

Figure S1. Hematoxylin and eosin staining of malignant pleural effusion. A total of 6 representative MPE samples are shown compared with a control. Patient 18050707 MPE displayed numerous cancer cells exhibiting striate chromatin, uneven or irregular nucleoli, abundant cytoplasm, uneven staining, and vesicle-like and well-defined malignant cells. Patient 18051504 MPE displayed clusters of abnormal cells with abnormal nuclei, abundant cytoplasm and uneven staining. Patient 18051602 MPE displayed scattered abnormal cells in mass, with uneven cytoplasm staining and a cloudy cytoplasm, and morphologically malignant clear cells. Patient 18052526 MPE displayed abnormal cells of different sizes, clusters of cells, abnormal nuclei, different shades of cytoplasm and cells with a notably malignant appearance. Patient 18052605 MPE displayed abnormal cells differing in size, including clumpy and scattered cells, large abnormal nuclei, rich cytoplasm, and the morphology of malignant cells was clear. Patient 18062543 MPE displayed clusters of abnormal cells differing in size, with heterogeneous nuclei, uneven cytoplasm coloring and a notably malignant appearance.

H&E staining of malignant pleural effusion

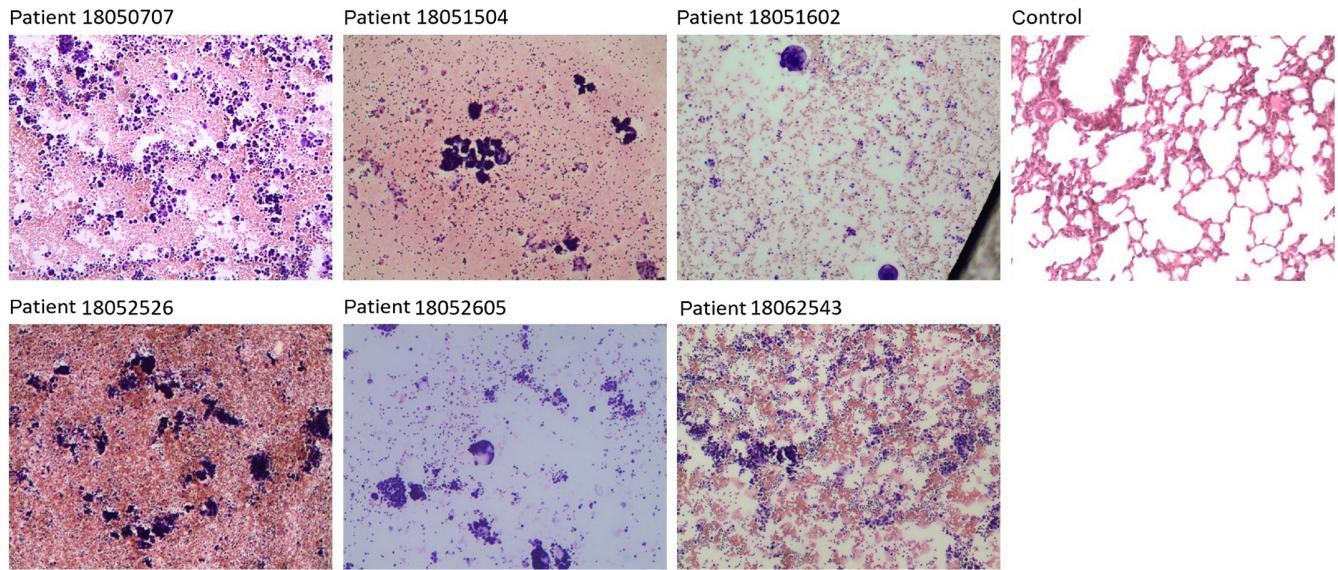


Figure S2. Amplification refractory mutation system PCR result of a gene fusion of ALK + EML4 based on the RNA level from patient 18040535. The Ct values were 9.98 and 16.89.

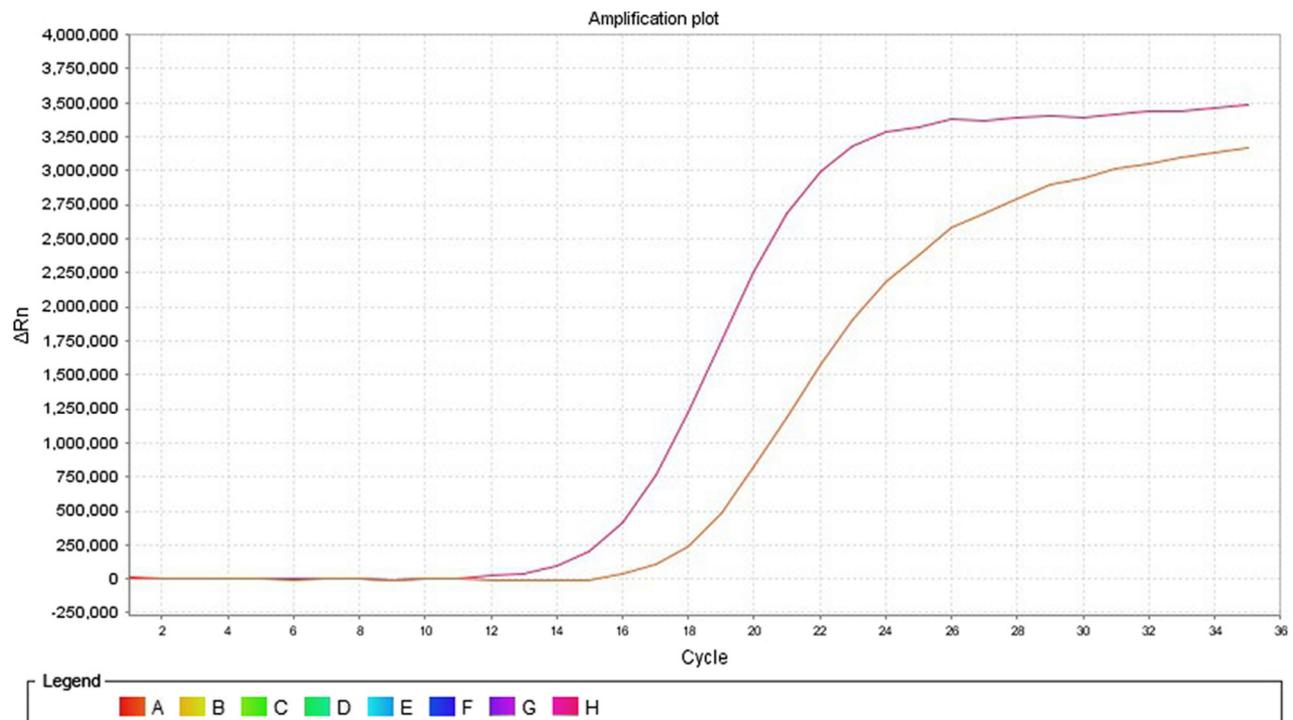


Figure S3. Amplification refractory mutation system PCR result of a BRAF N581I mutation from patient 18042126. The Ct values were 11.86 and 17.36.

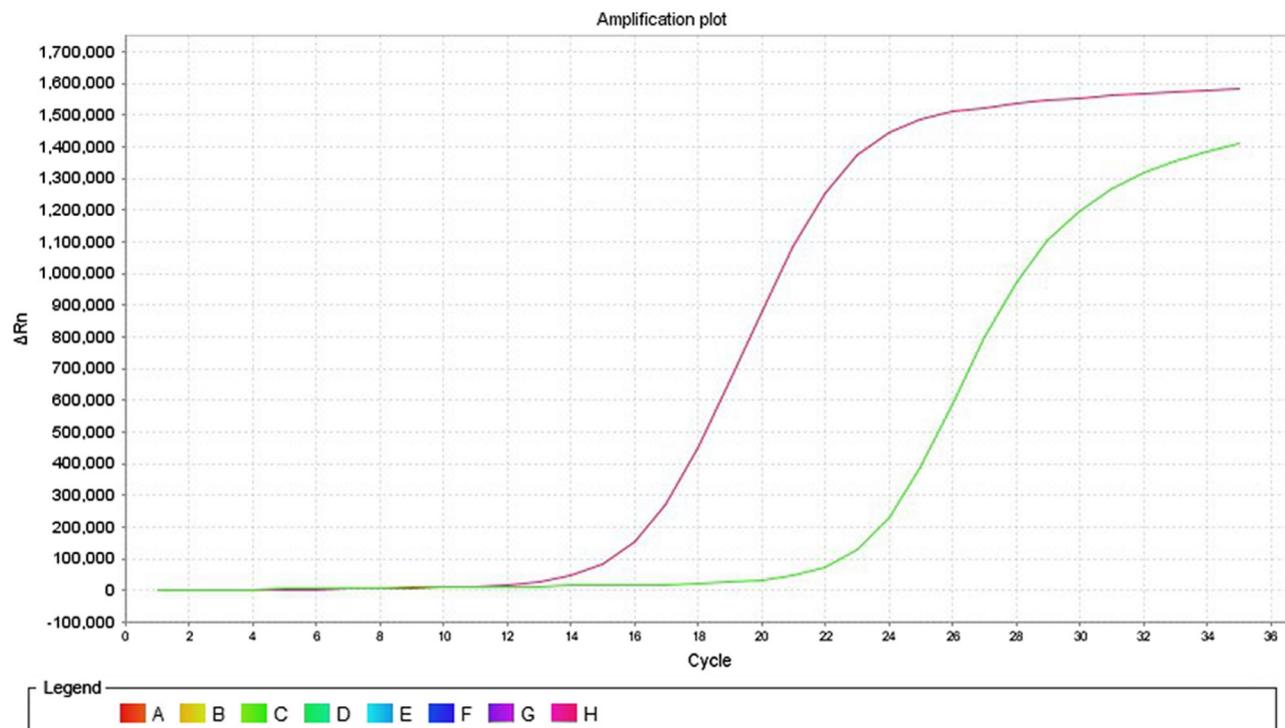


Figure S4. Amplification refractory mutation system PCR result of an EGFR exon 19 deletion mutation from patient 18062314. The Ct values were 13.66 and 16.46.

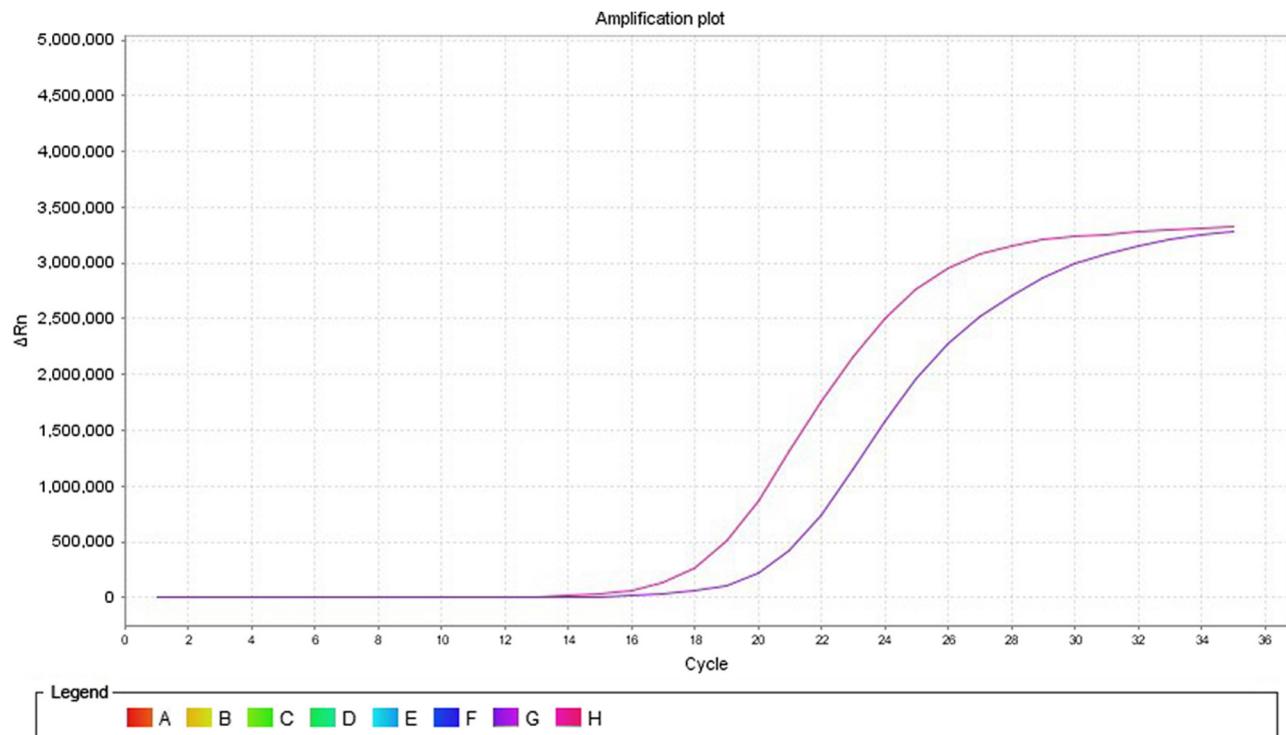


Figure S5. Amplification refractory mutation system PCR result of an EGFR exon 20 insertion mutation from patient 18011709. The Ct values were 11.66 and 20.12.

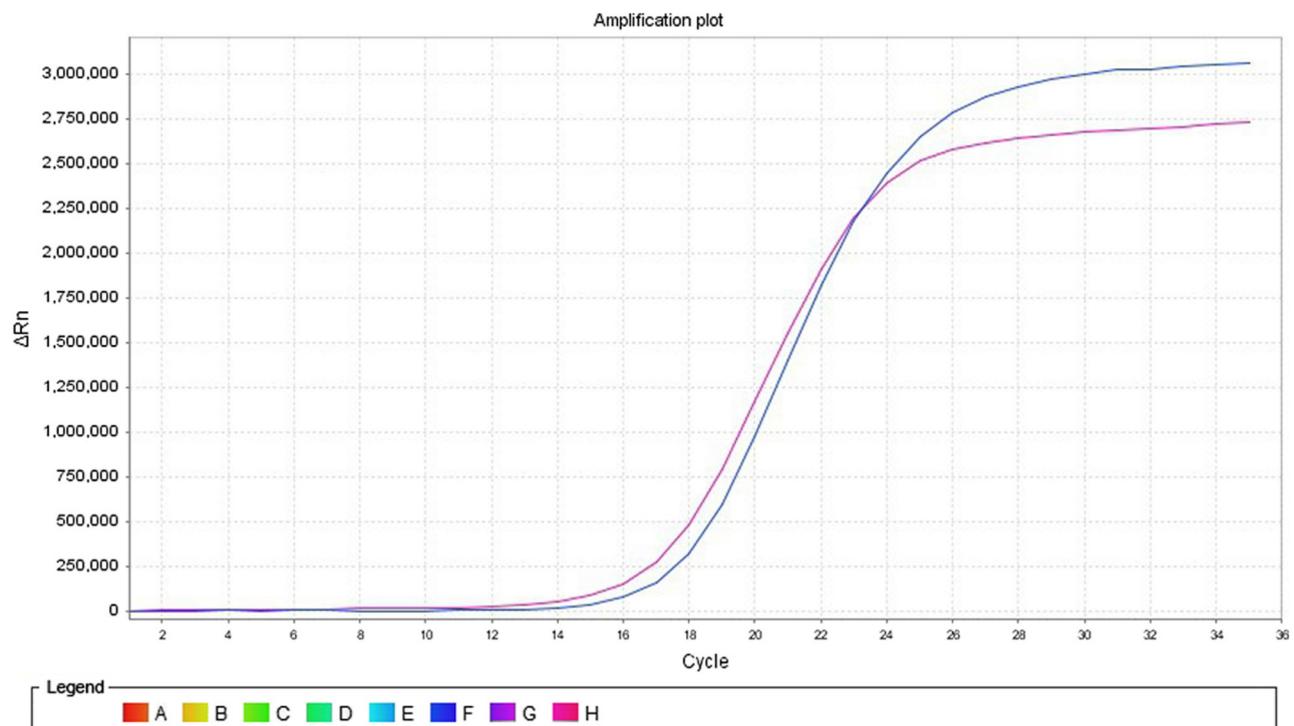


Figure S6. Amplification refractory mutation system PCR result of an EGFR G719A mutation from patient 18012202. The Ct values were 14.71 and 17.02.

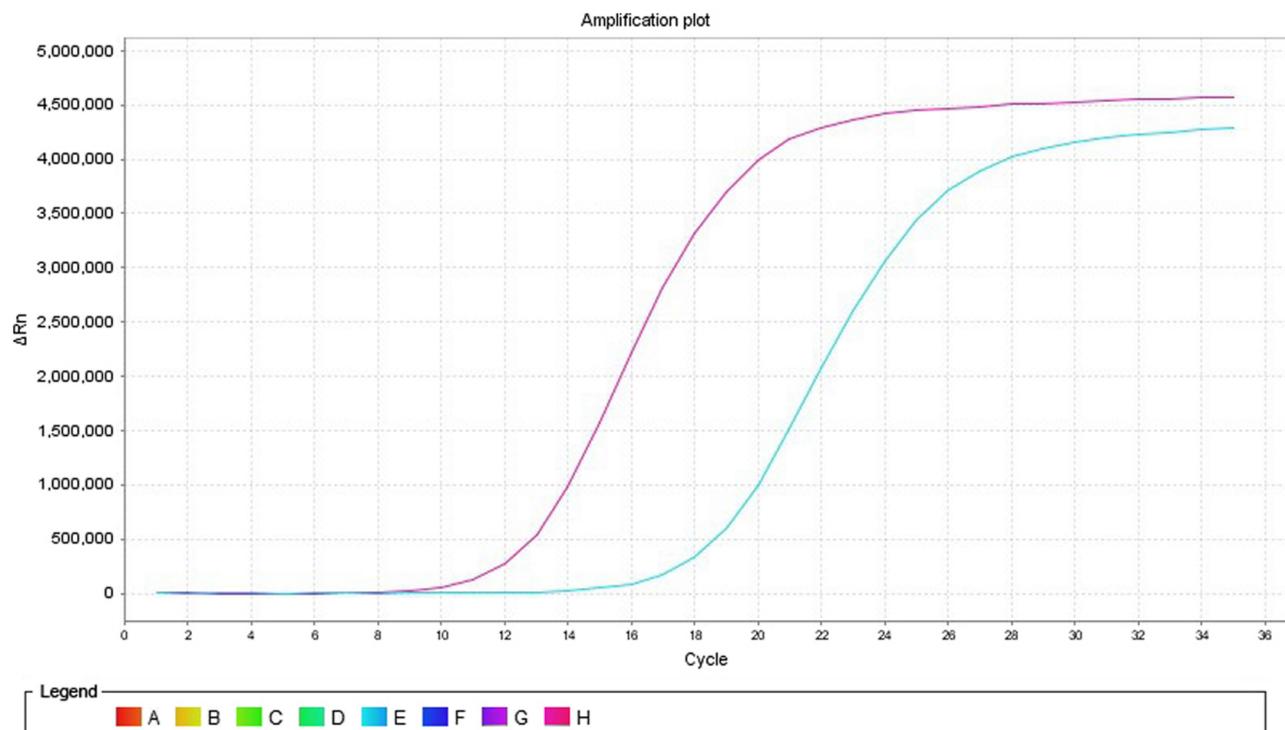


Figure S7. Amplification refractory mutation system PCR result of a KRAS G12S mutation from patient 18060717. The Ct values were 11.85 and 18.26.

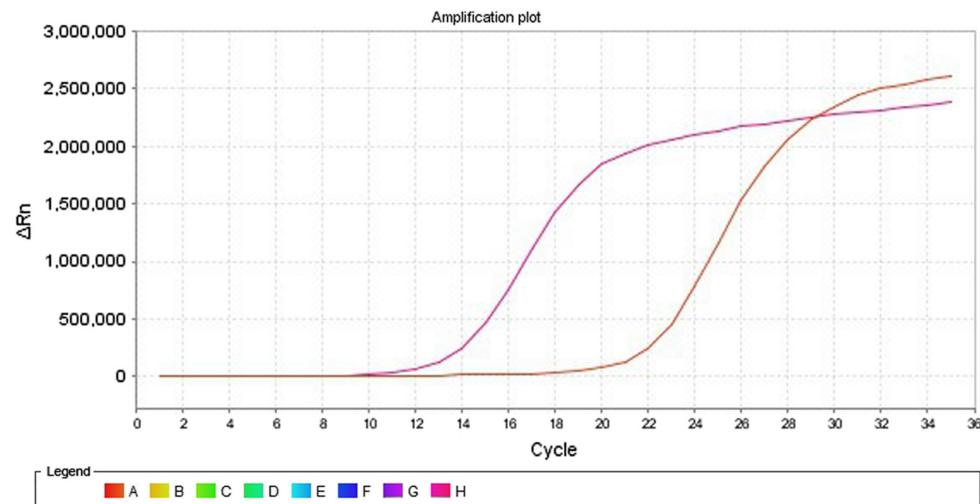


Figure S8. Amplification refractory mutation system PCR result of an EGFR L858R mutation from patient 18050707. The Ct values were 12.96 and 13.56.

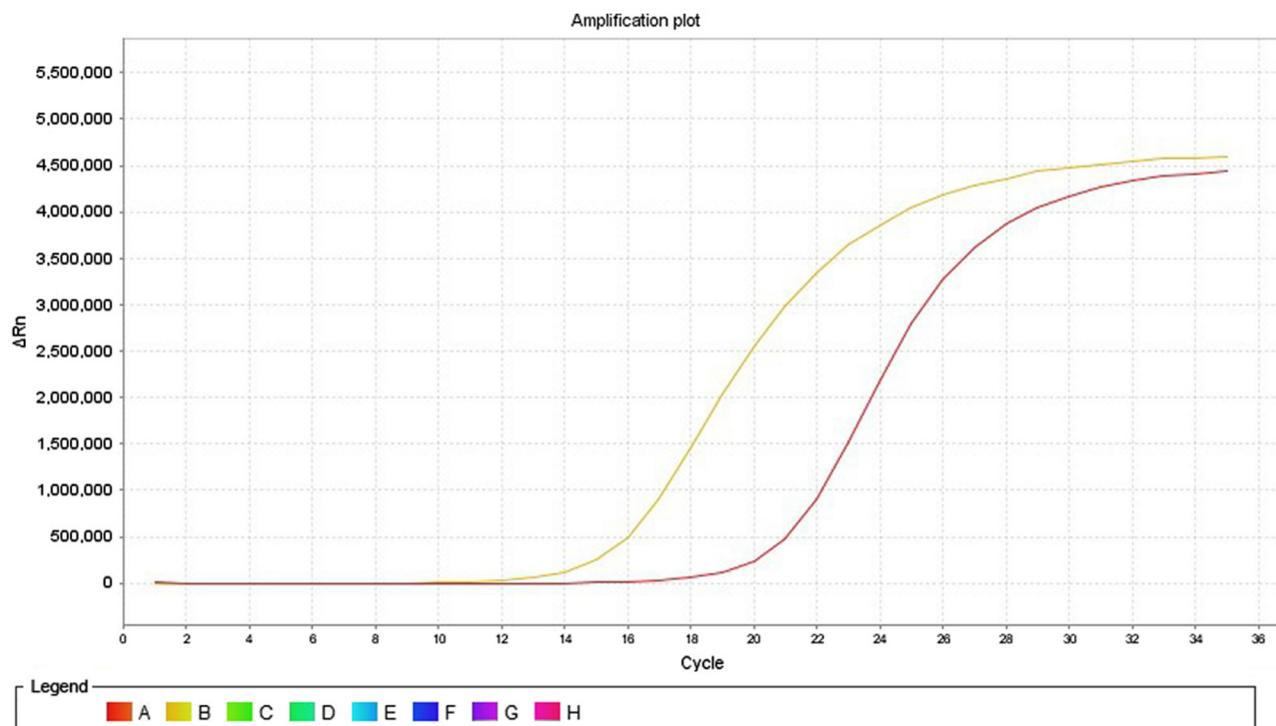


Figure S9. Amplification refractory mutation system PCR result of an EGFR L861Q mutation from patient 18011513. The Ct values were 12.22 and 15.64.

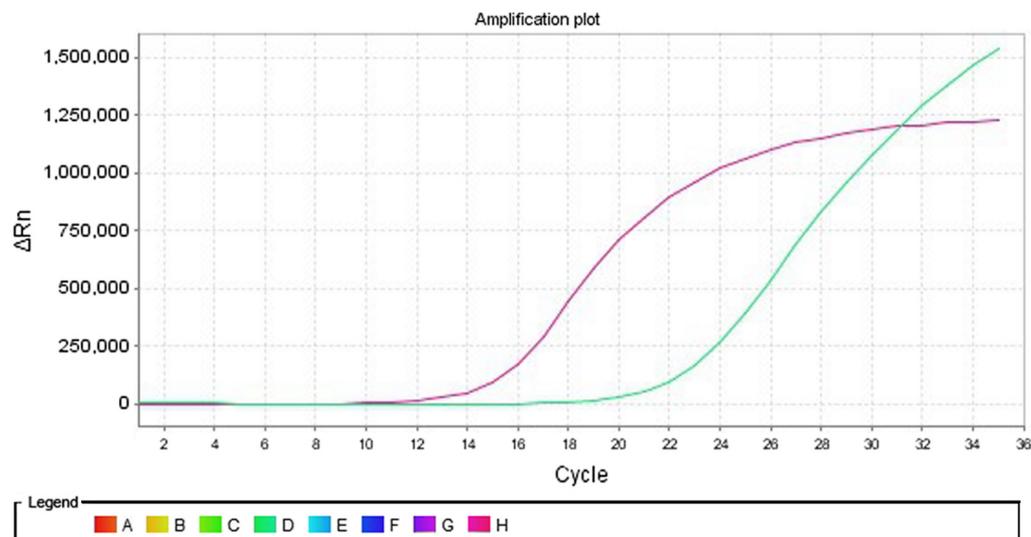


Figure S10. Amplification refractory mutation system PCR result of an EGFR T790M mutation from patient 18051017. The Ct values were 9.36 and 14.01.

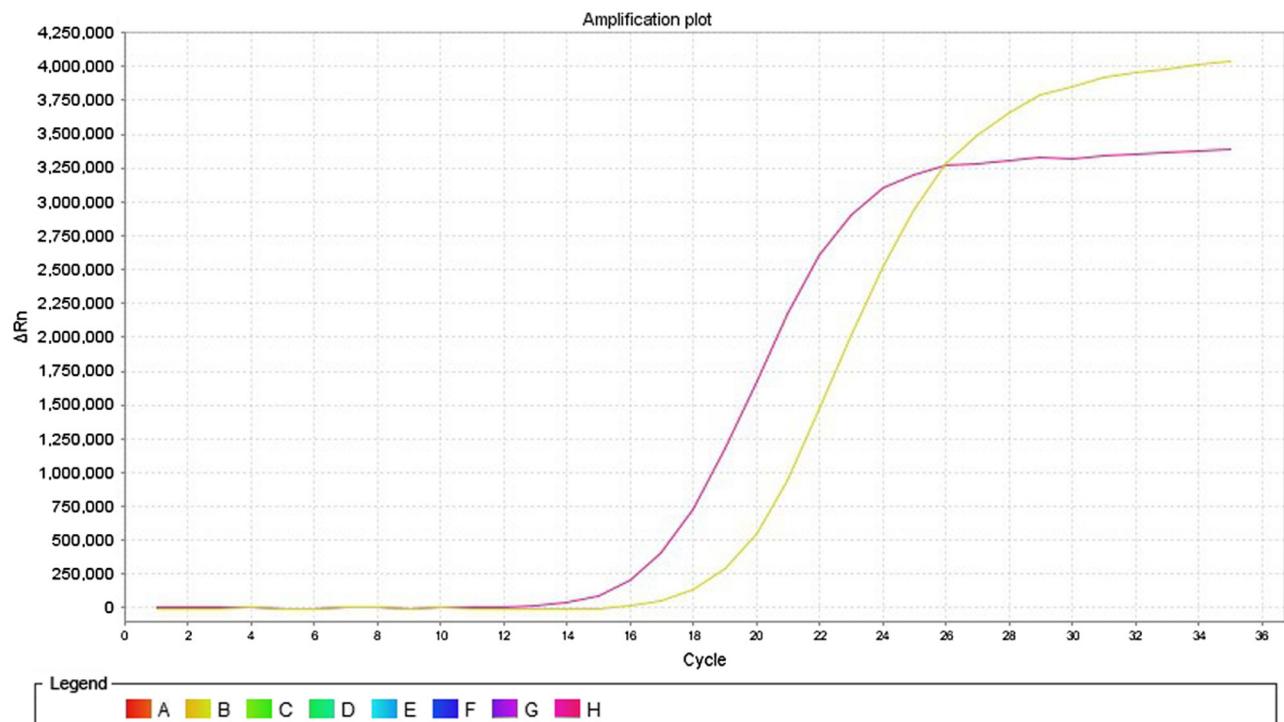


Table SII. NGS results of 108 patients with MPE.

Table SII. Continued.

A	B	Subtype	C	D	E	F1	G	H	I	J	K	L	M1	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	A	B	C1	D	E	F	G	H	I	J	K	L	M	N
1806	F	AC	86										7	6								1	L85		2	3		1	E5		1									
1120																							8R																	
1806	M	AC	66										7	3																										
1121																																								
1806	F	AC	83										15	1	1	1	E74																							
2314																																								
1806	M	AC	62										3	2																									1	
2317																																								
1806	F	Lung cancer	71																																					
2543																																								
1806	M	AC																																						
2601																																								
1807	F	AC	45	1									5	2	1	2																						2		
0609																																								
1807	M	AC	67										4	1	1	2																								
0713																																								
1807	F	LC	64										1	1	1	1	L74																				2			
0901																																								
1807	NA	NA											1	1	1		7_T7																				1			
0904																																								
1807	M	AC	76										1	1	1		E74																				3			
0908																																								
1807	F	AC	30																																				1	
0919																																								
1807	F	AC	55																																					
1002																																								
1807	M	AC	88	1																																			2	
1006																																								

NGS, next generation sequencing; MPE, malignant pleural effusion. Table SII. Results of NGS, ‘lung cancer’ means NSCLC, unconfirmed if adenocarcinoma. ‘adenocarcinoma’ means lung subtype adenocarcinoma. The meanings of capital letters in line 1 are: A, No. of patients; B, gender; C, age; D, ALK fusion; E, BIM (2903bp deletion); F, female; G, ERBB2 amplification; H, FGFR1 amplification; I, MET (exon4 deletion); J, MET amplification; K, RET fusion; L, ROS1 fusion; M, male; M1, ERBB2 other mutation; N, ERBB2 special mutation; O, MET other mutation; P, TP53 mutation; Q, ALK other mutation; R, EGFR exon 18 mutation; S, EGFR exon 19 mutation; T, EGFR exon 19 special mutation; U, EGFR exon 20 special mutation; V, EGFR exon 20 mutation; W, EGFR exon 20 special mutation; X, EGFR exon 21 mutation; Y, EGFR exon 21 special mutation; Z, DDR2 mutation; AA, NRAS mutation; AB, HRAS mutation; AC, adenocarcinoma; AC1, EGFR other mutation; AD, RET other mutation; AE, PTEN mutation; AF, AKT1 mutation; AG, BRAF mutation; AH, BRAF special mutation; AI, PIK3CA mutation; AJ, PIK3CA special mutation; AK, KRAS mutation; AL, KRAS special mutation; AM, ROS1 other mutation; AN, FGFR1 other mutation; LC, lung cancer. The numbers in lists D-AN indicate the number mutations observed at each site. NA, not available.