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# Correction: Mapping of quantitative trait locus for resistance to anthracnose in a population derived from genotypes PI 527538 and Ervilha of common bean

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## KEYWORDS

*Colletotrichum lindemuthianum*, quantitative trait locus, candidate genes, recombinant inbred lines, race

## A Correction on

### Mapping of quantitative trait locus for resistance to anthracnose in a population derived from genotypes PI 527538 and Ervilha of common bean

By Mulube M, Hamabwe S, Kuwabo K, Chinji M, Nkandela M, Botha J, Mwense B, Tembo L, Lungu D, Mukuma C, Parker T and Kamfwa K (2025). *Front. Plant Sci.* 16:1691703. doi: 10.3389/fpls.2025.1691703

There was a mistake in [Figure 1](#) and [2](#) as published. The figure images were swapped, but the captions were correct. That is, the linkage map figure originally presented as [Figure 1](#) should be [Figure 2](#), while the frequency table distribution graph presented as [Figure 2](#) should be [Figure 1](#).

The corrected [Figure 1](#) and [Figure 2](#) appear below.

The original version of this article has been updated.

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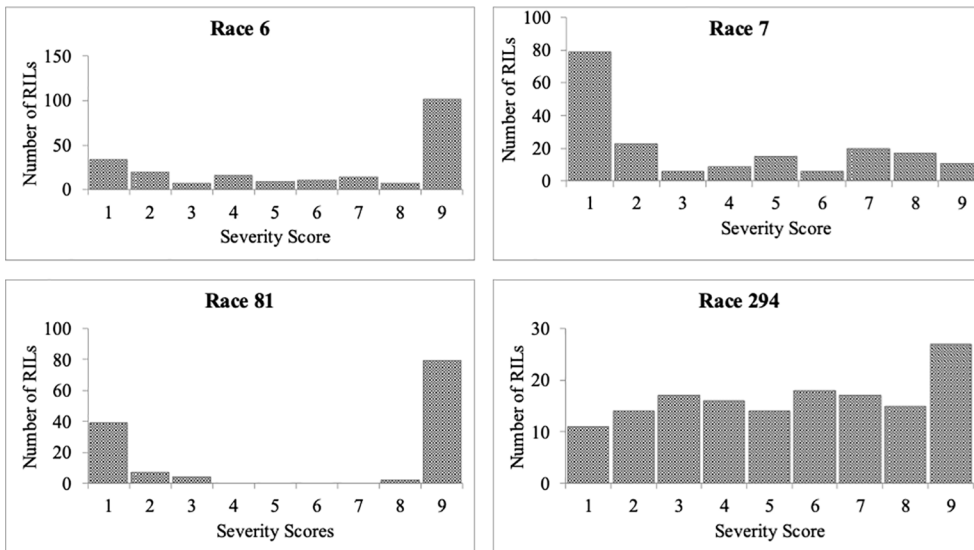


FIGURE 1 Frequency distributions of severity scores for *Colletotrichum lindemuthianum* races 6, 7, 81, and 294 inoculated on recombinant inbred lines derived from a cross of PI 527538 and Ervilha. This should be the bare graph (also known as histogram).

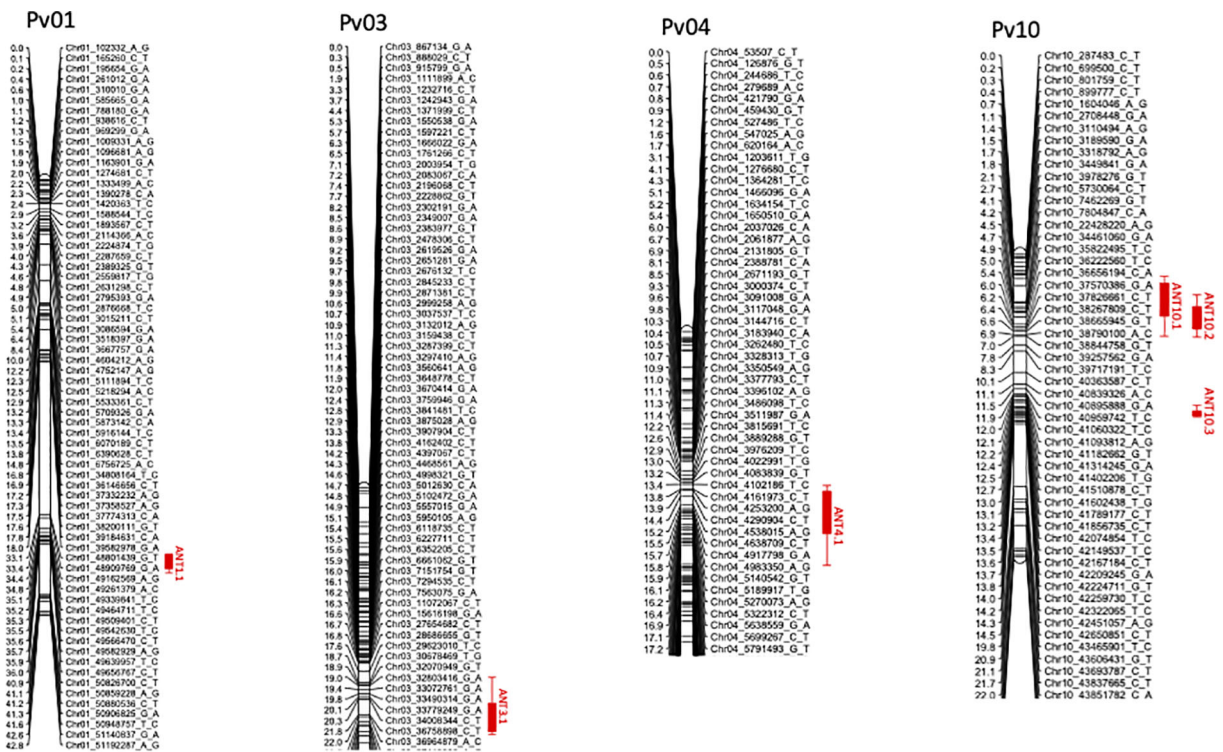


FIGURE 2 Linkage maps for chromosomes Pv01, Pv03, Pv04, and Pv10 with positions for QTL ANT1.1 (for races 6 and 81), ANT3.1 (race 294), ANT4.1 (race 6), ANT10.1 (race 6), ANT10.2 (race 7), and ANT10.3 (race 294). This should be the linkage maps.