

TABLE S1

Species	Proportion of isolates^a
<i>B. longum</i> subsp. <i>longum</i>	57.28%
<i>B. breve</i>	27.18%
<i>B. animalis</i> subsp. <i>lactis</i>	7.28%
<i>B. bifidum</i>	6.80%
<i>B. dentium</i>	0.97%
<i>B. tsurumiense</i> (<i>low confidence</i>)	0.49%

^aThere were 206 isolates of *Bifidobacterium*; internal transcribed spacer region (ITS) sequences were identified using BLAST in the NCBI database.