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Comparative transcriptomic analysis of the roots of intercropped peanut and maize reveals novel insights into peanut iron nutrition

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## ACCEPTED MANUSCRIPT

- 1 **Abbreviations**: Fe, iron; SSH, suppression subtractive hybridization; ACC,
- 2 aminocyclopropane-1-carboxylic acid synthase; MAs, mugineic acid family
- 3 phytosiderophores; TOM1, transporter of mugineic acid 1; YS1, yellow stripe 1
- 4 transporter; NRAMP, natural resistance-associated macrophage protein; Zn, zinc; P,
- 5 phosphorus; K, potassium; DMT, divalent metal ion transporter genes; Cd, cadmium;
- 6 M, monocropping; I, intercropping; MTP, metal tolerance protein; Cu, copper;
- 7 NRT/PRT, peptide/nitrate transporter; IAA, Indole-3-acetic acid; ABA, abscisic acid;
- 8 OPTs, oligopeptide transporters; GSH, glutathione; ABC, ATP□binding cassette
- 9 transporter; ERFs, ethylene-response factors; JA, jasmonic acid; JAs, jasmonates; GO,
- 10 gene ontology

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