

Genus	Average	Median	Source
Acetanaerobacterium	7.7	8.0	rmDB [O]
Acetivibrio	7.7	8.0	rmDB [O]
Acidaminobacter	7.7	8.0	rmDB [O]
Acidaminococcus	7.7	8.0	rmDB [O]
Acinetobacter	6.2	6.0	rmDB [G]
Actinomyces	3.0	3.0	rmDB [G]
Adlercreutzia	2.3	3.0	rmDB [F]
Akkermansia	3.0	3.0	rmDB [G]
Alistipes	5.8	6.0	rmDB [O]
Allisonella	7.7	8.0	rmDB [O]
Allobaculum	6.3	6.0	rmDB [P]
Anaerococcus	7.7	8.0	rmDB [O]
Anaerofilum	7.7	8.0	rmDB [O]
Anaerofustis	5.0	5.0	rmDB [F]
Anaerophaga	6.0	6.0	rmDB [F]
Anaerosinus	7.7	8.0	rmDB [O]
Anaerospobacter	7.7	8.0	rmDB [O]
Anaerostipes	7.7	8.0	rmDB [O]
Anaerotruncus	7.7	8.0	rmDB [O]
Anaerovirgula	7.7	8.0	rmDB [O]
Anaerovorax	7.7	8.0	rmDB [O]
Bacteroides	6.0	6.0	rmDB [G]
Barnesiella	5.5	5.5	rmDB [F]
Bifidobacterium	3.6	4.0	rmDB [G]
Blautia	7.7	8.0	rmDB [O]
Brachybacterium	3.0	3.0	rmDB [G]
Brachyspira	1.0	1.0	rmDB [G]
Bradyrhizobium	1.3	1.0	rmDB [G]
Bryantella	9.0	9.0	rmDB [F]
Bulleidia	6.3	6.0	rmDB [P]
Butyricoccus	7.7	8.0	rmDB [O]
Butyricimonas	5.5	5.5	rmDB [F]
Butyrivibrio	7.7	8.0	rmDB [O]
Caenispirillum	3.3	4.0	rmDB [F]
Caloranaerobacter	9.0	9.0	rmDB [F]
Caminicella	9.0	9.0	rmDB [F]
Campylobacter	2.9	3.0	rmDB [G]
Catenibacterium	6.3	6.0	rmDB [P]
Cetobacterium	5.0	5.0	rmDB [F]
Clostridiisalibacter	9.0	9.0	rmDB [F]
Clostridium	9.0	9.0	rmDB [F]
Collinsella	2.3	3.0	rmDB [F]
Coprobacillus	6.3	6.0	rmDB [P]
Coprococcus	7.7	8.0	rmDB [O]
Corynebacterium	4.6	5.0	rmDB [G]
Cupriavidus	5.0	5.0	rmDB [G]
Dehalobacter	5.3	5.0	rmDB [F]
Desulfifibacter	5.3	5.0	rmDB [F]
Desulfitobacterium	6.0	6.0	rmDB [G]
Desulfovibrio	4.5	4.5	rmDB [G]
Dialister	7.7	8.0	rmDB [O]
Dorea	7.7	8.0	rmDB [O]
Eggerthella	3.0	3.0	rmDB [G]
Enterobacter	7.0	7.0	rmDB [G]
Enterococcus	4.0	4.0	rmDB [G]
Escherichia	7.0	7.0	rmDB [G]
Ethanoligenens	7.7	8.0	rmDB [O]
Eubacterium	5.0	5.0	rmDB [G]
Faecalibacterium	7.7	8.0	rmDB [O]
Fusibacter	7.7	8.0	rmDB [O]
Fusobacterium	5.0	5.0	rmDB [G]
Garciella	5.0	5.0	rmDB [F]
Gemella	5.3	5.0	rmDB [F]
Geopsychrobacter	2.4	2.0	rmDB [F]
Geosporobacter	9.0	9.0	rmDB [F]
Gordonibacter	2.3	3.0	rmDB [F]
Haemophilus	5.8	6.0	rmDB [G]
Haloplasma	1.7	2.0	rmDB [C]
Helicobacter	2.0	2.0	rmDB [G]
Herbaspirillum	2.0	2.0	rmDB [F]
Hespella	9.0	9.0	rmDB [F]
Holdemania	1.0	1.0	NCBI
Howardella	7.7	8.0	rmDB [O]
Hydrogenoanaerobacterium	7.7	8.0	rmDB [O]

Genus	Average	Median	Source
Kerstersia	3.0	3.0	rmDB [F]
Kiloniella	2.3	2.0	rmDB [C]
Klebsiella	8.0	8.0	rmDB [G]
Lachnobacterium	7.7	8.0	rmDB [O]
Lachnospira	7.7	8.0	rmDB [O]
Lactobacillus	5.4	5.0	rmDB [G]
Lactococcus	6.0	6.0	rmDB [G]
Lactonifactor	7.7	8.0	rmDB [O]
Laribacter	4.4	4.0	rmDB [F]
Leptonema	2.0	2.0	rmDB [G]
Leptospira	2.0	2.0	rmDB [G]
Limnobacter	4.9	5.0	rmDB [F]
Lutispora	7.7	8.0	rmDB [O]
Mahella	3.2	3.5	rmDB [F]
Marvinbryantia	7.7	8.0	rmDB [O]
Megasphaera	7.7	8.0	rmDB [O]
Mesorhizobium	2.0	2.0	rmDB [G]
Methylobacterium	5.5	5.0	rmDB [G]
Microvirgula	4.4	4.0	rmDB [F]
Mitsuokella	7.7	8.0	rmDB [O]
Mogibacterium	7.7	8.0	rmDB [O]
Neisseria	4.0	4.0	rmDB [G]
Novosphingobium	3.0	3.0	rmDB [G]
Odoribacter	5.5	5.5	rmDB [F]
Oribacterium	7.7	8.0	rmDB [O]
Oscillibacter	7.7	8.0	rmDB [O]
Oxalobacter	2.0	2.0	rmDB [F]
Oxobacter	9.0	9.0	rmDB [F]
Pannonibacter	2.9	3.0	rmDB [F]
Papillibacter	7.7	8.0	rmDB [O]
Parabacteroides	7.0	7.0	rmDB [G]
Paraprevotella	5.8	6.0	rmDB [O]
Parasporobacterium	9.0	9.0	rmDB [F]
Parasutterella	3.0	3.0	rmDB [F]
Parvimonas	7.7	8.0	rmDB [O]
Paucimonas	4.9	5.0	rmDB [F]
Pelotomaculum	2.0	2.0	rmDB [G]
Peptococcus	5.3	5.0	rmDB [F]
Peptostreptococcus	7.7	8.0	rmDB [O]
Phascolarctobacterium	7.7	8.0	rmDB [O]
Prevotella	5.8	6.0	rmDB [O]
Prosthecomicrobium	2.7	3.0	rmDB [F]
Pseudobutyrvibrio	7.7	8.0	rmDB [O]
Ralstonia	4.4	4.0	rmDB [G]
Raoultella	6.5	7.0	rmDB [F]
Rhodanobacter	2.3	2.0	rmDB [F]
Rikenella	5.8	6.0	rmDB [O]
Robinsoniella	7.7	8.0	rmDB [O]
Roseburia	7.7	8.0	rmDB [O]
Roseospirillum	2.7	3.0	rmDB [O]
Rothia	2.7	3.0	rmDB [C]
Ruminococcus	7.7	8.0	rmDB [O]
Shigella	7.0	7.0	rmDB [G]
Slackia	2.3	3.0	rmDB [F]
Solobacterium	6.3	6.0	rmDB [P]
Sporacetigenium	7.7	8.0	rmDB [O]
Sporobacter	7.7	8.0	rmDB [O]
Staphylococcus	5.4	5.0	rmDB [G]
Streptococcus	5.4	6.0	rmDB [G]
Subdoligranulum	7.7	8.0	rmDB [O]
Succiniclasticum	7.7	8.0	rmDB [O]
Succinivibrio	9.5	9.5	rmDB [O]
Sutterella	3.0	3.0	rmDB [F]
Syntrophococcus	5.3	6.0	rmDB [O]
Tannerella	5.5	5.5	rmDB [F]
Terasakiella	2.7	3.0	rmDB [O]
Thermotalea	9.0	9.0	rmDB [F]
Turicibacter	6.3	6.0	rmDB [P]
Veillonella	7.7	8.0	rmDB [O]
Verrucomicrobium	3.0	3.0	rmDB [F]
Victivallis	3.0	3.0	NCBI
Wautersia	4.9	5.0	rmDB [F]
Yersinia	6.9	7.0	rmDB [G]
Zymophilus	7.7	8.0	rmDB [O]

Source:
 rmDB - rmDB database
 [G] - genus level statistics
 [F] - family level statistics
 [O] - order level statistics
 [C] - class level statistics
 NCBI - NCBI microbial genomes