

ERRATUM

ParABS Systems of the Four Replicons of *Burkholderia cenocepacia*: New Chromosome Centromeres Confer Partition Specificity

Nelly Dubarry, Franck Pasta, and David Lane

*Laboratoire de Microbiologie et Génétique Moléculaire, Centre National de Recherche Scientifique, 118 route de Narbonne,
31062 Toulouse, France*

Volume 188, no. 4, p. 1489–1496, 2006. Page 1495, column 1: The sixth sentence of the first full paragraph should read as follows: The weak partition activity of the c2, c3, and p1 systems may be partly a consequence of intrinsic inefficiency, sensed even in *Burkholderia*; whereas the efficient c1 *parAB* is normally supplemented with only two isolated *parS* sites, those of c2, c3, and p1 are accompanied by three to four nearby *parS* sites in a clustered arrangement which may serve to compensate for a lower effectiveness of its protein partners.

Page 1495, column 2: The fourth sentence should read as follows: The second chromosome of *Vibrio cholerae* also betrays a plasmid ancestor through the phylogeny of its ParA (6) as well as by the nature of its replication origin (5).