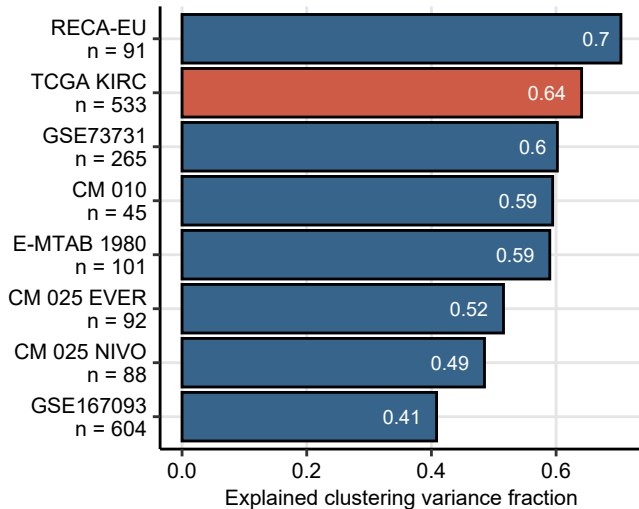
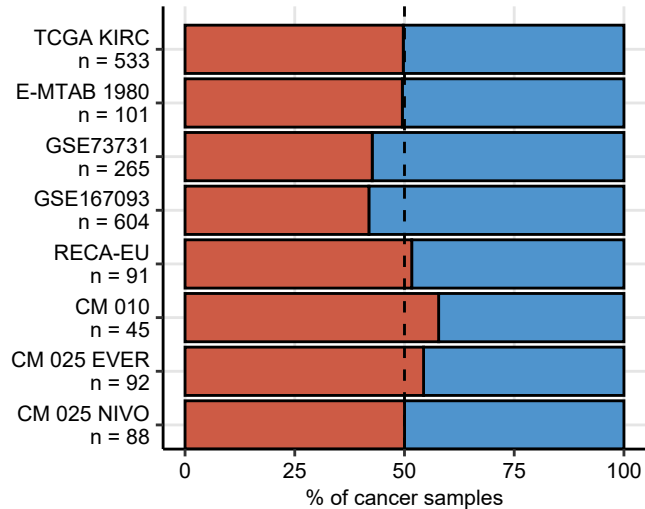
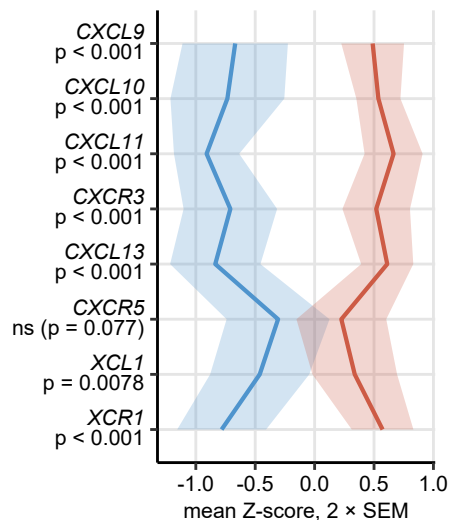
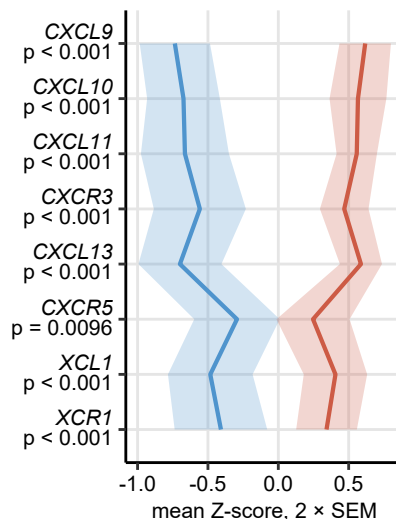


A**Semi-supervised clustering by lymphokine genes****Distribution of cancer samples****B****CM 010**

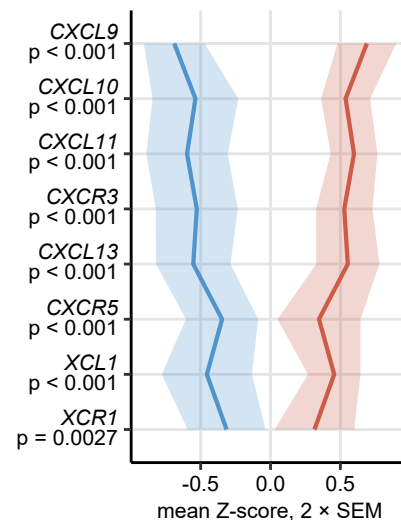
low: n = 19, high: n = 26

**CM 025 EVER**

low: n = 42, high: n = 50

**CM 025 NIVO**

low: n = 44, high: n = 44



Lymphokine cluster chemox low chemox high