



Figure S4. RT-qPCR analyses of RirA targets. Free-living cultures under iron replete (37 μM FeCl_3) or deficient (0.37 μM FeCl_3) conditions, and rhizobia within soybean nodules (28 dpi). The transcription level relative to 16S rRNA gene is shown. Significant differences are indicated (mean \pm SE based on three independent experiments; *t* test, *, P-value < 0.05; **, P-value < 0.01; ***, P-value < 0.001).