Figure: Horizontal stacked bar chart showing the number of eQTLs above PIP threshold across cell types and tissues, colored by eVariant–eGene distance.

Legend: **eVariant–eGene distance**
- 0–10 Kb
- 10–100 Kb
- 100–250 Kb
- 250–1000 Kb
- > 1000Kb

Y-axis (top to bottom): T_CD4_pos, blood, monocyte, LCL, Treg, neutrophil, T_CD8_pos, NK, iPSC, spleen, macrophage, B, monocyte_CD16_pos, Tfh, T, T_CD4_pos_naive, B_memory, monocyte_CD14_pos, dendritic

X-axis: Number of eQTLs above PIP threshold (0, 1000, 2000, 3000)