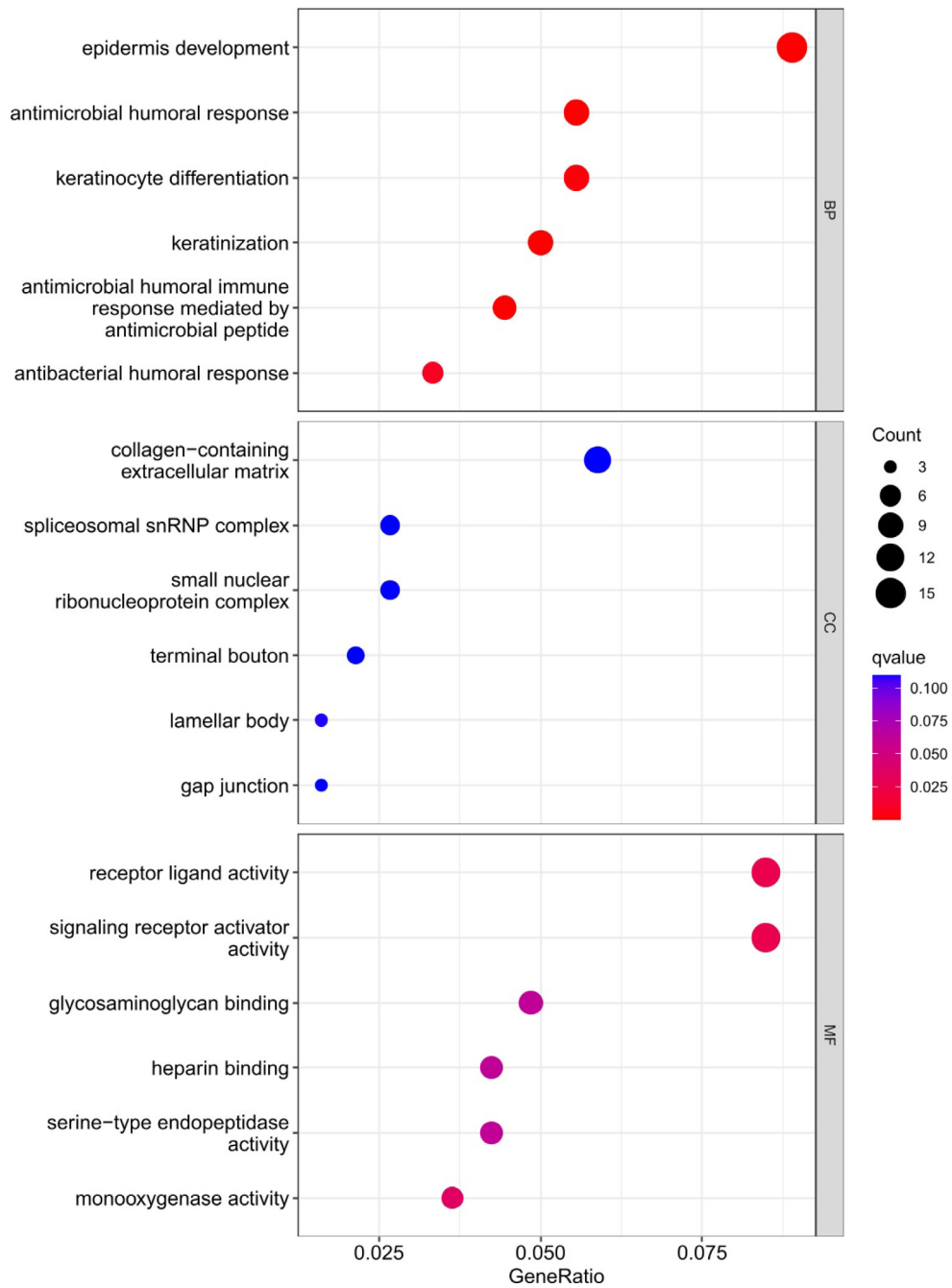


**Table S1** Clinical characteristics of NSCLC patients involved in the study

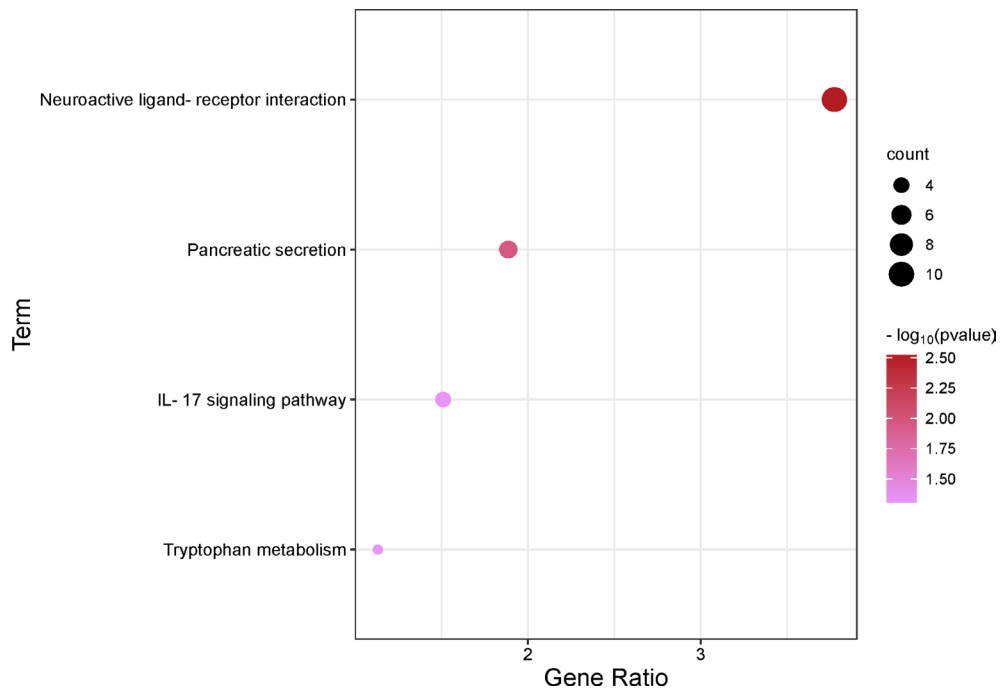
Characteristics	Testing cohort	Training cohort	TCGA cohort
Age, years			
≤65	213	215	428
>65	279	279	558
Unknown	9	7	16
Gender			
Female	190	211	401
Male	311	290	601
Stage			
I	256	258	514
II	144	135	279
III	81	83	164
IV	13	20	33
Unknown	7	5	12

**Table S2** Sources of cuproptosis-related genes

Cuproptosis-related genes	References
ATP7B	(40)
ATP7A	(41)
NFE2L2	(43)
NLRP3	(44)
SLC31A1, DBT, GCSH and DLST	(42)
FDX1, LIAS, LIPT1, LIPT2, DLD, DLAT, PDHA1, PDHB, MTF1, GLS and CDKN2A	(23)



**Figure S1** The GO enrichment analysis in the high- and low-risk groups of NSCLC. GO, Gene Ontology; NSCLC, non-small cell lung cancer.



**Figure S2** The KEGG enrichment analysis in the high- and low-risk groups of NSCLC. KEGG, Kyoto Encyclopedia of Genes and Genomes; NSCLC, non-small cell lung cancer.