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Supporting information for article:

Structure of the GDP-bound state of the SRP GTPase FlhF

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Figure S1  Plasmid map employed for this study. B. Sequence alignment of *S. putrefaciens* and *Bacillus subtilis* FlhF.
Figure S2  B-factors of the GDP bound NG-FlhF published with this study. B-factors range from 44 (yellow, thin) to 91 (red, thick).