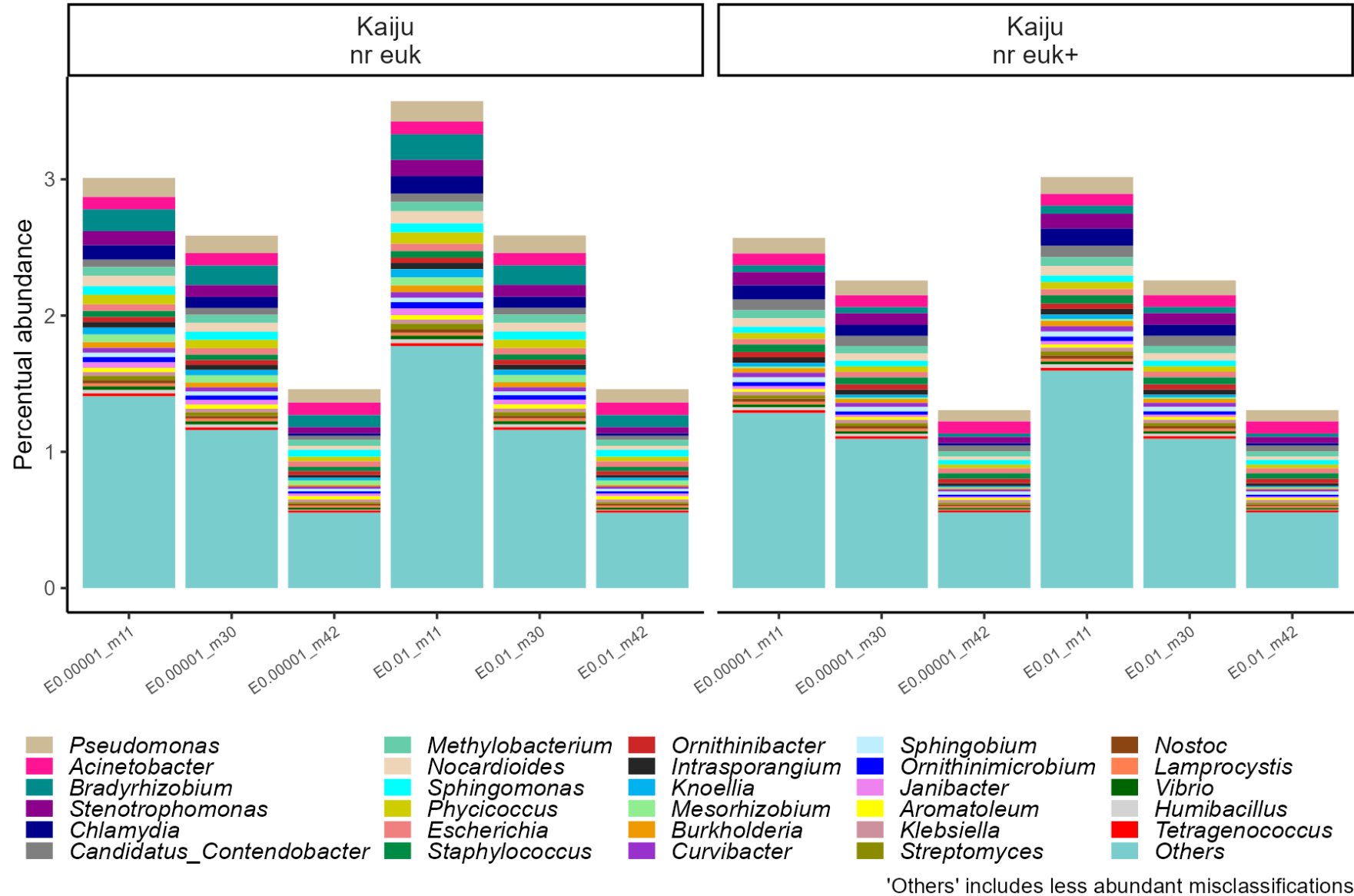




Supplementary figure 2: Bar plots illustrating the relative abundances of the most abundant misclassified genera across classifications obtained using Kaiju with various parameter settings. The displayed relative abundances account for the correct classification counts (not shown in this plot), thereby representing the total extent of misclassifications. The x-axis represents the classification types: “E” denotes the E-value threshold while “m” indicates the coverage threshold.