| Gene  | Group | Gene  | Group | Gene  | Group | Gene  | Group |
|-------|-------|-------|-------|-------|-------|-------|-------|
| abrB  | 1     | rpeE  | 1     | yflH  | 3     | pthC  | 1     |
| rplL  | 1     | rpmD  | 1     | yflD  | 3     | ylaI  | 4     |
| rplM  | 1     | rplO  | 1     | cspB  | 1     | ylaJ  | 5     |
| rplN  | 1     | rplM  | 1     | yncN  | 2     | divIVA| 3     |
| rplO  | 1     | rplO  | 1     | prsA  | 1     | rpmB  | 1     |
| rplP  | 1     | yfbQ  | 3     | yflD  | 1     | flhJ  | 2     |
| rplP  | 1     | ycdA  | 3     | cotW  | 3     | cotE  | 5     |
| rplP  | 1     | ycmE  | 1     | ykwD  | 1     | ynzH  | 2     |
| rplF  | 1     | ydcN  | 4     | ykuT  | 5     | ynzC  | 3     |
| rplR  | 1     | ydcN  | 2     | ykzG  | 2     | yonK  | 2     |
| ypzA  | 4     | eno   | 1     | rpsT  | 1     | cotG  | 1     |
| yokF  | 2     | yttA  | 3     | cspD  | 1     | ytlB  | 4     |
| ypeD  | 5     | yvcE  | 1     | fer   | 1     | yvzB  | 1     |
| yscF  | 5     | ywhB  | 3     | rpmI  | 1     | yxeE  | 3     |
| yscB  | 5     | alpC  | 1     | rpsD  | 1     |        |       |

Table 3: Repartition of the genes employed to gauge the Codon Adaptation Index [5], as in [17], among the clusters identified in *B. subtilis*. Note the highly significant concentration in the first cluster. Genes used to gauge the CAI index for *E. coli* are all concentrated in the first cluster.