recurrent CDI
allogenic
responder
n.r.
ulcerative colitis
allogenic
responder
non-responder
autologous
r.
non-resp
Crohn's disease
allogenic
resp.
non-resp.
ESBL
allogenic
resp.
non-resp.
metabolic syndrome
allogenic
autologous

Bacteroides ovatus (A0023)
Bacteroides stercoris (r0275)
Bacteroides cellulosilyticus (r0692)
Bacteroides thetaiotaomicron (r0897)
Bacteroides uniformis (r0899)
Bacteroides xylanisolvens (r1072)
Bacteroides ovatus (r1073)
Bacteroides caccae (r1382)
Phocaeicola massiliensis (r0455)
Phocaeicola vulgatus (r0898)
Prevotella copri (r4448)
Barnesiella intestinihominis (r4880)
Odoribacter splanchnicus (r4846)
Alistipes onderdonkii (r0775)
Alistipes finegoldii (r1481)
Alistipes obesi (r1825)
Alistipes butredinis (r4266)
Alistipes shahii (r4873)
Parabacteroides distasonis (r1074)
Parabacteroides merdae (r1378)
Bifidobacterium adolescentis (A0399)
Bifidobacterium longum (r0150)
Bifidobacterium adolescentis (r1156)
Collinsella aerofaciens (r1383)
Adlercreutzia celatus_A (r3198)
Eggerthella lenta (r0642)
Bilophila sp900550745 (A3477)
Bilophila wadsworthia (r1149)
Parasutterella excrementihominis (r1405)
Sutterella wadsworthensis (r0767)
Erysipelatoclostridium sp000752095 (r3608)
Holdemanella sp002299315 (m7589)
Streptococcus salivarius (r0199)
Streptococcus thermophilus (r0219)
Oscillospiraceae (A1608)
CAG-103 sp900317855 (A0333)
CAG-103 sp900317855 (A0334)
CAG-103 sp000432375 (m7180)
CAG-110 sp003525905 (m5396)
CAG-170 sp000432135 (m6348)
CAG-170 sp002404795 (m6801)
CAG-83 sp001916855 (A0311)
CAG-83 sp000431575 (m5805)
CAG-83 sp000435975 (m5944)
CAG-83 sp001916855 (m6272)
ER4 sp000765235 (A0294)
ER4 sp900546295 (A0297)
ER4 sp900317525 (m6795)
ER4 sp000765235 (r3624)
Flavonifractor plautii (r1377)
Intestinimonas butyriciproducens (r2968)
Lawsonibacter (A0066)
Lawsonibacter sp900553135 (A1972)
Lawsonibacter asaccharolyticus (m6419)
Oscillibacter (A0040)
Oscillibacter (A1530)
Oscillibacter sp900548505 (A2315)
Oscillibacter sp001916835 (m6676)
Oscillibacter welbionis (r0858)
Faecalibacterium sp900539885 (A0336)
Faecalibacterium prausnitzii_J (A0341)
Faecalibacterium prausnitzii_A (A0343)
Faecalibacterium prausnitzii_F (A0344)
Faecalibacterium (m5779)
Faecalibacterium prausnitzii_I (m5815)
Faecalibacterium sp900539945 (m6631)
Faecalibacterium prausnitzii_E (m7143)
Faecalibacterium prausnitzii (r1379)
Faecalibacterium prausnitzii_C (r4211)
Faecalibacterium prausnitzii_G (r4875)
Faecalibacterium prausnitzii_D (r4910)
Fournierella sp002160145 (A2320)
Fournierella sp004558145 (A2532)
Gemmiger (A2183)
Gemmiger sp900539695 (m6478)
Gemmiger qucibialis (m6557)
Gemmiger sp900540775 (m6850)
Gemmiger formicilis (m6905)
Ruminiclostridium_E siraeum (r1387)
Ruminococcus_D bicirculans (r2358)
Ruthenibacterium lactatiformans (r4738)
SFJ001 sp004555865 (A2586)
1XD42-69 (A2476)
Acetatifactor sp003447295 (A0202)
Acetatifactor sp900066365 (m5669)
Acetatifactor sp900066565 (m5978)
Agathobacter faecis (m5354)
Agathobacter rectalis (r1416)
Anaerobutyricum hallii_A (m6657)
Anaerobutyricum hallii (r4207)
Anaerostipes hadrus (r1309)
Bariatricus comes (r4313)
Blautia_A (A2141)
Blautia_A sp900066355 (A2800)
Blautia_A sp003478765 (m6013)
Blautia_A sp000285855 (m6525)
Blautia_A sp900548245 (m6716)
Blautia_A sp900066335 (m6834)
Blautia_A sp900066145 (m7018)
Blautia_A sp000436615 (m7076)
Blautia_A sp900066165 (m7130)
Blautia_A sp900066505 (m7306)
Blautia_A wexlerae (r0466)
Blautia_A massiliensis (r0859)
Blautia_A obeum (r4202)
Blautia_A sp900066355 (r4719)
CAG-127 (A0140)
CAG-127 sp900319515 (m5364)
CAG-45 sp900066395 (m6916)
CAG-81 sp900066785 (m6808)
CAG-81 sp900066535 (r3825)
Clostridium_Q sp003024715 (m6832)
Coprococcus eutactus_A (m5379)
Coprococcus_A sp900548825 (A3431)
Coprococcus_A catus (r4874)
Dorea sp900550865 (A0100)
Dorea sp900066555 (m5826)
Dorea sp000433215 (m7455)
Dorea formicigenerans (r0973)
Dorea longicatena_B (r2893)
Dorea longicatena (r4203)
Eisenbergiella sp900066775 (m6088)
Enterocloster sp900541315 (m5712)
Enterocloster sp000431375 (m6792)
Eubacterium_F sp003491505 (m6883)
Eubacterium_I ramulus (r2795)
Fusicatenibacter saccharivorans (m6937)
GCA-900066135 sp900543575 (m6852)
GCA-900066575 sp900553635 (A2429)
Hungatella_A sp003478355 (m5603)
KLE1615 sp900066985 (m6632)
Lachnospira rogosae (m6721)
Lachnospira sp000437735 (m7668)
Lachnospira eligens_B (r4389)
Mediterraneibacter torques (r1376)
Mediterraneibacter faecis (r4718)
Roseburia intestinalis (r1427)
Roseburia hominis (r4572)
Roseburia inulinivorans (r4632)
Ruminococcus_A sp003011855 (m5411)
Ruminococcus_B gnavus (r0280)
UBA11774 sp003507655 (m6875)
UBA7160 (A3024)
UBA7182 sp003480725 (m7031)
UBA9502 sp003480315 (m7014)
UMGS1375 sp900066615 (m6787)
Anaerotignum sp000436415 (m5947)
CAG-41 sp900066215 (m6602)
Clostridium_A leptum (r4234)
Ruminococcus_E bromii_B (r4720)
Ruminococcus_H sp003531055 (m7093)
Agathobaculum butyriciproducens (m5437)
UBA1191 sp900549125 (m5477)
Intestinibacter bartlettii (r4268)
Romboutsia timonensis (m7331)
SFFH01 sp900548125 (m7415)
SFFH01 sp900542445 (m7782)
UBA11524 sp000437595 (m6452)
UMGS1975 sp900546685 (m5735)

FMT outcome
rejection
colonisation
novel strains
novel str. (consp.)
colonisation (consp.)
coexistence
persistence (consp.)
persistence
species loss