Claims

1. Use of a pharmaceutical composition comprising a plurality of bacterial isolates in manufacture of a medicament for treating or preventing inflammatory bowel disease (IBD) in a subject in need thereof, wherein the plurality of bacterial isolates comprises *Bacteroides stercoris and Bacteroides cellulosilyticus*, and at least one of *Bacteroides uniformis*, *Subdoligranulum variabile*, *Anaerostipes hadrus*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Faecalibacterium prausnitzii*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*, wherein the *Bacteroides stercoris* comprises a 16S ribosomal ribonucleic acid (rRNA) sequence that has at least 95%, or at least 97%, or at least 98%, or at least 99% sequence identity with the nucleotide sequence of SEQ ID NO: 13.

2. The use of claim 1, wherein at least two of the plurality of bacterial isolates are isolated from a stool of different human donors.

3. The use of claim 1 or 2, wherein the plurality of bacterial isolates comprises at least two of *Bacteroides uniformis*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Anaerostipes hadrus*, *Faecalibacterium prausnitzii*, *Subdoligranulum variabile*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*.

4. The use of claim 1 or 2, wherein the plurality of bacterial isolates comprises at least three of *Bacteroides uniformis*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Anaerostipes hadrus*, *Faecalibacterium prausnitzii*, *Subdoligranulum variabile*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*.

5. The use of claim 1 or 2, wherein the plurality of bacterial isolates comprises at least four of *Bacteroides uniformis*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Anaerostipes hadrus*, *Faecalibacterium prausnitzii*, *Subdoligranulum variabile*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*.

6. The use of claim 1 or 2, wherein the plurality of bacterial isolates comprises at least five of *Bacteroides uniformis*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Anaerostipes hadrus*, *Faecalibacterium prausnitzii*, *Subdoligranulum variabile*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*.

7. The use of claim 1 or 2, wherein the plurality of bacterial isolates comprises at least six of *Bacteroides uniformis*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Anaerostipes hadrus*, *Faecalibacterium prausnitzii*, *Subdoligranulum variabile*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*.

8. The use of claim 1 or 2, wherein the plurality of bacterial isolates comprises at least seven of *Bacteroides uniformis*, *Odoribacter splanchnicus*, *Roseburia faecis*, *Anaerostipes hadrus*, *Faecalibacterium prausnitzii*, *Subdoligranulum variabile*, *Akkermansia muciniphila*, *Alistipes shahii*, and *Eubacterium rectale*.

9. The use of any one of claims 1-8, wherein the plurality of bacterial isolates comprises one or more 16S rRNA sequences that have at least 95%, or at least 97%, or at least 98%, or at least 99% sequence identity with one or more nucleotide sequences selected from SEQ ID NOs: 1, 2, 3, 7, 8, 11, 14, 18, 19, 20, 22 and 23.

10. The use of any one of claims 1-9, wherein the plurality of bacterial isolates comprises at least two bacterial isolates comprising *Faecalibacterium prausnitzii*, wherein the at least two bacterial isolates comprise different 16S rRNA sequences.

11. The use of claim 10, wherein the at least two bacterial isolates comprising *Faecalibacterium prausnitzii* comprise 16S rRNA sequences that have at least 95%, or at least 97%, or at least 98%, or at least 99% sequence identity with nucleotide sequences of SEQ ID NOs: 1 and/or 7.

12. The use of claim 10 or 11, wherein the at least two bacterial isolates comprising *Faecalibacterim prausnitzii* are isolated from a stool of different human donors.

13. The use of any one of claims 1-12, wherein the plurality of bacterial isolates comprises lyophilized bacteria.

14. The use of any one of claims 1-13, wherein the plurality of bacterial isolates does not include *Escherichia coli*.