

Supplemental Table 6. Comparison of Motif characteristics in tumor samples.

	QSRANV		PIW		GTFR		KYTG		RHH		P-value ¹
	Baseline	Week 5	Baseline	Week 5	Baseline	Week 5	Baseline	Week 5	Baseline	Week 5	
Count, Mean (SD)	4.7 (2.4)	3.4 (2.6)	0.6 (1.1)	0.3 (0.9)	0.3 (0.8)	0.3 (1.1)	2.4 (1.9)	2.2 (1.9)	0.4 (1.3)	0.2 (0.7)	<0.01**
Proportion (of all clones), Mean (SD)	2E-3 (4E-3)	3E-3 (7E-3)	7.9E-5 (1E-4)	8.6E-5 (2E-4)	1.8E-5 (4.2E-5)	6.1E-5 (1E-4)	6.8E-4 (1.7E-3)	2.7E-3 (6.3E-3)	6.5E-5 (2.9E-4)	5.3E-5 (1.5E-4)	<0.01**
Patients' motif presence at baseline, N(%)											0.54 ⁺
Presence	40 (97.6)	-	13 (31.7)	-	9 (22.0)	-	33 (80.5)	-	7 (17.1)	-	
Absence	1 (2.4)	-	28 (68.3)	-	32 (78.0)	-	8 (19.5)	-	34 (82.9)	-	
Change in proportion from baseline[#], N(%)											0.35 ⁺
Increase	-	29 (70.7)	-	5 (29.4)	-	7 (50.0)	-	28 (71.8)	-	5 (45.5)	
Decrease	-	12 (29.3)	-	12 (70.6)	-	7 (50.0)	-	11 (28.2)	-	6 (54.5)	

¹ Overall p-value comparing all motifs

⁺ Comparing baseline values across all motifs

^{*} Significant P-value

[#] Patients with no motifs present at baseline and week 5 where excluded from the analysis