FIGURE 5

The graph illustrates the percentage of gene pairs across various ranges of average gene length, with three categories: Inparalogs, Outparalogs, and Inparalogs + Outparalogs. Each category is represented by a different color: blue for Inparalogs, green for Outparalogs, and brown for Inparalogs + Outparalogs. The x-axis represents ranges of average gene length, while the y-axis shows the percentage of gene pairs.