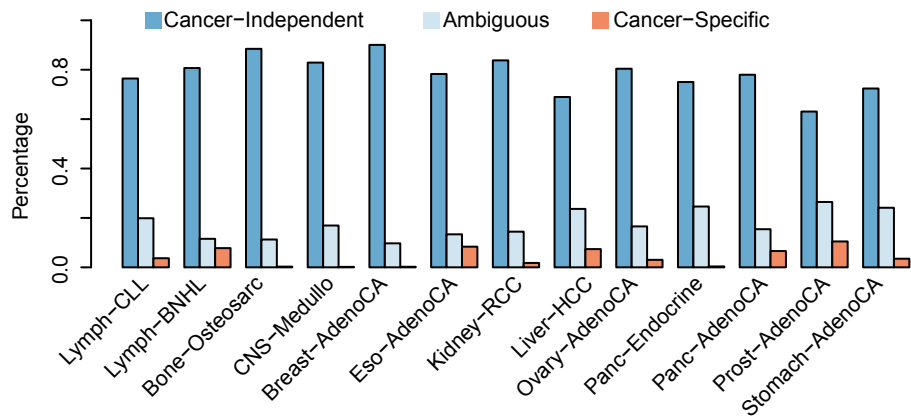
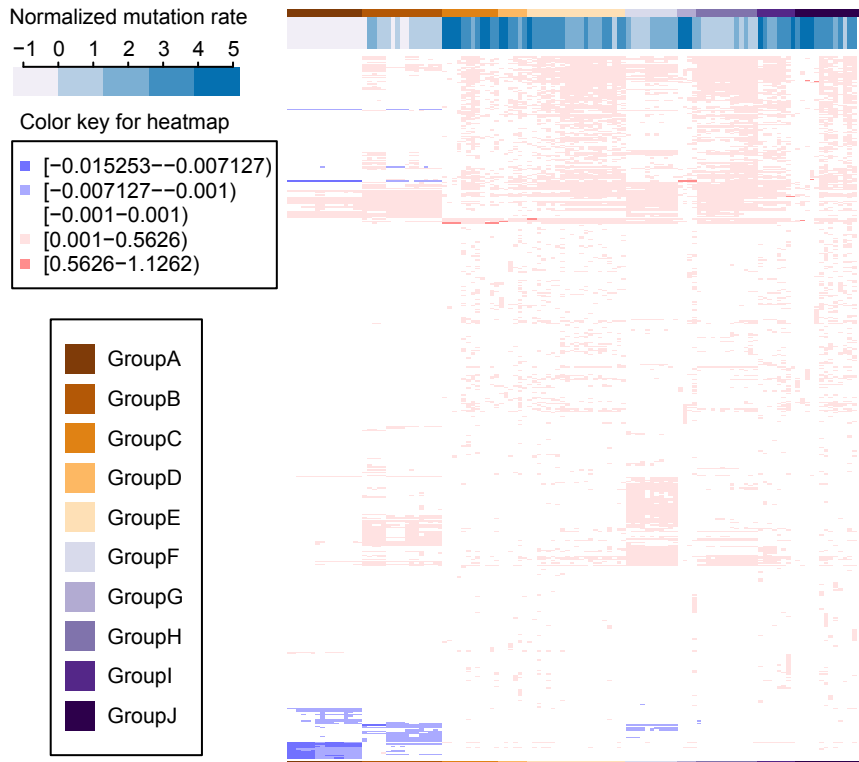


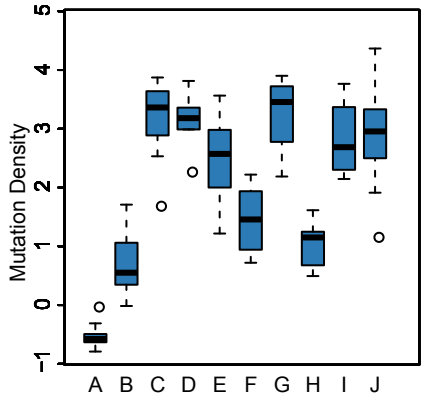
a Percentages of regions in each category



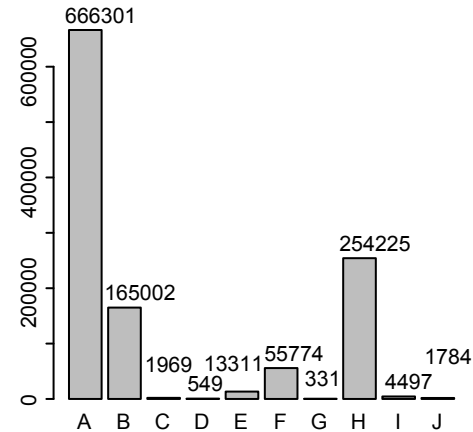
b Contextual weights of the motifs



c Normalized mutation rates in groups



d The numbers of regions in groups



e Fold changes in group A

