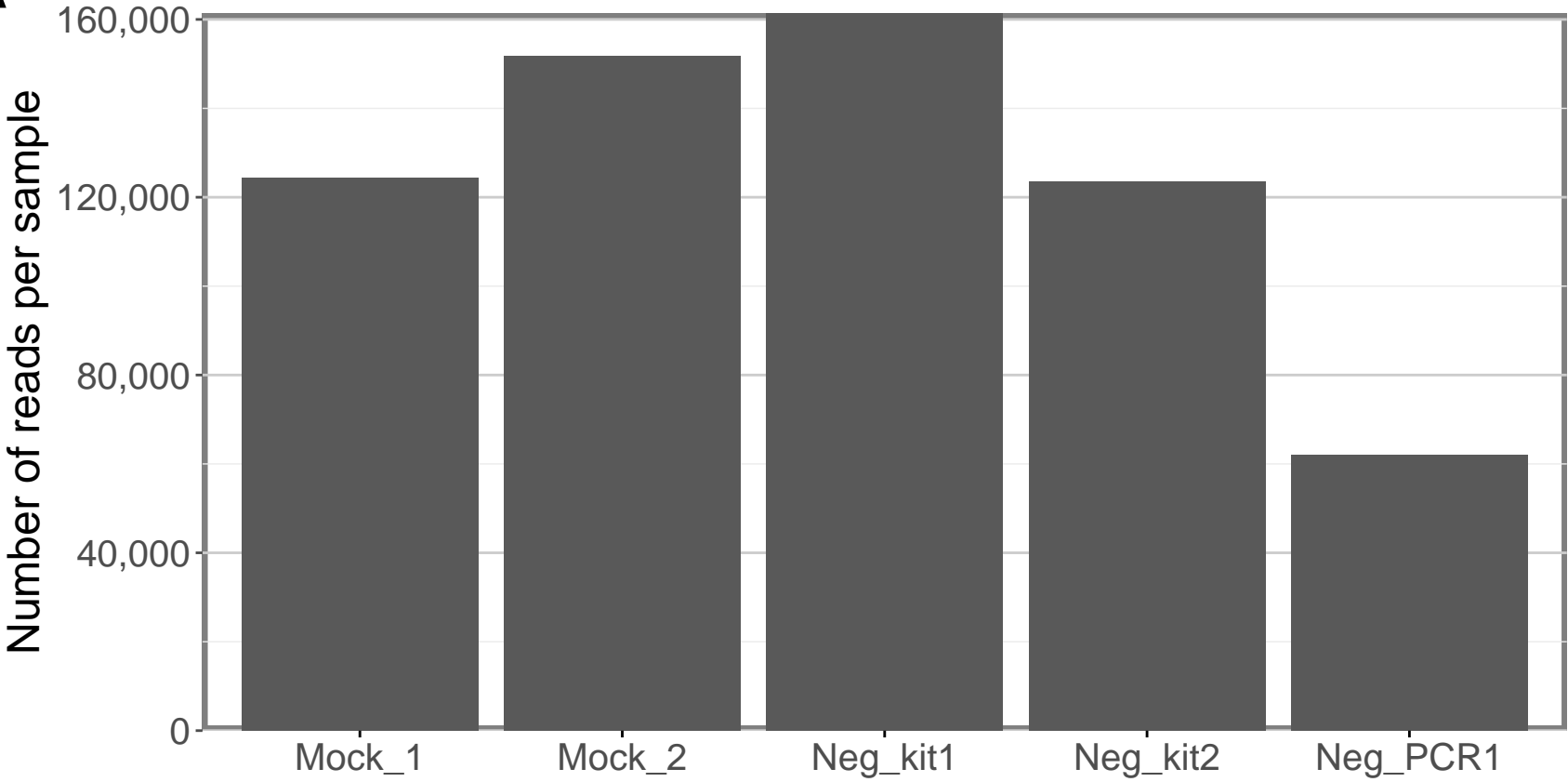
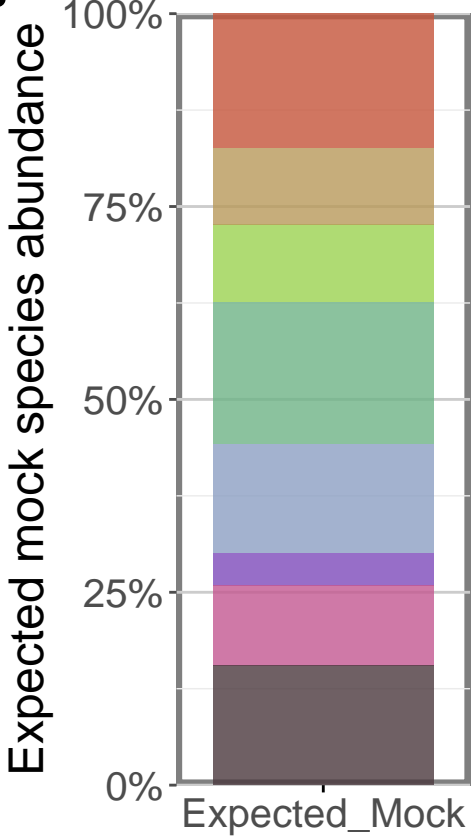


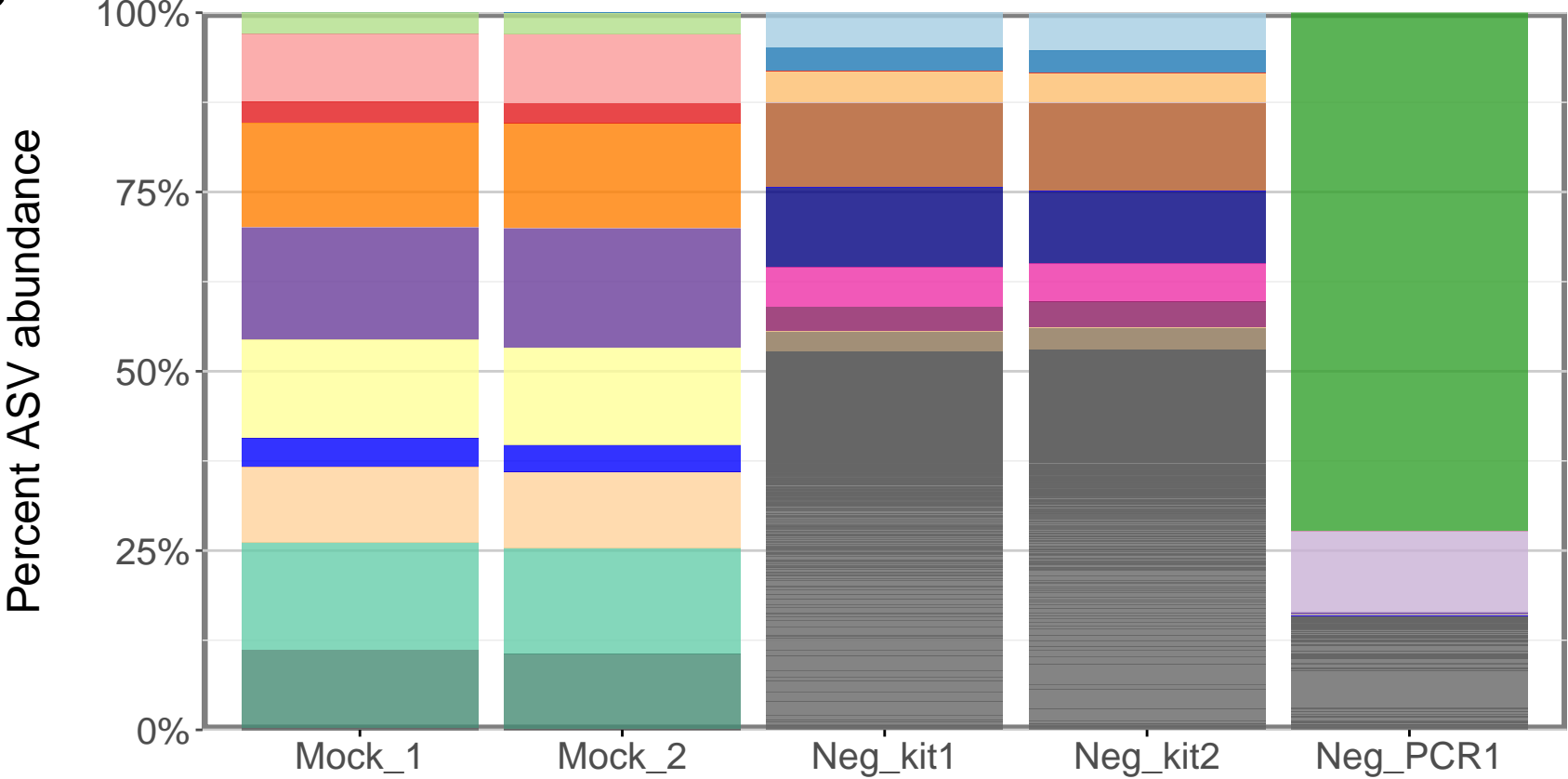
A



B



C



Mock species

- Bacillus subtilis
- Enterococcus faecalis
- Escherichia coli
- Lactobacillus fermentum
- Listeria monocytogenes
- Pseudomonas aeruginosa
- Salmonella enterica
- Staphylococcus aureus

ASV

- 19d57d86ee1d5ea49fe9f4256bfa9d22: d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Flavobacterium
- 1c34565a5c5d3d9fa3953518e9a4b59c: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Burkholderiales; f__Oxalobacteraceae
- 3019073bed0ceecaec9f446607c003a: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacterales; f__Enterobacteriaceae; g__Salmonella
- 33731ac0a1051571d808de3e91b11356: d__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhizobiales; f__Beijerinckiaceae; g__Methylobacterium-Methylobacterium; s__Methylobacterium-jeotgali
- 456f366888eac2ce0809986d3ed1b237: d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales
- 5c9b89d5acf8e0020d9de19b53d21663: d__Bacteria; p__Firmicutes; c__Bacilli; o__Staphylococcales; f__Staphylococcaceae; g__Staphylococcus
- 7a6da1a5e2be4eadd87d67705eb68f4a: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Alteromonadales; f__Alteromonadaceae; g__Rheinheimera
- 94efb59163996601f8a20a8f9b0ac0d9: d__Bacteria; p__Firmicutes; c__Bacilli; o__Bacillales; f__Bacillaceae; g__Bacillus
- a64d8357b6a2010ba13a1ebfc7df7c9b: d__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodobacterales; f__Rhodobacteraceae; g__Amaricoccus; s__Amaricoccus-kaplicensis
- b08e8a180eba5a20a70b190bf5f9ad2: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacterales; f__Enterobacteriaceae
- d02f4c8904ba636c08bbdf8b65b5d7d0: d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Listeriaceae; g__Listeria
- d23c811ea38d6cb704bb13445f43d599: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
- d3962c483c45f0765b9b7656ec7f43a0: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria
- e1a72c614544de9ecfc0b0cd7cc24738: d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Flavobacterium
- e6242a18b930fae98ab395c927393aa3: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Burkholderiales; f__Comamonadaceae
- ebb95142c2d5fb1d9a1e0e779348b6f1: d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Flavobacterium
- f0090fc348a4c45135f858d776e30dd0: d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Lactobacillaceae; g__Lactobacillus
- f0af66950b4ed3e76bd78e09b28e1f02: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales; f__Moraxellaceae; g__Acinetobacter
- f9eed79f2bd69b0da2385144b13658a7: d__Bacteria; p__Firmicutes; c__Bacilli; o__Staphylococcales; f__Staphylococcaceae
- fdaab5e761766ddd1297bd5f49d1cfb3: d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacterales
- Others