

## Supplementary materials

Supplementary Table S1. Relative abundance at phylum levels.

| Phylum                  | Control |         |         | Treatment |         |         |
|-------------------------|---------|---------|---------|-----------|---------|---------|
|                         | Donor 1 | Donor 2 | Donor 3 | Donor 1   | Donor 2 | Donor 3 |
| <i>Firmicutes</i>       | 21.3986 | 24.9092 | 50.0761 | 31.5851   | 28.2239 | 49.5107 |
| <i>Proteobacteria</i>   | 78.3565 | 73.7341 | 49.1644 | 68.2224   | 70.1170 | 47.7291 |
| <i>Bacteroidetes</i>    | 0.0125  | 0.0314  | 0.0104  | 0.0108    | 0.0107  | 0.0000  |
| <i>Actinobacteria</i>   | 0.0177  | 0.7823  | 0.4790  | 0.0614    | 1.4547  | 2.3280  |
| <i>Verrucomicrobia</i>  | 0.0000  | 0.0015  | 0.0005  | 0.0000    | 0.0000  | 0.0010  |
| <i>Fusobacteria</i>     | 0.0000  | 0.3156  | 0.0047  | 0.0000    | 0.0076  | 0.0000  |
| <i>Gemmatimonadetes</i> | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Synergistetes</i>    | 0.0000  | 0.0000  | 0.0016  | 0.0000    | 0.0000  | 0.0000  |

Supplementary Table S2. Relative abundance at family levels.

| Family                       | Control |         |         | Treatment |         |         |
|------------------------------|---------|---------|---------|-----------|---------|---------|
|                              | Donor 1 | Donor 2 | Donor 3 | Donor 1   | Donor 2 | Donor 3 |
| <i>Acidaminococcaceae</i>    | 9.5747  | 2.4803  | 4.6785  | 19.1722   | 2.1647  | 1.0408  |
| <i>Clostridiaceae</i> 1      | 7.3435  | 0.5876  | 3.8234  | 4.4148    | 0.0041  | 20.5760 |
| <i>Desulfovibrionaceae</i>   | 0.0000  | 0.0000  | 0.0026  | 0.0026    | 0.0000  | 0.0025  |
| <i>Enterococcaceae</i>       | 2.4730  | 17.8295 | 33.4064 | 3.7032    | 7.5518  | 26.1675 |
| <i>Erysipelotrichaceae</i>   | 0.3198  | 0.0582  | 0.1434  | 1.3464    | 0.1973  | 0.1061  |
| <i>Eubacteriaceae</i>        | 0.0000  | 0.0000  | 0.2244  | 0.0000    | 0.0000  | 0.0000  |
| <i>Lachnospiraceae</i>       | 0.7014  | 2.8625  | 0.3465  | 1.2060    | 8.6140  | 0.6646  |
| <i>Lactobacillaceae</i>      | 0.1731  | 0.1724  | 0.0010  | 0.1068    | 0.0540  | 0.0025  |
| <i>Peptoniphilaceae</i>      | 0.0000  | 0.0000  | 0.0052  | 0.0021    | 0.0000  | 0.0000  |
| <i>Peptostreptococcaceae</i> | 0.0484  | 0.06432 | 5.8831  | 0.0408    | 9.4250  | 0.0652  |
| <i>Ruminococcaceae</i>       | 0.5189  | 0.1133  | 0.3668  | 0.8200    | 0.0668  | 0.0096  |
| <i>Selenomonadaceae</i>      | 0.0000  | 0.0096  | 0.0005  | 0.0000    | 0.0280  | 0.0000  |
| <i>Streptococcaceae</i>      | 0.0530  | 0.0592  | 0.0208  | 0.1316    | 0.0301  | 0.0152  |
| <i>Veillonellaceae</i>       | 0.0915  | 0.0000  | 0.6229  | 0.5460    | 0.0102  | 0.0854  |
| <i>Enterobacteriaceae</i>    | 77.7955 | 72.9281 | 48.0886 | 67.1206   | 69.2898 | 46.8461 |
| <i>Morganellaceae</i>        | 0.0031  | 0.2311  | 0.0000  | 0.0098    | 0.0836  | 0.0000  |
| <i>Pasteurellaceae</i>       | 0.0395  | 0.0000  | 0.0000  | 0.2193    | 0.0000  | 0.0000  |
| <i>Phyllobacteriaceae</i>    | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0005  | 0.0000  |
| <i>Rhodobacteraceae</i>      | 0.0031  | 0.0091  | 0.0016  | 0.0057    | 0.0036  | 0.0000  |
| <i>Sphingomonadaceae</i>     | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Sutterellaceae</i>        | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Muribaculaceae</i>        | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Odoribacteraceae</i>      | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Prevotellaceae</i>        | 0.0000  | 0.0000  | 0.0010  | 0.0010    | 0.0005  | 0.0000  |

| Family                      | Control |         |         | Treatment |         |         |
|-----------------------------|---------|---------|---------|-----------|---------|---------|
|                             | Donor 1 | Donor 2 | Donor 3 | Donor 1   | Donor 2 | Donor 3 |
| <i>Rikenellaceae</i>        | 0.0000  | 0.0000  | 0.0016  | 0.0000    | 0.0000  | 0.0000  |
| <i>Eggerthellaceae</i>      | 0.0042  | 0.0030  | 0.0026  | 0.0191    | 0.0025  | 0.0091  |
| <i>Micrococcaceae</i>       | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Nakamurellaceae</i>      | 0.0000  | 0.0000  | 0.0000  | 0.0010    | 0.0056  | 0.0015  |
| <i>Porphyromonadaceae</i>   | 0.0000  | 0.0000  | 0.0000  | 0.0010    | 0.0000  | 0.0000  |
| <i>Pseudonocardiaceae</i>   | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Solirubrobacteraceae</i> | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Rubritaleaceae</i>       | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Fusobacteriaceae</i>     | 0.0000  | 0.3155  | 0.0047  | 0.0000    | 0.0076  | 0.0000  |
| <i>Gemmatimonadaceae</i>    | 0.0000  | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Synergistaceae</i>       | 0.0000  | 0.0000  | 0.0016  | 0.0000    | 0.0000  | 0.0000  |

Supplementary Table S3. Percentage change at genus levels.

| Genus                    | Control  |          |          | Treatment |          |          |
|--------------------------|----------|----------|----------|-----------|----------|----------|
|                          | Donor 1  | Donor 2  | Donor 3  | Donor 1   | Donor 2  | Donor 3  |
| <i>Adlercreutzia</i>     | -0.0332  | -0.0061  | -0.0052  | 0.0401    | 0.0036   | 0.0094   |
| <i>Terrisporobacter</i>  | -0.0302  | -0.0056  | -0.0690  | 0.0424    | 0.0010   | 0.0673   |
| <i>Promicromonospora</i> | -0.5617  | -0.2125  | -1.5741  | 1.2150    | 0.3586   | 1.9098   |
| <i>Pseudonocardia</i>    | -0.7932  | -1.5335  | -1.4444  | 2.1024    | 3.2836   | 1.6846   |
| <i>Anaerostipes</i>      | -1.9218  | -0.1260  | -0.2782  | 3.0414    | 0.2375   | 0.2779   |
| <i>Akkermansia</i>       | -0.0042  | -0.0097  | -0.0197  | 0.0070    | 0.0272   | 0.0328   |
| <i>Alistipes</i>         | -0.0286  | -0.0466  | -0.8382  | 0.0316    | 0.0646   | 0.8999   |
| <i>Parabacteroides</i>   | -16.3417 | -15.4376 | -26.1212 | 13.5192   | 12.4244  | 23.1782  |
| <i>Ruminococcus 2</i>    | -1.4958  | -14.1600 | -18.5971 | 2.2853    | 10.2125  | 16.5730  |
| <i>Roseburia</i>         | -7.3000  | -0.1716  | -1.4926  | 5.0034    | 0.2746   | 1.4147   |
| <i>Megasphaera</i>       | -0.5343  | -1.5168  | -3.0740  | 1.0407    | 1.7551   | 3.6659   |
| <i>Weissella</i>         | -0.0064  | -0.0020  | -0.0249  | 0.0190    | 0.0056   | 0.0407   |
| <i>Neglecta</i>          | -6.7556  | 0.0000   | -0.06885 | 5.9144    | 0.0000   | 0.0965   |
| <i>Romboutsia</i>        | -1.0271  | -3.0153  | -1.3982  | 1.5801    | 2.2438   | 1.1298   |
| <i>Enterocloster</i>     | -3.8984  | -1.5035  | -3.2172  | 5.3252    | 2.6369   | 3.1128   |
| <i>Peptacetobacter</i>   | 0.0000   | -6.5689  | -0.7458  | 0.0000    | 12.1658  | 1.2034   |
| <i>Collinsella</i>       | -11.2280 | -2.7956  | -0.0431  | 6.8622    | 1.3445   | 0.0448   |
| <i>Paraclostridium</i>   | -0.0916  | -0.2868  | -0.8999  | 0.2199    | 0.7557   | 1.1106   |
| <i>Duncaniella</i>       | -1.7761  | -0.0102  | -1.2393  | 1.4411    | 0.0195   | 1.0158   |
| <i>Streptococcus</i>     | -0.8264  | -0.1352  | -0.0379  | 1.0430    | 0.1934   | 0.0423   |
| <i>Gillisia</i>          | -1.9951  | -1.2536  | -0.0099  | 2.1649    | 0.6211   | 1.0158   |
| <i>Lawsonibacter</i>     | 6.6491   | 0.0000   | 2.1445   | -6.5535   | 0.0000   | -2.1697  |
| <i>Proteus</i>           | 2.4677   | 17.6293  | 33.3981  | -2.4674   | -17.4997 | -33.3996 |
| <i>Butyricicoccus</i>    | 2.2504   | 1.2140   | 21.3012  | -1.8365   | -0.9277  | -20.3182 |

| Genus                        | Control  |          |         | Treatment |         |         |
|------------------------------|----------|----------|---------|-----------|---------|---------|
|                              | Donor 1  | Donor 2  | Donor 3 | Donor 1   | Donor 2 | Donor 3 |
| <i>Bifidobacterium</i>       | 0.0260   | 0.0025   | 0.0000  | -0.0225   | -0.0015 | -0.0005 |
| <i>Fusobacterium</i>         | 0.4532   | 2.0020   | 2.0617  | 0.2838    | -1.9637 | -1.9650 |
| <i>Flavonifractor</i>        | 0.0016   | 0.5628   | 0.0000  | -0.0011   | -0.5103 | 0.0000  |
| <i>Lachnospira</i>           | 0.0000   | -0.0307  | -0.2258 | 0.0000    | 0.0159  | 0.1893  |
| <i>Tyzzerella</i>            | 0.0000   | -0.0031  | -0.0862 | 0.0000    | 0.0056  | 0.0751  |
| <i>Holdemania</i>            | 0.0000   | 0.0000   | 0.1263  | 0.0000    | 0.0000  | -0.1285 |
| <i>Prevotella</i>            | 0.0000   | 0.0000   | -0.0114 | 0.0000    | 0.0005  | 0.0057  |
| <i>Mesorhizobium</i>         | 0.0000   | 0.0000   | -0.0078 | 0.0000    | 0.0000  | 0.0177  |
| <i>Phoea</i>                 | 0.0000   | 0.0000   | -0.0182 | 0.0000    | 0.0005  | 0.0302  |
| <i>Allisonella</i>           | 0.0005   | -22.2866 | -0.5276 | -0.0005   | 18.3060 | 0.5421  |
| <i>Acidaminococcus</i>       | -0.0087  | -0.0169  | -0.3078 | 0.0100    | 0.0210  | 0.3603  |
| <i>Coprococcus</i>           | -0.0001  | 0.0000   | 0.0000  | -0.0005   | 0.0000  | 0.0000  |
| <i>Gemella</i>               | -0.0432  | 0.0000   | 0.0000  | 0.0901    | 0.0000  | 0.0000  |
| <i>Longibaculum</i>          | 0.0000   | -0.0118  | -0.0145 | 0.0000    | 0.0031  | 0.0167  |
| <i>Rothia</i>                | 0.0000   | -5.7517  | -2.1885 | 0.0000    | 6.4206  | 2.8079  |
| <i>Sulfitobacter</i>         | 0.0000   | 0.0471   | 0.0000  | 0.0000    | -0.0455 | 0.0000  |
| <i>Lactiplantibacillus</i>   | 0.0000   | 0.0000   | -0.0109 | 0.0000    | 0.0000  | 0.0224  |
| <i>Bilophila</i>             | 0.0000   | -0.0118  | -0.0311 | 0.0000    | 0.0087  | 0.0235  |
| <i>Rubritalea</i>            | 3.9880   | 0.5789   | 3.2816  | -3.7752   | -0.5727 | -3.2968 |
| <i>Bacteroides</i>           | -0.0017  | 0.0000   | -0.0353 | 0.0015    | 0.0000  | 0.0746  |
| <i>Faecalibacterium</i>      | 0.0000   | -0.0015  | -0.3130 | 0.0000    | 0.0000  | 0.4395  |
| <i>Blautia</i>               | -0.0064  | 0.0000   | 0.0000  | 0.0045    | 0.0000  | 0.0000  |
| <i>Parasutterella</i>        | -0.0017  | 0.0000   | -0.0016 | 0.0015    | 0.0005  | 0.0010  |
| <i>Sphingobium</i>           | 0.0000   | 0.0000   | -0.0628 | 0.0000    | 0.0000  | 0.0464  |
| <i>Turicibacter</i>          | 0.0000   | -0.0046  | -0.0477 | 0.0005    | 0.0051  | 0.0542  |
| <i>Agathobacter</i>          | -0.0234  | -0.0082  | -0.0156 | 0.0726    | 0.0046  | 0.0162  |
| <i>Catenibacterium</i>       | 0.0000   | 0.0000   | -0.0036 | 0.0000    | 0.0000  | 0.0021  |
| <i>Intestinimonas</i>        | -0.0006  | -0.0031  | 0.0587  | 0.0000    | 0.0041  | -0.0508 |
| <i>Sutterella</i>            | 0.0000   | 0.0000   | -0.4565 | 0.0000    | 0.0000  | 0.7792  |
| <i>Nakamurella</i>           | -0.0834  | 0.0000   | -0.0389 | 0.0811    | 0.0000  | 0.0323  |
| <i>Eggerthella</i>           | -0.0006  | 0.0000   | 0.0052  | 0.0000    | 0.0000  | -0.0052 |
| <i>Odoribacter</i>           | -0.0017  | 0.0000   | -0.0358 | 0.0050    | 0.0005  | 0.0334  |
| <i>Odoribacter</i>           | 0.0000   | -0.0092  | -0.0462 | 0.0000    | 0.0041  | 0.0615  |
| <i>Dialister</i>             | 0-0.8265 | -0.4224  | -1.4672 | 0.9225    | 0.6484  | 1.7373  |
| <i>Enterococcus</i>          | 0.0000   | 0.0504   | -0.4240 | 0.0000    | -0.0463 | 0.5417  |
| <i>Phascolarctobacterium</i> | -0.0111  | -0.0026  | -0.0031 | 0.0140    | 0.0031  | 0.0068  |
| <i>Plesiomonas</i>           | -2.6932  | -1.3939  | -0.2154 | 3.2903    | 1.1242  | 0.2091  |
| <i>Roseisolibacter</i>       | 0.0056   | 0.0075   | -0.0026 | 0.0018    | -0.0116 | 0.0000  |
| <i>Dysosmobacter</i>         | -0.3448  | -0.0118  | 0.0000  | 0.3611    | 0.0169  | 0.0005  |

| Genus                                | Control |         |         | Treatment |         |         |
|--------------------------------------|---------|---------|---------|-----------|---------|---------|
|                                      | Donor 1 | Donor 2 | Donor 3 | Donor 1   | Donor 2 | Donor 3 |
| <i>Paraprevotella</i>                | 0.0000  | -2.2520 | -6.1622 | 0.0000    | 2.8584  | 6.4513  |
| <i>Solirubrobacter</i>               | 0.0000  | -0.0005 | -0.0130 | 0.0000    | -0.0005 | 0.0099  |
| <i>Faecalibacillus</i>               | 0.0000  | -0.1301 | -0.0903 | 0.0000    | 0.2411  | 0.1330  |
| <i>Dielma</i>                        | 0.0000  | 0.0000  | -0.0031 | 0.0000    | 0.0000  | 0.0016  |
| <i>Morganella</i>                    | -0.7788 | -0.0246 | -0.2979 | 0.7933    | 0.0380  | 0.3003  |
| <i>Veillonella</i>                   | -0.6101 | -0.1761 | -0.4411 | 0.3576    | 0.1518  | 0.5657  |
| <i>Oscillibacter</i>                 | -0.2183 | 0.0000  | -0.0047 | 0.3368    | 0.0000  | 0.0042  |
| <i>Escherichia/Shigella</i>          | -0.0012 | 0.0000  | -0.0016 | 0.0010    | 0.0000  | 0.0010  |
| <i>Anaerotruncus</i>                 | 0.0320  | -0.0719 | -0.0343 | -0.0238   | 0.1024  | 0.0391  |
| <i>Flintibacter</i>                  | -0.0006 | 0.0000  | -0.0078 | 0.0000    | 0.0005  | 0.0162  |
| <i>Eisenbergiella</i>                | 0.0000  | -0.0015 | -0.0042 | 0.0000    | 0.0010  | 0.0026  |
| <i>Senegalimassilia</i>              | 0.0000  | 0.0000  | -0.0031 | 0.0000    | 0.0000  | 0.0052  |
| <i>Clostridium XVIII</i>             | 0.0000  | 0.0000  | -0.0052 | 0.0000    | 0.0000  | 0.0052  |
| <i>Sediminicola</i>                  | -0.0017 | 0.0000  | -0.8745 | 0.0145    | 0.0000  | 1.0245  |
| <i>Lactacaseibacillus</i>            | 0.0000  | 0.0000  | 0.0000  | 0.0085    | 0.0000  | 0.0000  |
| <i>Anaerofilum</i>                   | 0.0000  | 0.1451  | 0.0000  | 0.0000    | -0.1431 | 0.0000  |
| <i>Schaalia</i>                      | 0.0000  | -0.0010 | -0.0016 | 0.0005    | 0.0000  | 0.0000  |
| <i>Lachnospiracea incertae sedis</i> | 0.0000  | -0.0031 | -1.0255 | 0.0000    | 0.0036  | 0.9057  |
| <i>Burkholderia</i>                  | 0.0000  | -0.0154 | -0.5097 | 0.0000    | 0.0185  | 0.6846  |
| <i>Clostridium IV</i>                | 0.0000  | 0.0000  | -0.0042 | 0.0000    | 0.0000  | 0.0047  |
| <i>Lactonifactor</i>                 | 0.0000  | 0.0000  | -0.0119 | 0.0000    | 0.0000  | 0.0182  |
| <i>Gemmiger</i>                      | -0.0087 | -0.0246 | -0.6161 | 0.0170    | 0.0339  | 0.5673  |
| <i>Haemophilus</i>                   | 0.0174  | -0.0056 | 5.2598  | -0.0178   | 0.0077  | -5.2592 |
| <i>Phocaeicola</i>                   | 0.0000  | 0.0000  | -0.0602 | 0.0000    | 0.0000  | 0.0454  |
| <i>Eubacterium</i>                   | 0.0000  | -0.0184 | -0.0036 | 0.0000    | 0.0154  | 0.0115  |
| <i>Amedibacillus</i>                 | -0.0001 | 0.0000  | 0.0000  | 0.0165    | 0.0000  | 0.0000  |
| <i>Finegoldia</i>                    | -0.5827 | -0.0502 | -0.1474 | 0.5148    | 0.0308  | 0.0673  |
| <i>Clostridium sensu stricto</i>     | 0.0000  | 0.0480  | 0.3138  | 0.0000    | -0.0475 | -0.3230 |
| <i>Megamonas</i>                     | 1.3008  | 0.0005  | 0.0005  | -1.3015   | -0.0005 | -0.0005 |
| <i>Ruminococcus</i>                  | 0.0000  | 0.0000  | 0.0000  | 0.0090    | 0.0000  | 0.0000  |
| <i>Sellimonas</i>                    | 0.0031  | 0.0051  | 0.0000  | -0.0031   | -0.0051 | 0.0000  |
| <i>Winogradskyella</i>               | 0.0000  | 0.0687  | 0.0000  | 0.0000    | -0.0698 | 0.0000  |
| <i>Massiliimalia</i>                 | 0.0000  | 0.0000  | 0.0000  | 0.0611    | 0.0000  | 0.0000  |
| <i>Providencia</i>                   | 0.0000  | 0.0000  | -0.0254 | 0.0000    | 0.0000  | 0.0261  |
| <i>Intestinibacter</i>               | -0.0140 | -0.0041 | 0.0000  | 0.0135    | 0.0000  | 0.0000  |
| <i>Anaerotignum</i>                  | -0.0087 | 0.0000  | 0.0000  | 0.0000    | 0.0000  | 0.0000  |
| <i>Raoultibacter</i>                 | 0.0000  | 0.0000  | -2.9624 | 0.0000    | 0.0000  | 3.4849  |
| <i>Anaerobutyricum</i>               | -0.0023 | -0.0010 | -0.2242 | 0.0000    | 0.0000  | 0.2398  |

| Genus                         | Control |         |         | Treatment |         |         |
|-------------------------------|---------|---------|---------|-----------|---------|---------|
|                               | Donor 1 | Donor 2 | Donor 3 | Donor 1   | Donor 2 | Donor 3 |
| <i>Cloacibacillus</i>         | -0.0321 | -0.0077 | -0.0083 | 0.0761    | 0.0077  | 0.0141  |
| <i>Dorea</i>                  | -0.0006 | -0.0016 | 0.0000  | 0.0015    | -0.0015 | 0.0000  |
| <i>Ruthenibacterium</i>       | -0.0052 | 0.0000  | 0.0000  | 0.0100    | 0.0000  | 0.0000  |
| <i>Hungatella</i>             | 0.0000  | -0.2628 | -0.2694 | 0.0000    | 0.2151  | 0.2884  |
| <i>Lactococcus</i>            | 0.0000  | -0.1470 | 0.0000  | 0.0000    | 0.1211  | 0.0000  |
| <i>Mediterraneibacter</i>     | -0.8749 | -0.6175 | -1.3193 | 1.0552    | 0.8454  | 1.0652  |
| <i>Fusicatenibacter</i>       | 0.0000  | 0.0000  | 0.0000  | 0.0300    | 0.0000  | 0.0000  |
| <i>Pediococcus</i>            | 0.0000  | 0.0000  | 0.0000  | 0.0336    | 0.0000  | 0.0000  |
| <i>Actinomyces</i>            | 0.0000  | 0.0000  | 0.0000  | 0.0125    | 0.0000  | 0.0000  |
| <i>Negativibacillus</i>       | 0.0000  | -0.0020 | -0.0062 | 0.0000    | 0.0015  | 0.0021  |
| <i>Desulfovibrio</i>          | 0.0000  | -0.1341 | -0.0109 | 0.0000    | 0.0898  | 0.0099  |
| <i>Erysipelatoclostridium</i> | -0.0010 | 0.0045  | 0.0005  | 0.0064    | -0.0070 | -0.0016 |
| <i>Citrobacter</i>            | 0.0000  | -0.1070 | -0.0042 | 0.0005    | 0.0539  | 0.0052  |
| <i>Slackia</i>                | -0.0105 | 0.0000  | -1.0161 | 0.0050    | 0.0000  | 1.1035  |
| <i>Faecalimonas</i>           | 0.0198  | -0.0072 | 0.4068  | 0.0207    | 0.0021  | -0.4068 |
| <i>Butyricimonas</i>          | 0.0010  | 0.0025  | -0.0010 | -0.0011   | -0.0030 | 0.0000  |