

1 **SUPPLEMENTAY METHODS**
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4 **Sedimentation velocity.** Sedimentation velocity analysis of FtsZ-YFP-mts was performed
5 at 0.5 g/L protein concentration in 50 mM Tris-HCl, 150 mM KCl, 5 mM MgCl₂, pH 7.5
6 buffer. The experiments were carried out at 38,000 rpm and at 20 °C in an XL-I analytical
7 ultracentrifuge (Beckman-Coulter Inc.) equipped with a UV-VIS detection system, an An-
8 50 Ti rotor and 12 mm double-sector centerpieces. Sedimentation profiles were registered
9 every 5 min. The sedimentation coefficient distributions were calculated by least-squares
10 boundary modeling of sedimentation velocity data using the c(s) method as implemented
11 in the SEDFIT program (Schuck, P., 2000). These S-values were corrected to standard
12 conditions (water, 20 °C and infinite dilution) to get the corresponding standard s_{20,w}
13 using the software SEDNTERP (Holde, v.E.v.,1985). The frictional ratio, f/f₀, was
14 calculated for the monomer using this software.

15 **GTPase activity.** FtsZ GTPase activity was determined using the BIOMOL GREEN
16 reagent for phosphate detection (Enzo). In brief FtsZ at 5 μM concentration is measured
17 every 20 seconds after adding 1 mM GTP for a total of 7 time points. After 15 minutes of
18 incubation with Biomol Green, the samples are measured OD_{620nm}. The data is later fit to a
19 standard curve of 40 μM phosphate.
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