## **Description of Additional Supplementary Files:**

**Supplementary Data 1.** Spreadsheet summarizing the frequency of all persistent (>10% of the last 500 ns of simulation) hydrogen bonds formed at important interfaces in the complex. Each sheet corresponds to a different interface, and hydrogen bond frequencies are listed for each system simulated.

**Supplementary Data 2.** Directory containing coordinates for predicted structures and conformers after 1 microsecond of MD simulation, both for the protein and for both protein and membrane.

**Supplementary Movie 1.** Epifluorescence time-lapse imaging of single Halo-FtsB molecule in M9-glucose, corresponding to Figure 1A (top). Scale bar,  $0.5 \mu m$ .

**Supplementary Movie 2.** Epifluorescence time-lapse imaging of single Halo-FtsB molecule in M9-glucose, corresponding to Figure 1A (bottom). Scale bar,  $0.5 \mu m$ .

**Supplementary Movie 3.** One-microsecond simulation of the FtsQLBWI complex.

**Supplementary Movie 4.** One-microsecond simulation of FtsWI embedded in a lipid bilayer. Phosphorus (dark purple) and oxygen (light purple) atoms of the polar head groups are shown.

**Supplementary Movie 5.** One microsecond simulation of the FtsQLBWI complex with bound activator FtsN.