Fig. S1

(a) In the figure, patients affected by various types of mutations are shown. The y-axis represents the percentage of patients affected, and the x-axis lists different types of diseases: AML, MM, ALL, and CLL.

(b) The bar chart illustrates the copy number variation of m6A regulatory genes in AML. The bars are color-coded to represent different types of variations: black for point mutation, red for deep deletion, blue for amplification, green for shallow deletion or gain. The x-axis shows the number of events, and the y-axis represents the number of patients.