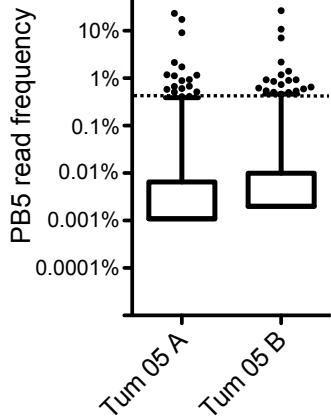
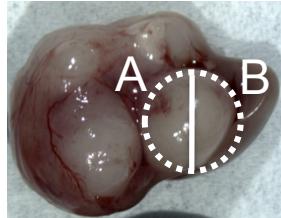


A

Tumor 05, liver Formalin fixed tissue		
	Insertions	PB5 Reads
Tumor 05, A	1857	90406
Tumor 05, B	1902	50160

**Tumor 05 A**

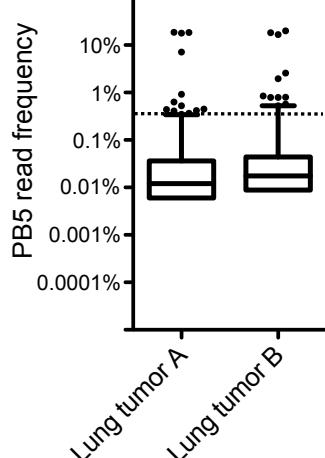
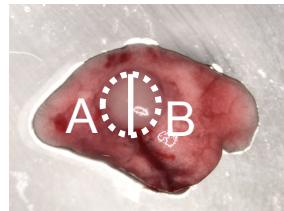
Chr	Position	PB5 f	
chr11	41573884	23.2%	intergenic
chr18	16209166	17.4%	intergenic
chr1	87855454	9.1%	Dgkd
chr8	38677940	2.2%	intergenic
chr19	34319719	1.7%	Fas
chr19	32786662	1.2%	Pten
chr8	100880553	1.2%	intergenic
chr13	69768968	1.1%	intergenic
chr6	120080572	0.9%	intergenic
chr1	156608148	0.9%	Abi2
chr3	100720626	0.7%	intergenic
chr9	114777333	0.7%	Cmtm7
chr1	181237712	0.6%	intergenic
chr16	23230498	0.6%	St6gal1
chr1	134962318	0.5%	intergenic
chr6	131293588	0.5%	Magohb
chr13	32829461	0.4%	intergenic
chr2	158268767	0.4%	Bpi1
chr9	98589940	0.4%	Mrps22

Tumor 05 B

Chr	Position	PB5 f	
chr11	41573884	26.5%	intergenic
chr18	16209166	10.7%	intergenic
chr1	87855454	7.1%	Dgkd
chr8	38677940	2.2%	intergenic
chr19	34319719	1.4%	Fas
chr19	32786662	1.2%	Pten
chr1	156608148	0.9%	Abi2
chr1	97039868	0.9%	intergenic
chr6	120080572	0.9%	intergenic
chr9	114777333	0.9%	Cmtm7
chr11	22125225	0.7%	Ehbp1
chr9	63572479	0.7%	Igch
chr6	131293588	0.6%	Magohb
chr8	100880553	0.6%	intergenic
chr13	69768968	0.5%	intergenic
chr16	95849795	0.5%	1600002D24Rik
chr9	5699470	0.5%	intergenic
chr9	98589940	0.5%	Mrps22
chr2	11637292	0.5%	intergenic
chr8	87954812	0.5%	Zfp423
chr3	100720626	0.4%	intergenic
chr8	28543227	0.4%	intergenic
chr12	53713078	0.4%	Npas3
chr15	39478446	0.4%	Rims2

B

Animal ID #4180 R26-PBase; ATP1-S2; INK+/- Age: 42 weeks		
	Insertions	PB5 Reads
Lung tumor, A	1327	16314
Lung tumor, B	1109	11394

**Lung tumor A**

Chr	Position	PB5 f	
chr11	80623112	18.6%	Myo1d
chr9	89968709	18.4%	Rasgrf1
chr12	69422766	17.8%	intergenic
chr10	56904704	7.1%	intergenic
chr18	14015898	0.9%	intergenic
chr10	15213909	0.6%	intergenic
chr10	32264523	0.5%	Nkain2
chr10	67016841	0.4%	Reep3
chr10	86146164	0.4%	Syn3
chr6	17017596	0.4%	intergenic
chr4	54976180	0.4%	Zfp462

Lung tumor B

Chr	Position	PB5 f	
chr12	69422766	19.9%	intergenic
chr11	80623112	18.2%	Myo1d
chr9	89968709	16.6%	Rasgrf1
chr10	56904704	2.6%	intergenic
chr7	143292394	2.0%	Kcnq1
chr2	52623038	0.8%	Cacnb4
chr18	14015898	0.8%	intergenic
chr10	15061726	0.8%	intergenic
chrX	89274343	0.8%	intergenic
chr10	86146164	0.6%	Syn3
chr2	76150368	0.5%	Pde11a
chr10	119901129	0.5%	Grip1
chr10	32264523	0.5%	Nkain2
chr10	67016841	0.5%	Reep3
chr2	49096760	0.4%	Mbd5
chr10	15091000	0.4%	intergenic