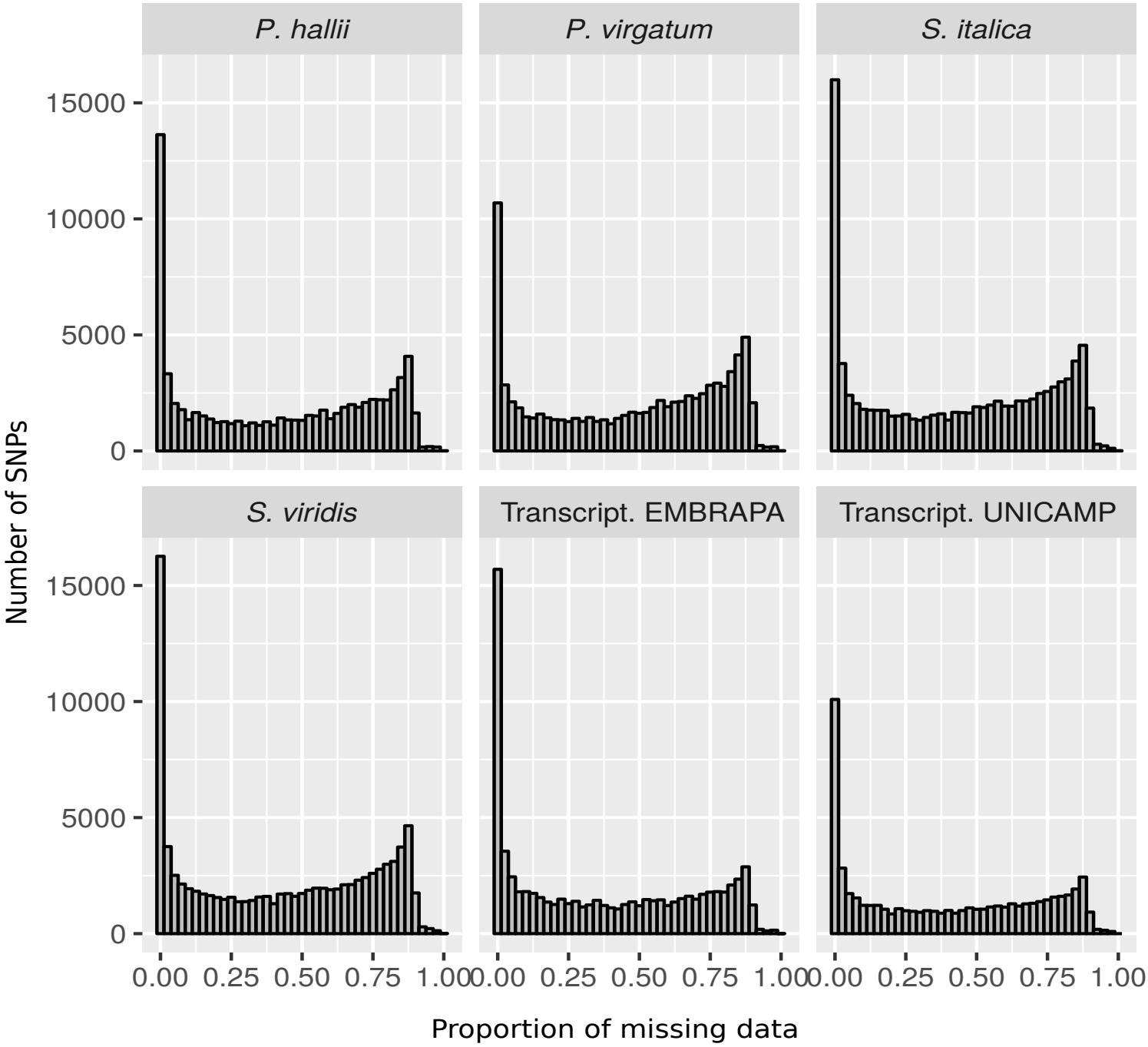

Figure S2 Proportion of missing data in GBS markers using *Panicum hallii*, *Panicum virgatum*, *Setaria italica*, *Setaria viridis*, and two transcriptomes of *Panicum maximum* (obtained by EMBRAPA and UNICAMP research groups) as reference genomes.