





















Variable	N	Hazard ratio	p	
Age	546		1.03 (0.99, 1.06)	0.115
Stage	T2 188		Reference	
	T3 293		3.37 (0.63, 18.04)	0.156
	T4 10		0.00 (0.00, Inf)	0.998
GleasonScore	498		2.97 (1.34, 6.59)	0.007
n_SNV	543		1.00 (1.00, 1.00)	0.766
n_driver	546		1.10 (0.96, 1.27)	0.176
Ti_fraction	541		0.98 (0.95, 1.00)	0.056
Tv_fraction	541		1.03 (1.00, 1.05)	0.056
cnaBurden	546		1.04 (1.00, 1.07)	0.058
n_CNV	546		1.01 (1.00, 1.01)	<0.001
n_Amp	546		1.01 (1.00, 1.01)	<0.001
n_Del	546		1.02 (1.00, 1.04)	0.036
ploidy	546		1.07 (0.88, 1.30)	0.502
TD score	546		3.21 (1.89, 5.43)	<0.001
Chromothripsis score	546		1.01 (1.00, 1.02)	0.089
MACN score	543		1.06 (0.93, 1.22)	0.383
MATH	543		1.00 (0.98, 1.01)	0.802
n_mutation_cluster	543		1.35 (1.04, 1.75)	0.022
purity	546		7.72 (2.12, 28.21)	0.002