**A novel predictive model for anoikis-related gene markers in squamous cell carcinoma of the lung based on machine learning and single-cell analysis, revealing their immune relevance.**

**Supplementary figures**

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**Fig S1**

network diagram. (A) Network diagram of 45 prognostic genes interacting with anoikis closely related genes. (B) Protein Interaction PPI Network Diagram.



**Fig. S2**

LUSC drug sensitivity.



**Fig. S3**

LUSC drug sensitivity



**Fig. S4**

LUSC drug sensitivity.



**Fig. S5**

LUSC drug sensitivity.



**Fig. S6**

LUSC drug sensitivity.



**Fig. S7**

LUSC drug sensitivity.



**Fig. S8**

LUSC drug sensitivity.



**Fig. S9**

LUSC drug sensitivity.



**Fig. S10**

LUSC drug sensitivity.