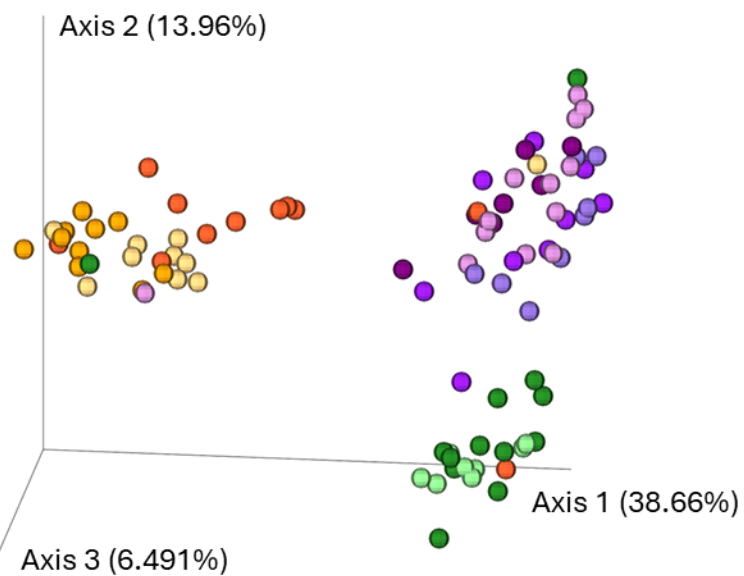
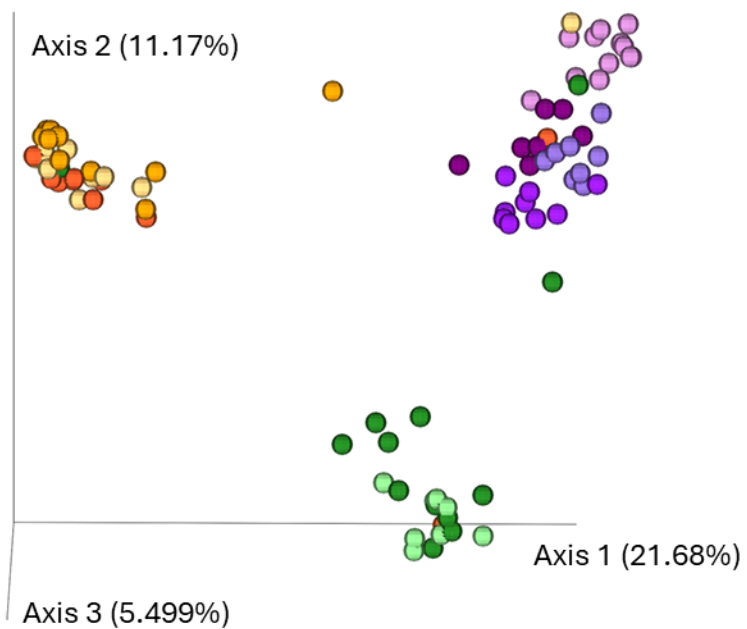
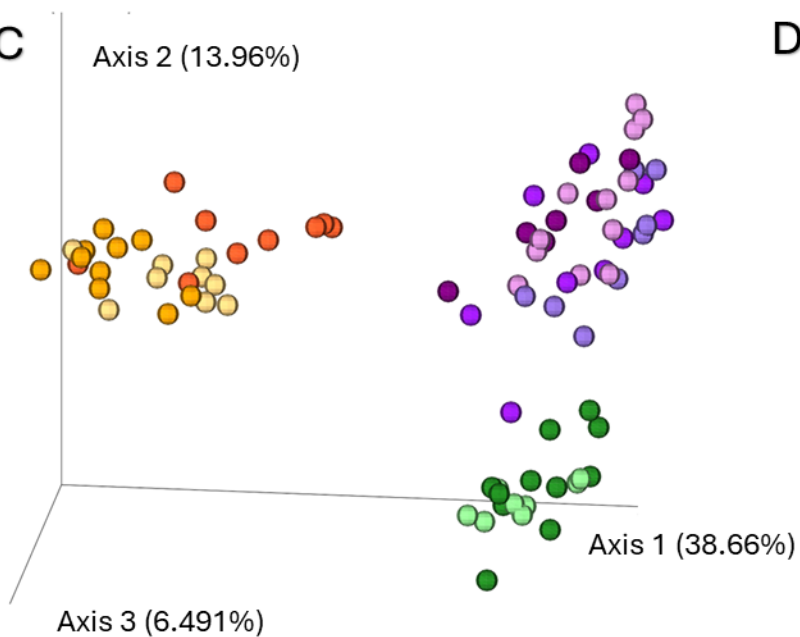
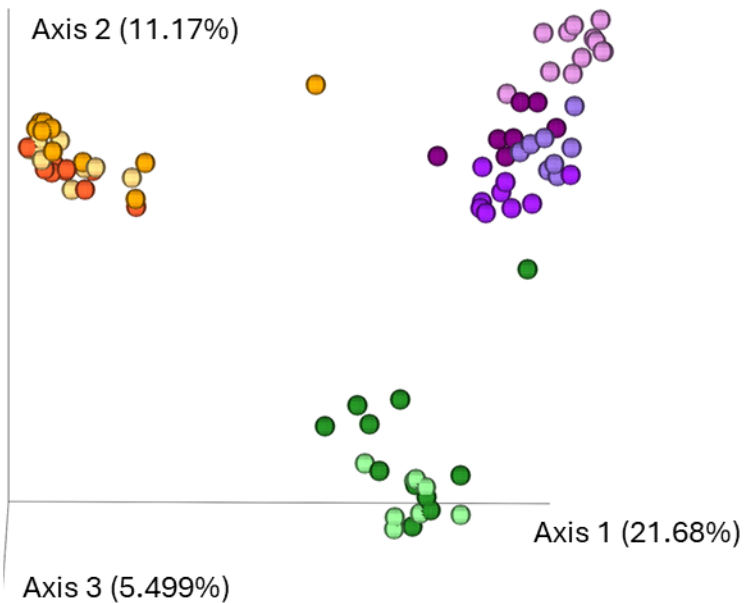
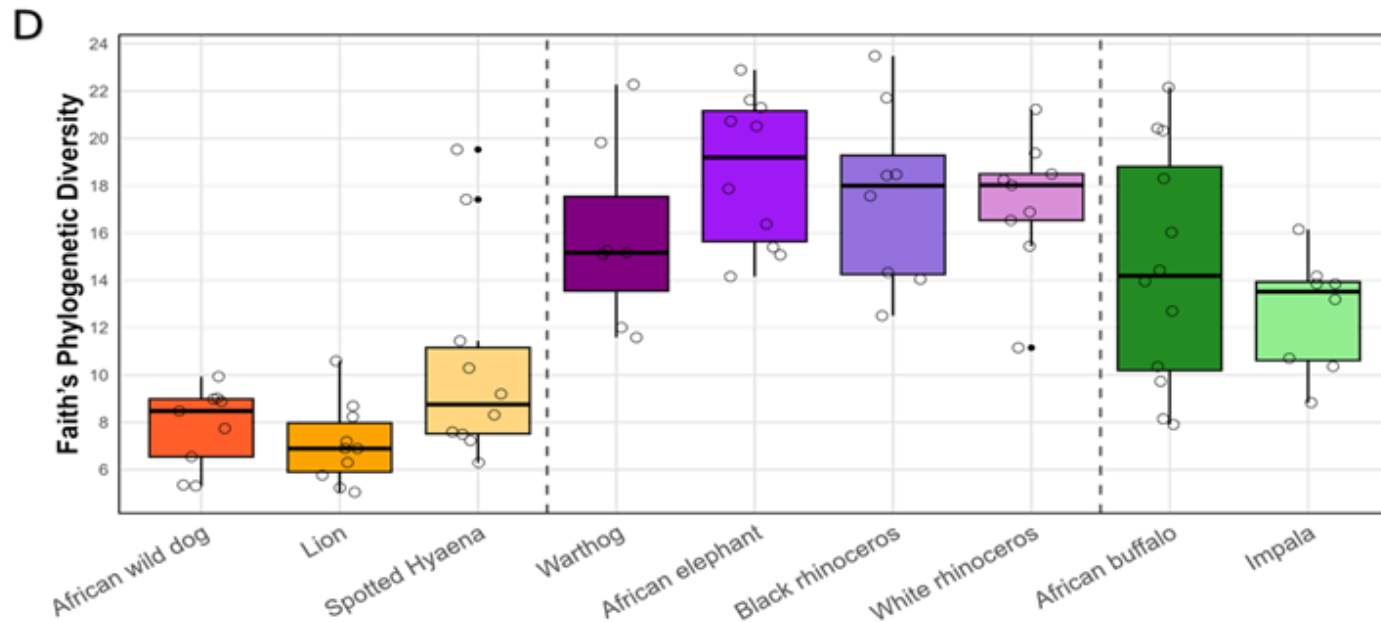
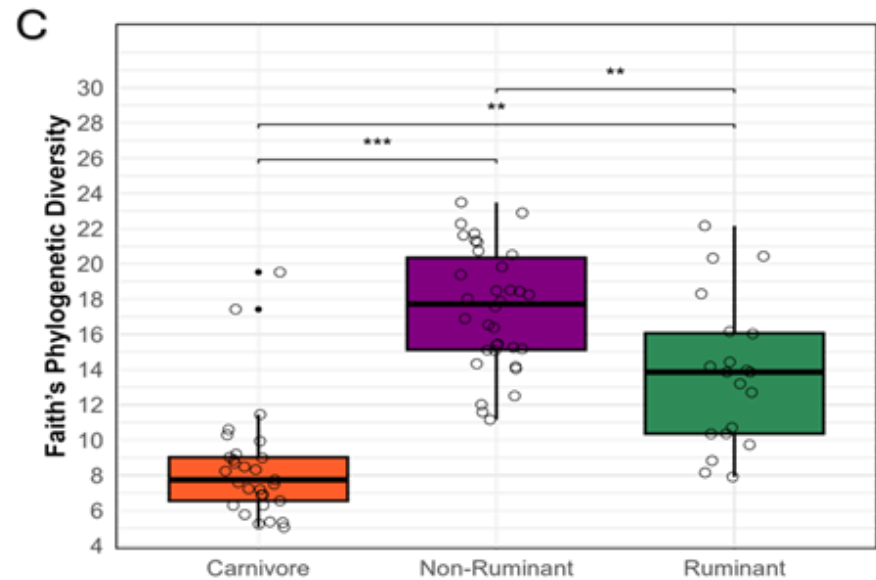
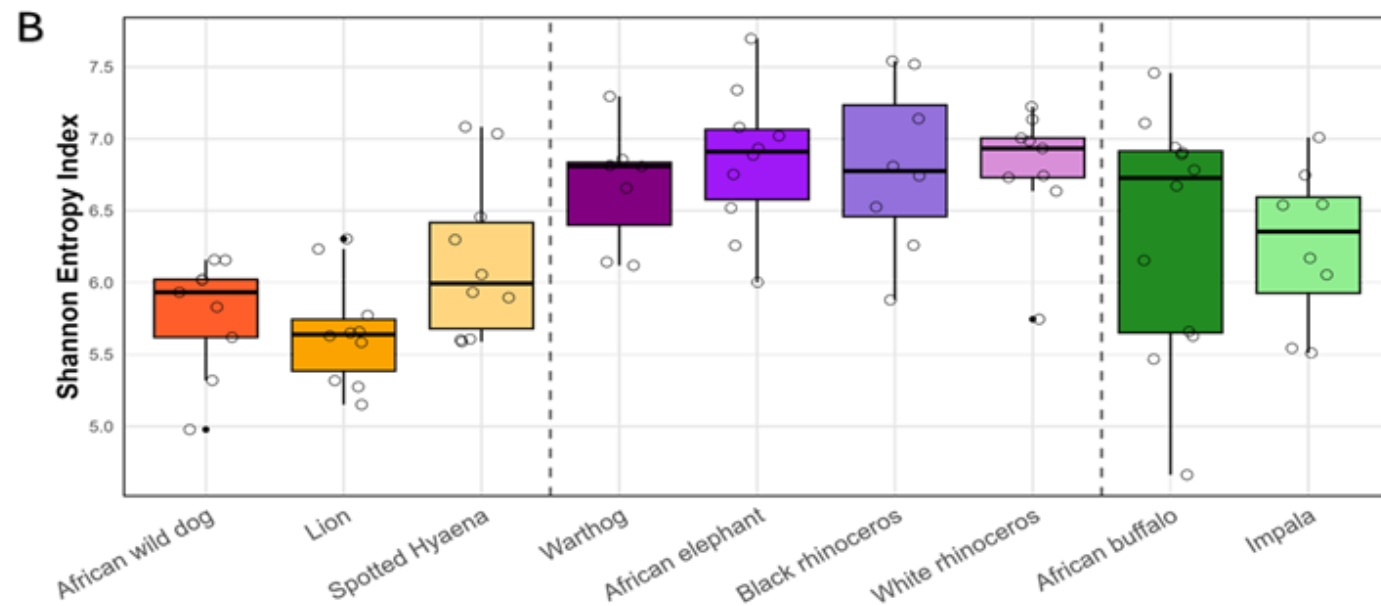
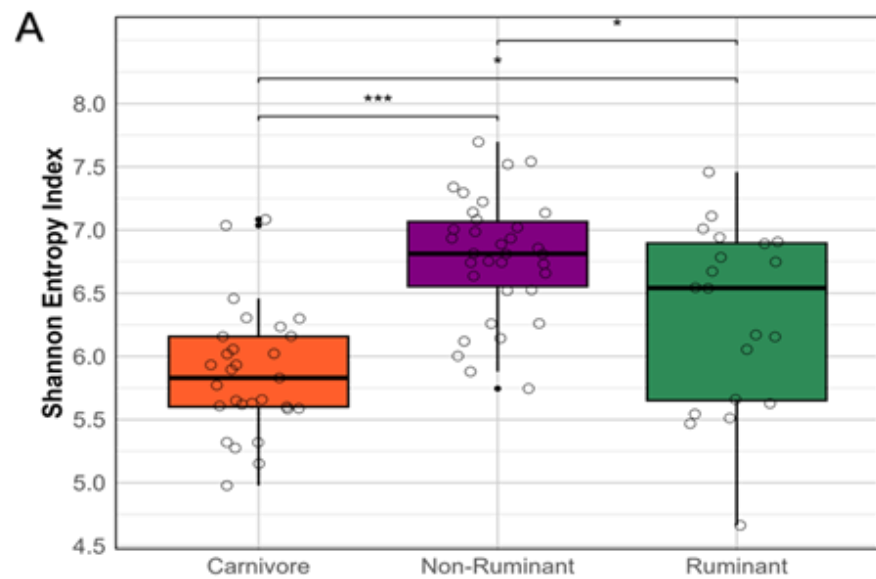
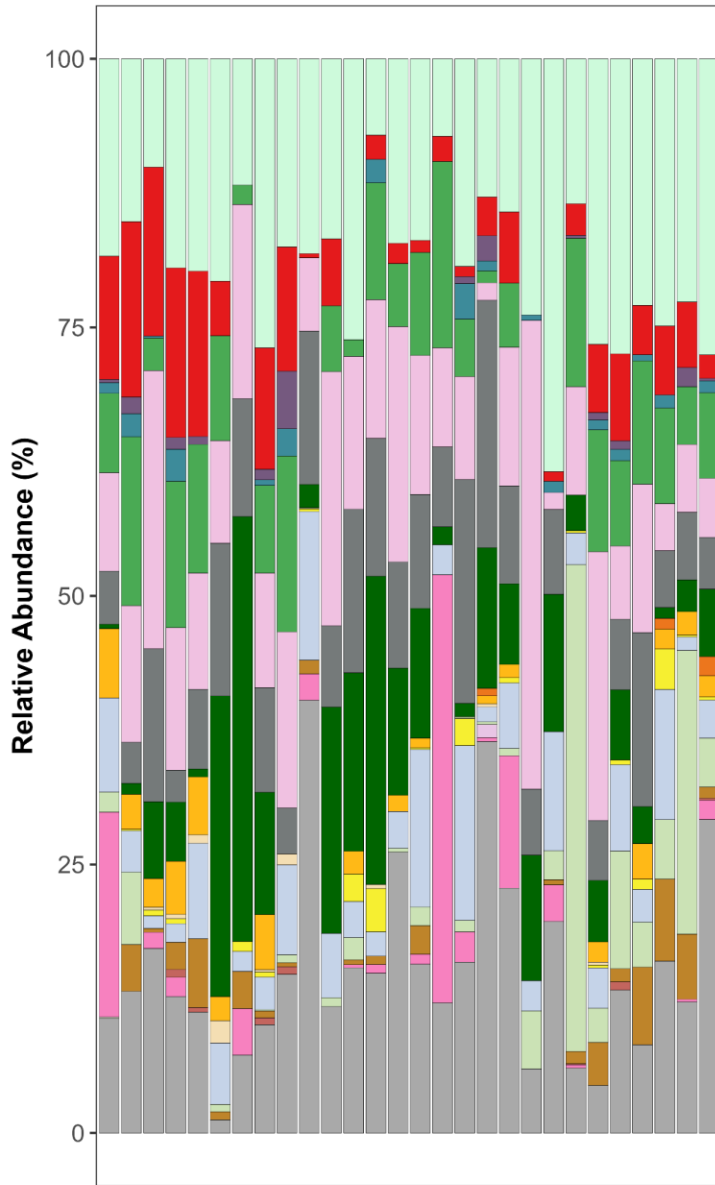


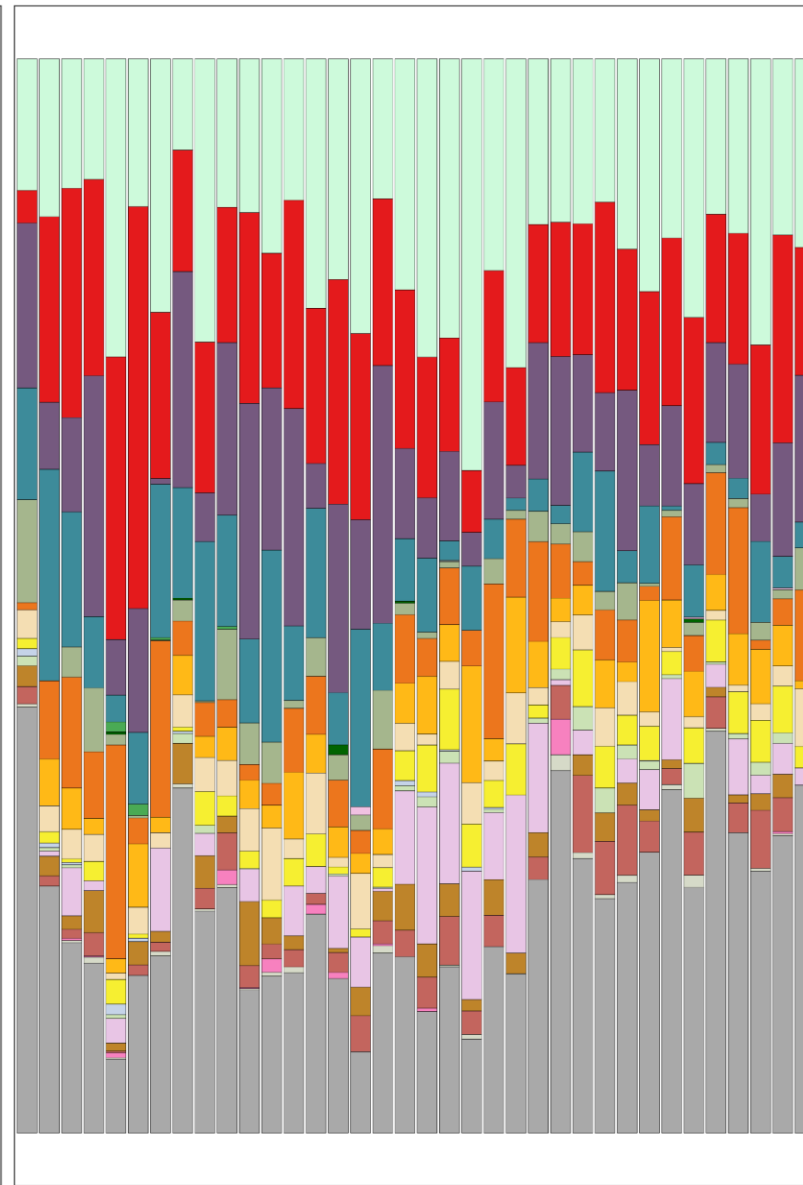
A**B****C****D**



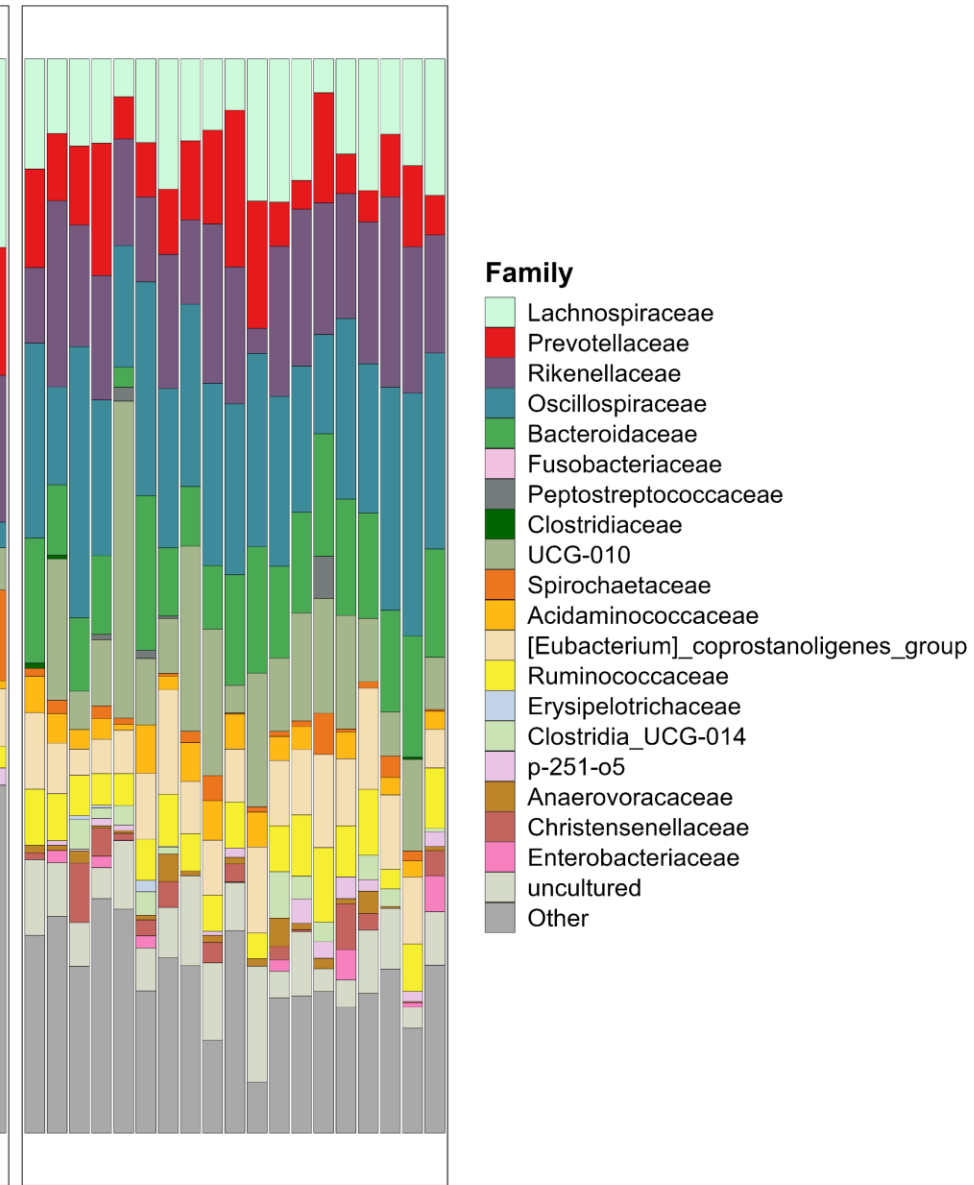
Carnivore



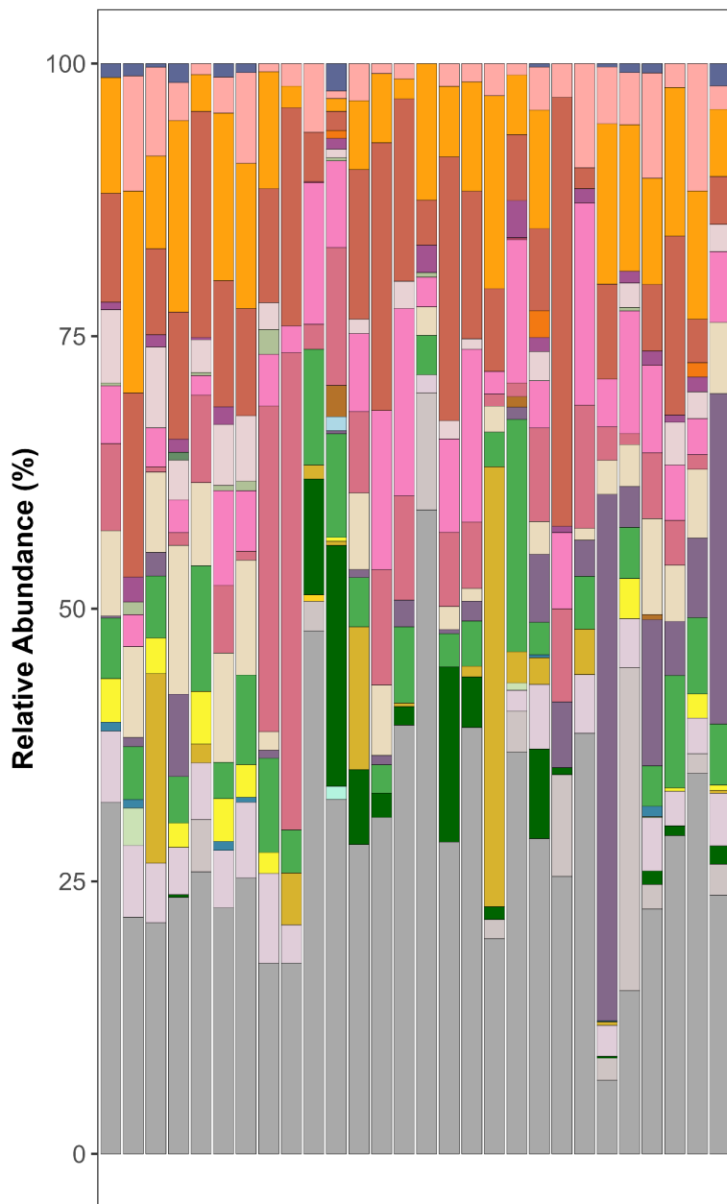
Non-Ruminant



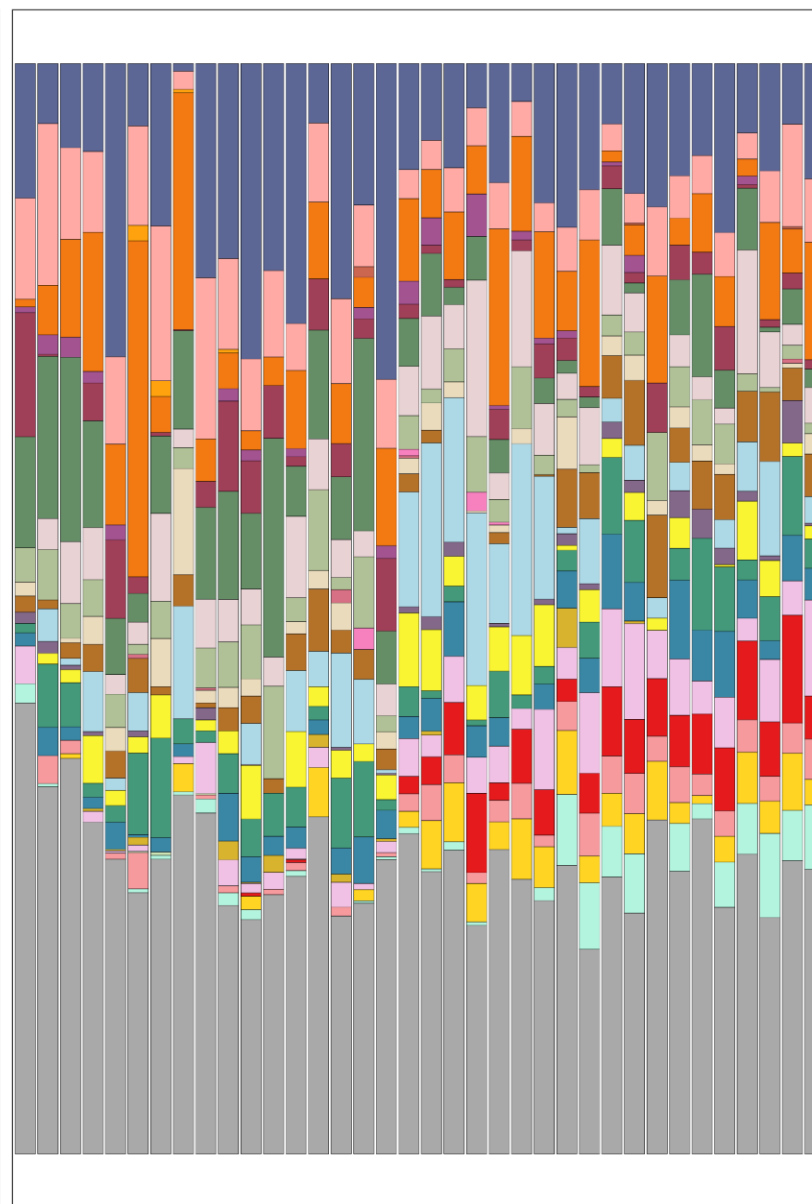
Ruminant



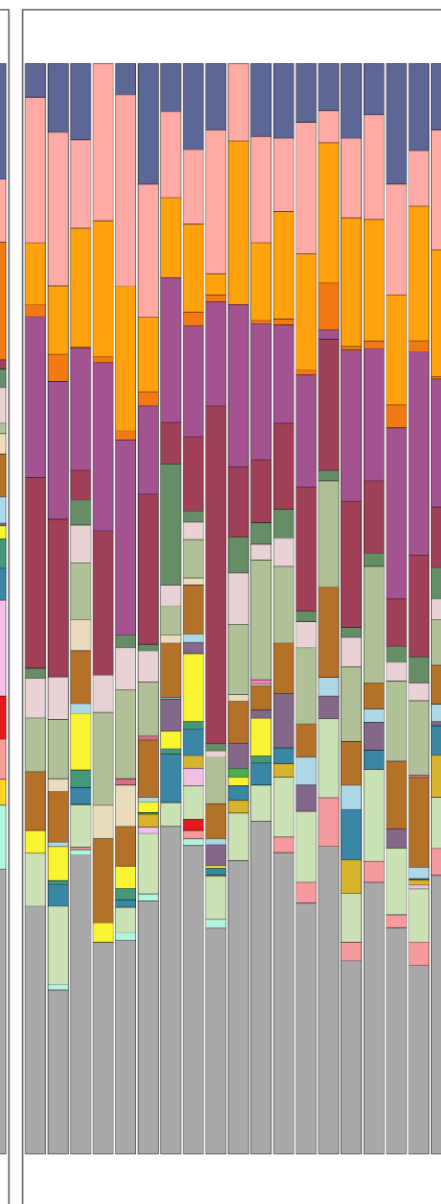
Carnivore



Non-Ruminant

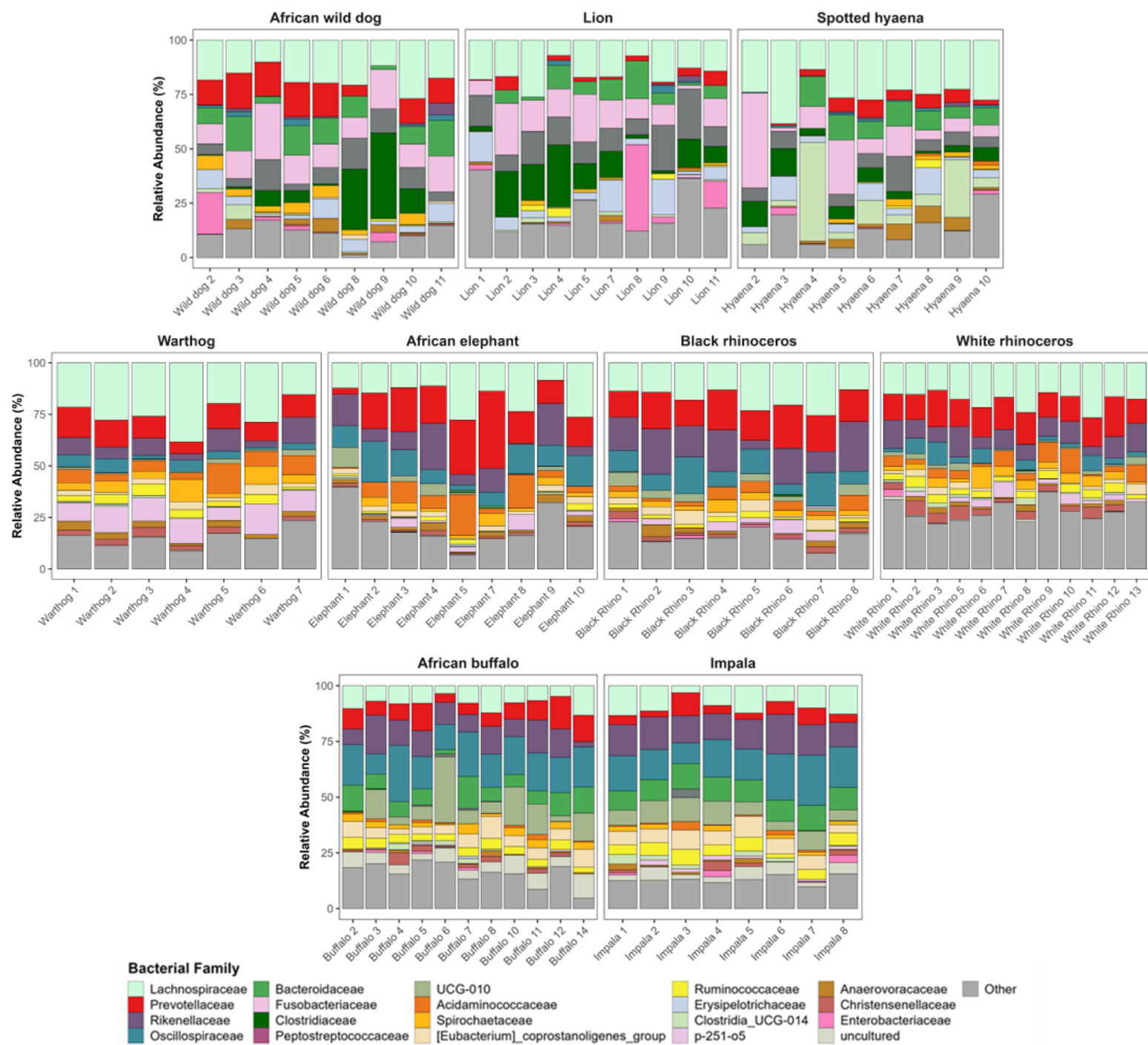


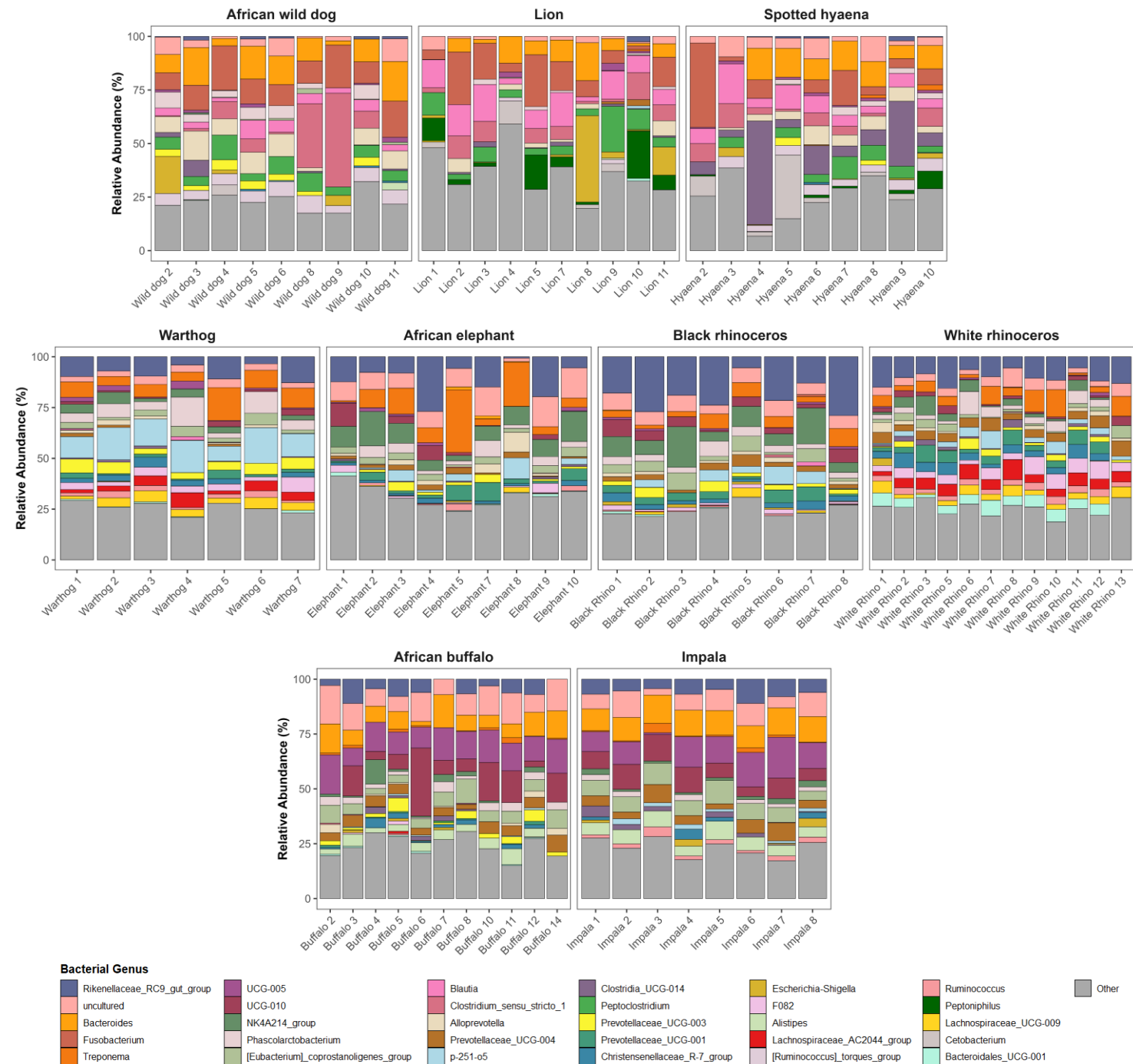
Ruminant



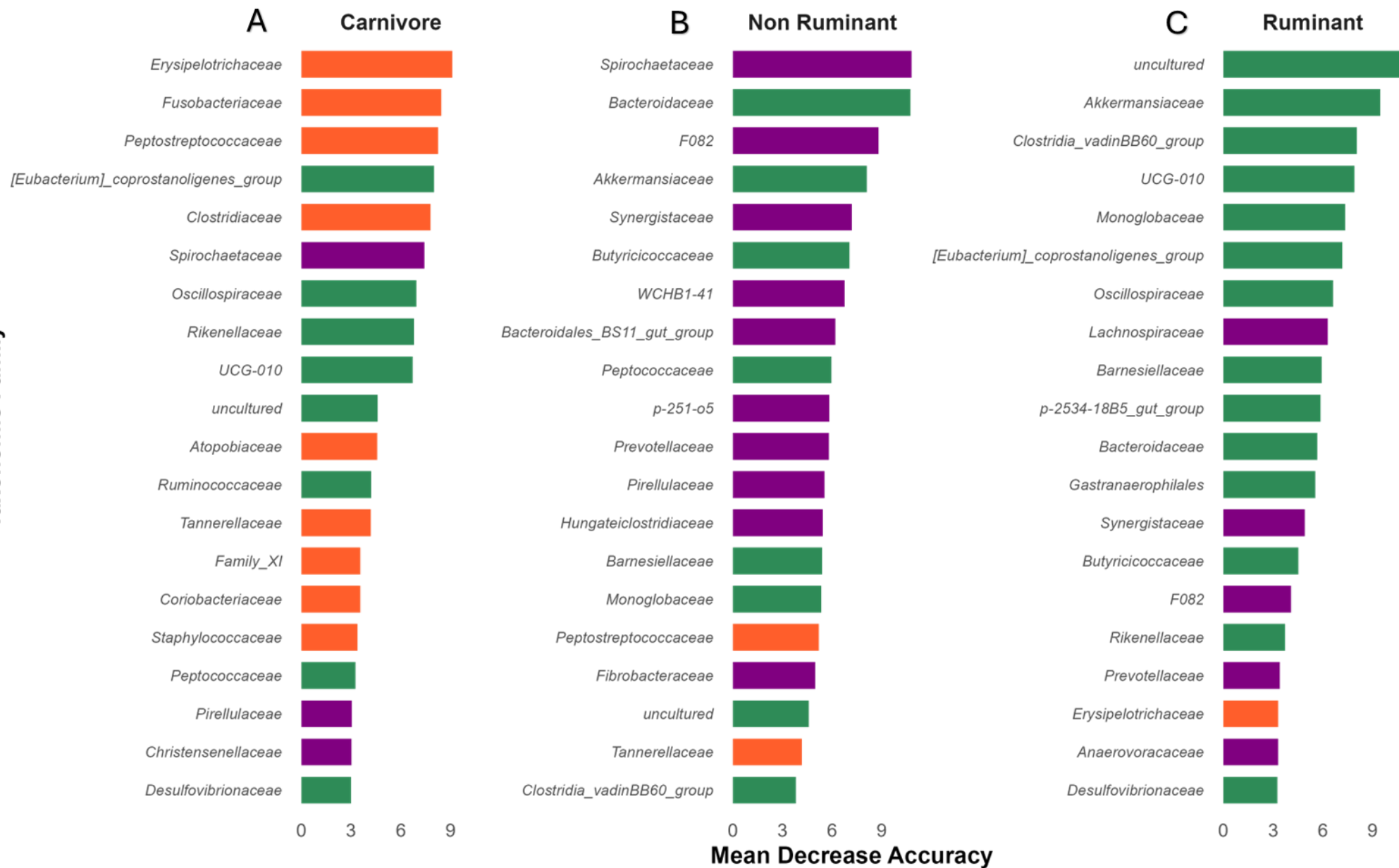
Genera

- Rikenellaceae_RC9_gut_group
- uncultured
- Bacteroides
- Fusobacterium
- Treponema
- UCG-005
- UCG-010
- NK4A214_group
- Phascolarctobacterium
- [Eubacterium]_coprostanoligenes_group
- Blautia
- Clostridium_sensu_stricto_1
- Alloprevotella
- Prevotellaceae_UCG-004
- p-251-o5
- Clostridia_UCG-014
- Peptoclostridium
- Prevotellaceae_UCG-003
- Prevotellaceae_UCG-001
- Christensenellaceae_R-7_group
- Escherichia-Shigella
- F082
- Alistipes
- Lachnospiraceae_AC2044_group
- [Ruminococcus]_torques_group
- Ruminococcus
- Peptoniphilus
- Lachnospiraceae_UCG-009
- Cetobacterium
- Bacteroidales_UCG-001
- Other





Taxonomic Family



Highest CLR Abundance ■ Carnivore ■ Non Ruminant ■ Ruminant

Taxonomic Genera

