

Erratum to: Genome sizes in diploid and allopolyploid *Arachis* L. species (section *Arachis*)

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We regret but noticed the following corrections went required;

Herewith please find a renewed Fig. 1 to replace the former published which contained an inadequately marked distribution border line.

Secondly, in the Materials and methods, Flow cytometry section, the name “*Paspalum notatum* Chirú byotype” should have read “*Paspalum dilatatum* Chirú Q4081”.

Furthermore, the last article section “The constancy in the Cx values suggests that the hybridization and chromosome doubling events that occurred during the origin of the cultivated peanut did not

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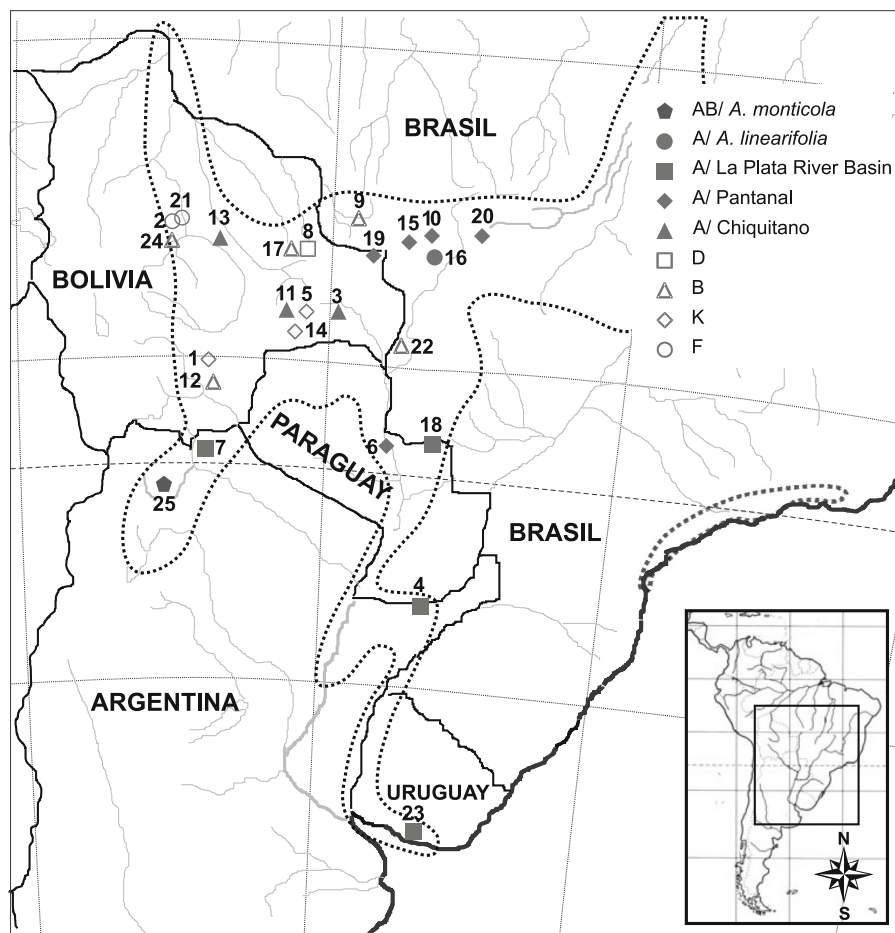


Fig. 1 Geographic distribution of the localities in which the accessions of the wild *Arachis* species analysed in this study were collected. For location details and species code see

Table 1. The *dash-dotted line* denotes the distribution of the *Arachis* section. Each genome and karyotype group (of the A genome) is indicated with different symbol

induced significant changes in genome size.” should read “The constancy in the Cx values suggests that the hybridization and chromosome doubling events

that occurred during the origin of the cultivated peanut did not induce significant changes in genome size.”