



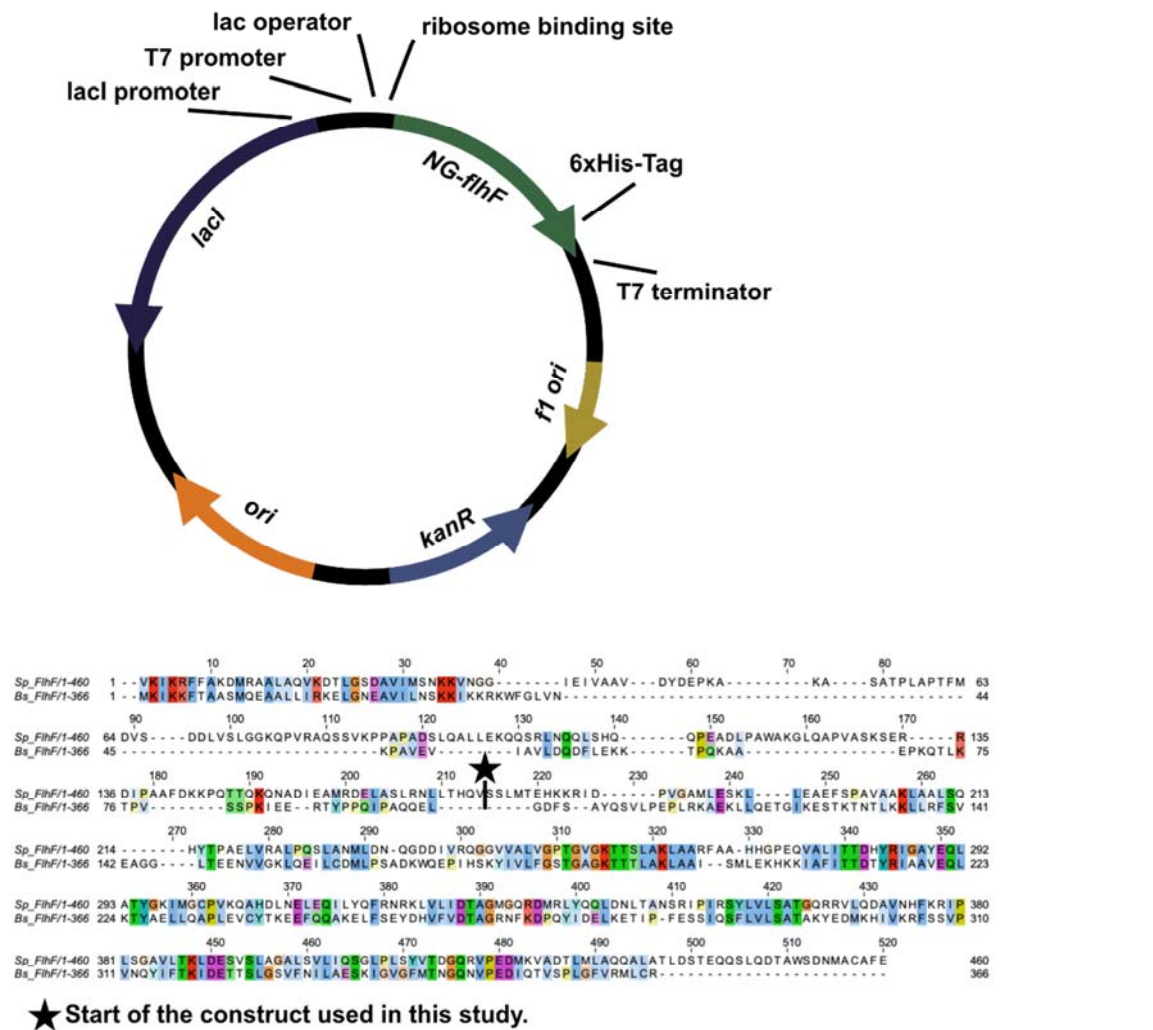
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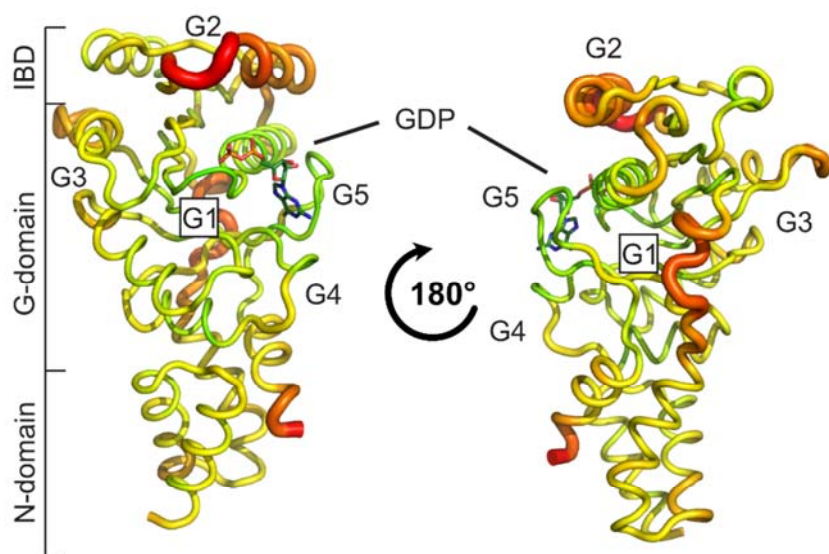
**Supporting information for article:**

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

**Anita Dornes, Christopher-Nils Mais and Gert Bange**



**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).