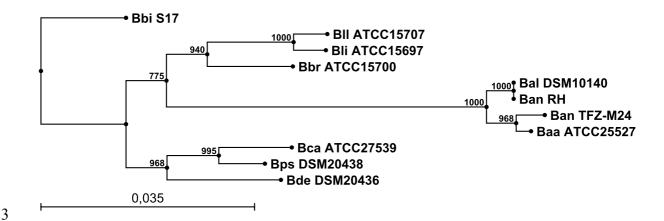
## 1 S1 Figure



Supplementary Figure S1: Taxonomic analysis of the *Bifidobacterium sp.* strain isolated from the animal facility at Ulm University. A phylogenetic tree was calculated with the 16S rRNA gene sequence of the isolated strain (Ban TFZ-M24) obtained by PCR (described in Materials and Methods) and the corresponding sequences of *B. bifidum* S17 (Bbi S17), *B. longum* subsp. *longum* ATCC15707 (Bll ATCC15707), *B. longum* subsp. *infantis* ATCC15697 (Bli ATCC15697), *B. breve* ATCC15700 (Bbr ATCC15700), *B. animalis* subsp. *lactis* DSM10140 (Bal DSM10140), *B. animalis* RH (Ban RH), *B. animalis* subsp. *animalis* ATCC25527 (Bal ATCC25527), *B. catenulatum* ATCC27539 (Bca ATCC27539), *B. pseudocatenulatum* DSM20438 (Bps DSM20438), and *B. dentium* DSM20436 (Bde DSM20436). Numbers indicate bootstrap values from n=1000 iterations.