



Figure 2. Strain library identification at each body site. (A) Individual pie charts represent the community structure at each site. Pie labels indicate body site, font colors indicate body site type (sebaceous, moist, rarely moist), and the n-value of each pie chart is shown in parentheses. Bacterial composition is shown within the pie charts. Pie charts can be read by starting at the 12-o'clock position going in the clockwise direction to align with the legend. The large, light pink wedge seen in all charts except for the nares corresponds to *Micrococcus*. **(B)** Relative abundance of taxa across eight body sites comparing cultured versus metagenomic datasets. **(C)** Scatter plot depicts relative abundance of individual genera as derived from metagenomic (y-axis) and cultured datasets (x-axis). Each dot represents a specific genus, with its position reflecting the relative abundance across the two dataset types. Genera are color-coded according to their respective phyla.