

Table S2

	FapC	CsgA
M_0	50 μM	5.3 μM
P_0	39 nM	8 nM
n_c	1.31	1.21
k_n	$1.07 \times 10^{-4} \text{ M}^{-n_c+1} \text{ h}^{-1}$	$5.06 \times 10^{-4} \text{ M}^{-n_c+1} \text{ h}^{-1}$
k_+	$1.11 \times 10^7 \text{ M h}^{-1}$ (3100 $\text{M}^{-1} \text{ s}^{-1}$)	$7.47 \times 10^7 \text{ M h}^{-1}$ (21000 $\text{M}^{-1} \text{ s}^{-1}$)
Mean residual error of fit	0.00067	0.00074

Table S2: Kinetic parameters for global fitting of the experimental data for seeded aggregation of FapC and CsgA to a nucleation-elongation model, where M_0 is the concentration of fibril seeds, P_0 is the number concentration of seed fibrils, n_c is the reaction order of the primary nucleation process, k_n is the rate constant for the primary nucleation process and k_+ is the rate constant for the elongation of existing fibrils. The values for the elongation rates are within a factor of 2 of those determined from the initial gradient analysis.