

Table S5. 454-sequencing and newbler-assembly data of the genomes sequenced in this study.

	<i>B. clarridgeiae</i>	<i>B. rochalimae</i>	<i>B. sp. 1-1C</i>	<i>B. sp. AR 15-3</i>	<i>B. schoenbuchensis</i>
Final length (bp)	1,522,743	1,540,290	1,566,739	1,590,248	1,667,713
Number of 454-reads	212,905	287,512	253,303	254,535	173,358
Average read length (bp)	247	211	241	236	285
Number of Newbler contigs (> one read)	43	78	74	93	220
Number of contigs after manual assembly	1	13	19	17	18
454-sequence coverage	35x	39x	39x	37x	29x