

	Group 1 : <3 cores, <3mm (n=89)	Group 2 : <3 cores, <50% (n=112)	Group 3 : <33% positive cores (n=177)	Comparisons between groups (p values)		
				1 vs 2	2 vs 3	1 vs 3
Mean PSA density (ng/ml/gr)	0.165	0.166	0.172	0.925	0.516	0.487
<i>Pathological features on biopsies</i>						
Mean percent of cancer in any core (%)	8.0	11.7	15.8	0.004	0.006	<0.001
No of positive cores	1.30	1.35	2.21	0.504	<0.001	<0.001
Total tumor length (mm)	1.2	2.0	5.2	<0.001	<0.001	<0.001
<i>Pathological features on RP specimens</i>						
Gleason score:						
6	46	52	79	0.640	0.971	0.589
7	42	58	95			
8 or more	1	2	3			
Primary Gleason 4	8	13	22	0.547	0.835	0.403
pTNM:						
T0	1	1	1	0.457	0.499	0.160
T2	77	93	142			
T3-T4	11	18	34			
Positive surgical margins:	16	22	37	0.765	0.796	0.573
Extracapsular extension:	10	16	31	0.522	0.456	0.175
Seminal vesicle invasion:	1	2	2	1.00	0.643	1.00
Overall unfavourable*	19	27	49	0.644	0.501	0.264
Overall unfavourable**	12	19	36	0.497	0.476	0.170
PSA failure	5	7	8	0.851	0.518	0.766

* Defined as ECE, SVI and/or primary Gleason score 4

** Defined as ECE, SVI and/or Gleason score ≥8