

## Supplementary

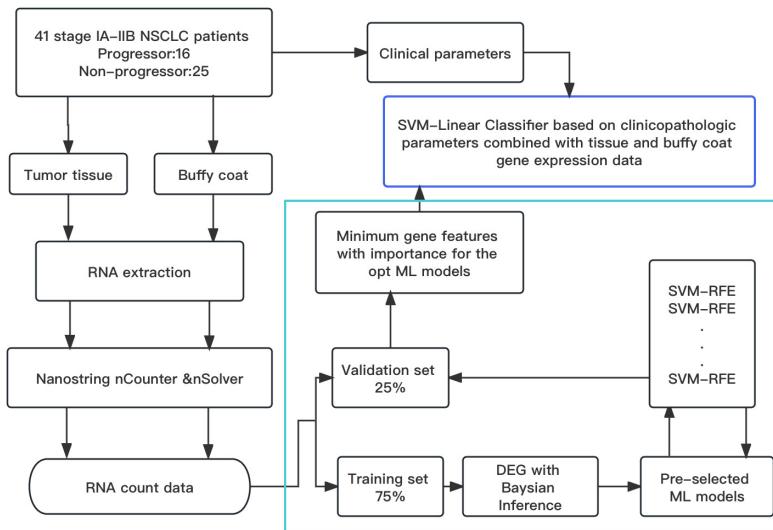
**Table S1** List of housekeeping genes

Housekeeping genes
<i>ABCF1</i>
<i>AGK</i>
<i>ALAS1</i>
<i>AMMECR1L</i>
<i>CC2D1B</i>
<i>CNOT10</i>
<i>CNOT4</i>
<i>COG7</i>
<i>DDX50</i>
<i>DHX16</i>
<i>DNAJC14</i>
<i>EDC3</i>
<i>EIF2B4</i>
<i>ERCC3</i>
<i>FCF1</i>
<i>G6PD</i>
<i>GPATCH3</i>
<i>GUSB</i>
<i>HDAC3</i>
<i>HPRT1</i>

**Table S1** (continued)

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Housekeeping genes
<i>MRPS5</i>
<i>MTMR14</i>
<i>NOL7</i>
<i>NUBP1</i>
<i>POLR2A</i>
<i>PPIA</i>
<i>PRPF38A</i>
<i>SAP130</i>
<i>SDHA</i>
<i>SF3A3</i>
<i>TBP</i>
<i>TLK2</i>
<i>TMUB2</i>
<i>TRIM39</i>
<i>TUBB</i>
<i>USP39</i>
<i>ZC3H14</i>
<i>ZKSCAN5</i>
<i>ZNF143</i>
<i>ZNF346</i>

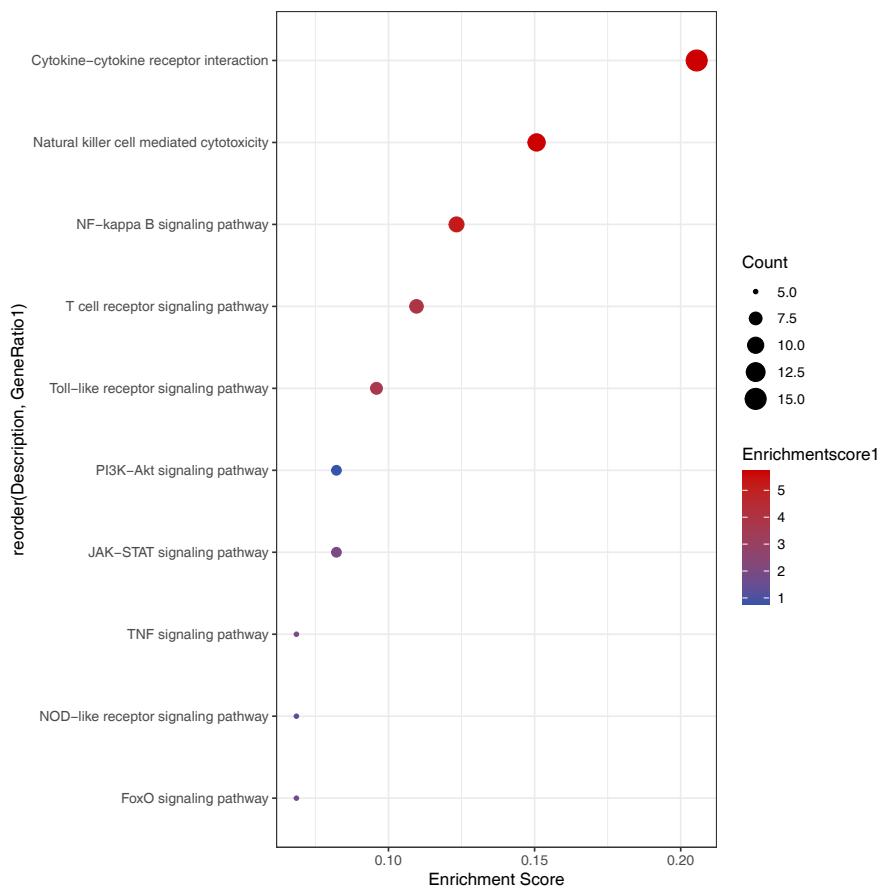


**Figure S1** SVM-RFE data analysis flow chart. NSCLC, non-small cell lung cancer; SVM-Linear, SVM with the linear kernel; SVM, support vector machine; ML, machine learning; SVM-RFE, SVM with recursive feature elimination; DEG, differentially expressed gene.

**Table S2** Top 20 in network string\_interactions.tsv ranked by MNC method

Rank	Name	Score
1	<i>TNF</i>	64
2	<i>IL6</i>	61
3	<i>CD8A</i>	59
4	<i>GZMB</i>	40
5	<i>CXCL8</i>	39
6	<i>TBX21</i>	37
7	<i>PRF1</i>	34
8	<i>KLRK1</i>	30
9	<i>IRF4</i>	29
10	<i>CD247</i>	28
10	<i>KLRD1</i>	28
10	<i>CD34</i>	28
10	<i>TLR9</i>	28
10	<i>CD1C</i>	28
15	<i>LAG3</i>	27
16	<i>ZAP70</i>	25
16	<i>STAT4</i>	25
18	<i>IDO1</i>	24
19	<i>BCL6</i>	23
19	<i>FCGR2B</i>	23

MNC, maximum neighborhood component.



**Figure S2** KEGG pathway analysis of DEGs. KEGG, Kyoto Encyclopedia of Genes and Genomes; DEGs, differentially expressed genes.