Adenocarcinoma; Tumor (n=155), non-tumor lung (n=117)

Tumor FCM RNA-seq n=85 n=142

13 72 70

Cases in which FCM was performed (n=85)

	Done	Not-done
IHC	65	20
WES	69	16
TCR repertoire	55	30
Metabolome	29	56

FCM; flow cytometry IHC; immunohistochemistry WES; whole exome seq TCR; T cell receptor

Squamous cell carcinoma; Tumor (n=80), non-tumor lung (n=65)

FCM RNA-seq n=50 n=74

6 44 30

Cases in which FCM was performed (n=50)

	Done	Not-done
IHC	44	6
WES	45	5
TCR repertoire	36	14
Metabolome	24	26

Other lung cancer; Tumor (n=46), non-tumor lung (n=37)

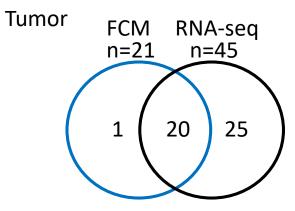


Figure S1. The number of patients and analyses in LUAD and LUSQ. The immunohistochemistry of tissue microarray, whole exome sequencing, TCR repertoire analysis, and metabolome analysis were performed in the cases FCM was conducted.