

recA

Supplementary Figure S3. The single most parsimonious tree inferred from the *recA* data for *Burkholderia kirstenboschensis* and related *Burkholderia* species. The tree depicted is 571 steps long with consistency, retention and homoplasy indices of 0.39, 0.66 and 0.61, respectively. Bootstrap values ($\geq 60\%$) are indicated at the branches and type strains of the species *Burkholderia cepacia* and *B. vietnamiensis* were used outgroup. The scale bar indicates the number of nucleotide changes per site. Type strain information and sequence accession numbers are indicated as in Supplementary Figure S1.

