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.