

Table S1. Detected genera that are linked to butyrate producers. For functional gene IDs associated with individual taxa see fungene (<http://fungene.cme.msu.edu>) and Vital et al (2013). The applied gene-association should not be regarded as strict for those taxa as several genera contain both *but* and *buk* as well as non-butyrate producing members (see main text). Thus, this type of analysis cannot substitute for a functional gene-targeted approach, but is used here for global comparisons. Individual (gene) categories are indicated: butyryl-CoA:acetate CoA-transferase (*but*), butyrate kinase (*buk*), alternative transferases (*other*) and amino acids fed butyrate-producing pathways (*aa*). Brachyspir. (*Brachyspiraceae*), Clost.IS_XI (*Clostridiales_Incertae_Sedis_XI*), Oxalobact. (*Oxalobacteraceae*).

<p>but</p> <p>Lachnospiraceae <i>Clostridium XIVa</i> <i>Roseburia</i> <i>Anaerostipes</i> <i>Butyrivibrio</i> <i>Pseudobutyrvibrio</i> <i>Lachnospiraceae_incertae_sedis</i></p> <p>Ruminococcaceae <i>Butyricoccus</i> <i>Anaerococcus</i> <i>Faecalibacterium</i></p> <p>Acidaminococcaceae <i>Acidaminococcus</i></p> <p>Veillonellaceae <i>Megasphaera</i></p> <p>Eubacteriaceae <i>Anaerofustis</i> <i>Eubacterium</i> <i>Pseudaminobacter</i></p> <p>Spirochaetaceae <i>Treponema</i></p> <p>Brachyspir./Clost.IS_XI/Oxalobact. <i>Peptoniphilus</i> <i>Brachyspira</i> <i>Oscillibacter</i></p>	<p>other</p> <p>Erysipelotrichaceae <i>Clostridium XVIII</i> <i>Erysipelotrichaceae_incertae_sedis</i></p> <p>Carnobacteriaceae <i>Carnobacterium</i></p> <p>buk</p> <p>Clostridiaceae <i>Clostridium sensu stricto</i></p> <p>Peptostreptococcaceae <i>Clostridium XI</i></p> <p>Ruminococcaceae <i>Subdoligranulum</i> <i>Anaerotruncus</i></p> <p>Lachnospiraceae <i>Coprococcus</i> <i>Shuttleworthia</i></p> <p>aa</p> <p>Porphyromonadaceae <i>Odoribacter</i> <i>Porphyromonas</i> <i>Butyricimonas</i> <i>Propionibacterium</i></p> <p>Fusobacteriaceae <i>Fusobacterium</i></p> <p>Rikenellaceae <i>Alistipes</i></p>
---	---

Table S2. Main ingredients and major enrichments of animal diets.

SampleID	Diet	Main Ingredients	Major Enrichments
BaldEagle	C	Natural Balance® Meat eating bird diet, fish, mice, rat, rabbit	
Duck	H	Corn, scratch, Mazuri® Waterfowl Maintenance	Crickets
Mousebird	H	Fruit and Mazuri® Softbill	
Hawk	C	Nebraska Brand® Bird of Prey diet	Prey items
Peafowl	O	Corn, scratch, Mazuri® Waterfowl Maintenance, bugs, worms	Produce, bugs, worms
KingVulture	C	Fish, Nebraska Brand® Premium Meat, rabbit	Raw eggs
Penguin	C	Fish	
Ostrich	O	Ratite pellets	Produce, corn, starch, browsw
Tenrec1	C	Insecticore diet, meal worms, cricket, Science Diet® Feline Adult Lite, apple	
Tenrec2	C	Insecticore diet, meal worms, cricket, Science Diet® Feline Adult Lite, apple	
GuineaHog	H	Grass, hay, Mazuri® Mini-pig Elder	Omolene® Sweet Feed, oats, scratch, mealworms, eggs, bread, cereal
Dog1	O	Wellness® natural dry dog food	
Dog2	O	Wellness® natural dry dog food	
ArcticFox1	C	Natural Balance® Carnivore Diet, Science Diet® K-9 Small Bites dry, rib bone, dog food	Frozen fish, live fish, deer meat, eggs, dog treats, peanut butter tubes
ArcticFox2	C	Natural Balance® Carnivore Diet, Science Diet® K-9 Small Bites dry, rib bone, dog food	Frozen fish, live fish, deer meat, eggs, dog treats, peanut butter tubes
AfricanLion	C	Carnivore diet, rib bone, femur bone	Herring, rabbits, venison, chicken, melon, pumpkin
Tiger	C	Nebraska Brand® Exotic Feline Diet	
Meerkat	O	Insectivore diet, cat food, crickets, mealworms, hard boiled eggs, fruit, veggies, greens, fuzzy mice	Nuts, seeds, grains, cereal, pasta, dried fruit, honey
Ferret	C	Meat, eggs, pinky mice, bugs	
RiverOtter	C	Fish, Science Diet® Active Longevity	Earthworms, mice, carrots, apples, pumpkin
RedPanda	H	Banana, apple, Marion™ Leafeater diet, Marion™ Leafeater Lemur Biscuits, bamboo	
GrizzlyBear	C	Mazuri® Omnivore 5635, produce, bones	
GiantPanda1	H	Bamboo	
GiantPanda2	H	Bamboo	
Binturong	O	Mazuri® Leaf Eater Sticks, dry dog food, banana, fruit, half egg	
Armadillo	O	Insectivore pellets, potato, apple, grape, waxworm	Crickets
Opossum	O	Veggies, fruit, Science Diet® Canine Small Bites, adult mouse whole	Produce
Rabbit1	H	Oxbow basix bunny/T pellets, hay, fresh produce	
Rabbit2	H	Oxbow basix bunny/T pellets, hay, fresh produce	
Donkey	H	Hay, mineral pellet	
CottontopTamarin	O	Mazuri® Marmoset Jelly, banana, Mazuri® High Protein Primate Stick, fruit, bugs	
GoldenLionTamarin	O	Tamarin diet, waxworms, mealworms	Peanut butter, honey, dried fruit, cereal, cooked pasta, popcorn, yogurt
Human1	O	North american omnivore diet	
Human2	O	North american omnivore diet	
Mandril	O	Mazuri® Primate Browse Biscuits, glucosamine	
MongooseLemur	O	Mazuri® Primate Browse Biscuits, greens	<i>Peanut butter, honey, dried fruit, cereal, cooked pasta, popcorn, yogurt,</i>
RedruffeledLemur	O	Mazuri® Primate Browse Biscuits, greens	<i>crickets, earthworms, bread, mealworms, waxworms</i>
RingtailedLemur	O	Mazuri® Primate Browse Biscuits, greens, browse, flowers, fruit, veggies	<i>(for all Lemurs and SpiderMonkey)</i>
SpiderMonkey	O	Marion™ Leafeater Lemur Biscuit, Mazuri® New World Primate diet, fruit, veggie	
GuineaPig	H	Hay, Kaytee® Timothy Complete Fiber diet	
PatagonianCavy	H	Grass, hay, Mazuri® Primate Browse Biscuits, greens, veggies, guinea pig diet, seeds, scratch, rodent chow, fruit	
Chinchilla	H	Hay, Mazuri® Chin Chow, cheerios, vegetables, fresh and dried fruit, grains	
Porcupine	H	Mazuri® Primate Browse Biscuits, Science Diet® Active Longevity	Browse, pumpkins, melons, corn stalks, corn on the cob, scents, spices
Mouse	O	Kaytee® healthy support diet	
Rat	O	Kaytee® healthy support diet	
Treeshrew	O	Insectivore diet, cat food, mixed fruit, veggies, bugs	
BeardedDragon	O	Greens topped with fruit or vegetables, insects	Fuzzy mice
MilkSnake	C	Dead Fuzzies (mice)	
RatSnake	C	Dead Fuzzies (mice)	
Chuckwalla	H	Greens, carrot, potato, beet, and seed	
Skink	O	Produce, crickets, worms, pinky mouse	
BoxTurtle	O	Fruit, vegetables, earthworms, crickets	
Tortoise1	H	Diced greens, carrot, potato, tortoise chow pellets, hay/grass mix	Fruits
Tortoise2	H	Diced greens, carrot, potato, tortoise chow pellets, hay/grass mix	Fruits

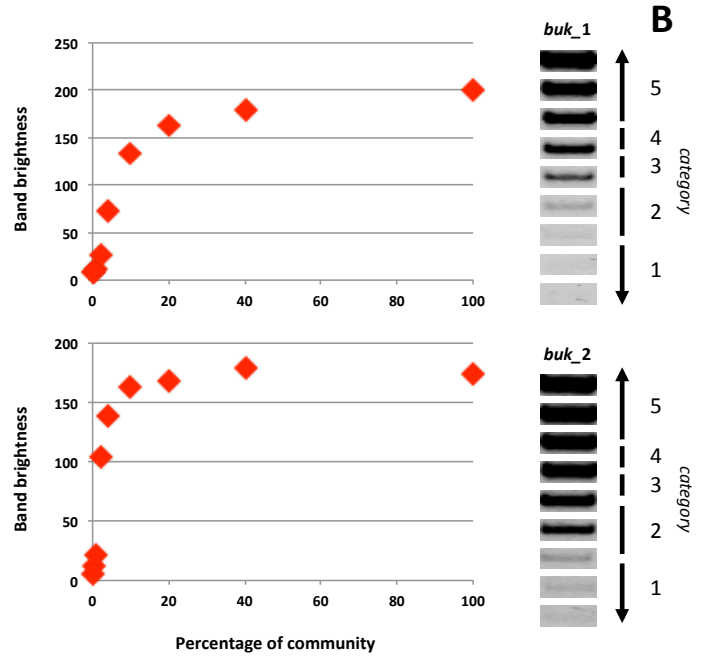
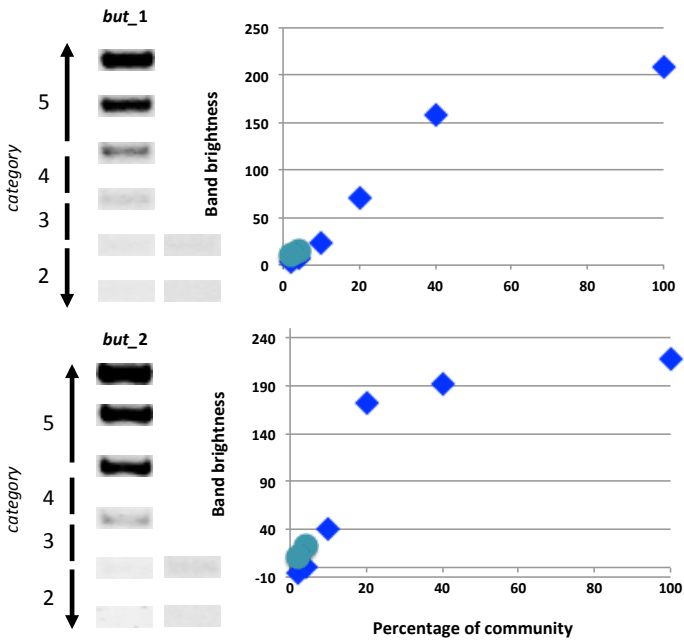
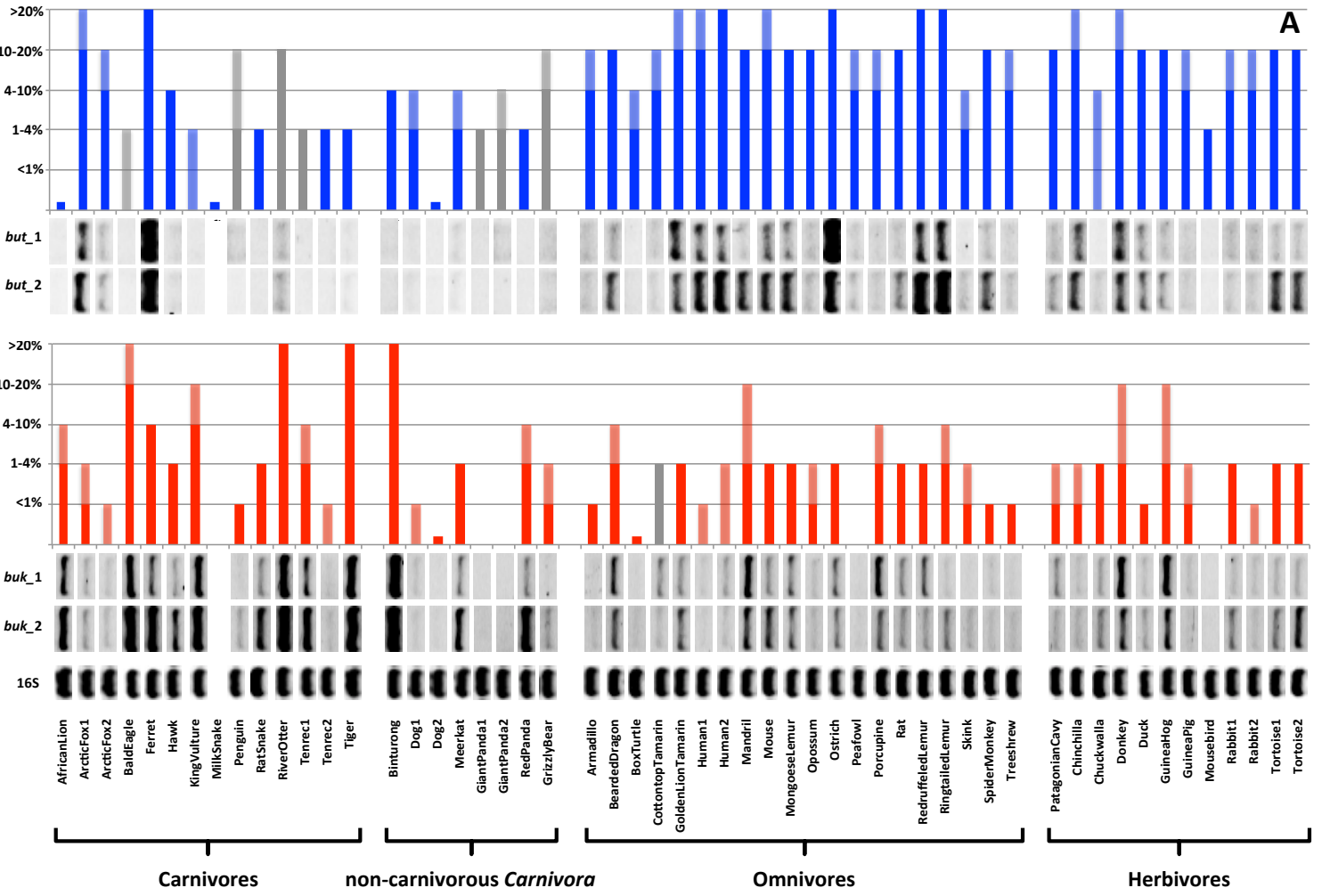


Figure S1. Estimating gene abundances of butyryl-CoA:acetate CoA-transferase (blue) and butyrate kinase (red) for all samples (panel A). Band intensities of PCR products were categorized into six distinct brightness groups (0-5) based on established standard curves from reference genomes (shown in panel B; gel background was subtracted from intensity values). Two experiments were performed for each gene. In order to increase detection confidence for *but* at low intensities two reference samples were analyzed at lowest concentrations and median values were used for calculations. Intensities of samples that were categorized as zero, but yielded sequences after re-amplification, are indicated as well (bars exhibiting 0.2 intensity). Samples where the majority of reads ($\geq 97\%$) were filtered out during the pipelining (see material and methods) are marked grey (also grizzly bear is marked grey as only 12 % of reads were considered as real *but*). For more details see materials and methods. No data is available for milk snake. Since no bands were visible for both genes in previous experiments (16S *rRNA* amplification displayed a bright band) we categorized estimated gene abundance as 0 % for both genes in this animal.

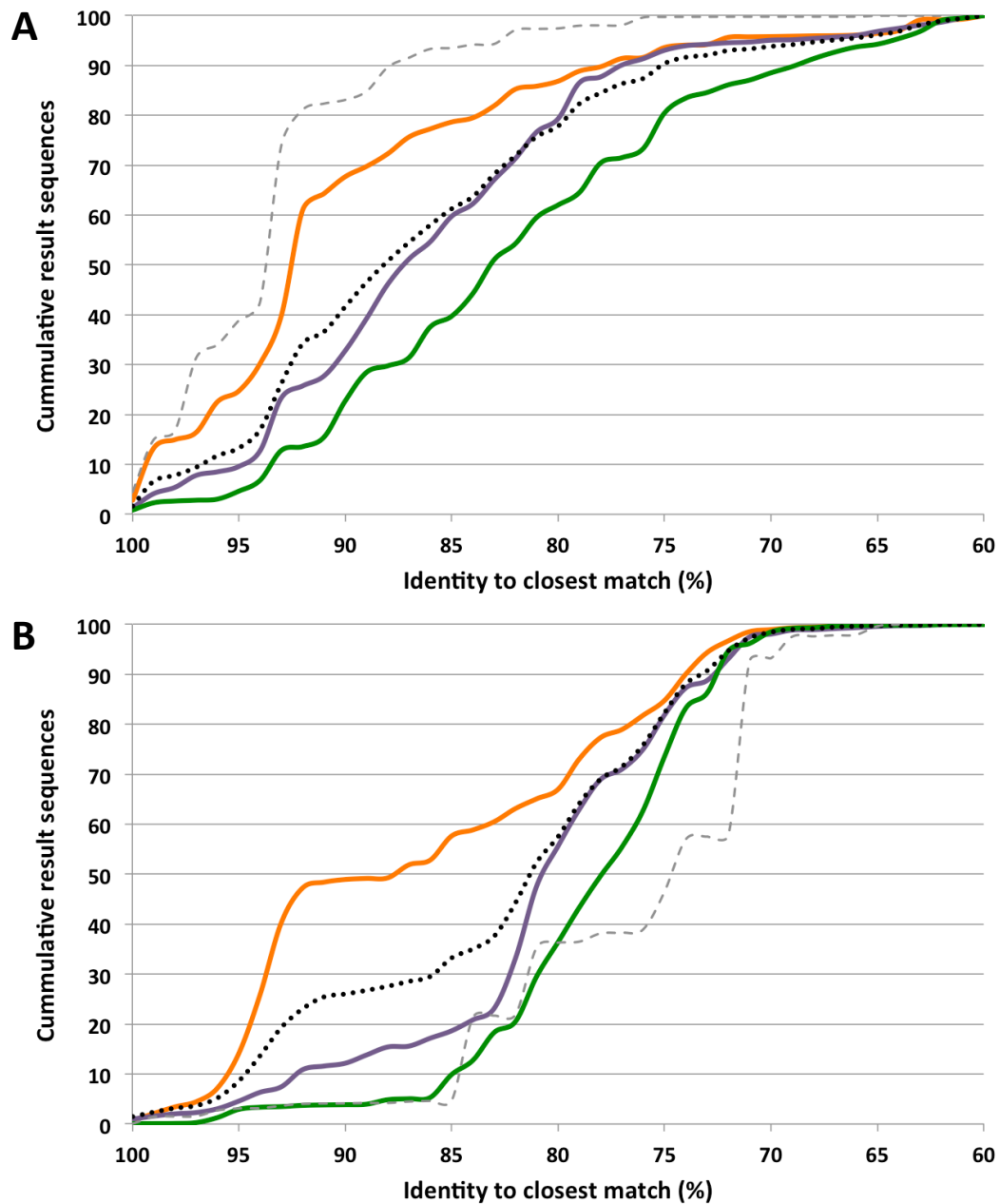


Figure S2. Identity values of all obtained sequences to a reference (on protein level) for butyryl-CoA:acetate CoA-transferase (A) and butyrate kinase (B) is shown. Individual diet groups are indicated as orange (carnivores including non-carnivorous *Carnivora*), violet (omnivores), green (herbivores), grey-dashed line (human) and black-dotted line (all groups together). The data is displayed as a cumulative percentage of sequence identities to a reference gene in our database.

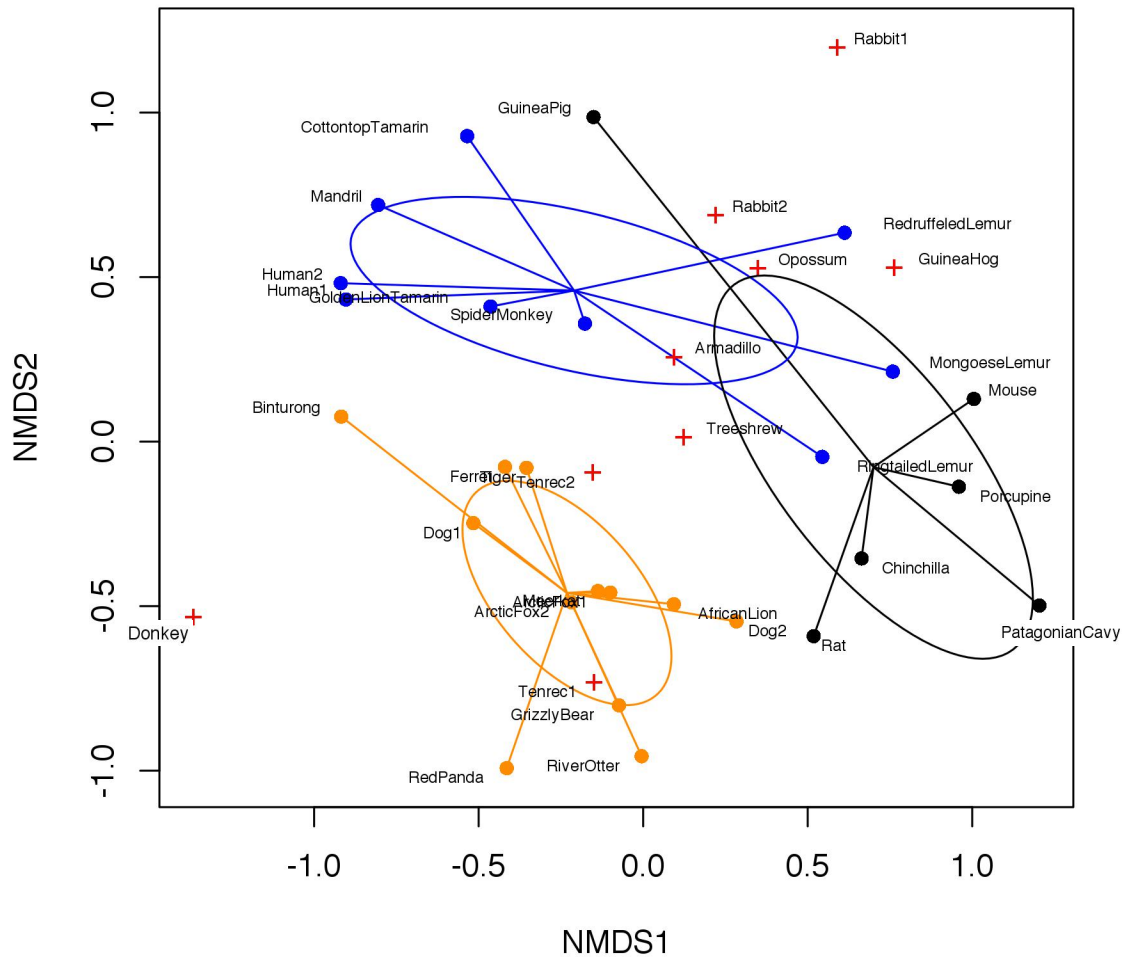
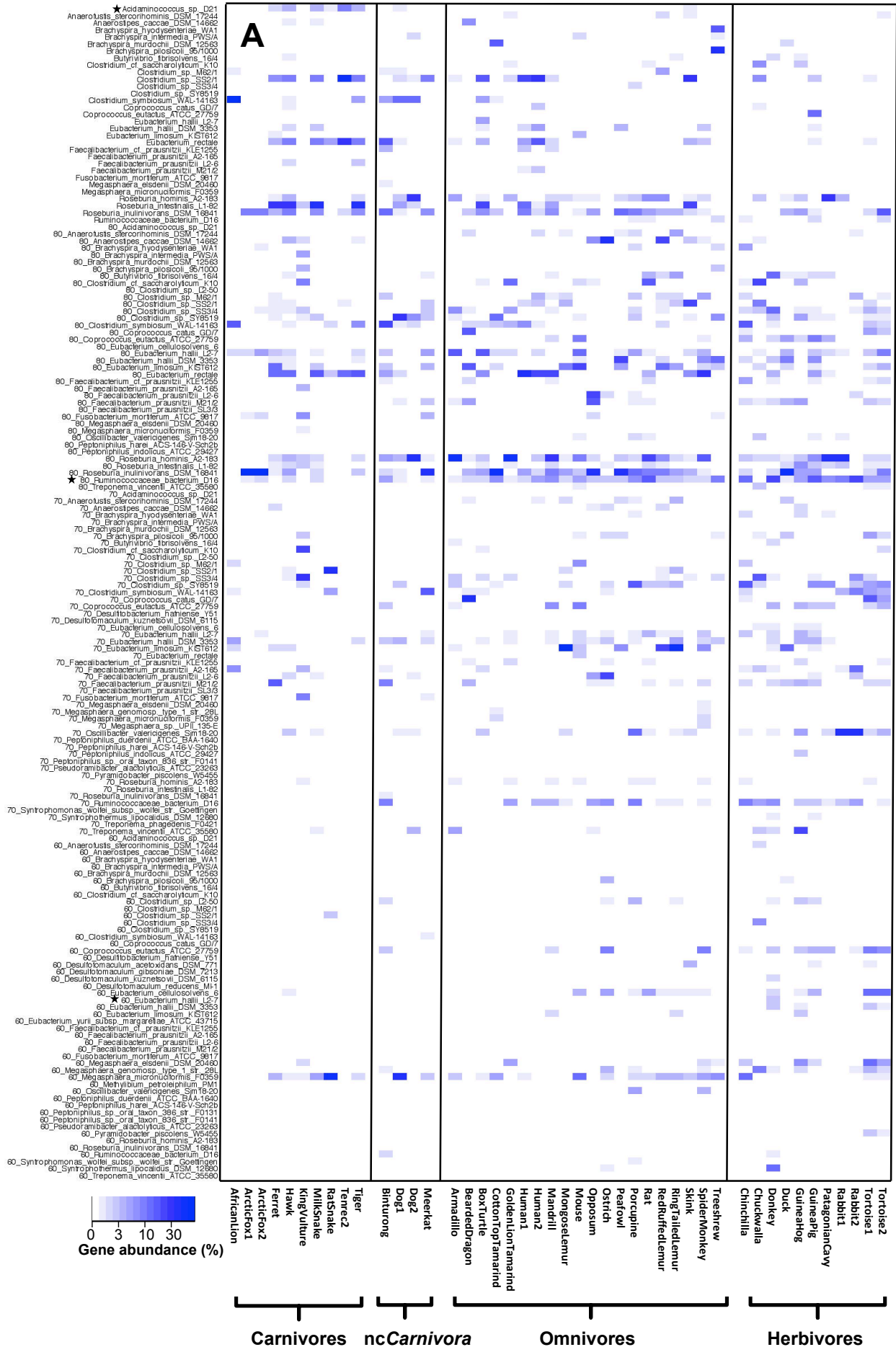
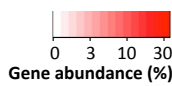


Figure S3. Non-metric multidimensional scaling (NMDS) of combined gene data (butyryl-CoA:acetate CoA-transferase and butyrate kinase) based on multi-linkage clustering (5% cut-off on protein level) for *Mammalia* only is shown. Individual orders are indicated as orange (*Carnivora*), blue (*Primates*) and black (*Rodentia*). Other subjects are displayed as well.



- Clostridium_acetobutylicum
- Clostridium_bartlettii_DSM_16795
- Clostridium_beijerinckii_NCIMB_8052
- Clostridium_botulinum_B_str_Eklund_17B
- Clostridium_botulinum_H04402_065
- Clostridium_butyricum
- Clostridium_cellulovorans_743B
- Clostridium_hiranonis_DSM_13275
- ★ Clostridium_perfringens
- Clostridium_sp._L2-50
- Clostridium_tetani_E88
- Coproccoccus_sp._ART55/1
- Eubacterium_ventriosum_ATCC_27560
- Lachnospiraceae_bacterium_3_1_57FAA_CT1
- Subdoligranulum_variabile_DSM_15176
- 80_Butyrvibrio_crossotus_DSM_2876
- 80_Butyrvibrio_proteoclasticus_B316
- 80_Clostridium_bartlettii_DSM_16795
- 80_Clostridium_beijerinckii_NCIMB_8052
- 80_Clostridium_botulinum_A3_str_Loch_Maree
- 80_Clostridium_botulinum_B_str_Eklund_17B
- 80_Clostridium_botulinum_E1_str_BoNT_E_Beluga
- 80_Clostridium_botulinum_F_str_230613
- 80_Clostridium_botulinum_H04402_065
- 80_Clostridium_butyricum
- 80_Clostridium_carboxidivorans_P7
- ★ 80_Clostridium_difficile_R20291
- 80_Clostridium_hiranonis_DSM_13275
- ★ 80_Clostridium_perfringens
- 80_Clostridium_sp._L2-50
- 80_Clostridium_sporogenes_ATCC_15579
- 80_Clostridium_tetani_E88
- 80_Coproccoccus_sp._ART55/1
- 80_Eubacterium_ventriosum_ATCC_27560
- 80_Lachnospiraceae_bacterium_3_1_57FAA_CT1
- 80_Peptostreptococcus_stomatis_DSM_17678
- 80_Shuttleworthia_satelles_DSM_14600
- 80_Subdoligranulum_variabile_DSM_15176
- 70_Butyrvibrio_crossotus_DSM_2876
- 70_Butyrvibrio_proteoclasticus_B316
- 70_Clostridium_acetobutylicum
- 70_Clostridium_bartlettii_DSM_16795
- 70_Clostridium_beijerinckii_NCIMB_8052
- 70_Clostridium_botulinum_A3_str_Loch_Maree
- 70_Clostridium_botulinum_B_str_Eklund_17B
- 70_Clostridium_botulinum_Bf
- 70_Clostridium_botulinum_E1_str_BoNT_E_Beluga
- 70_Clostridium_botulinum_F_str_230613
- 70_Clostridium_botulinum_H04402_065
- 70_Clostridium_butyricum
- 70_Clostridium_carboxidivorans_P7
- 70_Clostridium_cellulovorans_743B
- 70_Clostridium_difficile_R20291
- 70_Clostridium_hiranonis_DSM_13275
- 70_Clostridium_perfringens
- 70_Clostridium_sp._L2-50
- 70_Clostridium_sporogenes_ATCC_15579
- 70_Clostridium_tetani_E88
- 70_Coproccoccus_sp._ART55/1
- 70_Eubacterium_ventriosum_ATCC_27560
- 70_Lachnospiraceae_bacterium_3_1_57FAA_CT1
- 70_Peptostreptococcus_stomatis_DSM_17678
- 70_Shuttleworthia_satelles_DSM_14600
- 70_Subdoligranulum_variabile_DSM_15176
- 60_Butyrvibrio_crossotus_DSM_2876
- 60_Butyrvibrio_proteoclasticus_B316
- 60_Clostridium_acetobutylicum
- 60_Clostridium_bartlettii_DSM_16795
- 60_Clostridium_beijerinckii_NCIMB_8052
- 60_Clostridium_botulinum_A3_str_Loch_Maree
- 60_Clostridium_botulinum_B_str_Eklund_17B
- 60_Clostridium_botulinum_F_str_230613
- 60_Clostridium_botulinum_H04402_065
- 60_Clostridium_butyricum
- 60_Clostridium_carboxidivorans_P7
- 60_Clostridium_cellulovorans_743B
- 60_Clostridium_difficile_R20291
- 60_Clostridium_hiranonis_DSM_13275
- 60_Clostridium_perfringens
- 60_Clostridium_sp._L2-50
- 60_Clostridium_sporogenes_ATCC_15579
- 60_Clostridium_tetani_E88
- 60_Coproccoccus_sp._ART55/1
- 60_Eubacterium_ventriosum_ATCC_27560
- 60_Lachnospiraceae_bacterium_3_1_57FAA_CT1
- 60_Shuttleworthia_satelles_DSM_14600
- 60_Subdoligranulum_variabile_DSM_15176



Tortoise2
 Tortoise1
 Rabbit2
 Rabbit1
 PatagonianCavey
 GuineaPig
 GuineaHog
 Duck
 Donkey
 Chinchilla
 TreeShaw
 SpiderMonkey
 Skink
 RingTailedLemur
 Rat
 Porcupine
 Ostrich
 Opposum
 Mouse
 MongooseLemur
 Mandrill
 Human2
 Human1
 GoldenlionTamarin
 BoxTurtle
 BeardedDragon
 Armadillo
 GrizzlyBear
 Redpanda
 Meerkat
 Dog2
 Dog1
 Binturong
 Tiger
 Tenrec2
 Tenrec1
 RiverOtter
 RedTailedHawk
 Ratsnake
 Peruvian
 KingVulture
 Ferret
 BaldEagle
 ArcticFox2
 ArcticFox1
 Africanlion

Carnivores **ncCarnivora** **Omnivores** **Herbivores**

Figure S4. FrameBot closest match analysis of butyryl-CoA:acetate CoA-transferase (A: blue) and butyrate kinase (B: red) for all samples is shown. Closest match results were separated into distinct categories based on their percent identity to a reference gene (bins containing 10% identity ranges were created, where the number preceding the reference name specifies the lowest identity in that group; e.g. “70_reference name” is the combined percentage of all sequences that show 70%-79% identity to that reference; no number indicates categories from 90%-100% identity). nc*Carnivora*: non-carnivorous *Carnivora*. A black star indicates taxa significantly different ($p < 0.05$) between diet groups (non-carnivorous *Carnivora* were excluded from the analysis).

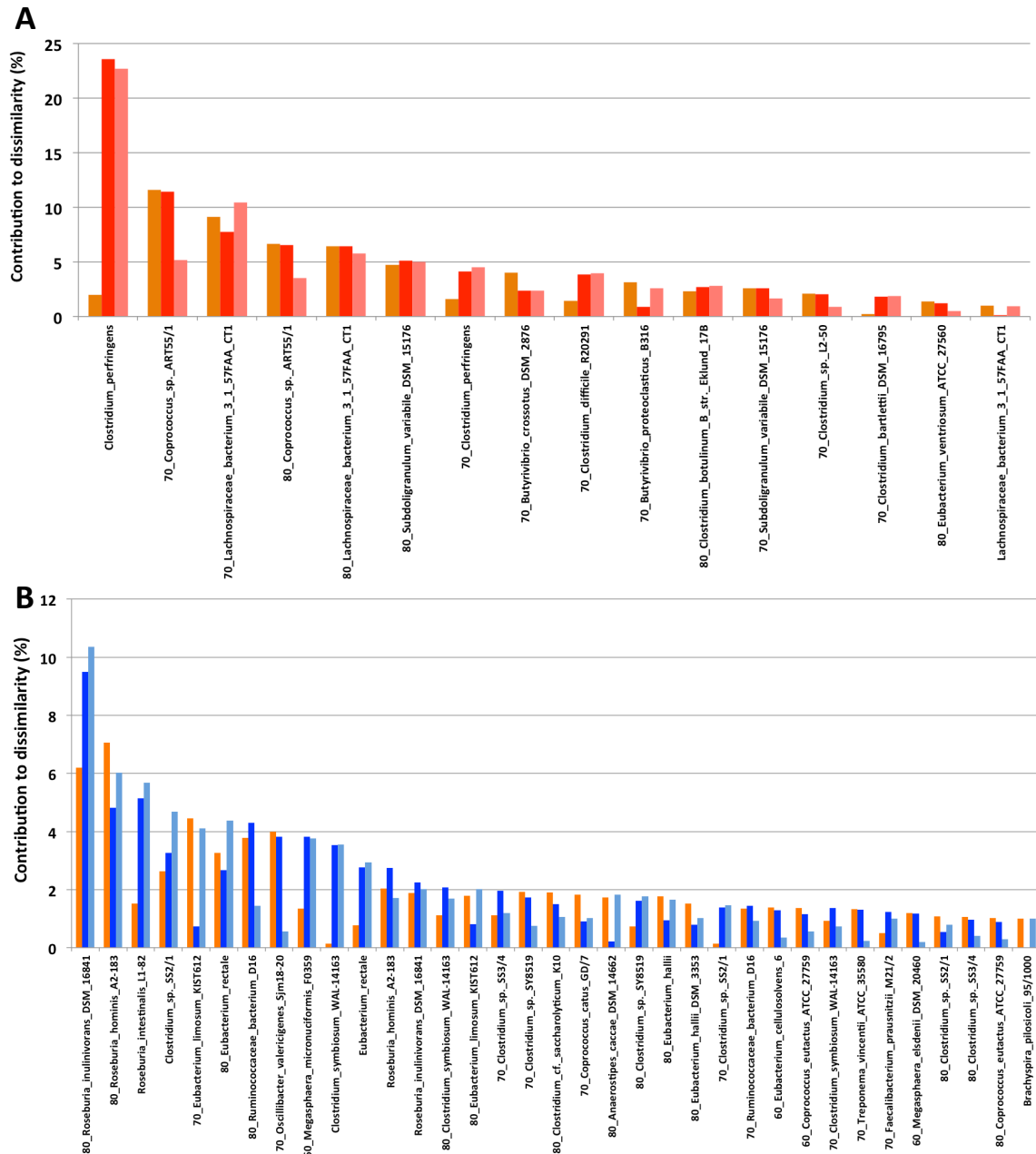


Figure S5. Simper analysis of butyryl-CoA:acetate CoA-transferase (A) and butyrate kinase (B) is shown. Analysis comparing herbivores/omnivores (orange bar), carnivores/herbivores (red, blue) and carnivores/omnivores (light red, light blue) are indicated. Only taxa contributing more than 1 % are displayed. Non-carnivorous *Carnivora* were excluded from the analysis.

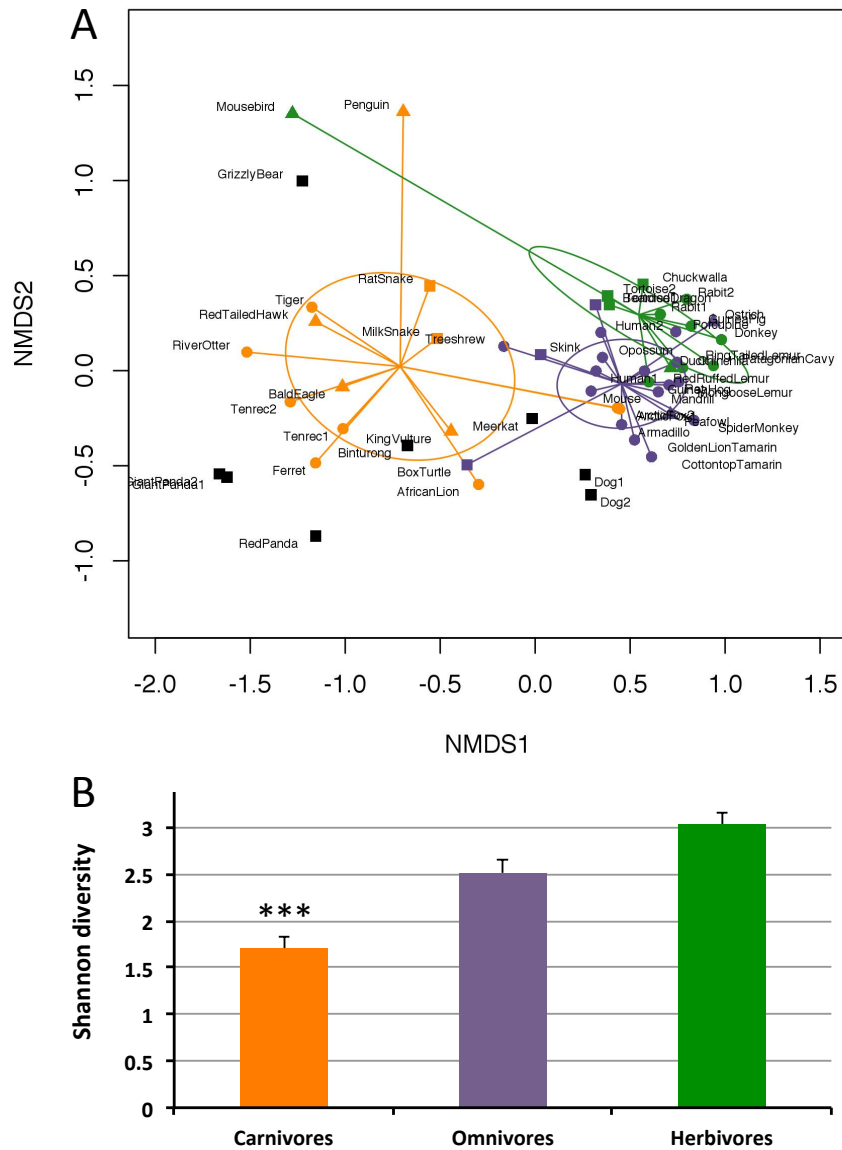


Figure S6. In panel A non-metric multidimensional scaling (NMDS) of entire bacterial 16S *rRNA* data is shown (stress = 0.13). Individual diet groups are indicated as orange (carnivores), violet (omnivores) and green (herbivores). Distinct *Vertebrate* classes are specified with symbols. Non-carnivorous *Carnivora* are presented as black squares. Ellipses represent the standard deviations of points. Diet was revealed as a significant factor ($p <$

0.01 based on permutational anova analysis). Shannon diversity is indicated in panel B where error bars indicate standard error of the mean. ***: $p < 0.01$.

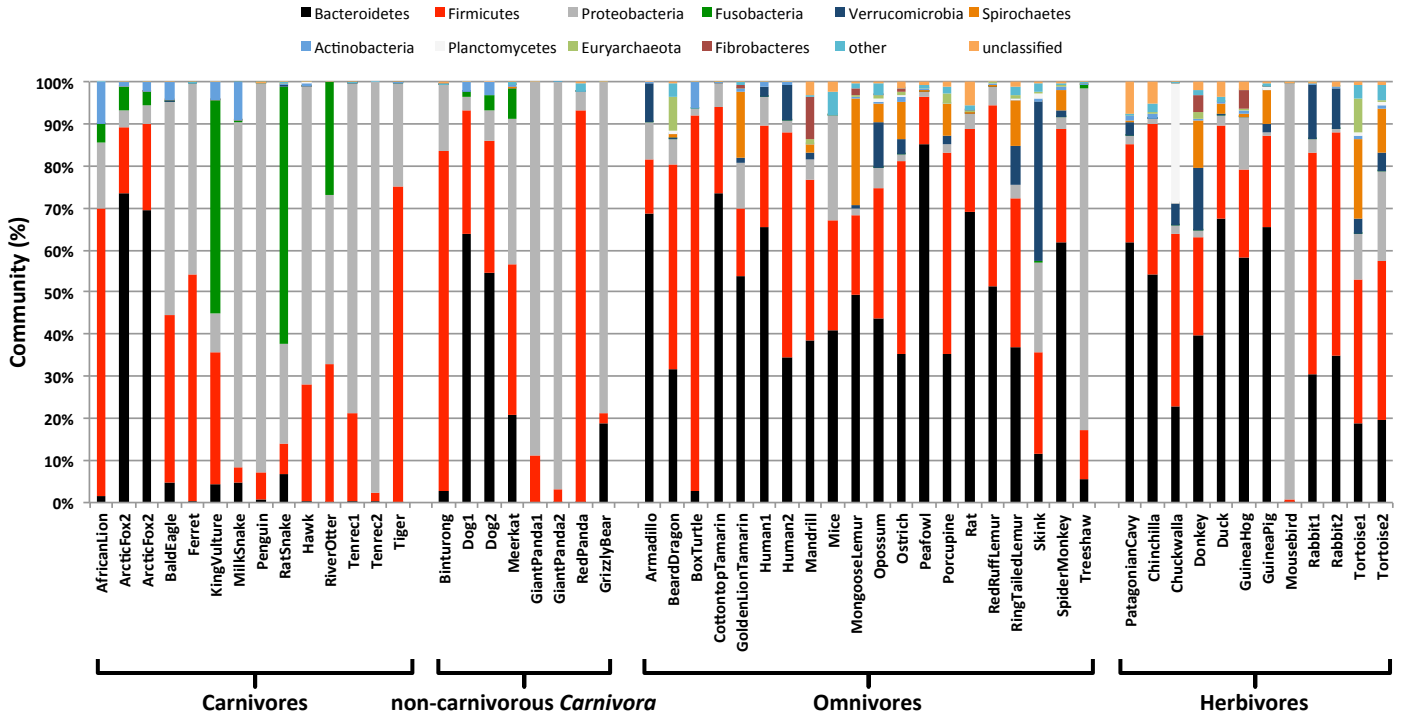


Figure S7 16S rRNA analysis on phylum level for all samples is shown.