

Supplemental Information

Biochemical validation of a second class of tetrahydrofolate riboswitches in bacteria

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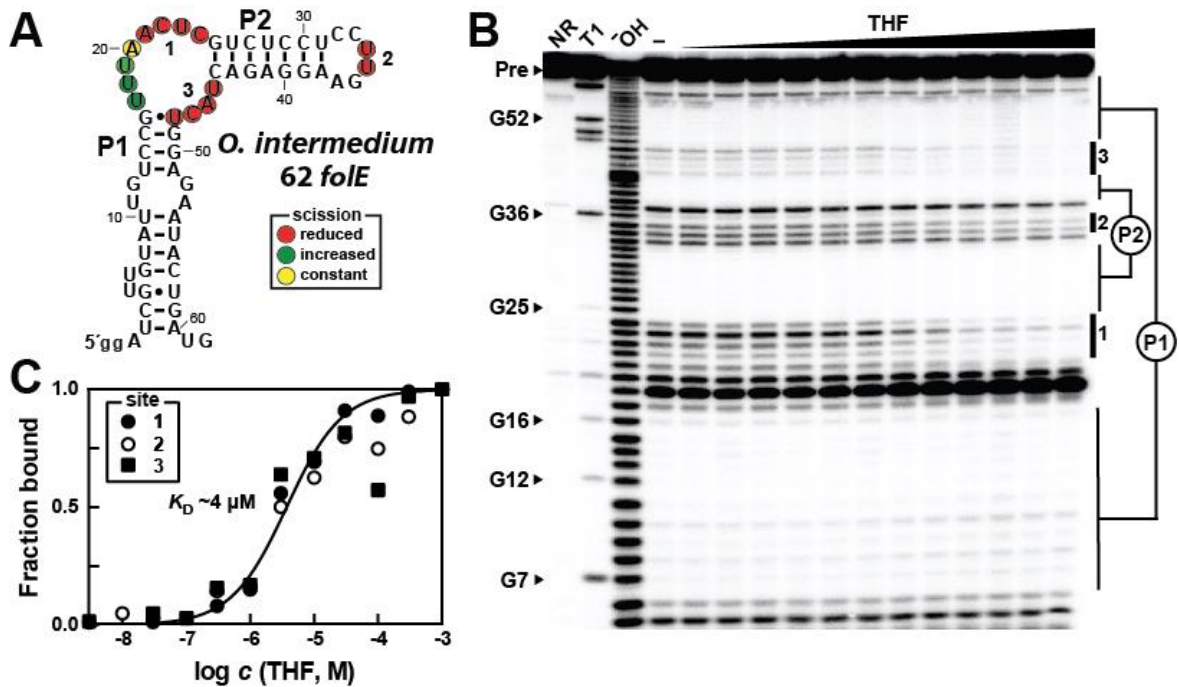


FIGURE S1. THF binding by the 62 *folE* RNA construct derived from *Ochrobactrum intermedium*. (A) Sequence and secondary structure model of the *O. intermedium* 62 *folE* RNA construct. Additional notations are as described in the legend to **Fig. 2A**. (B) PAGE analysis of the *O. intermedium* 62 *folE* RNA subjected to in-line probing assays in the absence (–) or presence of THF concentrations ranging from 10 nM to 1 mM. Additional annotations are as described in the legend to **Fig. 2B**. (C) Plot of fraction of RNA bound to THF versus the logarithm of the molar concentration of THF. The data is derived from B. Additional annotations are as described in the legend to **Fig. 2C**.

Supplemental Table S1. Sequences of synthetic DNAs used in this study.

| Name | Sequence (5' to 3') | Annotation |
|-------|---|--|
| folE1 | CATAAAGGCGTCTCCCGTA GTTCTCTCGAGGAGGAAC GAGTTGAACGGACCACGA CCTTTCCTATAGTGAGTCG TATTA | The template for transcription of <i>M. loti</i> 62 <i>folE</i> RNA |
| folE2 | CATAAAGGCGTCTCCCGTA GTTCTCTCGAGGAGGAAC GAGTTGTACGGACCACGAC CCTTTCCTATAGTGAGTCGT ATTA | The template for transcription of M1 <i>M. loti</i> 62 <i>folE</i> RNA |
| folE3 | CATAAAGGCGTCTCCCGTA GTTCTCTCGAGGAGGAAC GACTTGAACGGACCACGAC CCTTTCCTATAGTGAGTCGT ATTA | The template for transcription of M2 <i>M. loti</i> 62 <i>folE</i> RNA |
| folE4 | CATAAAGGCGTCTCCCGCA GTTCTCTCGAGGAGGAAC GAGTTGAACGGACCACGA CCTTTCCTATAGTGAGTCG TATTA | The template for transcription of M3 <i>M. loti</i> 62 <i>folE</i> RNA |
| folE5 | CATCAGTATTCTCCAGTAG TCTCCTTCAAGGAGGAGAC GAGTTAAACGGACAATAC AACGATCCTATAGTGAGTC GTATTA | The template for transcription of <i>O. intermedium</i> 62 <i>folE</i> RNA |