

P #	PSA at diagnosis	PSA Post-RP	GS Primary	GS Secondary	Gleason	T Stage	Margin	Stratification	Notes
P5004		0.17	4	4	8	T3b	-	High Risk	
P5006	4	0.36	3	4	7	T3a	-	Low Risk	
P5007	10	0.26	4	3	7	T3a	-	Low Risk	
P5008	10	0.12	4	3	7	T2c	-	Low Risk	
P5010	18	0.04	3	4	7	T3b	-	High Risk	
P5011	5	0.03	4	4	8	T3a	+	High Risk	
P5013	6	0.02	4	3	7	T3a	+	High Risk	
P5014	17	0.48	4	5	9	T3a	+	High Risk	
P5015	140	0.03	3	4	7	T3a	+	High Risk	
P5018	17	0.20	4	3	7	T3a	+	High Risk	
P5019	17	0.01	4	3	7	T2c	+	Low Risk	
P5021	9	0.35	3	4	7	T3a	+	High Risk	
P5022	19	2.28	4	5	9	T3b	-	High Risk	
P5026	6	0.13	3	4	7	T2c	+	Low Risk	
P5027	9	0.01	4	3	7	T3b	-	High Risk	
P5028	7	0.02	4	5	9	T3a	-	High Risk	
P5029	7	0.00	4	5	9	T3a	-	High Risk	
P5030	7	0.04	4	3	7	T2c	+	Low Risk	
P5031	22	0.10	4	5	9	T3b	+	High Risk	
P5034	10	0.59	3	4	7	T3a	-	Low Risk	
P5035	20	0.99	4	5	9	T3b	+	High Risk	
P5036	5	0.11	5	3	8	T3a	-	High Risk	
P5037	11	0.10	4	3	7	T3b	+	High Risk	
P5039	5	0.24	3	4	7	T3b	+	High Risk	
P5048	5	0.16	4	4	8	T3a	+	High Risk	
P5049	67	16.59	5	4	9	T3b	+	High Risk	
P5051		0.02	3	4	7	T2c	-	Low Risk	
P5054	5	0.10	4	4	8	T3a	-	High Risk	
P5056	7	0.00	3	3	6	T3a	-	Low Risk	
P5057	10	0.48	4	5	9	T3b	+	High Risk	
P5060	4	0.02	3	4	7	T2	-	Low Risk	
P5063	12	0.02	4	5	9	T3b	-	High Risk	
P5064	16	0.00	4	3	7	T3a	+	High Risk	

P5066	13	0.00	4	3	7	T3b	+	High Risk	
P5067	10	0.09	4	3	7	T2c	+	Low Risk	
P5068	7	0.00	3	4	7	T2c	-	Low Risk	
P5069	5	0.00	4	3	7	T3a	+	High Risk	
P5072		0.06	4	3	7	T3b	+		*
P5073	5	0.88	3	3	6	T2	-	Low Risk	
P5077	14	2.94	4	5	9	T3b	-	High Risk	
P5078	20	0.00	4	5	9	T3b	+	High Risk	
P5080	7	0.27	3	4	7	T2	+	Low Risk	
P5090	11	0.00	3	4	7	T3b	-	High Risk	
P5095		0.00	3	4	7	T3a	-		*
P5097	14	0.31	5	4	9	T3b	+	High Risk	
P5098	13	0.44	4	3	7	T4	+	High Risk	
P5099	10	0.10	3	4	7	T3b	-	High Risk	
P5101		0.38	3	3	6	T2c	+	Low Risk	
P5102	6	0.00	4	5	9	T2c	+	High Risk	
P5103	16	0.18	4	3	7	T3a	+	High Risk	
P5104	7	0.00	3	4	7	T2c	-	Low Risk	
P5105	12	0.31	4	4	8	T2a	-	High Risk	
P5106	4	0.00	3	4	7	T2c	-	Low Risk	
P5108	9	3.80	4	5	9	T3a	+	High Risk	
P5110		0.00	3	4	7	T2	-	Low Risk	
P5112	5	0.03	4	4	8	T3a	+	High Risk	
P5113	9	0.52	3	4	7	T2a	-	Low Risk	
P5115	6	0.04	4	5	9	T3a	+	High Risk	
P5116	8	0.00	3	4	7	T2c	+	Low Risk	
P5117		0.99	3	4	7	T2	-	Low Risk	
P5118	6	0.00	4	3	7	T2c	+	Low Risk	
P5119	4	0.44	4	3	7	T2	-	Low Risk	
P5120	2	0.03	4	4	8	T3a	+	High Risk	
P5121	5	0.26	3	4	7	T2	+	Low Risk	
P5122	10	0.08	4	3	7	T3a	-	Low Risk	
P5123		0.18	4	3	7	T2c	+		*
P5125	6	0.27	4	3	7	T3a	-	Low Risk	

P5126	6	0.06	3	4	7	T3a	+	High Risk	
P5127	9	5.08	3	5	8	T2	-	High Risk	
P5128	12	0.14	3	4	7	T3a	+	High Risk	
P5129		1.72	3	4	7	T2	+	Low Risk	
P5130	5	0.00	4	5	9	T3b	+	High Risk	
P5131	11	0.66	4	3	7	T3a	+	High Risk	
P5132	5	0.14	3	4	7	T2	+	Low Risk	
P5133	4	0.15	4	3	7	T3a	-	Low Risk	
P5135	10	0.00	3	4	7	T3a	+	High Risk	
P5137	6	0.11	3	4	7	T2	-	Low Risk	
P5139	4	0.15	3	4	7	T3a	-	Low Risk	

High risk	44
Low risk	31

75

* denotes insufficient RNA quality/quantity and sample excluded