## **Supplementary Figure**

## Fig. S1. Zur-Box consensus region in *E.coli znuA*, and *S.* Typhimurium *znuA*, *zinT* and *zupT*

Ec znuA	GA <b>AATGTTATAATATCACATTTC</b>
St znuA	AG <u>AATGTTATAATATCACATTTC</u>
St zinT	GC <u>AATGTTATAATAT</u> A <u>ACA</u> A <u>T</u> CA
St zupT	AC <u>A</u> T <u>T</u> T <u>T</u> GT <u>TAA</u> C <u>ATCAC</u> CA <u>T</u> AC

**Fig. S2**. Growth curves of *S*. Typhimurium wild type ( $\blacksquare$ ), *mntH sitABCD* ( $\blacktriangle$ ) and *fepA/entF feoB* ( $\blacktriangledown$ ) mutant strains in VB-MM. Panel A: growth in standard VB-MM; Panel B: growth in VB-MM supplemented with 3  $\mu$ M MnCl<sub>2</sub>; Panel C growth in VB-MM supplemented with 3  $\mu$ M FeSO<sub>4</sub>.



**Fig. S3**. Growth curves of *S*. Typhimurium *znuABCzupT* in standard VB-MM ( $\blacksquare$ ), in VB-MM supplemented with 3  $\mu$ M ZnSO<sub>4</sub>( $\blacktriangle$ ) or in VB-MM supplemented with 3  $\mu$ M MnCl<sub>2</sub>( $\blacklozenge$ ).



Fig S3