

Supplementary Table 1: TSC1/TSC2 mutation findings in angiomyolipoma subject to H3K27ac and MITF ChIP-Seq.

Tumor ID	Gene	Nucleotide change (hg19)	Protein Change	Variant Classification	Allele fraction	Tumor characterization
AML-1	TSC1	c.2183_2186dupCTCT	p.E730Sfs*5	Frameshift duplication	0.66	2-hit tumor/ Copy neutral Loss of heterozygosity
AML-2	TSC2	c.421G>T	p.E141*	Nonsense	0.06	2-hit tumor
		c.5238delC	p.H1746Qfs*80	Frameshift deletion	0.06	
AML-3		c.100A>T	p.K34*	Nonsense	0.19	2-hit tumor
		c.481+1G>A	p.A161_splice	Splice	0.19	
AML-4		c.265delT	p.L89Cfs*16	Frameshift deletion	0.35	2-hit tumor
		c.3097_3098insA	c.Y1033*	Frameshift insertion	0.38	