

SI Table 4: Comparison of Rate Constants of folding at 0.54 M GuHCl monitored by intrinsic fluorescence and ANS fluorescence.

| CTPRan | ^a <i>k</i> Intrinsic Fluorescence (s ⁻¹) | ^b <i>k</i> ANS Fluorescence (s ⁻¹) |
|--------|---|---|
| 2 | ^c 450 ± 50 | No observed binding to ANS |
| 3 | 350 ± 40 | No observed binding to ANS |
| 4 | 220 ± 23 | 162 ± 44 |
| 5 | 180 ± 14 | 151 ± 13 |
| 6 | 180 ± 12 | 156 ± 24 |
| 8 | 170 ± 14 | 143 ± 34 |
| 10 | 140 ± 5 | 123 ± 14 |

All rate constants were determined from refolding traces that had a final concentration 0.54 M GuHCl except CTPRa2 (0.73 M). ^aMonitored by intrinsic fluorescence which were fitted to a single exponential equation. ^bMonitored by ANS fluorescence which were fitted to a double exponential equation. The *k* listed is the faster rate with larger amplitude. The second phase was due to photolysis as decreasing protein concentration decreased the first phase but not the second (data not shown). ^cRate constant monitored at 0.73 M GuHCl as 0.54 M was too rapid for the stopped flow instrument. Errors reported are ± 1 standard deviation of the experimental data.