nM (F).

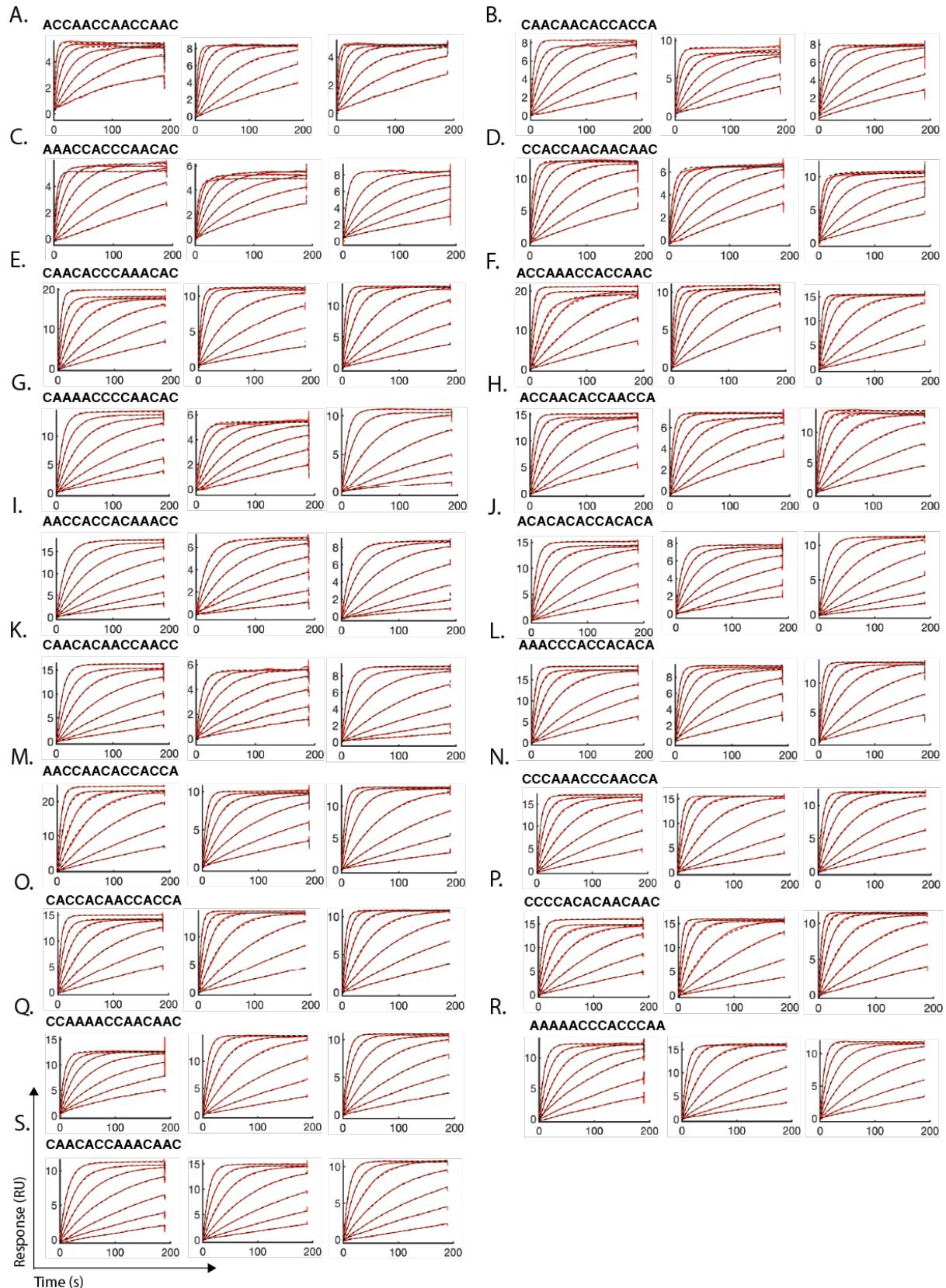


Figure S2. Replicate SPR sensorgrams for all non-repetitive 14 base AC sequences. Raw sensorgrams are shown in red with mono-exponential fits overlaid in dashed black). Concentrations are serial dilutions from 50 nM to 0.78125 nM.

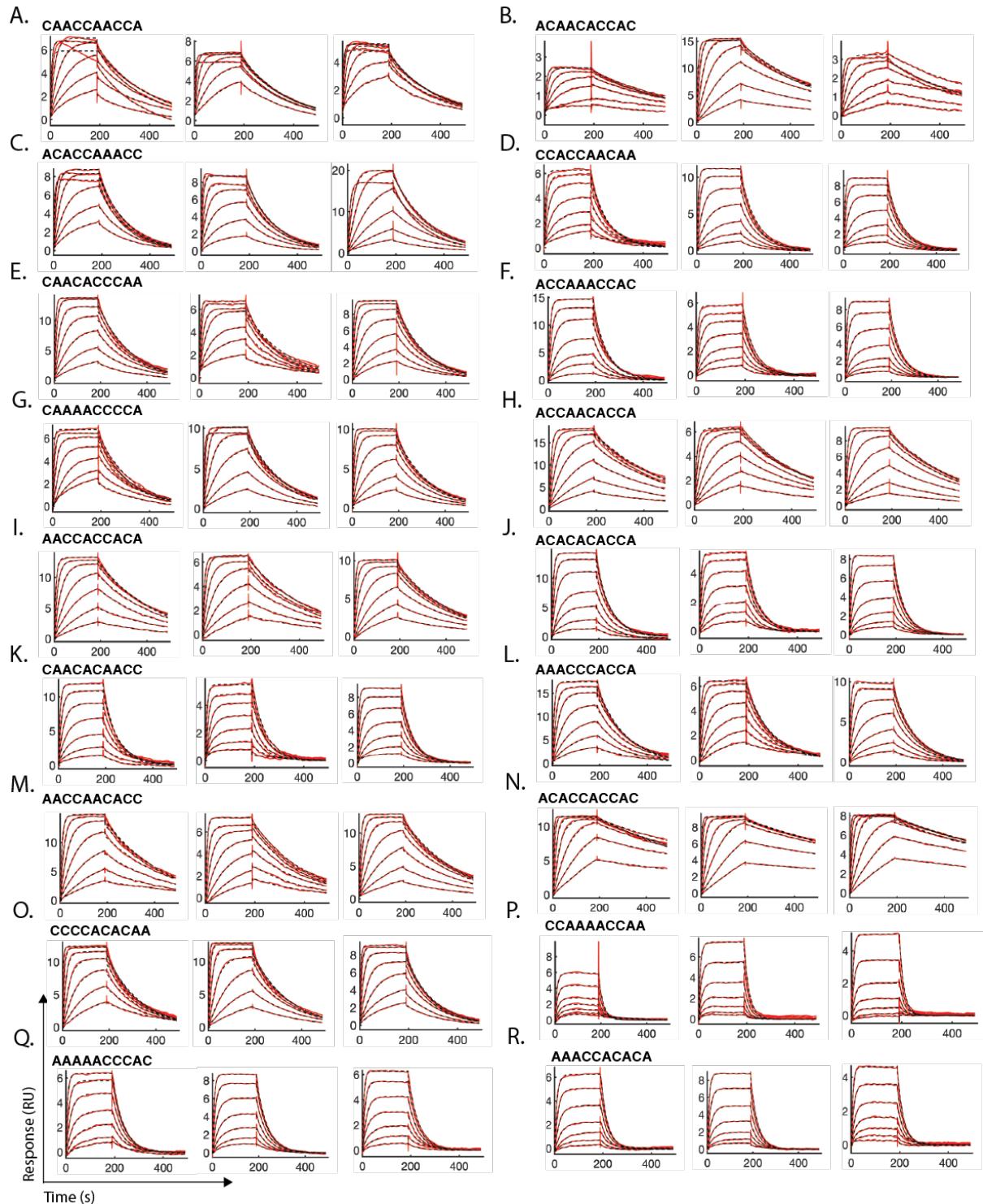


Figure S3. Replicate SPR sensorgrams for all non-repetitive 10 base AC sequences. Raw sensorgrams are shown in red with mono-exponential fits for association and dissociation overlaid in dashed black). Concentrations are serial dilutions from 50 nM to 0.78125 nM.

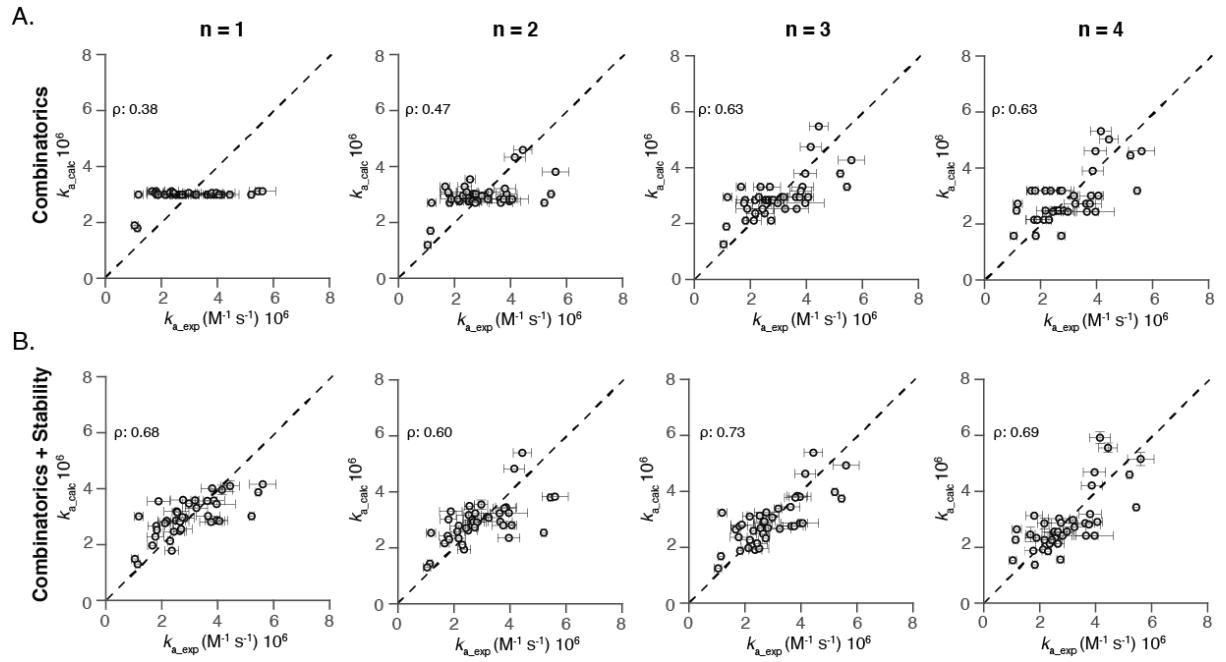


Figure S4. Model vs experimentally measured hybridisation rates without repetitive 10AC and 14AC sequences. (A) shows fits from combinatorics alone and (B) from the full model, which includes stability. Errors are standard deviations from at least three independent measurements.

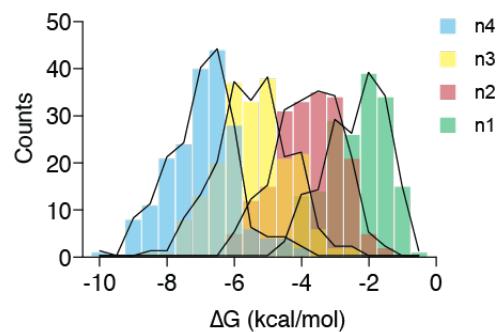


Figure S5. Distribution of binding energies at different nucleation lengths for all unique nucleation states in this study.

Supplementary Table 1: Comparison of equilibrium dissociation constants determined from kinetic rate constants and from steady state measurements for all 10 base DNA sequences in this study.

Number	Name	Sequence 5' > 3'	K _D (nM) kinetics	K _D (nM) steady state
4	10NR	GTTCGGTCTA	7.7 ± 0.2	6.7 ± 0.1
5	10AC	ACACACACAC	1.6 ± 0.1	1.3 ± 0.2
6	10AG	TCTCTCTCTC	NA	212 ± 15
50 % GC content				
20		CAACCAACCA	1.4 ± 0.1	1.4 ± 0.2
21		ACAACACCAC	0.9 ± 0.02	0.6 ± 0.06
22		ACACCAAACCC	3.7 ± 0.3	3.9 ± 0.4
23		CCACCAACAA	5.1 ± 0.9	6.3 ± 1.8
24		CAACACCCAA	7.0 ± 0.4	7.8 ± 3.2
25		ACCAAACCCAC	1.5 ± 0.2	1.4 ± 0.4
26		CAAAACCCCA	6.8 ± 0.6	7.1 ± 2.3
27		ACCAACACCA	3.2 ± 0.5	3.1 ± 1.1
28		AACCACCCACA	2.0 ± 0.2	1.9 ± 0.3
29		ACACACACCA	1.7 ± 0.1	1.5 ± 0.1
30		CAACACAAACC	1.5 ± 0.2	1.4 ± 0.05
31		AAACCCACCA	6.6 ± 0.8	5.8 ± 0.7
32		AACCAACACC	1.6 ± 0.1	1.4 ± 0.5
57 % GC content				
36		ACACCACCCAC	0.2 ± 0.03	NA
37		CCCCACACAA	1.3 ± 0.2	1.4 ± 0.4
42 % GC content				
41		CCAAAACCAA	24.9 ± 6.3	28.1 ± 5.1
42		AAAAACCCAC	8.9 ± 0.6	8.4 ± 1.8
43		AAACCACACA	18.8 ± 3.8	16.0 ± 4.8

Supplementary Table 2: Statistics of the correlation coefficient, ρ , between experimental data and model predictions, for the null distribution sampled by random permutations, and resultant statistics of p -value estimates.

Model	n	Outliers excluded [†]	Data	mean (std-dev)	95% CI	mean (std-dev)	95% CI	mean (std-dev)	95% CI	p-value
Combinatorics	1	N	0.368 (0.0189)	[0.332, 0.406]	-4.71×10^{-5} (0.158)	[-0.329, 0.279]	0.00223 (0.00169)	[0.000250, 0.00654]	[0.000250, 0.00654]	
Comb. + length	1	N	0.371 (0.0187)	[0.335, 0.408]	0.0777 (0.161)	[0.262, 0.362]	0.0218 (0.00819)	[0.00920, 0.04111]	[5.13 $\times 10^{-6}$, 2.50 $\times 10^{-5}$]	
Stability	1	N	0.688 (0.0254)	[0.636, 0.736]	0.0508 (0.165)	$-0.295, 0.345$	$1.02 \times 10^{-5}, 5.53 \times 10^{-6}$	[5.19 $\times 10^{-5}$, 4.39 $\times 10^{-5}$]	[5.19 $\times 10^{-5}$, 4.39 $\times 10^{-5}$]	
Stab. + length	1	N	0.690 (0.0253)	[0.639, 0.738]	0.118 (0.163)	[0.227, 0.406]	1.39 $\times 10^{-5}$ (1.18 $\times 10^{-5}$)			
Combinatorics	2	N	0.495 (0.0364)	[0.421, 0.563]	0.000317 (0.158)	[-0.300, 0.315]	0.00107 (0.00115)	[7.79 $\times 10^{-5}$, 0.00406]	[7.79 $\times 10^{-5}$, 0.00406]	
Comb. + length	2	N	0.504 (0.0384)	[0.426, 0.576]	0.0399 (0.162)	[-0.271, 0.359]	0.00205 (0.00215)	[0.000184, 0.00781]	[0.000184, 0.00781]	
Stability	2	N	0.600 (0.0363)	[0.525, 0.667]	0.0554 (0.160)	$-0.256, 0.365$	0.00192 (0.00305)	[2.58 $\times 10^{-6}$, 0.00999]	[2.58 $\times 10^{-6}$, 0.00999]	
Stab. + length	2	N	0.602 (0.0367)	[0.527, 0.670]	0.0886 (0.162)	[0.230, 0.400]	0.000447 (0.000623)	[2.78 $\times 10^{-5}$, 0.00207]	[2.78 $\times 10^{-5}$, 0.00207]	
Combinatorics	3	N	0.559 (0.0389)	[0.478, 0.630]	0.000282 (0.158)	[-0.292, 0.320]	0.000257 (0.000405)	[1.61 $\times 10^{-5}$, 0.00132]	[1.61 $\times 10^{-5}$, 0.00132]	
Comb. + length	3	N	0.569 (0.0414)	[0.484, 0.646]	0.0278 (0.162)	[-0.273, 0.352]	0.000391 (0.000666)	[1.31 $\times 10^{-5}$, 0.00215]	[1.31 $\times 10^{-5}$, 0.00215]	
Stability	3	N	0.635 (0.0366)	[0.578, 0.721]	0.0158 (0.159)	$-0.279, 0.355$	$2.85 \times 10^{-5}, 3.87 \times 10^{-5}$	[5.96 $\times 10^{-5}$, 0.00105]	[5.96 $\times 10^{-5}$, 0.00105]	
Stab. + length	3	N	0.658 (0.0371)	[0.580, 0.725]	0.0428 (0.162)	[0.262, 0.386]	4.07 $\times 10^{-5}$ (9.42 $\times 10^{-5}$)	[5.39 $\times 10^{-6}$, 0.000263]	[5.39 $\times 10^{-6}$, 0.000263]	
Combinatorics	4	N	0.543 (0.0397)	[0.461, 0.616]	0.000473 (0.158)	[-0.292, 0.320]	0.000412 (0.000595)	[2.19 $\times 10^{-5}$, 0.00202]	[2.19 $\times 10^{-5}$, 0.00202]	
Comb. + length	4	N	0.563 (0.0437)	[0.474, 0.645]	0.0242 (0.161)	[-0.275, 0.349]	0.000441 (0.000764)	[8.55 $\times 10^{-6}$, 0.00249]	[8.55 $\times 10^{-6}$, 0.00249]	
Stability	4	N	0.617 (0.0397)	[0.534, 0.689]	0.0454 (0.161)	$-0.263, 0.362$	0.000134 (0.000268)	[7.97 $\times 10^{-6}$, 0.000776]	[7.97 $\times 10^{-6}$, 0.000776]	
Stab. + length	4	N	0.623 (0.0419)	[0.536, 0.700]	0.0810 (0.160)	[0.230, 0.392]	0.000210 (0.000450)	[1.22 $\times 10^{-5}$, 0.00122]	[1.22 $\times 10^{-5}$, 0.00122]	
Combinatorics	1	Y	0.376 (0.0192)	[0.339, 0.414]	0.000210 (0.162)	[-0.342, 0.281]	0.00204 (0.00152)	[0.000229, 0.000601]	[0.000229, 0.000601]	
Comb. + length	1	Y	0.378 (0.0190)	[0.341, 0.415]	0.0801 (0.165)	$-0.272, 0.388$	0.0219 (0.00812)	[0.000772, 0.0409]	[0.000772, 0.0409]	
Stability	1	Y	0.675 (0.0253)	[0.624, 0.723]	0.0529 (0.170)	$-0.306, 0.352$	$1.37 \times 10^{-5}, 9.96 \times 10^{-6}$	[5.25 $\times 10^{-6}$, 3.64 $\times 10^{-5}$]	[5.25 $\times 10^{-6}$, 3.64 $\times 10^{-5}$]	
Stab. + length	1	Y	0.679 (0.0250)	[0.629, 0.727]	0.122 (0.167)	[0.237, 0.416]	3.29 $\times 10^{-5}$ (2.72 $\times 10^{-5}$)	[7.40 $\times 10^{-6}$, 8.98 $\times 10^{-5}$]	[7.40 $\times 10^{-6}$, 8.98 $\times 10^{-5}$]	
Combinatorics	2	Y	0.466 (0.0330)	[0.400, 0.529]	0.000394 (0.162)	[-0.315, 0.318]	0.00186 (0.00157)	[0.000232, 0.000602]	[0.000232, 0.000602]	
Comb. + length	2	Y	0.473 (0.0326)	[0.408, 0.535]	0.0502 (0.167)	$-0.276, 0.374$	0.00495 (0.00367)	[0.000926, 0.0146]	[0.000926, 0.0146]	
Stability	2	Y	0.602 (0.0325)	[0.536, 0.663]	0.0606 (0.163)	$-0.258, 0.377$	0.00235 (0.00289)	[2.69 $\times 10^{-5}$, 0.000991]	[2.69 $\times 10^{-5}$, 0.000991]	
Stab. + length	2	Y	0.604 (0.0324)	[0.538, 0.665]	0.103 (0.166)	[0.223, 0.421]	0.000609 (0.000701)	[5.19 $\times 10^{-5}$, 0.00251]	[5.19 $\times 10^{-5}$, 0.00251]	
Combinatorics	3	Y	0.631 (0.0336)	[0.563, 0.694]	$-5.16 \times 10^{-5}, 0.162$	[-0.306, 0.325]	$4.44 \times 10^{-5}, 6.24 \times 10^{-5}$	[6.14 $\times 10^{-6}$, 0.000210]	[6.14 $\times 10^{-6}$, 0.000210]	
Comb. + length	3	Y	0.643 (0.0319)	[0.578, 0.703]	0.0409 (0.167)	$-0.276, 0.373$	$9.01 \times 10^{-5}, 0.000107$	[1.03 $\times 10^{-5}$, 0.000359]	[1.03 $\times 10^{-5}$, 0.000359]	
Stability	3	Y	0.731 (0.300)	[0.669, 0.786]	0.0197 (0.167)	$-0.297, 0.350$	$1.09 \times 10^{-5}, 6.19 \times 10^{-6}$	[5.15 $\times 10^{-6}$, 2.68 $\times 10^{-5}$]	[5.15 $\times 10^{-6}$, 2.68 $\times 10^{-5}$]	
Stab. + length	3	Y	0.734 (0.290)	[0.674, 0.788]	0.0581 (0.171)	$-0.268, 0.397$	$1.28 \times 10^{-5}, 1.15 \times 10^{-5}$	[5.19 $\times 10^{-6}$, 3.61 $\times 10^{-5}$]	[5.19 $\times 10^{-6}$, 3.61 $\times 10^{-5}$]	
Combinatorics	4	Y	0.624 (0.0355)	[0.551, 0.690]	$-7.38 \times 10^{-5}, 0.163$	[-0.309, 0.323]	$5.15 \times 10^{-5}, 7.59 \times 10^{-5}$	[6.07 $\times 10^{-6}$, 0.000238]	[6.07 $\times 10^{-6}$, 0.000238]	
Comb. + length	4	Y	0.666 (0.0305)	[0.603, 0.722]	0.0367 (0.166)	$-0.277, 0.366$	$4.46 \times 10^{-5}, 5.56 \times 10^{-5}$	[5.36 $\times 10^{-6}$, 0.000198]	[5.36 $\times 10^{-6}$, 0.000198]	
Stability	4	Y	0.692 (0.0389)	[0.609, 0.761]	0.0237 (0.165)	$-0.290, 0.353$	$3.83 \times 10^{-5}, 5.52 \times 10^{-5}$	[5.89 $\times 10^{-6}$, 0.000154]	[5.89 $\times 10^{-6}$, 0.000154]	
Stab. + length	4	Y	0.715 (0.0371)	[0.636, 0.780]	0.0677 (0.166)	$-0.250, 0.396$	$2.87 \times 10^{-5}, 4.11 \times 10^{-5}$	[6.57 $\times 10^{-6}$, 0.000122]	[6.57 $\times 10^{-6}$, 0.000122]	

[†] Yes (Y)/No (N) indicates exclusion/inclusion of sequences ‘ACACACACAC’ and ‘CACACACACACACA’, respectively.

Supplementary Table 3: Statistics of optimised parameters for various applications of the model on experimental data.

Model	n	Outliers excluded [†]	$\kappa (\times 10^6 \text{ s}^{-1})$	mean (std-dev)	95% CI	α	mean (std-dev)	95% CI	γ	mean (std-dev)	95% CI
Combinatorics	1	N	6.12 (0.103)	[5.92, 6.32]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	1	N	5.12 (1.29)	[3.06, 8.07]	-1.92 (0.0997)	[-2.11, -1.72]	-	-	- ∞ [‡] (-)	-	-
Stability	1	N	28.4 (5.35)	[20.6, 41.3]	-2 [‡] (-)	-	-	-	6.56 (0.320)	[5.96, 7.22]	
Stab. + length	1	N	33.2 (9.18)	[18.8, 54.4]	-2.06 (0.105)	[-2.26, -1.85]	-	-	6.55 (0.325)	[5.95, 7.22]	
Combinatorics	2	N	10.3 (0.178)	[10.0, 10.7]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	2	N	6.63 (1.49)	[4.19, 10.0]	-1.80 (0.0936)	[-1.99, -1.62]	-	-	- ∞ [‡] (-)	-	-
Stability	2	N	26.2 (5.38)	[13.0, 35.0]	-2 [‡] (-)	-	-	-	7.50 (0.753)	[5.15, 8.29]	
Stab. + length	2	N	25.1 (11.2)	[7.68, 50.2]	-1.96 (0.117)	[-2.18, -1.73]	-	-	7.40 (0.867)	[5.08, 8.32]	
Combinatorics	3	N	13.6 (0.243)	[13.1, 14.1]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	3	N	8.01 (1.60)	[5.34, 11.6]	-1.76 (0.0876)	[-1.93, -1.59]	-	-	- ∞ [‡] (-)	-	-
Stability	3	N	19.4 (0.653)	[18.1, 20.7]	-2 [‡] (-)	-	-	-	8.17 (0.143)	[7.87, 8.44]	
Stab. + length	3	N	16.3 (3.49)	[10.5, 24.1]	-1.92 (0.0895)	[-2.09, -1.74]	-	-	8.17 (0.146)	[7.86, 8.43]	
Combinatorics	4	N	14.7 (0.271)	[14.2, 15.2]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	4	N	6.73 (1.19)	[4.70, 9.34]	-1.63 (0.0821)	[-1.79, -1.47]	-	-	- ∞ [‡] (-)	-	-
Stability	4	N	46.4 (5.33)	[36.6, 56.7]	-2 [‡] (-)	-	-	-	13.2 (0.499)	[12.7, 13.6]	
Stab. + length	4	N	27.6 (6.77)	[16.1, 42.0]	-1.74 (0.0881)	[-1.91, -1.56]	-	-	13.2 (1.10)	[12.7, 13.8]	
Combinatorics	1	Y	5.99 (0.104)	[5.79, 6.19]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	1	Y	5.57 (1.45)	[3.25, 8.90]	-1.96 (0.103)	[-2.16, -1.76]	-	-	- ∞ [‡] (-)	-	-
Stability	1	Y	23.0 (3.41)	[17.8, 31.1]	-2 [‡] (-)	-	-	-	6.18 (0.283)	[5.65, 6.77]	
Stab. + length	1	Y	30.7 (9.07)	[16.8, 51.9]	-2.11 (0.106)	[-2.31, -1.90]	-	-	6.16 (0.282)	[5.65, 6.75]	
Combinatorics	2	Y	10.6 (0.186)	[10.3, 11.0]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	2	Y	7.79 (1.79)	[4.85, 11.8]	-1.86 (0.0948)	[-2.04, -1.67]	-	-	- ∞ [‡] (-)	-	-
Stability	2	Y	20.4 (2.92)	[15.5, 26.7]	-2 [‡] (-)	-	-	-	6.76 (0.449)	[5.85, 7.57]	
Stab. + length	2	Y	19.6 (6.09)	[5.83, 7.58]	-1.97 (0.0988)	[-2.16, -1.78]	-	-	6.75 (0.456)	[5.83, 7.58]	
Combinatorics	3	Y	15.2 (0.263)	[14.7, 15.7]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	3	Y	9.55 (1.89)	[6.36, 13.8]	-1.79 (0.0852)	[-1.96, -1.62]	-	-	- ∞ [‡] (-)	-	-
Stability	3	Y	20.1 (0.795)	[18.6, 21.8]	-2 [‡] (-)	-	-	-	7.75 (0.211)	[7.32, 8.15]	
Stab. + length	3	Y	16.9 (3.61)	[10.9, 24.9]	-1.91 (0.0872)	[-2.08, -1.74]	-	-	7.74 (0.217)	[7.29, 8.15]	
Combinatorics	4	Y	17.4 (0.303)	[16.8, 18.0]	-2 [‡] (-)	-	-	-	- ∞ [‡] (-)	-	-
Comb. + length	4	Y	6.92 (1.18)	[4.88, 9.51]	-1.57 (0.0767)	[-1.72, -1.42]	-	-	- ∞ [‡] (-)	-	-
Stability	4	Y	22.2 (2.72)	[18.4, 29.3]	-2 [‡] (-)	-	-	-	9.96 (0.772)	[8.32, 11.4]	
Stab. + length	4	Y	10.4 (2.62)	[6.15, 16.2]	-1.62 (0.0826)	[-1.78, -1.46]	-	-	9.95 (1.35)	[7.28, 12.0]	

[†] Yes (Y)/No (N) indicates exclusion/inclusion of sequences ‘ACACACACAC’ and ‘CACACACACACACACA’, respectively.

[‡] These coefficients are fixed and are not allowed to freely vary. In the case of γ , the value should be considered in the appropriate limit such that all complementary binding states possess a probability of hybridisation of 1 and all mis-matched sites possess a probability of 0.

Supplementary Table 4: Point estimates for ρ , R^2 , and model parameters using mean experimental rates as point estimates.

Model	n	Outliers excluded [†]	ρ	R^2	$\kappa (\times 10^6 \text{ s}^{-1})$	α	γ
Combinatorics	1	N	0.381	0.122	6.12	-2 [‡]	- ∞ [‡]
Comb. + length	1	N	0.383	0.124	4.97	-1.92	- ∞ [‡]
Stability	1	N	0.712	0.502	27.5	-2 [‡]	6.55
Stab. + length	1	N	0.712	0.503	31.5	-2.06	6.54
Combinatorics	2	N	0.514	0.249	10.3	-2 [‡]	- ∞ [‡]
Comb. + length	2	N	0.521	0.259	6.46	-1.80	- ∞ [‡]
Stability	2	N	0.622	0.340	26.9	-2 [‡]	7.70
Stab. + length	2	N	0.622	0.340	25.0	-1.97	7.68
Combinatorics	3	N	0.580	0.163	13.6	-2 [‡]	- ∞ [‡]
Comb. + length	3	N	0.589	0.180	7.85	-1.76	- ∞ [‡]
Stability	3	N	0.679	0.309	19.4	-2 [‡]	8.18
Stab. + length	3	N	0.681	0.312	15.9	-1.91	8.18
Combinatorics	4	N	0.564	-0.0878	14.7	-2 [‡]	- ∞ [‡]
Comb. + length	4	N	0.583	-0.0397	6.63	-1.63	- ∞ [‡]
Stability	4	N	0.638	-0.153	46.3	-2 [‡]	13.2
Stab. + length	4	N	0.644	-0.130	27.3	-1.74	13.3
Combinatorics	1	Y	0.389	0.127	59.9	-2 [‡]	- ∞ [‡]
Comb. + length	1	Y	0.390	0.128	5.40	-1.96	- ∞ [‡]
Stability	1	Y	0.699	0.478	22.4	-2 [‡]	6.16
Stab. + length	1	Y	0.701	0.481	29.2	-2.11	6.15
Combinatorics	2	Y	0.484	0.234	10.6	-2 [‡]	- ∞ [‡]
Comb. + length	2	Y	0.489	0.239	7.60	-1.86	- ∞ [‡]
Stability	2	Y	0.623	0.384	20.1	-2 [‡]	6.78
Stab. + length	2	Y	0.622	0.384	18.6	-1.97	6.76
Combinatorics	3	Y	0.655	0.429	15.2	-2 [‡]	- ∞ [‡]
Comb. + length	3	Y	0.665	0.443	9.36	-1.79	- ∞ [‡]
Stability	3	Y	0.758	0.574	20.1	-2 [‡]	7.76
Stab. + length	3	Y	0.759	0.576	16.5	-1.91	7.75
Combinatorics	4	Y	0.647	0.400	17.4	-2 [‡]	- ∞ [‡]
Comb. + length	4	Y	0.689	0.470	68.3	-1.57	- ∞ [‡]
Stability	4	Y	0.717	0.458	21.6	-2 [‡]	9.98
Stab. + length	4	Y	0.738	0.507	9.95	-1.64	9.99

[†] Yes (Y)/No (N) indicates exclusion/inclusion of sequences ‘ACACACACAC’ and ‘CACACACACACACA’, respectively.

[‡] These coefficients are fixed and are not allowed to freely vary. In the case of γ , the value should be considered in the appropriate limit such that all complementary binding states possess a probability of hybridisation of 1 and all mis-matched sites possess a probability of 0.