

Figure S1. Typical EPR spectra of the homogenate. The arrows show the magnetic field for $g = 4.3$ and the peak to peak intensity (I_s) used to measure the relative amount of labile iron.

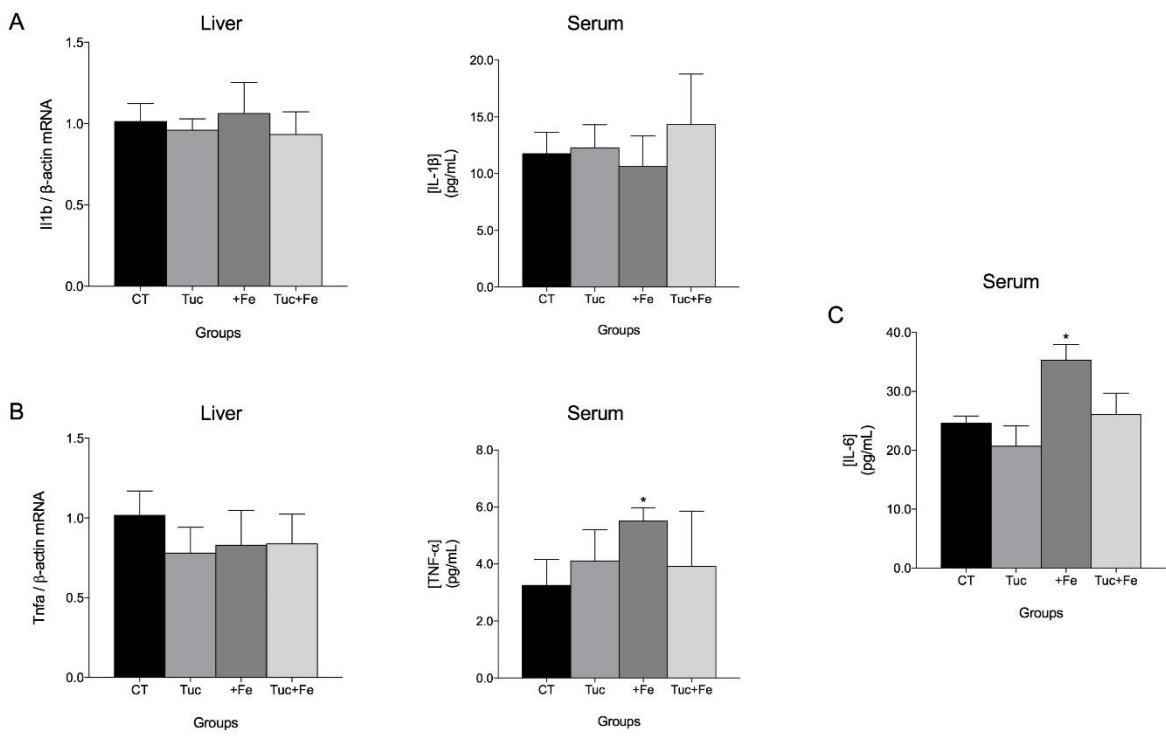


Figure S2. Hepatic mRNA levels quantification of interleukin 1 beta (Il1b) and tumor necrosis factor alpha (Tnfa) mRNA levels and serum protein levels of $\text{IL-1}\beta$, IL-6 and $\text{TNF-}\alpha$ in rats fed a control diet, tucum-do-cerrado (Tuc), iron-enriched diet (+Fe) or tucum-do-cerrado iron-enriched (Tuc+Fe) diet, for 12 weeks. Data are the average \pm standard deviation ($n = 6$). Statistical differences compared to: * CT group, # Tuc group and § +Fe group ($p < 0.05$).

Table S1. Primer sequences used for Bmp6, Hamp, Hfe, Hfe2, Smad7, Trf1 and Actb real-time PCR assays.

| Gene | Primers sequences (5'- 3') | GenBank accession number |
|---------------|-----------------------------------|--------------------------|
| Bmp6 | GACAGCAGAGTCGCAATCG (forward) | NM_013107.1 |
| | AGCTCACGTAAAGCTCATGC (reverse) | |
| Hamp | TGATGCTGAAGCGAAGGGA (forward) | NM_053469 |
| | TGTGTTGAGAGGTCAGGAC (reverse) | |
| Hfe | TGGGCAAGATCACCTTGAATT (forward) | NM_053301.4 |
| | GGATCCTGTGCTCTTCCACT (reverse) | |
| Hfe2 | GTAGCATCGGGAGCCAAC (forward) | NM_001012080.1 |
| | TCAAAGG CTGCAGGAAGATT (reverse) | |
| Smad 7 | AGAGGCTGTGTTGCTGTG (forward) | NM_030858.1 |
| | CATCGGGTATCTGGAGTAAGG (reverse) | |
| Trf1 | GAGTTCACTGACATCATCAAGCA (forward) | NM_022712.1 |
| | TCCAGCCTCACGAGGAGTAT (reverse) | |
| Actb | GTCGTACCACTGGCATTGTG (forward) | NM_031144 |
| | CTCTCAGCTGTGGTGGTGAA (reverse) | |

Table S2. Dilutions and companies of primary antibodies used for protein immunoblotting.

| Protein | Dilution | Code/Company |
|----------------|----------|------------------------------------|
| pSMAD1/5/8 | 1:500 | #9511 / Cell Signaling Technology |
| pSTAT3 | 1:500 | #9131 / Cell Signaling Technology |
| β -actin | 1:1000 | #4967S / Cell Signaling Technology |