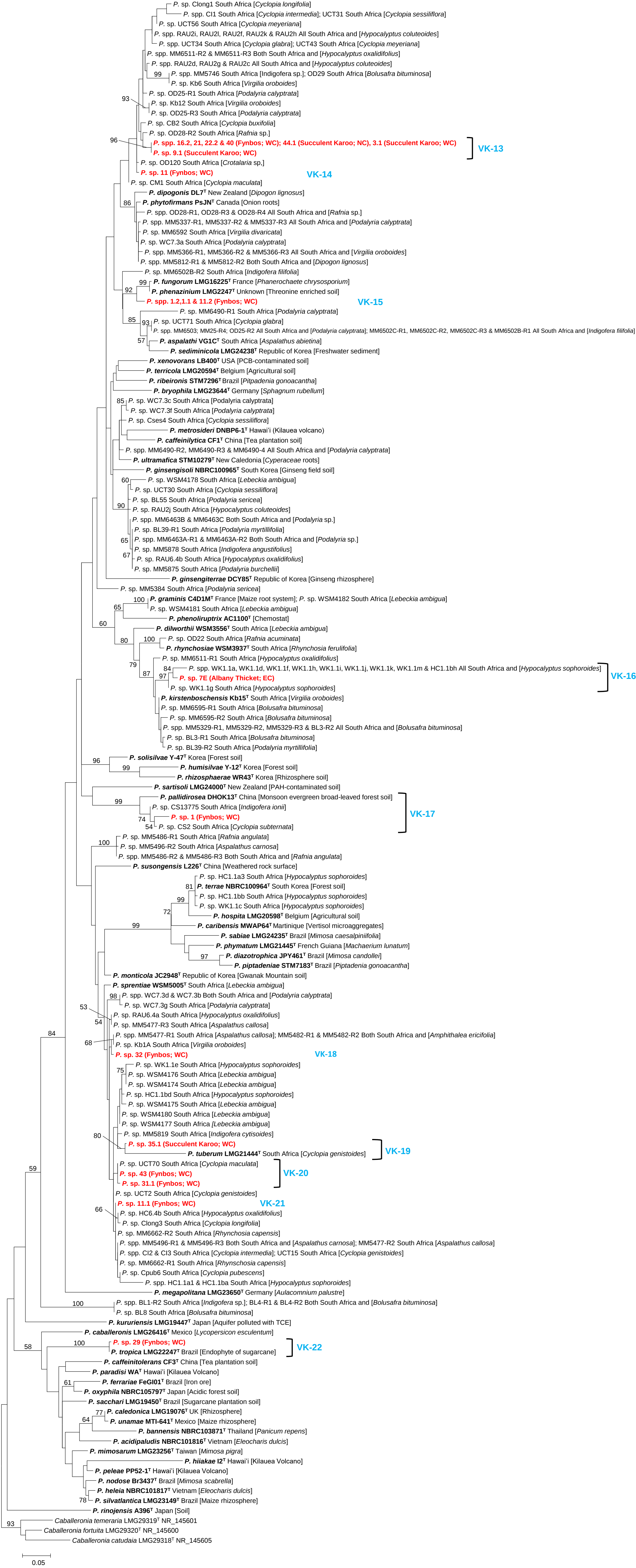


Paraburkholderia
recA



Suppl. Fig. S3 A *recA* maximum-likelihood phylogeny of the genus *Paraburkholderia*. Isolates from this study appear in red followed by information for the biome and province (as abbreviated for Table 1) from which the 'trapping' soil originated. *Paraburkholderia* type strains appear in bold, and all the isolates list information for their country of origin and host or source. Suppl. Table S3 contains the GenBank accessions and associated references for all the *Paraburkholderia* isolates included in this phylogeny. The delineated *Vachellia karroo*-associated lineages (VK-13 to VK-22) are listed in blue. Three species of the genus *Caballeronia* were used for an outgroup (their accession numbers are listed in the phylogeny). Nodes with $\geq 50\%$ bootstrap support are shown, while the scale bar indicates the number of nucleotide substitutions per site.