

Supplemental Material

Table S1. Primers, plasmids and strains used in experiments

| Primers | Sequence |
|-------------------------------|--------------------------------------------------------------------------------------------------|
| <i>BsnifZ</i> -PciI5 | 5'-GCGGACATGTTATATTTAGATAAATAG-3' |
| <i>BsnifZ</i> -BamHI3 | 5'-TAAAGGATCCATGTTCTACCTCATC-3' |
| <i>BsnifZ</i> '-BamHI3 | 5'-CTCGGATCCCTTAATTTTTTGGATGCC-3' |
| <i>BsthiI</i> -PciI5 | 5'-CACATGTATTACGATCATATATTAATTCG-3' |
| <i>BsthiI</i> -BglII3 | 5'-CGAGATCTACAAAGCTTTCAAAGTG-3' |
| <i>BsthiI</i> -NheI5 | 5'-CATGGCTAGCGATCATATTAATTCG-3' |
| <i>BsthiI</i> -KpnI3 | 5'-CTGAGGTACCTCCTTGTGAGTATAGAATGTGTC-3' |
| <i>BssufC</i> -NcoI3 | 5'-TACCATGGCTGCTTCAACATTAAC-3' |
| <i>BssufC</i> -BamHI5 | 5'-CAGGATCCGGAGTGATGTAG-3' |
| <i>Bssuf</i> -pro5 | 5'-GAGTCGACGTATCTGAATAAGACTGAAAC-3' |
| <i>Bssuf</i> -pro3 | 5'-GTCCCGGGTGAAGCAGCCATAT-3' |
| <i>EctRNA</i> ^{Met5} | 5'-AATTCCTGCAGTATACGACTCACTATAGGCTACGTA GCTCAGTTGGTTAGAGCACATCACTCATAATGATGGG GTCACAGG -3' |
| <i>EctRNA</i> ^{Met3} | 5'-mUmGGTGGCTACGACGGGATTCGAACCTGTGAC CCATCATTATG-3' |

| Plasmids | Relevant Gene Cloned | Vector | Reference |
|----------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------|-----------|
| pDS13 | <i>nifZ</i> PCR product using <i>BsnifZ</i> -PciI5 and <i>BsnifZ</i> -BamHI3 primers. It places PciI and BamHI sites at 5' and 3' sites flanking <i>nifZ</i> coding sequence (codons 2-382) | pCR2.1 TOPO | This work |
| pDS18 | <i>thil</i> ' PCR product using <i>BsthiI</i> -PciI5 and <i>BsthiI</i> -BglII3 primers. It places PciI and BglII sites at 5' and 3' sites flanking <i>thil</i> ' coding sequence (codons 1-358) | pCR2.1 TOPO | This work |
| pDS19 | 908 bp HindIII ' <i>thil</i> ' fragment (198-1101 bp) from pDS18 into the HindIII site of pMutin4 plasmid. | pMutin4 | This work |
| pDS20 | 1.1 kb PciI-BglII fragment of <i>thil</i> ' ligated into NcoI-BglII sites of pAra13 | pAra13 | This work |
| pDS21 | 1.15 kb PciI and BamHI <i>nifZ</i> fragment from pDS13 ligated <i>nifZ</i> gene into NcoI-BglII sites of pAra13. | pAra13 | This work |
| pDS47 | <i>nifZ</i> ' PCR product using <i>BsnifZ</i> -PciI5 and <i>BsnifZ</i> '-BamHI3 primers. It places PciI and BamHI sites at 5' and 3' sites of <i>nifZ</i> (codons 2-380). | pCR2.1 TOPO | This work |
| pDS54 | 650 bp BglII-BamHI ' <i>nifZ</i> ' fragment (codons 130-380) ligated in forward orientation into BamHI-digested pMutin4 plasmid. | pMutin4 | This work |
| pDS71 | 2.2 kB <i>nifZ</i> - <i>thil</i> ' PCR product amplified with primers <i>BsnifZ</i> -PciI5 and <i>BsthiI</i> -BglII3. | pCR2.1 TOPO | This work |
| pDS80 | 2.2 kB StuI-KpnI ' <i>nifZ</i> - <i>thil</i> ' fragment ligated into StuI-KpnI sites of pDS21 | pAra13 | This work |
| pDS101 | <i>thil</i> PCR product using <i>BsthiI</i> -NheI5 and <i>BsthiI</i> -KpnI3 primers. It places NheI and KpnI sites at 5' and 3' sites of <i>thil</i> (codons 1-403, and additional 90 bp downstream). | pCR2.1 TOPO | This work |
| pDS108 | 976 bp NruI-KpnI ' <i>thil</i> fragment from pDS101 ligated into NruI-KpnI sites of pDS20 | pAra13 | This work |
| pDS115 | 976 bp NruI-KpnI ' <i>thil</i> fragment from pDS101 ligated into NruI-KpnI sites of pDS80. | pAra13 | This work |
| pBH113 | <i>Escherichia coli thil</i> | pet15b | (3) |
| pDB943 | <i>Azotobacter vinelandii iscS</i> | pT7-7 | (6) |

| Strain | Relevant Genotype | Reference |
|--------------------------|-----------------------|--------------------------------|
| <i>B. subtilis</i> PS832 | wild type strain | (1) |
| <i>B. subtilis</i> DD5 | 'nifZ':pMutin4 | This work |
| <i>B. subtilis</i> DD14 | 'nifZ':pMutin4 | This work |
| <i>B. subtilis</i> 1A603 | thiA84::Tn197 | (4) |
| <i>E. coli</i> MG1655 | wild type strain | Laboratory stock |
| <i>E. coli</i> CL100 | Δ iscS | (2) |
| <i>E. coli</i> JLD26501 | thi::Kan ^r | P1(VJS2890)→TL524 ^a |

^a Constructed by phage P1-mediated transduction of strain TL524 (5) to kanamycin resistance using stain VSJ2890 (3) as P1 donor

References

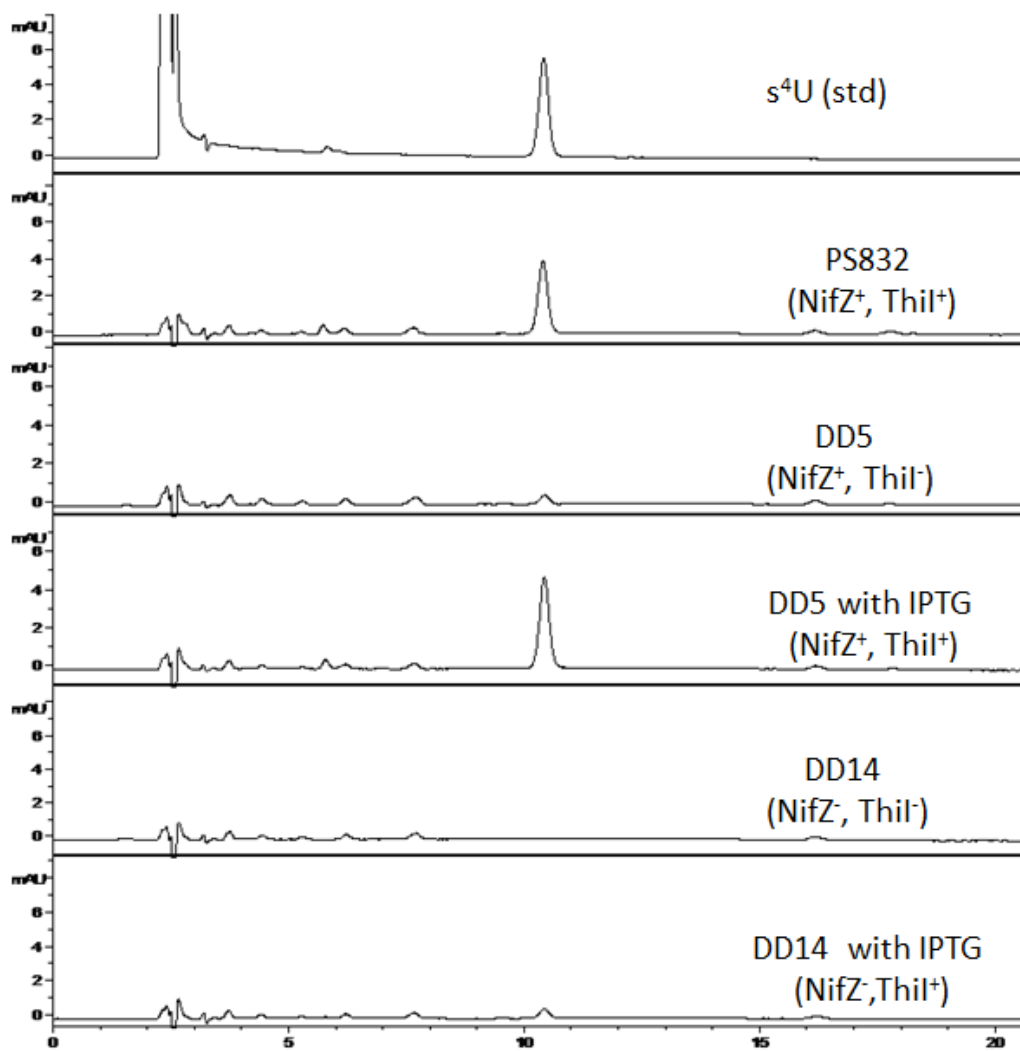
1. **Corfe, B. M., A. Moir, D. Popham, and P. Setlow.** 1994. Analysis of the expression and regulation of the gerB spore germination operon of *Bacillus subtilis* 168. *Microbiology* **140** (Pt 11):3079-3083.
2. **Lauhon, C. T., and R. Kambampati.** 2000. The *iscS* gene in *Escherichia coli* is required for the biosynthesis of 4-thiouridine, thiamin, and NAD. *J. Biol. Chem.* **275**:20096-20103.
3. **Mueller, E. G., C. J. Buck, P. M. Palenchar, L. E. Barnhart, and J. L. Paulson.** 1998. Identification of a gene involved in the generation of 4-thiouridine in tRNA. *Nucleic Acids Res.* **26**:2606-2010.
4. **Vandeyar, M. A., and S. A. Zahler.** 1986. Chromosomal insertions of Tn917 in *Bacillus subtilis*. *J. Bacteriol.* **167**:530-534.
5. **Wolfe, M. D., F. Ahmed, G. M. Lacourciere, C. T. Lauhon, T. C. Stadtman, and T. J. Larson.** 2004. Functional diversity of the rhodanese homology domain: the *Escherichia coli* *ybbB* gene encodes a selenophosphate-dependent tRNA 2-selenouridine synthase. *J. Biol. Chem.* **279**:1801-1809.
6. **Zheng, L., V. L. Cash, D. H. Flint, and D. R. Dean.** 1998. Assembly of iron-sulfur clusters. Identification of an *iscSUA-hscBA-fdx* gene cluster from *Azotobacter vinelandii*. *J. Biol. Chem.* **273**:13264-13272.

Figure S1. Detection of s⁴U8 in tRNA nucleosides isolated from cells lacking NifZ and/or Thil. (A) HPLC-MS chromatograms at Abs_{330nm} of tRNA nucleosides isolated from *B. subtilis* PS832, DD5 in the presence (*nifZ*⁺, *thil*⁺) and absence of IPTG (*nifZ*⁻, *thil*⁻) and DD14 culture in the presence (*nifZ*⁺, *thil*⁺) and absence (*nifZ*⁻, *thil*⁻) of IPTG. The peak corresponding to s⁴U was compared to the standard (shown on top), and to pseudouridine which was used as an internal control (not shown). (B) Negative mode mass spectrum of the peak corresponding to s⁴U.

Figure S2. Growth of *B. subtilis* strains in minimum medium in the presence (filled symbols) and in the absence (empty symbols) of 0.8 µg/mL thiamine. Cells were cultured in Spizizen's medium at 37°C, 300 RPM for PS832 (○) and DD14 in the absence (□) and presence (△) of IPTG. *B. subtilis* 1A603 (*thiA84::Tn917*) was cultured in Spizizen medium supplemented with tryptophan (◇). Growth patterns were monitored for 14 hours and optical density at 600 nm (A_{600nm}) was recorded as indicated.

Figure S1

A



B

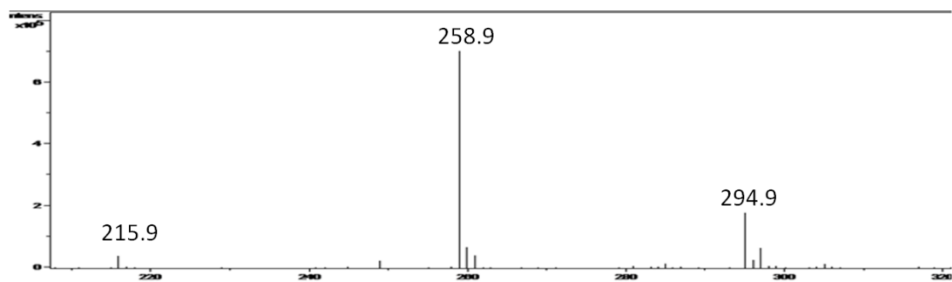


Figure S2

