

**Table S1** Primers used in real-time polymerase chain reaction assay

Gene name	Forward (5'-3')	Reverse (5'-3')
<i>SRC</i>	TGGCAAGATCACCAGACGG	GGCACCTTTCGTGGTCTCAC
<i>AIFM2</i>	GTGAGCGGGTGAGCAATCT	CTTGATGCCGGTGACAGAGAA
<i>SLC2A6</i>	CCGGACTACGACACCTTCC	GGATGTGTAGACCAGGGCATA
<i>HMOX1</i>	ATTCAGAAGGGCCAGGTGA	GGAAGTAGACAGGGGCGAAGA
<i>NGB</i>	ACAGTGGGTGAGTCTCTGCT	CCCGTAGAGTTGGCTCCAG
<i>MT1G</i>	CTTCTCGCTTGGGA ACTCTA	AGGGGTCAAGATTGTAGCAAA
<i>DRD4</i>	CCATCAGCGTGGACAGGTTCC	GCAGGGTAGGAAGAAGGAGCA
<i>AKR1C2</i>	CAACTTCAACCACAGGCTGC	GGGTCCACCCATGTTCTTC
<i>GAPDH</i>	AGCCTCAAGATCATCAGC	GAGTCCTTCCACGATACC

**Table S2** Antibodies used in Western blot assay

Antibodies	Source	Identifier
DRD4	Abcam	ab20424
SRC	Abcam	ab133283
AIFM2	Abcam	ab302673
AKR1C2	Abcam	ab179448

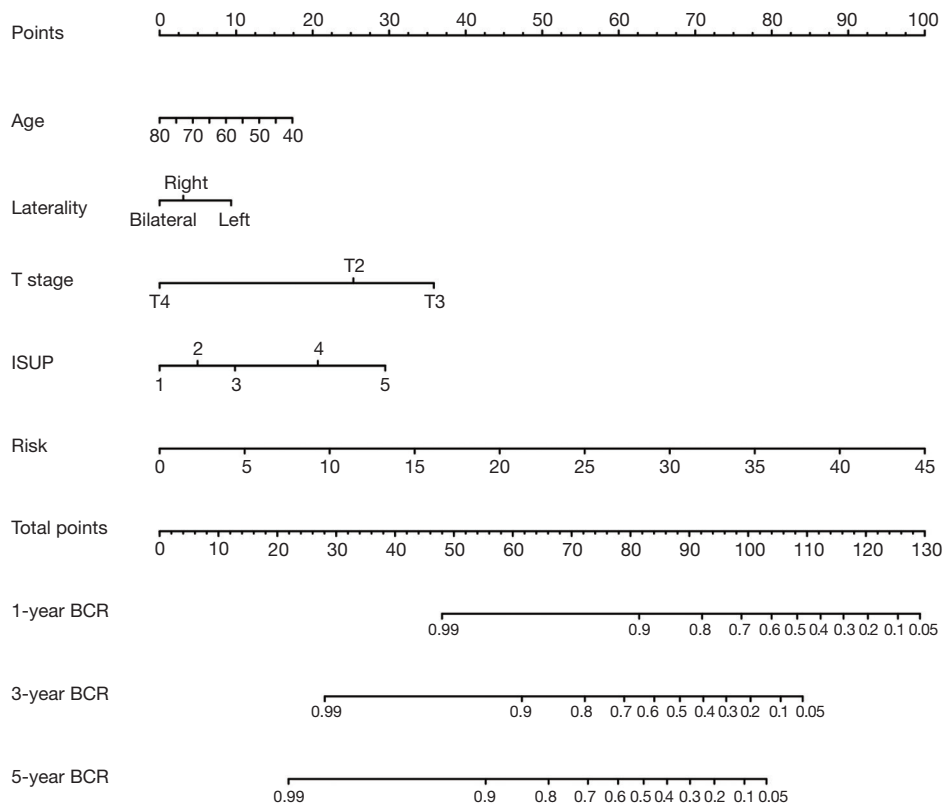
**Table S3** Clinical characteristics of the patients recruited for IHC

Disease	Sample ID	Age (years)	Sample types	PSA	Gleason score
BPH (N=6)	1	78	Biopsy	7.1	NA
	2	58	Biopsy	6.3	NA
	3	53	Biopsy	5.3	NA
	4	65	Biopsy	4.5	NA
	5	56	Biopsy	4.27	NA
	6	60	Biopsy	12.9	NA
PCa (N=10)	1	72	Biopsy	112	5+4=9
	2	69	Biopsy	56.4	4+4=8
	3	64	Biopsy	38.89	3+4=7
	4	61	Biopsy	33	4+4=8
	5	67	Biopsy	64	2+2=4
	6	77	Biopsy	8.5	3+3=6
	7	71	Biopsy	15	4+4=8
	8	63	Biopsy	7	5+4=9
	9	80	Biopsy	105	3+4=7
	10	56	Biopsy	17	5+5=10

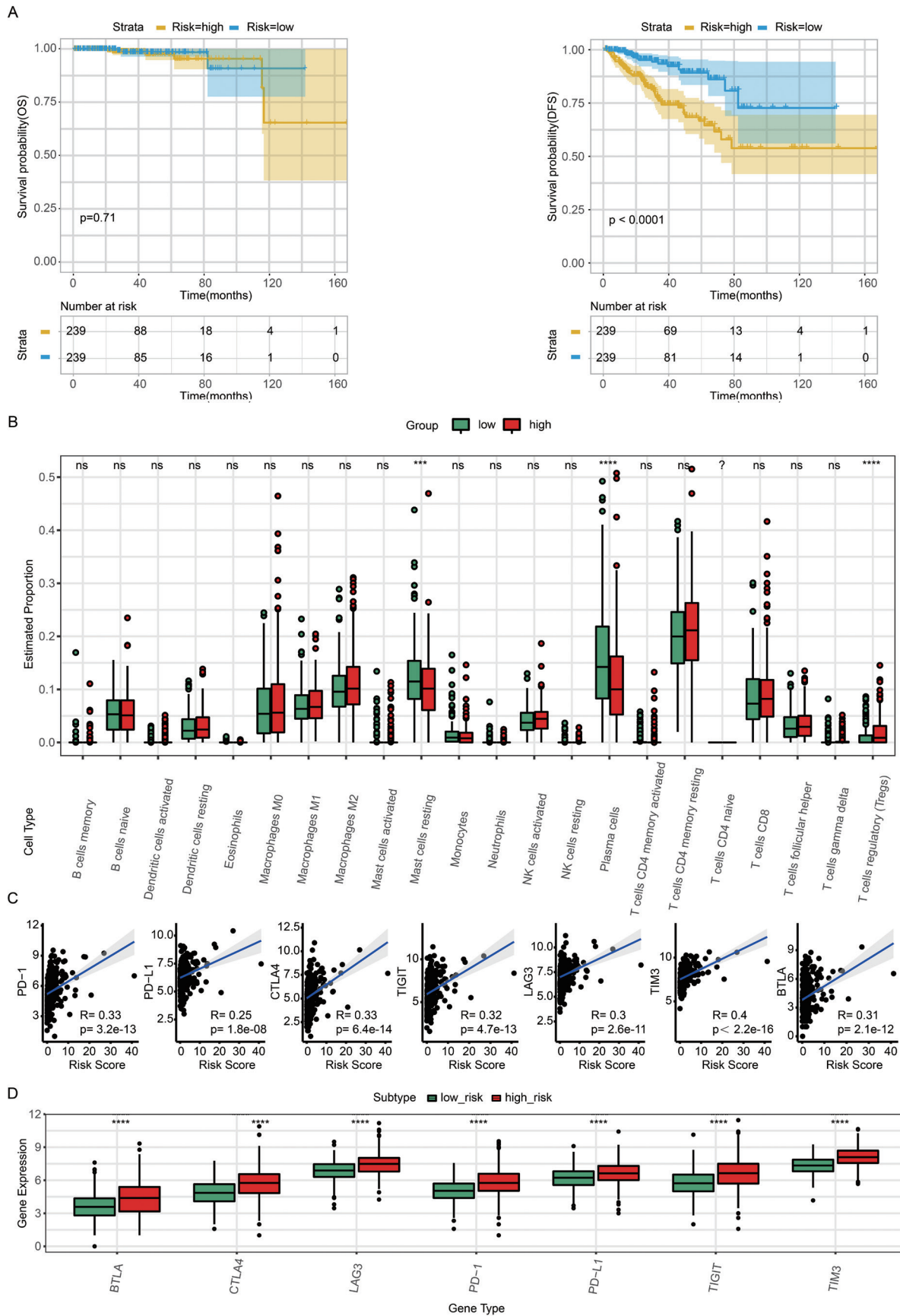
IHC, immunohistochemistry; PSA, prostate-specific antigen; BPH, benign prostatic hyperplasia; PCa, prostate cancer; NA, not applicable.



**Figure S1** Heatmap for the FRG prognostic signature and clinicopathological manifestations. BCR, biochemical recurrence; ISUP, International Society of Urological Pathology; FRG, ferroptosis-related gene.



**Figure S2** The nomogram combining clinicopathological variables and ferroptosis risk score was able to predict the 1-, 3-, and 5-year DFS of PCa patients. ISUP, International Society of Urological Pathology; BCR, biochemical recurrence; DFS, disease-free survival; PCa, prostate cancer.



**Figure S3** Gene expression profiles of immune cell populations. (A) Kaplan-Meier curves result of OS and DFS; (B) The proportion of immune infiltrating cells; (C) Correlation between immune checkpoint molecules and ferroptosis risk score; (D) The expression of immune checkpoint molecules. \*\*\*,  $P < 0.001$ ; \*\*\*\*,  $P < 0.0001$ ; ?, not be reliably quantified; ns, not significant. OS, overall survival; DFS, disease-free survival; PD-1, programmed cell death-1.