SUPPLEMENTARY MATERIAL

Supplementary Table 1: ddPCR probe specifications. Commercially available probes were purchased from Biorad and used following manufacturer's instructions for amplification cycles, with annealing temperatures as shown (in °C)

Probeset description	Assay ID	Annealing temperature
ddPCR [™] Mutation Assay: AKT3 p.E17K, Human	dHsaMDS121759946	55
ddPCR™ Mutation Assay: PIK3CA p.E542K, Human	dHsaMDV2010073	57
ddPCR™ Mutation Assay: PIK3CA p.E545K, Human	dHsaMDV2010075	57
ddPCR™ Mutation Assay: PIK3CA p.H1047R, Human	dHsaMDV2010077	55
ddPCR™ Mutation Assay: MTOR p.S2215F, Human	dHsaMDS365202467	55
ddPCR™ Mutation Assay: MTOR p.S2215Y, Human	dHsaMDS2514998	55

Supplementary Table 2: Complete list of samples and ddPCR results for the 17 mutation positive patients and comparison with orthogonal metods. Patients' de-identified code (Patient-ID), sex, diagnosis based on neuroimaging (Dx-MRI), Diagnosis based on neuropathology (Dx-NPath), genetic variant, sample type and results from the different variant identification methods (ddPCR, Oncoplex, smMIPs, Exome sequencing, and Amplicon sequencing are listed for each of the 17 mutation-positive patients. For ddPCR results, Variant allele fraction (VAF%), Delta confidence interval (DCI), and number of wild-type and mutant droplets (average of quadruplicate) are reported. Samples were considered positive for a given mutation via ddPCR when DCI>0.045 and VAF>0.05% (green text). Negative samples are indicated in red text. False positive results are indicated in green bolded text and *, while false negative results are indicated in red italic text and **. Empty cells represent lack of testing due to exhaustion of sample.

Patient ID LR16-470	Sex	Dx – MRI					IdPCR		Oncoplex	,		ID.	Exomes		Amplicon Sc	
		DX - MILL	Dx - NPATH	Genetic variant	Sample type	Mutant/total reads		DCI	Mutant/total reads	VAF %	smM Mutant/total reads	VAF %	Mutant/total reads	VAF %	Amplicon So Mutant/total reads	eq VAF %
	М	HMEG	NA	PIK3CA (p.E542K)	Saliva	70/4568	1.47	1.056	7/836	0.837						
LR13-197	F	DMEG	FCD 1b	PIK3CA (p.E545K)	Blood	4/9280	0.04	-0.077								
Latto tor	•	Dinto	10010	The en (plus on)	Saliva	792/3776	20.68	19.398								
LR15-251	м	HMEG	NA	PIK3CA (p.E545K)	Brain, FFPE Blood	246/1297 2/11069	18.86 0.02	16.691 -0.049	23/185 1/1241	12.432 0.081	0/205	0.00				
LICID-201	ivi	mand		тизся (р.1545к)	Biod	8/6472	0.02	-0.049	1/1241	0.081	0/164	0.00				
LR16-242	F	HMEG	NA	PIK3CA (p.E545K)	Saliva	1757/7865	21.76	20.780	169/1015	16.650	47/165	28.485				
					Brain	1827/8430	21.01	19.996			95/522	18.199				
LR16-413	М	HMEG	FCD	PIK3CA (p.E545K)	Peripheral blood						0/636	0.000				
					Skin fibroblasts	1377/5937	22.67	21.450	227/1007	20.902	0/582	0.000				
					Saliva Brain, anterior dysplasia	3/9796	0.03	21.459 -0.072	227/1086	20.902						
					Brain, superior temporal	10/14226	0.05	-0.072								
					Brain, temporal tip	17/11562	0.13	0.000								
					Brain, inferior temporal	44/14122	0.28	0.120	0/804**	0						
					Brain, anterior inferior	21/12662	0.15	0.010								
					Brain, middle temporal	10/9044	0.10	-0.040								
LR18-024	F	FCD	FCD 2a	PIK3CA (p.E545K)	Brain, amygdala Brain, hippocampus	21/15168 7/6259	0.12 0.11	0.000								
					Brain, cusa	2/4268	0.05	-0.072								
					Brain, FFPE, superior temporal	3/9620	0.03	-0.26								
					Brain, FFPE, inferior temporal 1	4/5482	0.07	0.00								
					Brain, FFPE, inferior temporal 2	9/6090	0.14	0.05								
					Brain, FFPE, inferior temporal 3	3/8068	0.03	-0.06	0/650	0						
LR12-251	F	FCD	FCD 2a	PIK3CA (p.H1047R)	Saliva Brain	574/11451	4.58	4.186	0000	U						
	~				Brain, frontal lobe	721/5114	13.76	12.771	i		104/1063	9.784	İ			1
ļ					Brain, operculum	1067/7604	13.51	12.703			92/932	9.871				
ļ					Brain, superior temporal	1404/7549	17.93	16.987			211/1172	18.003				
ļ					Brain, anterior temporal 1	1697/11371	14.06	13.352			192/1168	7.877				
LR11-443	F	HMEG	ECD 24	AKT2 (= E17V)	Brain, anterior temporal 2	1062/6515 88/8502	15.78	14.820								
LK11-443	£.	nated	FCD 2a	AKT3 (p.E17K)	Brain, dura Brain, parietal	88/8502 587/7862	0.95	0.668 6.437								+
					Brain, other	839/6792	11.80	10.963								
ļ					Brain, hippocampus	1039/6864	14.54	13.64			164/1090	15.046				
					Epidermis	84/11348	0.67	0.44	10/789	1.267						
					Cultured fibroblasts	4/7448	0.05	-0.09								
LR12-317	М	HMEG	NA	AKT3 (p.E17K)	Brain, FFPE Saliva	99/762 14/2347	12.93 0.58	10.461 0.208	3/1143	0.262						
					Brain	53/7955	0.58	0.208	3/1143	0.202						
LR13-351	F	FCD	FCD 2a	AKT3 (p.E17K)	Blood	3/9093	0.03	-0.023								
\neg					Brain, temporal lobe	57/3263	1.66	1.231								
					Brain, hippocampus	16/1254	1.26	0.639								
					Brain, insula-1	245/3262	7.26	6.384								
					Brain, insula-2 Brain, temporal tip	453/7056 80/3618	5.93 2.13	5.395 1.670								
LR16-313	F	FCD	FCD 2a	AKT3 (p.E17K)	Brain, FFPE-1	134/2048	6.44	5.315								
					Brain, FFPE-2	189/2621	7.08	6.033								
					Brain, FFPE-3	169/2018	8.25	6.990	38/497	7.646						
					Brain, FFPE-4	39/1576	2.44	1.608								
1012.244		mme	ECD A.	MEOD (- 02212E)	Skin	0/3688 19/10964	0.00	0.000	5/2879	0.002	0/48	0.000				
LR12-246	М	HMEG	FCD 2a	MTOR (p.S2215F)	Brain Brain, lateral frontal lobe	2/10948	0.16	0.047 -0.051	5/2879	0.002	0/48	0.000				
					Brain, medial frontal lobe	6/12085	0.02	-0.043				0.000				
					Brain, orbital frontal lobe	75/11983	0.57	0.388	11/1640**	0.0067	0/15**	0.000				
					Brain, FFPE, frontal lobe	8/12792	0.06	-0.034								
LR13-129	М	FCD	FCD 2b	MTOR (p.S2215F)	Brain, FFPE, parietal lobe	5/10161	0.05	-0.047								
					Brain, FFPE, posterior temporal lobe Brain, FFPE, occipital lobe	30/11250 4/11120	0.24	0.104 -0.051								
					Brain, FFPE, white matter	6/8714	0.06	-0.040								
					Brain, FFPE, superior frontal lobe	9/10528	0.08	-0.025								
					Brain, superior temporal	55/7789	0.65	0.390			0/351*	0.000			0/435*	0
ļ					Brain, inferior temporal	21/30292	0.05	-0.030			0/178	0.000			0/2798	0
ļ					Brain, middle temporal Brain, basal temporal	18/30290 6/15644	0.04	-0.030 -0.050			0/266 0/238	0.000			0/2416	0
ļ					Brain, amygdala	2/9287	0.03	-0.030			0/302	0.000			0/2941	0
ļ					Brain, hippocampus	4/8328	0.02	-0.047			3/246*	1.220			9/7583*	0.11868654
ļ					Brain, parietal lobe	243/9276	2.44	2.090			0/283**	0.000			0/560**	0
LR13-389	F	FCD	FCD 2a	MTOR (p.S2215F)	Brain, posterior temporal-1	1200/16993	6.22	5.824			15/158	9.494			76/1828	4.15754923
					Brain, posterior temporal-2	294/10678 16/9395	2.54 0.16	2.202 0.032			7/220 6/204	3.182 2.941			47/4016 54/2361	1.17031873
ļ					Brain, frontal operculum Brain, mid-frontal	4/8203	0.16	-0.032			0/204	2.941			34/2301	2.28716645
ļ					Brain, posterior frontal	421/9411	4.16	3.727								1
ļ					Brain, posterior temporal	143/8092	1.66	1.346								
ļ					Brain, inferior frontal	9/8096	0.10	-0.012								
ļ					Brain, orbital frontal	9/7531	0.11	-0.010								
LR14-046	М	FCD	FCD	MTOR (p.S2215F)	Saliva Brain, deep temporal	2/7167 323/7638	0.03 3.91	-0.047 3.379	75/2513	0.030						-
LA14+040	101	TCD	reb	MION (0.52215F)	Brain, deep temporal Brain, posterior frontal	323/7638 38/10187	0.34	0.121	6/1419	0.030						
LR14-155	F	FCD	FCD 2b	MTOR (p.S2215F)	Brain, anterior frontal	8/11312	0.06	-0.091	0/1417	0.000						
				- /	Saliva								0/91	0.00		
Ţ					Brain, lateral temporal lobe	601/7966	7.18	6.626			2/21	9.524				
ļ					Brain, inferior temporal lobe	89/7954	1.05	0.836			1/69	1.449				
ļ					Brain, temporal tip Brain, hippocampus	248/5042 2/6077	4.72 0.03	4.143 0.000			3/23 0/5	13.043 0.000				
ļ					Brain, hippocampus Brain, anterior frontal lobe	1/6009	0.03	0.000			0/102	0.000				1
ļ	м	HMEC	ECD 24	MTOR (P COLEV)	Brain, middle frontal lobe	877/6238	13.59	12.756			0/0	0.000				
1 8 16 004	М	HMEG	FCD 2a	MTOR (p.S2215Y)	Brain, parietal lobe	790/7623	9.88	9.224			5/75	6.667				
LR16-004			1		Brain, insula and operculum	798/8803	8.54	7.973			8/58	13.793	16/111	14.41		
LR16-004							1.60	1.341			3/53	5.660		1		
LR16-004					Brain, ventricle wall Brain, fronted basel contex	147/8551										
LR16-004					Brain, ventricle wall Brain, frontal basal cortex Brain, gyrus rectus	147/8551 233/6872 0/8727	3.20	2.798			4/29	13.793				-

Supplementary Table 3: Summary of clinical and neuroimaging features for the 17 mutation positive patients. Abbreviations: d, day; DMEG, dysplastic megalencephaly; FCD focal cortical dysplasia; HMEG, hemimegalencephaly; y, year; m, month; PQD, posterior quadrant dysplasia; N/A, not available. Data from two patients have been previously published, namely: LR13-389 (PMID 27159400); LR11-443 (PMID 28969385, 25722288, 27159400).

Supplementary Table 4: Data summary of patients and samples who tested negative for the 6 PI3K-AKT-MTOR hotspot mutations. Patients' de-identified code (Patient-ID), diagnosis based on neuroimaging (Dx-MRI), Diagnosis based on neuropathology (Dx-NPath), and sample type are listed for the mutation-negative patients (n=41). All samples were tested via ddPCR for all 6 hotspots in quadruplicates and in multiple independent runs. All samples had a DCI<0.045, thus they were considered negative (cells shaded in red). The average of each quadruplicate is reported here. Number of WT droplets is reported as measure of quality of the ddPCR run; samples with over 3000 were considered robust and subsequently analyzed (green shaded cells). If samples had less than 3000 WT droplets (yellow shaded cells) and VAF<0.1%, they were deemed negative. When multiple runs were performed for a given sample, results from the run yielding the highest number of wild-type droplets (WT droplets) is reported here. Patient LR12-245 was previously published in PMID: 27159400.

Sex Dx Starres - age of # of surgeries Sar (Neuropathology) onset (mo)	Seizures – age of # of surgeries onset (mo)	# of surgeries		Sample type	VAF%	ă	K3C4 p.	lets Mut droplets	VAF%	K3C	lets	Mut droplets VAF%	<u>^</u>	3C4 p-	ts Mut droplets	1	DCI DCI	MTOR p.S2115F WT Droplets	Mut droplets	VAF%	TOR	P.S2215Y WT Droplets Mut droplets	-	AKT3 DCI	AKT3 p.E17K	Mut drophets
F FCD 15 9.00 2 Binin 0.04 -0.083 4494 2	9.00 2 Brain 0.04 -0.083 4994 2	2 Brain 0.04 -0.083 4994 2	0.04 -0.083 4994 2	0.04 -0.083 4994 2	-0.083 4994 2	4994 2	2	0.01	+	-0.084	9127	1 0.02	+	+	c ,	0.03	0.000	6704	2	0.04	+	10584 5	0.01	-0.026	9337	
0.03 -0.093 5469 I 0.03	6.00 3 Barin modification 0.03 -0.093 -2469 1 0.03 Barin modification 0.03 -0.093 -2469 1 0.03	3 Saliva 0.03 -0.095 5469 I 0.03 Banin mudial 0.07 0.02 7813 7 0.05	Banin model and and a 0.03 -0.093 3469 I 0.03 Banin model and and a 0.03 -0.093 7915 7 0.05	Saltva 0.03 -0.093 3469 I 0.03 muddalamidad 0.02 0.093 2812 7 0.05	-0.095 5469 I 0.03	2469 I 0.03	1 0.03	+		+	5/6/	4 0.0	0.000	0152 78C0		000	0000	2955	0 -	0.00	0000	35/9 0	0.02	00070	35/3	
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M FCD2a 18.00 2 0.000 1.	18.00 1 Brain 0.02 0.000 10134 3	2 Dian 0.00 1000 10000 1 1	0.02 0.000 12154 3	0.02 0.000 12154 3	0.000 12154 3	12154 3		0.05	ľ	0.075	10954	6 0.0		12086	* ~	50.0	0.004	9732	n **	0.0		1839 0	0.03	-0.023	12903	• •
F FCD1a 105.00 2 Brain 0.03 0.000 10059 3	105.00 2 Brain 0.03 0.000 10059 3	2 Brain 0.03 0.000 10059 3	0.03 0.000 10059 3	0.03 0.000 10059 3	0.000 10059 3	10059 3	3	0			8979	0 0.0	H		2	0.03	0.000	8018	3	0.01		1 10126	0.01	-0.026	9025	1
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E F FCD 2a 182.50 1 Brain 0.07	182.50 I Brain 0.07 0.021	1 Brain 0.07 0.021	0.07 0.021	0.07 0.021	0.021		80	_	0.02		9182	2 0.03		10729	. 6	0.07	0.006	6653	5	0.05		9450 S	0.06	-0.012	10613	1
F FCD1a N/A 1 Saliva 0.04 0.000	N/A 1 Saliva 0.04 0.000	1 Saliva 0.04 0.000	0.04 0.000	0.04 0.000	0.000	+	4		-	-0.116	6434	3 0.0	+	7651	-	0.01	-0.049	8275	_	0.00	-	9010 0	0.03	-0.036	9656	9
FCD 1b 114.00 2 Brain 0.03 0.000	114.00 2 Brain 0.03 0.000	2 Brain 0.03 0.000	0.03 0.000	0.03 0.000	0.000	-	4		0.02	-0.116	8739	2 0.05	6 -0.050	9413	~	0.05	-0.044	10037	5	0.01		11206 1	0.02	-0.036	11883	2
M FCD 57.00 3 Brain 0.05 0.004	57.00 3 Brain 0.05 0.004	3 Brain 0.05 0.004	0.05 0.004	0.05 0.004	0.004		~		0.01	-0.116	8300	1 0.01	+	9711		0.05	-0.044	9126	~	0.02	0.000	1256 2	0.03	-0.036	1044	4 (
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73.00 1 Brain – Superficial cortex 0.01 0.000	73.00 1 Brain – Superficial cortex 0.01 0.000	1 Brain – Superficial cortex 0.01 0.000	- Superficial cortex 0.01 0.000	- Superficial cortex 0.01 0.000	0.000				0.04	-0.129	7407	3 0.0	G -0.058	6772	2	0.06	-0.071	6703	4	0.00	-0.051	7466 0	0.07	-0.099	7135	S
F FCD 3d 12.17 1 Saliva 0.00 0.000	12.17 1 Saliva 0.00 0.000	1 Saliva 0.00 0.000	0.00 0.00	0.00 0.00	0.000		0	+	0:00	-0.129	2653	0 0.0		2471	0	0.00	-0.071	2917	0	0.00	-0.051 2	2560 0	0.00	-0.105	2318	0
30.00 I Brain 0.03 -0.083	30.00 I Brain 0.03 -0.083	1 Brain 0.03 -0.083	0.03 -0.083	0.03 -0.083	-0.083		3		0.04	-0.084	9261	4 0.0.	1 -0.025	8971		0.04	0.000	7650	3	0.04	0.008	2374 6	0.04	-0.026	9661	4
0.04 0.000	5.00 1 Saliva 0.04 0.000	1 Saliva 0.04 0.000	0.04 0.000	0.04 0.000	0.000		2			-0.116	4096	1 0.02	2 0.000	5832	1	0.02	-0.049	5413	1	0.00	0.000 5	9976 0	0.06	-0.036	5011	3
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M N/A N/A Saliva 0.04 -0.059	N/A N/A Saliva 0.04 -0.059	N/A Saliva 0.04 -0.059	Saliva 0.04 -0.059	0.04 -0.059	-0.059		2		0.08	-0.116	4745	4 0.02	2 -0.055	5147	1	0.03	-0.049	5865	2	0.00	0.000 6	6581 0	0.00	-0.036	7332	0
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0.02 -0.059	0.02 -0.059	0.02 -0.059	0.02 -0.059	0.02 -0.059	-0.059			-	0.03	-0.114	6946	2 0.0	6 -0.044	8995	9	0.03	-0.051	8917	9	0.00		7535 0	0.01	-0.053	8425	-
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810.0- 0.00	0.00 -0.018	0.00 -0.018	0.00 -0.018	0.00 -0.018	-0.018	+	0	+	+	+	00/00	3 0.0.	+	+	0	0.03	0.000	138/6	4	10.0		0238 1	0.03	0.000	13156	4
Brain – Anterior deep 0.00 –0.024	Brain – Anterior deep 0.00 –0.024	Brain – Anterior deep 0.00 -0.024	Brain – Anterior deep 0.00 -0.024	0.00 -0.024	-0.024		0		0:00	-0.098	1215	0.0		1253	0	0.07	-0.053	1440	-	0.00	-0.014	1327 0	0.16	0.000	1217	2
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	Saliva 0.02 0.000	Saliva 0.02 0.000	Saliva 0.02 0.000	0.00 0.000	0000			_		-0.116	5459	3 0.00			0	0.01	-0,049	6324	_	0.03		5523 2	0.00	-0.036	7574	0
0.01 0.000	6.00 2 Brain 0.01 0.000	2 Rrain 0.01 0.000	Rrain 0.01 0.000	0.01 0.000	0.000		$\left \right $	-	$\left \right $	0116	8788	-00			"	0.03	-0.049	8000	~	10.0		1 09/2	100	920.0-	12599	-
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poral 0.0-	6000- 0000	6000- 0000	6000- 0000	6000- 0000	60.0-	+	+	-	+	-0.110	(435	10.0	+	+	4	s0:0	640.0-	0/0/	4	0.0		0 1000	0.0	050.0-	770/	0
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0.07 -0.052 6804	0.70 1 Brain – Hippocampus 0.07 –0.052 6804	1 Brain – Hippocampus 0.07 –0.052 6804	Brain – Hippocampus 0.07 -0.052 6804	0.07 -0.052 6804	-0.052 6804	6804		2		-0.114	7744	1 0.04		6750	9	0.00	-0.049	8790	0	0.01		7340 1	0.01	-0.036	8823	-
0.00 -0.059 5689	0.00 -0.059 5689	0.00 -0.059 5689	0.00 -0.059 5689	0.00 -0.059 5689	-0.059 5689	5689		0		-0.114	6945	1 0.0	0 -0.055		0	0.06	-0.043	7736	5	0.02		317 2	0.03	-0.036	8220	3
-0.059	0.01 -0.059	0.01 -0.059	0.01 -0.059	0.01 -0.059	-0.059		1	H	0.04	-0.114	8321	4 0.0			3	0.03	-0.049	8158	3	0.01	0.000	10484 1	0.01	-0.036	9288	1
0.00 -0.059	0.00 -0.059	0.00 -0.059	0.00 -0.059	0.00 -0.059	-0.059		•	-		-0.114	7339	3 0.06			1	0.04	-0.049	7581		0.00		4918 0	0.02	-0.036	9257	6

Supplementary Table 5: Neuropathological classification for cohort. The

neuropathological diagnosis for the 58 patients reported here, sub-classified by their primary clinical diagnosis, and the number of cases within each category. Abbreviations: FCD, focal cortical dysplasia; HIE, hypoxic-ischemic encephalopathy; MTS, mesial temporal sclerosis; Multifocal CD, multifocal cortical dysplasia; PMG, polymicrogyria

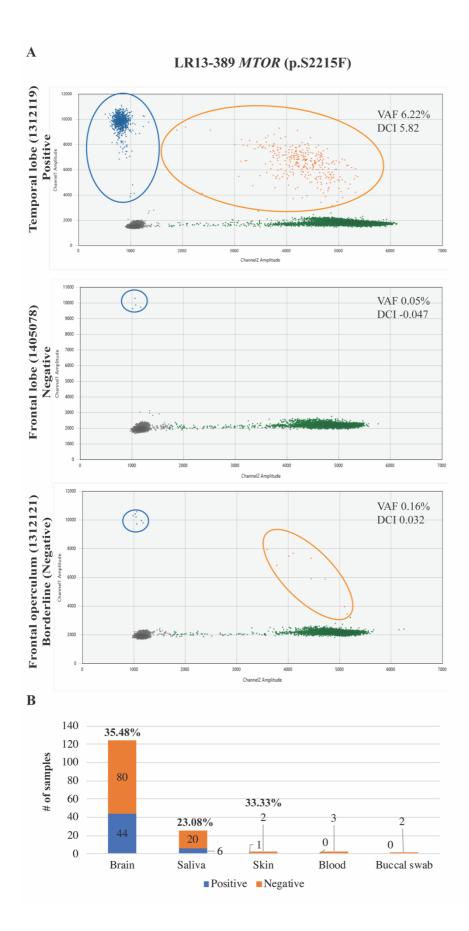
Clinical Diagnosis	Neuropathology	Cases
	FCD	5
	FCD 1	3
	FCD 1a	3
	FCD 1b	3
FCD/HMEG/DMEG	FCD 1c	1
	FCD 2	1
	FCD 2a	19
	FCD 2b	4
	FCD 3d	1
	PMG	2
MCD	Multifocal CD/ FCD 1c	1
	Multifocal CD/ FCD 2b	2
	Gliosis	3
	MTS	1
Other	Meningoangiomatosis	1
	Stroke	1
	HIE	1
	N/A	6
TOTAL		58

Supplementary Table 6: Genotype-neuropathology correlation for patient LR13-129; LR16-313 and LR18-024. Results from ddPCR analysis and neuropathology evaluation for these three representative patients are listed below. For neuropathologic evaluation, hematoxylin and eosin staining (H&E) was performed and cortical layering, as well as presence/absence of balloon/abnormal neurons was evaluated. The H&E results refer to the percentage of abnormal area as evaluated by the pathologist. When possible, pS6 staining was performed and presence of positive neurons (dark or light staining, percentage relative to the area of the section) was performed. For ddPCR results, cells shaded in gray represent brain regions that are positive for the relative hotspot mutation, while cells shaded in gray represent brain regions that do not present the mutation. For LR18-204, FFPE and fresh frozen samples are listed. Two representative brain regions (inferior temporal, positive; and superior temporal, negative) from patient LR18-024 were selected for a comparison of VAF% detected via ddPCR in fresh frozen vs FFPE samples. See Figure 5 for a schematic representation of the two LR18-024 specimens and relative neuropathology findings, and Supplementary Fig. 3 for neuropathology analysis for samples relative to LR13-129 and LR16-313. Blocks highlighted in Fig.5 and Supplementary Fig. 3 are highlighted here in yellow (positive blocks) and gray (negative blocks).

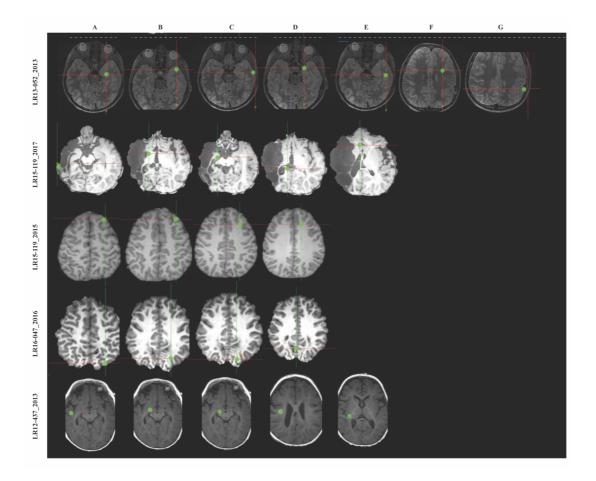
			(pp	ddPCR						Neuropathology			
							H&E				PS6+ pyramidal neurons	Suc	
	Sample description	VAF%	DCI	WT droplets	Mutant droplets	Dumbacha	Non Duarlantia	Not able to		Dark		Light	None
			-			Dysplastic	Non-Dysplastic	evaluate	%	Comment	%	Comment	%
	Left	0.02	-0.05	10946	2	50	50	0	5%	Dysplastic area	15%	Dysplastic area	80%
	Medial Frontal	0.04	-0.04	12079	9	0	0	100	5%		15%		80%
,	Inferior Frontal	0.57	0.39	11908	75	100	0	0	80%		20%		%0
671	Frontal	0.06	-0.03	12784	8	5%	80%	15%	5%	Mostly dysplastic area	%0		%0
-61	Parietal	0.05	-0.05	10156	5	20%	%SL	5%	25%	Diffusely distributed mostly in layer V	20%		55%
ชา	Posterior temporal	0.24	0.10	11220	30	%0	%0	100%	10%		80%		10%
	Occipital	0.03	-0.05	11116	4	20%	60%	20%	15%		30%		55%
	White matter	0.06	-0.04	8708	9	50%	%0	50%	20%	Concentrated in dysplastic area	20%	Concentrated in dysplastic area	9%09
	Superior Frontal	0.08	-0.02	10519	9	20%	30%	50%	15%		15%		70%
	Temporal lobe (A)					20	09	20	15%	Most prevalent in dysplastic foci and layer III	75%	Layers III and V	10%
	Temporal Lobe (B)	1.66	1.23	3206	57	50	40	10	40%	Most prevalent in dysplastic foci and layer III	55%	Layers III and V	5%
	Hippocampus	1.26	0.64	1238	16	25	10	50	%05	Most prevalent in dysplastic foci	45%		5%
EI	Insula (D1)	7.26	6.38	3017	245	70	20	10	50%	Most prevalent in dysplastic foci and layer III	45%		5%
E-9	Insula (D2)	5.93	5.39	6603	453	50	40	10	75%	Most prevalent in dysplastic foci and layer III	20%		5%
IN	Insula (D3)	6.44	5.32	3538	80	50	40	10	75%	Most prevalent in dysplastic foci and layer III	20%		5%
г	Insula (D4)	7.08	6.03	1914	134	50	40	10	%SL	Most prevalent in dysplastic foci and layer III	20%		5%
	Insula (D5)	8.25	6.99	2432	189	75	15	10	75%	Most prevalent in dysplastic foci and layer III	20%		5%
	temporal tip	2.44	1.61	1849	169	10	90	0	10%	Mostly scattered in layer V	25%		65%
	temporal tip	2.13	1.67	1537	39							N/A	
	Inferior temporal, FFPE, IT1	0.070	0.000	5478	4	10%	60%	30%	%0		0%		100%
	Inferior temporal, FFPE, IT2	0.140	0.046	6081	6	%0	50%	50%	3%		3%		95%
	E Inferior temporal, FFPE, IT3	0.030	0.000	8065	3	%0	60%	40%	%0		5%		95%
	E Superior temporal, FFPE, ST1, dysplastic	0.020	-0.260	5811	1								
	Superior temporal, FFPE, ST1, non-dysplastic	0.050	-0.260	7012	4	15	65	20	15	perfect correlation with dysplasia	85		0
1	Superior temporal, FFPE, ST1, full scroll	0.030	-0.260	9617	3								
70	Anterior Dysplasia	0.03	-0.072	9793	3		50%	50%					
-81	Superior Temporal	0.06	-0.050	14216	10	10%	30%	60%					
ชา	E Temporal Tip	0.13	0.000	11545	17	0	70	30					
	2 Inferior Temporal	0.28	0.120	14078	44	0%	75%	25%					
	Anterior Inferior	0.15	0.010	12641	21	10%	60%	30%					
	2 Middle Temporal	0.10	-0.040	9034	10	0	20	80					
	Amygdala	0.12	0.000	15147	21	0	20	80					
	Hippocampus	0.11	-0.050	6252	7	0	90	10					
	Cusa Specimen	0.05	-0.072	4266	2	too poorly o	too poorly oriented to evaluate for dysplasia	or dysplasia					

SUPPLEMENTARY FIGURES

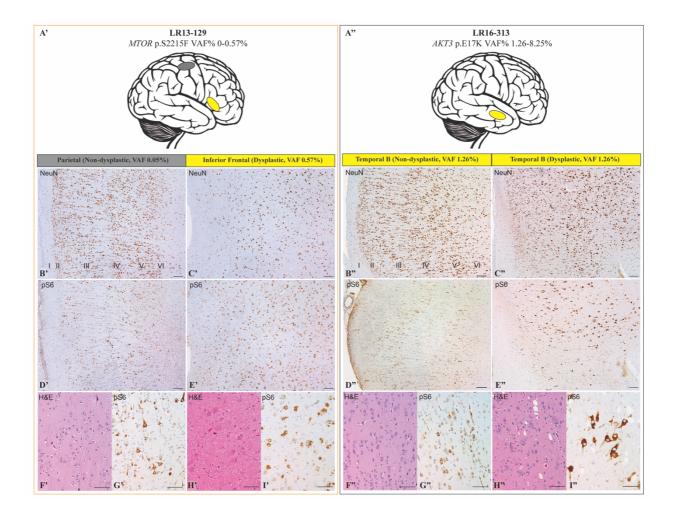
Supplementary Figure 1. ddPCR representative results. (A) Examples of 2D Quantasoft plots for brain tissues from patient LR13-389, carrying the MTOR p.S2215F mutation. Variant allele fractions (VAF%) and Delta confidence intervals (DCI) are shown for each sample. The temporal lobe (sample 1312119) shows a clear positive result, with mutant droplets (in the blue oval) clustering in the upper left quadrant and in higher number than the double positive droplets (in the orange oval), with wild-type droplets in green clustering in the lower right quadrant. A frontal lobe sample from the second surgical procedure (sample 1405078) was negative, with 4 mutant droplets (in the blue oval) detected which were also present in the wildtype control in the same run, as demonstrated by the negative DCI. A sample from the frontal operculum region (1312121) had a "borderline" result. Samples with VAF 0.16% and DCI=0.032 would be considered positive by the Biorad analysis protocol. However, as the plot shows, the number of mutant droplets (in the blue oval, n=7) are fewer than the double positive droplets (in the orange oval, n=9). The high number of double positive droplets leads to a false positive result, thus samples with DCI<0.045 are considered negative in this study. (B) The solve rate per type of tissue based on number of positive specimens vs. total. Hotspot mutations were not detectable in blood and buccal swabs.



Supplementary Figure 2: Brain MRI of four patients with FCD/HMEG/DMEG who tested negative for the 6 PI3K-AKT-MTOR hotspot mutations. Summary MRI images showing biopsy locations. For each image, cross-hairs reflect the sample site, with letters reflecting the multiple samples taken. Green dots represent the exact location of resection. The year in which the surgery was performed is indicated after the underscore, as patient LR15-119 underwent multiple brain surgeries. These samples were tested via ddPCR for the 6 hotspot mutations and were all negative.



Supplementary Figure 3: Histopathological findings and correlations with genotype in tissue samples from individuals LR13-129 and LR16-313. (A') Schematic representation of the resection locations for patient LR13-129. Immunohistochemical and ddPCR findings in the histologically non-dysplastic parietal (B',D',F',G') and dysplastic (FCD type IIB) inferior frontal (C', E', H', I') resection specimens from patient LR13-129. (B') NeuN labeling of neurons in a low magnification field from the parietal resection specimen shows intact sixlayered cortical gray matter as indicated by Roman numerals. The neurons appear cytologically normal (F'), but a significant subset show strong perikaryal pS6 immunoreactivity (D',G'). (C') In contrast, NeuN immunostaining shows marked disorganization of the cortical lamina in the inferior temporal region with balloon cells (H') and diffuse positive pS6 immunoreactivity in nearly the entire neuronal population (I'). Scale bars: B'-E', 200 µm; F'-I', 100 µm. (A''') Schematic representation of the resection location for patient LR16-313. Immunohistochemical and ddPCR findings in the histologically non-dysplastic (B", D", F", G") and dysplastic area (C", E", H", I") from the temporal lobe resection (block B) LR16-313. (B") NeuN labeling of neurons in a low magnification field from a portion of the temporal lobe resection specimen shows intact six-layered cortical gray matter as indicated by Roman numerals. The neurons appear cytologically normal (F"), despite having a detectable mosaic burden in this region (VAF 1.26%). In contrast, NeuN immunostaining shows disorganization of the cortical lamina in a portion of the same region (C") with atypical large neurons (H") and intense positive pS6 immunoreactivity in a subset of the neuronal population (I"). Scale bars: B-E, 200 µm; F-I, 100 µm. Overview of ddPCR results for all samples belonging to these three individuals with matching neuropathology evaluation is presented in Supplementary Table 6, with the negative blocks (red text) and positive blocks (green text) matching the ones represented in this figure.



Supplementary Figure 4: Comparison of number of surgeries and samples in the hotspot mutation positive and negative patients. (A) The number of surgeries per individual did not differ among the two populations. (B) The number of specimens per individual was significantly higher in the mutation-positive cohort (Kolmogorov-Smirnov test, pvalue=0.0068). Notably, our results do not rule out the possibility of a genetic mutation in the negative cases.

