

Table S1. Clinicopathologic characteristics of the study population

Characteristics	n (%)*	BCR		
		n of events	Median, months	P‡
All patients	320	116	53	
Age at diagnosis, years				
≤65	141 (44.6)	51	46	0.291
>65	175 (55.4)	63	64	
PSA at diagnosis, ng/mL				
≤10	132 (42.7)	28	NR	<0.001
>10	177 (57.3)	83	30	
Pathologic Gleason score				
2–6	122 (39.1)	33	NR	<0.001
7–10	190 (60.9)	79	30	
Pathologic stage†				
Localized	206 (65.6)	46	NR	<0.001
Locally advanced	108 (34.4)	65	10	
Surgical margin				
Negative	196 (70.5)	51	NR	<0.001
Positive	82 (29.5)	47	22	
Lymph node metastasis				
Negative	290 (93.2)	92	71	<0.001
Positive	21 (6.8)	19	2	

Abbreviations: BCR, biochemical recurrence; PSA, prostate-specific antigen; NR, the median survival was not reached.

*Column subtotals do not sum to 320 due to missing data.

†Tumor-node-metastasis staging by American Joint Committee on Cancer in 1997: Localized: T1/T2; Locally advanced: T3/T4.

‡P values were calculated using the log-rank test and P < 0.05 are in boldface.

Table S2. Regulatory annotation of variants linked with rs2394882

SNP ID	Chromosome	Position	LD (r ²)	Reference allele	Alternate allele	ASN frequency	Variant type	Promoter histone marks	Enhancer histone marks	DNase	Proteins bound	Motifs changed
rs2022084	6	31149276	0.8	A	G	0.63	intronic					Nkx2
rs3130498	6	31149842	0.8	T	C	0.63	intronic		LIV			Lmo2-complex,M tf1,Myf,NRSF,TC F12,ZEB1,Zbtb3
rs130078	6	31150788	0.8	C	G	0.63	synonymous		ESDR, IPSC, GI			Myf,p53
rs3130500	6	31152199	0.81	T	A	0.63	intronic					COMP1,HNF4,R XRA
rs2240059	6	31152837	0.84	C	T	0.63	intronic		ESDR			AP-2,ERalpha-a, HNF4,NRSF,Sin3 Ak-20,Znf143
rs3130450	6	31153736	0.82	C	T	0.63	intronic					
rs3130452	6	31155536	0.81	A	G	0.62	intronic					Irf,SP1,TATA,ZB RK1
rs3095239	6	31159013	0.8	A	G	0.54	5'-UTR	ESC, ESDR, LNG, IPSC, FAT, STRM, BRST, BLD, MUS, BRN, SKIN, VAS, LIV, GI, ADRL, HRT, KID, PANC, PLCNT, THYM, OVRY, CRVX, BONE	SPLN	ESC,ESDR,ESDR ,ESC,LNG,BRST, BLD,BLD,SKIN, SKIN,HRT,LNG, THYM,BLD,CR VX,MUS,VAS,B LD,SKIN		
rs3094187	6	31159167	0.81	C	T	0.54	5'-UTR	ESC, ESDR, LNG, IPSC, FAT, STRM, BRST, BLD, MUS, BRN, SKIN, VAS, LIV, GI, ADRL, HRT, KID, PLCNT, THYM, OVRY, PANC, CRVX, BONE	HRT, LNG, SPLN	ESC,ESDR,ESC, BRST,SKIN,HRT ,MUS,MUS,CRV X,LIV,VAS,BLD		
rs3130934	6	31160775	0.99	A	C	0.59	intronic	BLD	BLD			Fox,Foxa,Foxd3,F oxf1,Foxi1,Foxj1, Foxj2,Foxl1,Foxp 1,Foxq1,Irf,RREB -1,Sox,Zfp105

rs2073721	6	31161839	0.84 A	G	0.63	missense	IPSC, ESC, SKIN, BLD	IPSC	POL2	EBF,Irf,MZF1::1- 4,NF-kappaB,Nkx 2,STAT SREBP
rs2073723	6	31162301	0.84 T	C	0.63	intronic	ESC, IPSC, SKIN, BLD, CRVX	ESC		
rs1065461	6	31162725	0.84 T	C,G	0.63	3'-UTR		ESC	POL2,POL24H8	
rs3132528	6	31163792	0.84 C	T	0.63	3'-UTR	BLD	MUS, LNG, CRVX, BRST	CRVX	AIRE,ATF3,CAC D,Rad21,SP2 Myf
rs1061118	6	31164637	1 G	A	0.59	synonymous				
rs2394882	6	31164872	1 A	C	0.59	intronic	BLD	IPSC, BLD		DEC,E2A,Irf,Nan og,Pou5f1,TAL1
rs2106074	6	31165732	0.99 G	A	0.59	intronic		ESDR, ESC, IPSC, BLD		
rs3130932	6	31166166	0.99 C	A	0.59	5'-UTR	BLD	ESC, IPSC	BLD,MUS	Hand1
rs3130929	6	31167430	0.83 T	C	0.63	intronic	ESC, IPSC	ESC		Egr-1,Mtf1,Znf14 3
rs9263804	6	31167929	0.83 C	T	0.63	intronic	ESC, ESDR, IPSC	ESC, SKIN		AP-1,Hdx,Irf,Pou 3f2,TATA,TCF12
rs3130501	6	31168676	0.82 A	G	0.63	intronic	ESC, ESDR, IPSC	ESDR, BLD, SKIN, CRVX	ESC	PU.1,TCF4
rs3130502	6	31168889	0.83 A	G	0.63	intronic	ESC, ESDR, IPSC	ESDR, SKIN, ADRL, PLCNT, CRVX		E2F,ERalpha-a,N RSF
rs3132524	6	31168937	0.83 T	C	0.63	intronic	ESC, ESDR, IPSC	ESDR, SKIN, ADRL, PLCNT, CRVX		SIX5
rs3132523	6	31169055	0.83 T	C	0.63	intronic	ESC, ESDR, IPSC	ESDR, SKIN, PLCNT, CRVX, BLD	ESC	Bcl6b,GATA,HD AC2,RXRA,TAL 1
rs3132522	6	31169982	0.82 T	C	0.64	intronic	ESC, ESDR, IPSC	ESDR, BRST, SKIN, GI, PLCNT, SPLN, CRVX, BLD	POL2,POL24H8, AP2ALPHA,AP2 B1	ERalpha-a,SETD B1
rs3094192	6	31170027	0.82 C	G	0.64	intronic	ESC, ESDR, IPSC	ESDR, BRST, SKIN, GI, PLCNT, SPLN, CRVX, BLD	POL2,POL24H8, AP2GAMMA	BCL,E2A,GR,Irf, LBP-9,Nkx2,PU.1 ,Pax-5,RXRA,ST AT
rs3130504	6	31170060	0.82 A	T	0.64	intronic	ESC, ESDR, IPSC	ESDR, BRST, SKIN, GI, PLCNT, SPLN, CRVX, BLD	POL2,POL24H8, AP2GAMMA	SMC3,YY1