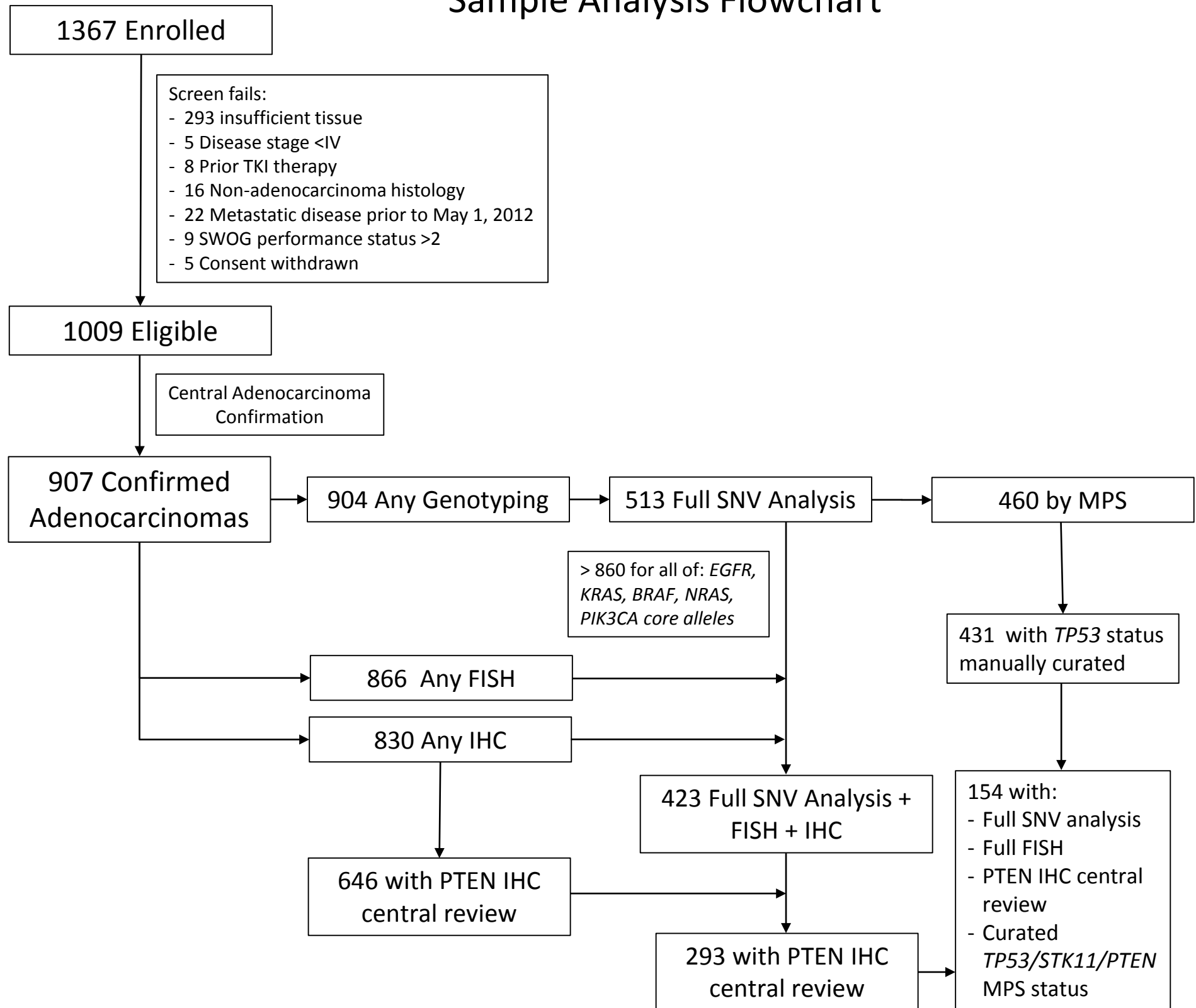


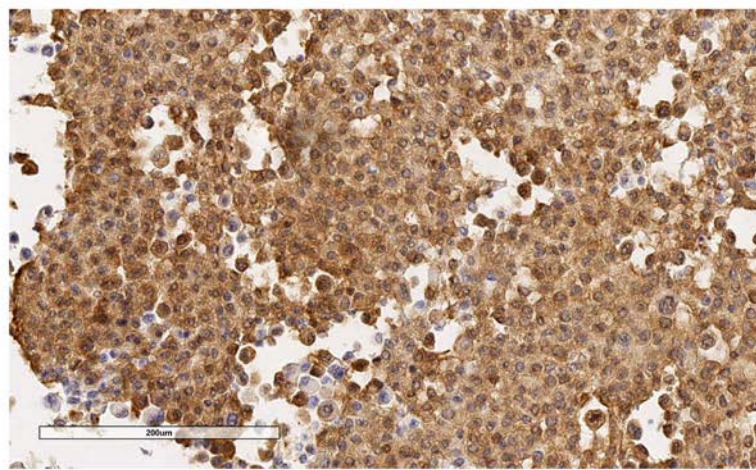
Sample Analysis Flowchart



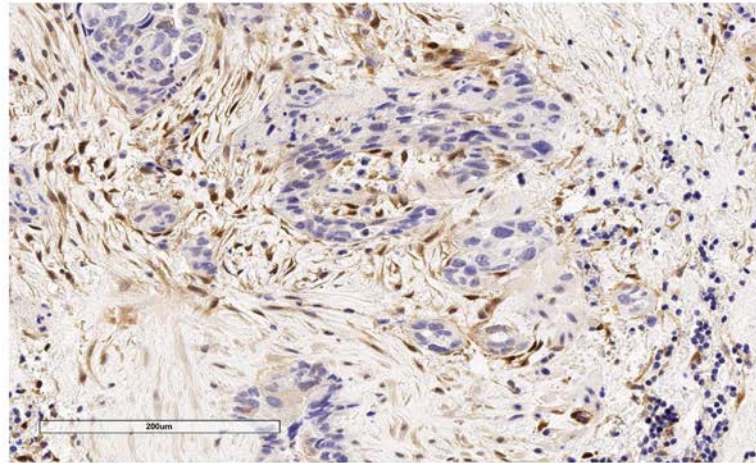
Supplemental Figure 3

PTEN Immunohistochemistry analysis of LCMC II samples.

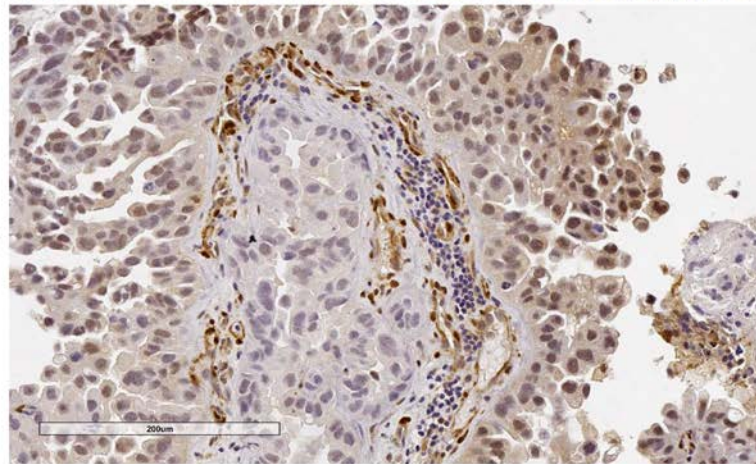
Intact



Lost

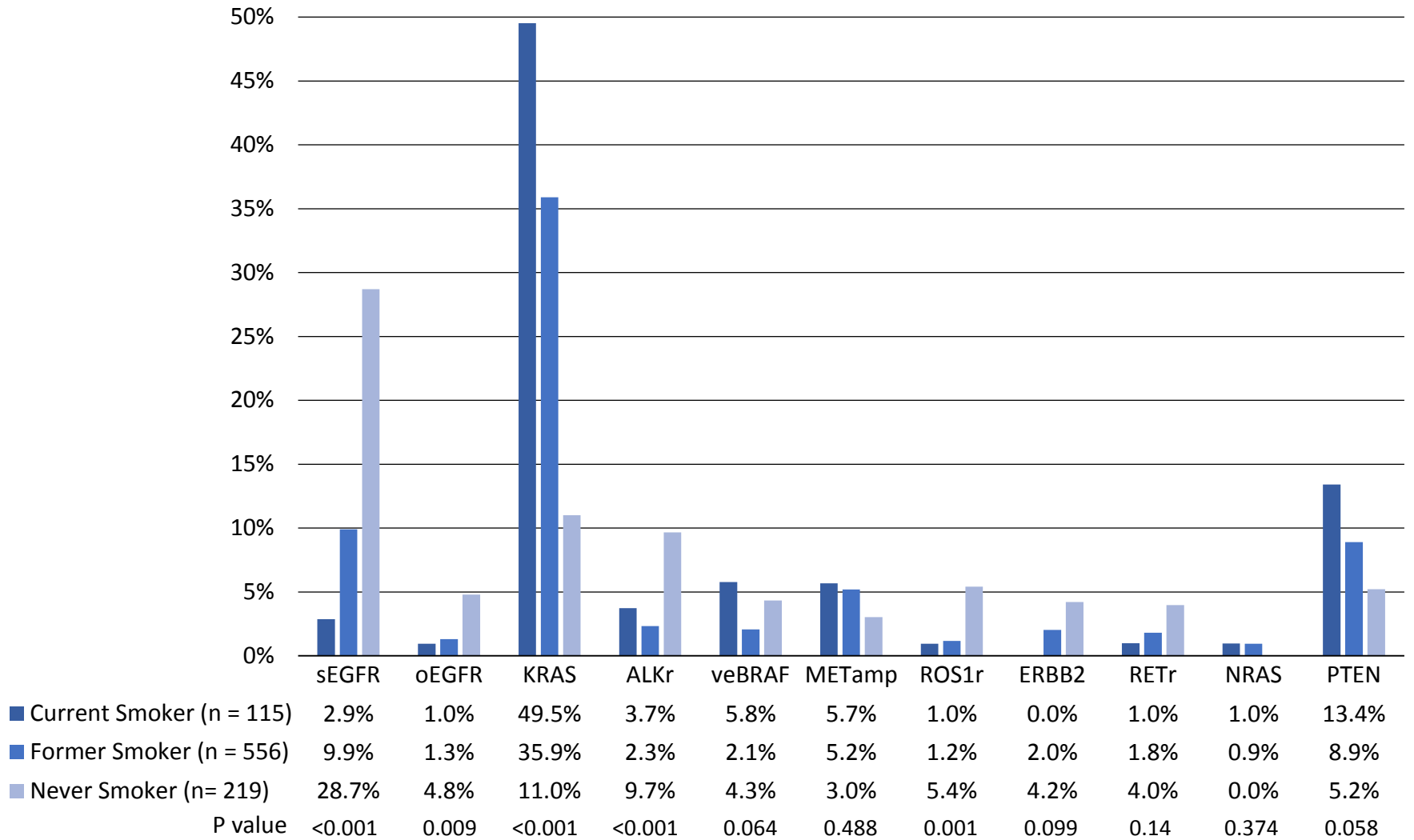


Heterogeneous



Supplemental Figure 4

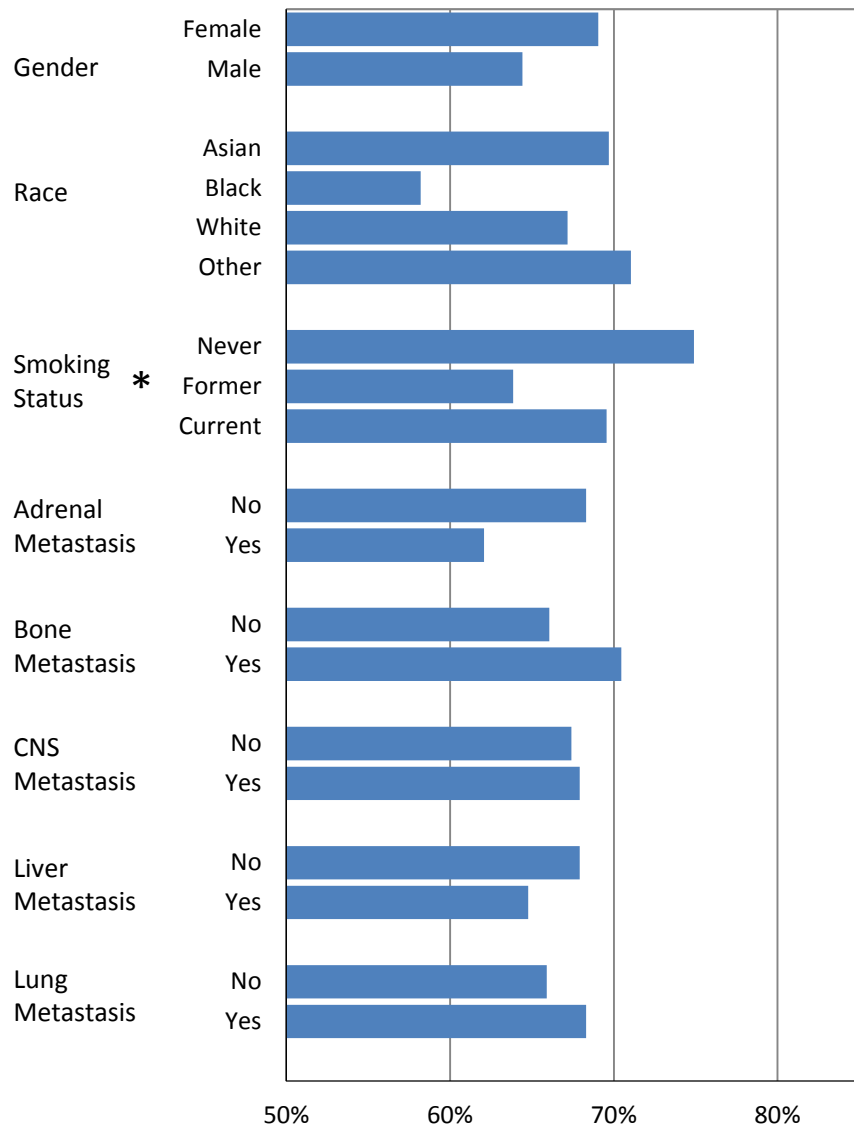
Mutation Frequency by Mutation and Smoking Status



Supplemental Figure 5

Gene: Any* (n=944)**

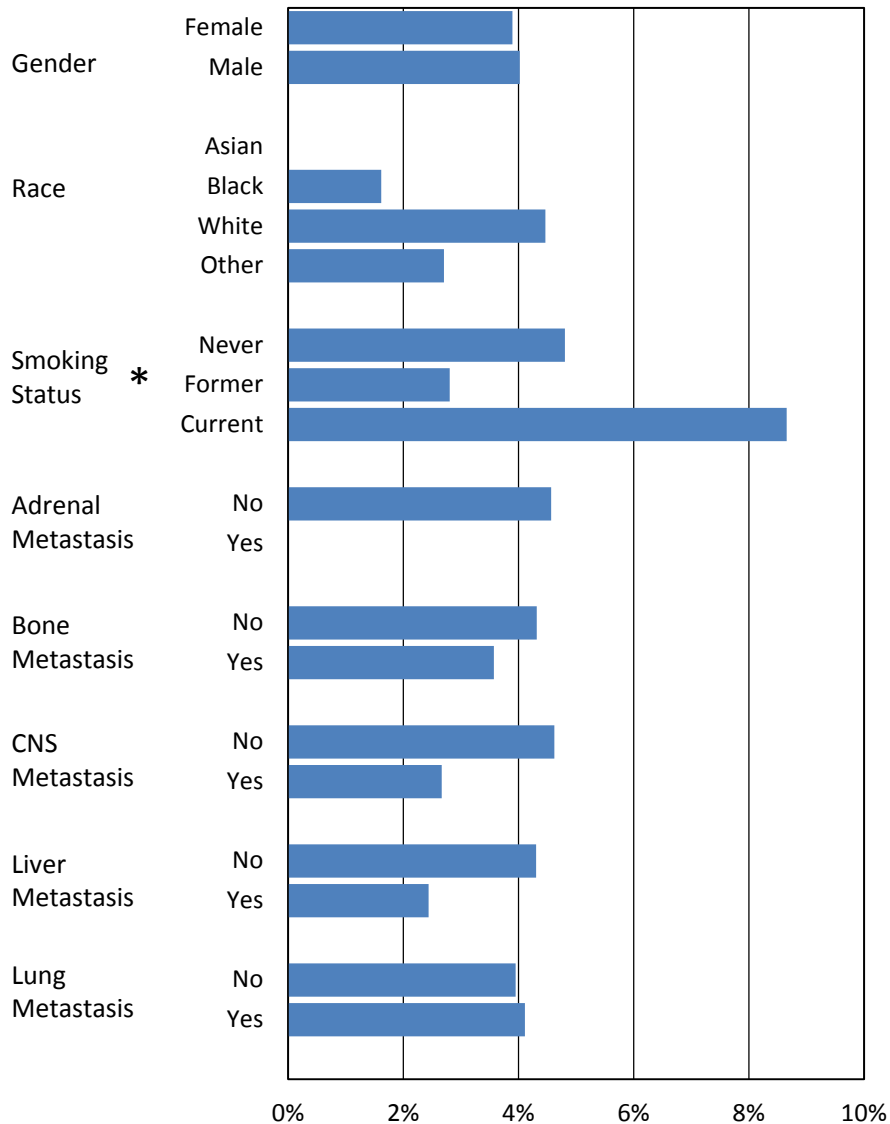
Any Gene, Mutation Positive (%)



Variable	N	WT (n=299)	Mut (n=605)	P
Sex	904			0.14
	Female	51% (151)	56% (337)	
	Male	49% (148)	44% (268)	
Race	904			0.27
	Amer Indian/Alaska Native	0% (0)	0% (3)	
	Asian	3% (10)	4% (23)	
	Black/African American	9% (28)	6% (39)	
	Native Hawaii/Pacific Isl.	0% (1)	0% (0)	
	White	80% (239)	81% (489)	
	Not disclosed	7% (21)	8% (51)	
Smoker	890			0.01
	Current	12% (35)	13% (80)	
	Former	69% (201)	59% (355)	
	Never	19% (55)	27% (164)	
Stage at diagnosis	887			0.06
	I	12% (36)	8% (49)	
	II	9% (26)	6% (34)	
	III	11% (34)	11% (65)	
	IV	68% (200)	75% (443)	
Adrenal metastasis	863			0.24
	No	88% (246)	91% (530)	
	Yes	12% (33)	9% (54)	
Bone metastasis	871			0.20
	No	73% (206)	68% (401)	
	Yes	27% (78)	32% (186)	
Brain metastasis	869			0.89
	No	73% (205)	72% (424)	
	Yes	27% (77)	28% (163)	
Liver metastasis	864			0.55
	No	89% (249)	90% (527)	
	Yes	11% (31)	10% (57)	
Lung metastasis	882			0.45
	No	46% (134)	44% (259)	
	Yes	54% (155)	56% (334)	
Histopathologic subtype	788			0.57
	Acinar predominant	29% (77)	35% (182)	
	Lepidic predominant	2% (5)	2% (11)	
	Micropapillary predom.	0% (0)	0% (1)	
	Mucinous	1% (2)	1% (4)	
	Papillary predominant	1% (2)	0% (2)	
	Solid predominant	68% (180)	62% (322)	
Age at diagnosis	887			0.04
	(Q1, median, Q3)	57.8 65.3 71.0	56.1 63.1 69.9	

Supplemental Figure 5

BRAF, Mutation Positive (%)

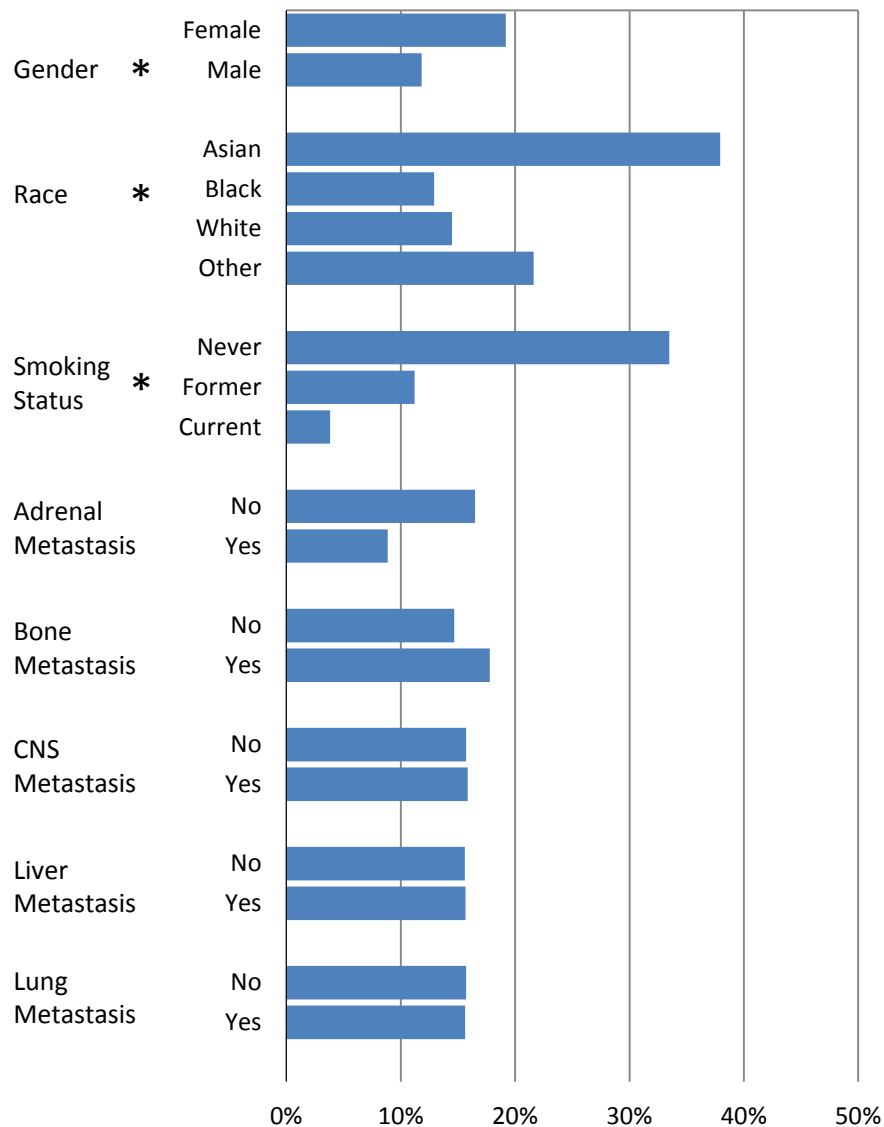


Gene: BRAF (n=860)

Variable	N	WT (n=826)	Mut (n=34)	P
Sex	904			0.93
Female		54% (444)	53% (18)	
Male		46% (382)	47% (16)	
Race	904			0.7
Amer Indian/Alaska Native		0% (3)	0% (0)	
Asian		4% (30)	0% (0)	
Black/African American		7% (61)	3% (1)	
Native Hawaii/Pacific Isl.		0% (1)	0% (0)	
White		80% (663)	91% (31)	
Not disclosed		8% (68)	6% (2)	
Smoker	890			0.02
Current		12% (95)	26% (9)	
Former		64% (520)	44% (15)	
Never		24% (198)	29% (10)	
Stage at diagnosis	887			0.92
I		10% (80)	12% (4)	
II		7% (55)	9% (3)	
III		11% (90)	9% (3)	
IV		72% (585)	71% (24)	
Adrenal metastasis	863			0.05
No		90% (711)	100% (34)	
Yes		10% (79)	0% (0)	
Bone metastasis	871			0.62
No		70% (554)	74% (25)	
Yes		30% (243)	26% (9)	
Brain metastasis	869			0.21
No		73% (578)	82% (28)	
Yes		27% (219)	18% (6)	
Liver metastasis	864			0.42
No		90% (711)	94% (32)	
Yes		10% (80)	6% (2)	
Lung metastasis	882			0.90
No		45% (365)	44% (15)	
Yes		55% (443)	56% (19)	
Histopathologic subtype	788			0.07
Acinar predominant		34% (244)	21% (6)	
Lepidic predominant		2% (16)	0% (0)	
Micropapillary predom.		0% (1)	0% (0)	
Mucinous		1% (5)	4% (1)	
Papillary predominant		0% (3)	4% (1)	
Solid predominant		63% (458)	71% (20)	
Age at diagnosis	887			0.59
(Q1, median, Q3)		56.4 63.8 70.4	58.7 63.5 70.1	

Supplemental Figure 5

EGFR, Mutation Positive (%)

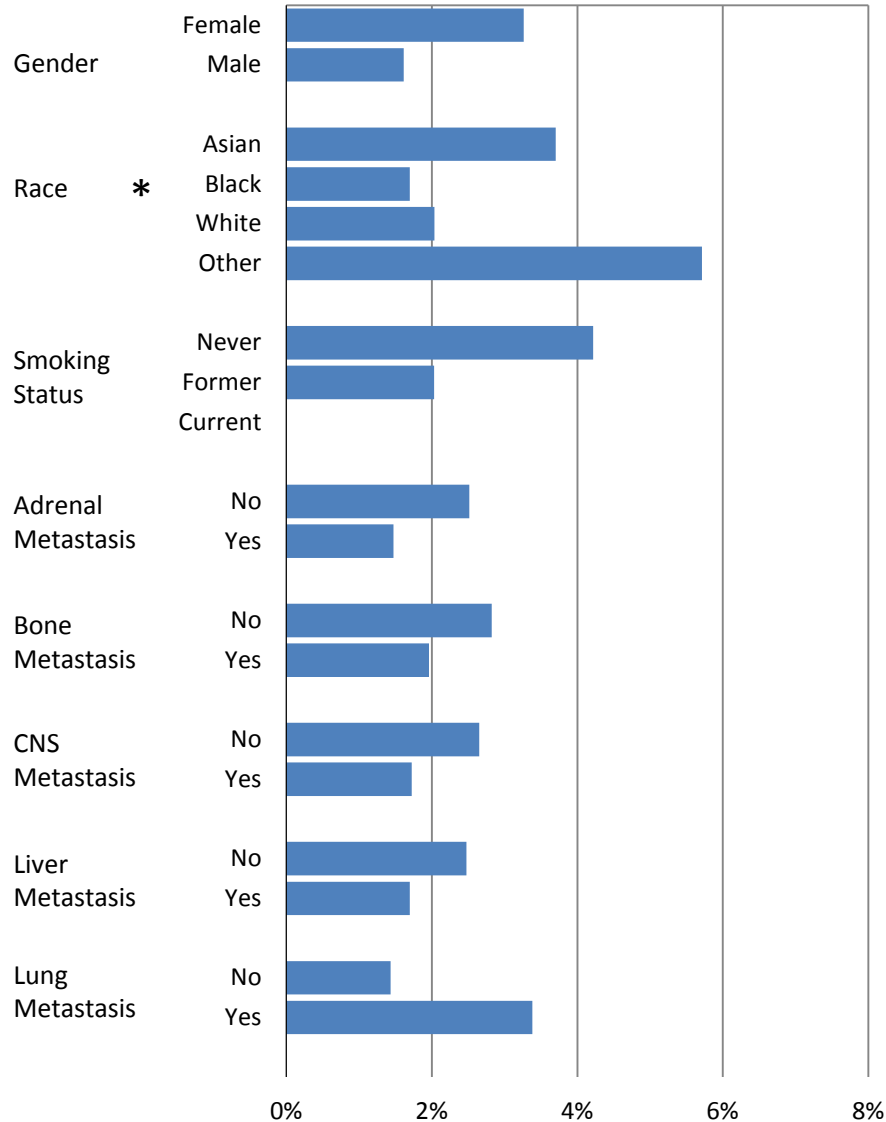


Gene: EGFR (n=862)

Variable	N	WT (n=726)	Mut (n=136)	P
Sex	904			0.003
	Female	52% (375)	65% (89)	
	Male	48% (351)	35% (47)	
Race	904			0.009
	Amer Indian/Alaska Native	0% (3)	0% (0)	
	Asian	2% (18)	8% (11)	
	Black/African American	7% (54)	6% (8)	
	Native Hawaii/Pacific Isl.	0% (1)	0% (0)	
	White	82% (596)	74% (101)	
	Not disclosed	7% (54)	12% (16)	
Smoker	890			<0.001
	Current	14% (101)	3% (4)	
	Former	66% (475)	45% (60)	
	Never	19% (139)	52% (70)	
Stage at diagnosis	887			0.002
	I	11% (79)	4% (5)	
	II	8% (54)	2% (3)	
	III	11% (82)	8% (11)	
	IV	70% (499)	85% (112)	
Adrenal metastasis	863			0.08
	No	90% (623)	95% (123)	
	Yes	10% (72)	5% (7)	
Bone metastasis	871			0.26
	No	70% (494)	65% (85)	
	Yes	30% (208)	35% (45)	
Brain metastasis	869			1
	No	73% (510)	73% (95)	
	Yes	27% (191)	27% (36)	
Liver metastasis	864			1
	No	90% (627)	90% (116)	
	Yes	10% (70)	10% (13)	
Lung metastasis	882			1
	No	45% (322)	45% (60)	
	Yes	55% (389)	55% (72)	
Histopathologic subtype	788			0.48
	Acinar predominant	32% (203)	41% (47)	
	Lepidic predominant	2% (14)	2% (2)	
	Micropapillary predom.	0% (1)	0% (0)	
	Mucinous	1% (5)	1% (1)	
	Papillary predominant	1% (4)	0% (0)	
	Solid predominant	65% (415)	57% (65)	
Age at diagnosis	887			0.17
	(Q1, median, Q3)	56.7 63.8 70.7	55.3 63.0 68.8	

Supplemental Figure 5

ERBB2, Mutation Positive(%)

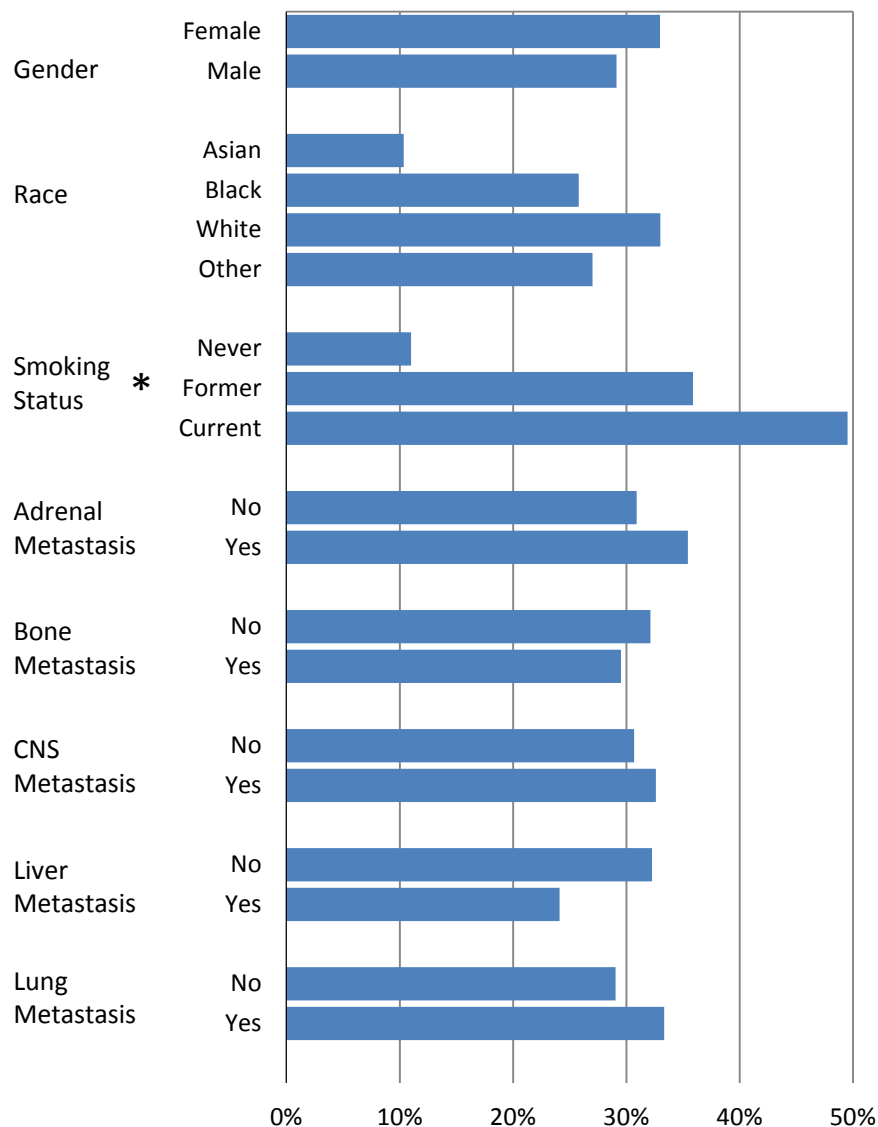


Gene: ERBB2 (n=647)

Variable	N	WT (n=631)	Mut (n=16)	P
Sex	904			0.18
Female		52% (326)	69% (11)	
Male		48% (305)	31% (5)	
Race	904			0.02
Amer Indian/Alaska Native		0% (2)	6% (1)	
Asian		4% (26)	6% (1)	
Black/African American		9% (58)	6% (1)	
Native Hawaii/Pacific Isl.		0% (1)	0% (0)	
White		76% (481)	62% (10)	
Not disclosed		10% (63)	19% (3)	
Smoker	890			0.1
Current		13% (80)	0% (0)	
Former		62% (386)	53% (8)	
Never		25% (159)	47% (7)	
Stage at diagnosis	887			0.26
I		8% (51)	19% (3)	
II		5% (32)	6% (1)	
III		12% (76)	0% (0)	
IV		74% (459)	75% (12)	
Adrenal metastasis	863			0.6
No		89% (543)	93% (14)	
Yes		11% (67)	7% (1)	
Bone metastasis	871			0.52
No		67% (413)	75% (12)	
Yes		33% (200)	25% (4)	
Brain metastasis	869			0.5
No		72% (441)	80% (12)	
Yes		28% (171)	20% (3)	
Liver metastasis	864			0.71
No		90% (552)	93% (14)	
Yes		10% (58)	7% (1)	
Lung metastasis	882			0.12
No		44% (275)	25% (4)	
Yes		56% (343)	75% (12)	
Histopathologic subtype	788			0.82
Acinar predominant		31% (169)	50% (6)	
Lepidic predominant		2% (11)	0% (0)	
Micropapillary predom.		0% (1)	0% (0)	
Mucinous		1% (6)	0% (0)	
Papillary predominant		0% (2)	0% (0)	
Solid predominant		65% (357)	50% (6)	
Age at diagnosis	887			0.43
(Q1, median, Q3)		56.6 63.0 69.5	54.7 68.0 71.9	

Supplemental Figure 5

KRAS, Mutation Positive (%)

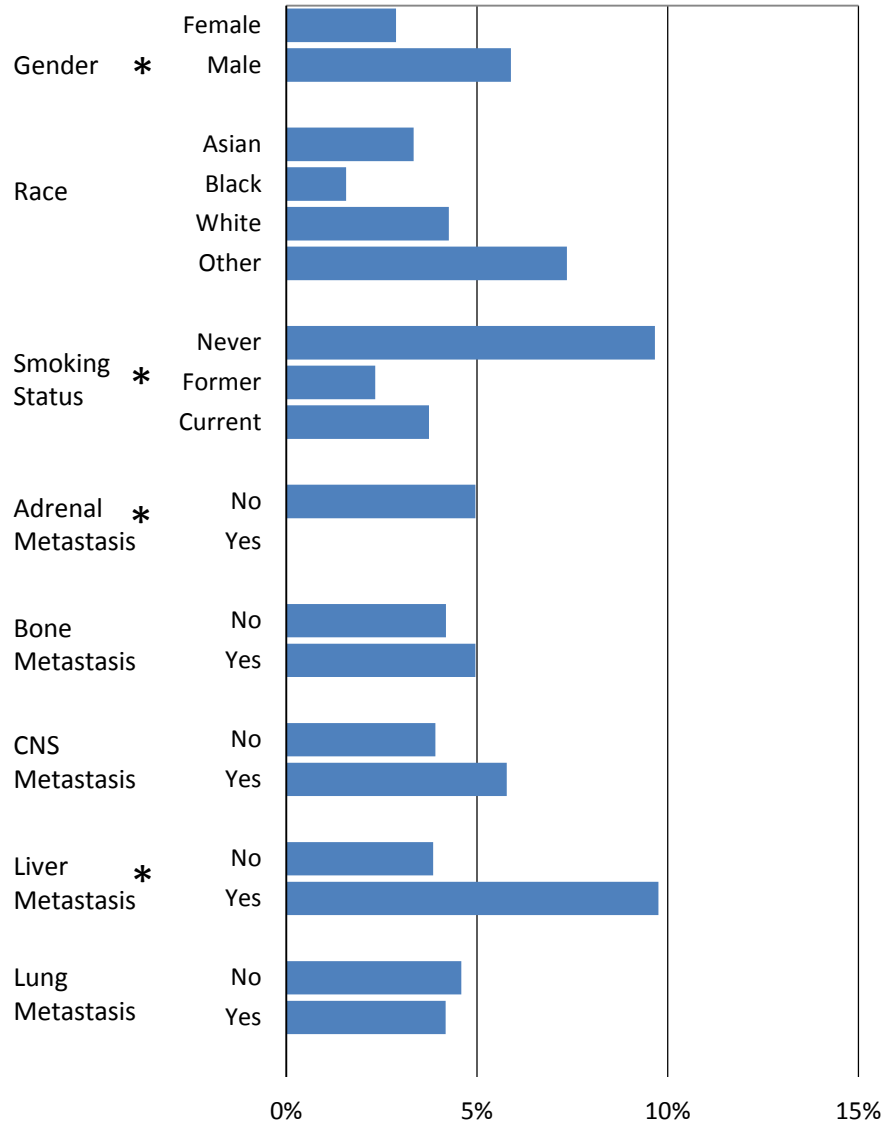


Gene: KRAS (n=862)

Variable	N	WT (n=593)	Mut (n=269)	P
Sex	904			0.23
Female		52% (311)	57% (153)	
Male		48% (282)	43% (116)	
Race	904			0.05
Amer Indian/Alaska Native		0% (1)	1% (2)	
Asian		4% (26)	1% (3)	
Black/African American		8% (46)	6% (16)	
Native Hawaii/Pacific Isl.		0% (1)	0% (0)	
White		79% (467)	86% (230)	
Not disclosed		9% (52)	7% (18)	
Smoker	890			<0.001
Current		9% (53)	19% (52)	
Former		59% (343)	72% (192)	
Never		32% (186)	9% (23)	
Stage at diagnosis	887			0.77
I		10% (56)	11% (28)	
II		6% (37)	8% (20)	
III		11% (62)	12% (31)	
IV		73% (428)	70% (184)	
Adrenal metastasis	863			0.41
No		91% (516)	89% (231)	
Yes		9% (51)	11% (28)	
Bone metastasis	871			0.46
No		69% (393)	71% (186)	
Yes		31% (179)	29% (75)	
Brain metastasis	869			0.6
No		73% (420)	72% (186)	
Yes		27% (153)	28% (74)	
Liver metastasis	864			0.13
No		89% (504)	92% (240)	
Yes		11% (63)	8% (20)	
Lung metastasis	882			0.18
No		47% (271)	42% (111)	
Yes		53% (308)	58% (154)	
Histopathologic subtype	788			0.06
Acinar predominant		31% (160)	37% (90)	
Lepidic predominant		1% (7)	4% (9)	
Micropapillary predom.		0% (1)	0% (0)	
Mucinous		1% (5)	0% (1)	
Papillary predominant		1% (3)	0% (1)	
Solid predominant		66% (338)	58% (142)	
Age at diagnosis	887			0.18
(Q1, median, Q3)		55.9 63.6 70.1	57.4 63.9 71.0	

Supplemental Figure 5

ALKr Positive, (%)

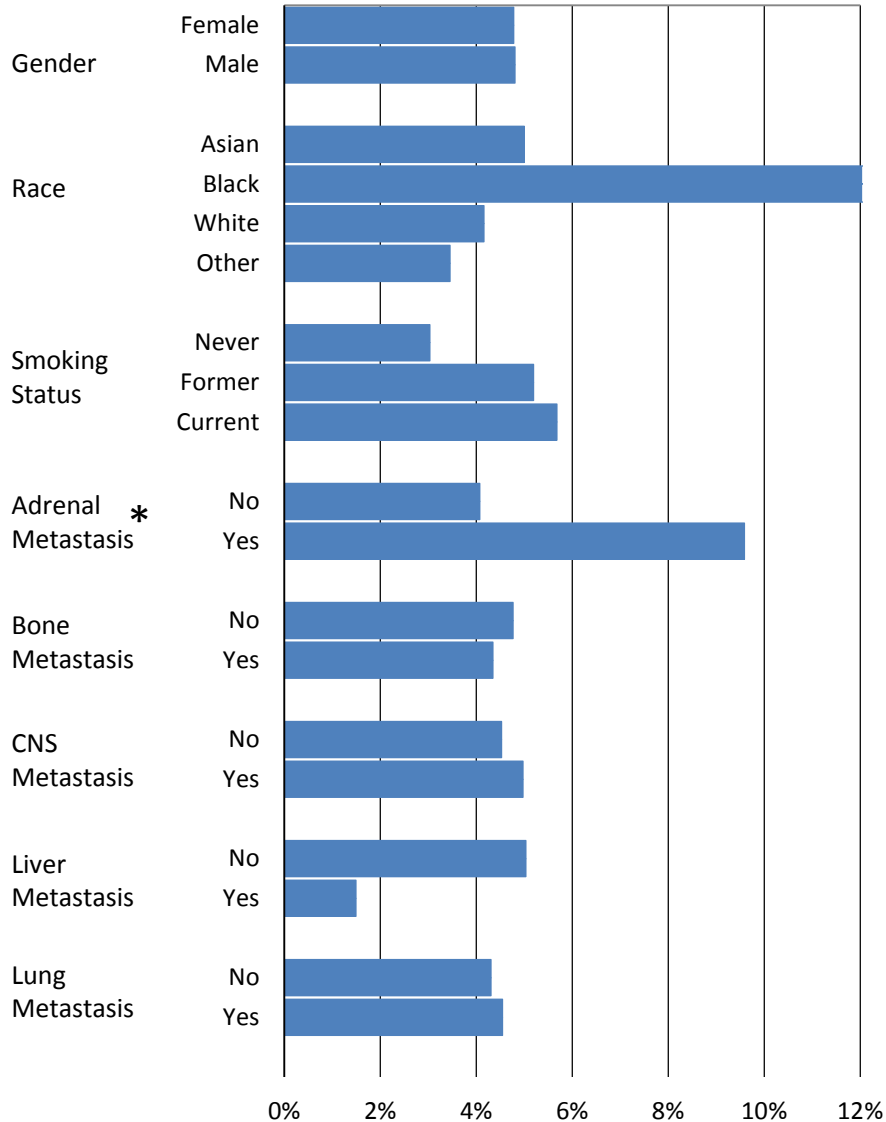


Gene: ALK (rearrangement) (n=843)

Variable	N	WT (n=807)	Mut (n=36)	P
Sex	904			0.03
		Female	54% (439)	36% (13)
		Male	46% (368)	64% (23)
Race	904			0.67
		Amer Indian/Alaska Native	0% (2)	0% (0)
		Asian	4% (29)	3% (1)
		Black/African American	8% (63)	3% (1)
		Native Hawaii/Pacific Isl.	0% (1)	0% (0)
		White	7% (60)	14% (5)
		Not disclosed	81% (652)	81% (29)
Smoker	890			<0.001
		Current	13% (103)	11% (4)
		Former	63% (504)	33% (12)
		Never	24% (187)	56% (20)
Stage at diagnosis	887			0.02
		I	10% (79)	0% (0)
		II	7% (58)	0% (0)
		III	11% (88)	23% (8)
		IV	72% (570)	77% (27)
Adrenal metastasis	863			0.04
		No	89% (690)	100% (36)
		Yes	11% (83)	0% (0)
Bone metastasis	871			0.62
		No	71% (550)	67% (24)
		Yes	29% (230)	33% (12)
Brain metastasis	869			0.25
		No	73% (566)	64% (23)
		Yes	27% (212)	36% (13)
Liver metastasis	864			0.01
		No	90% (700)	78% (28)
		Yes	10% (74)	22% (8)
Lung metastasis	882			0.78
		No	45% (354)	47% (17)
		Yes	55% (436)	53% (19)
Histopathologic subtype	788			0.91
		Acinar predominant	33% (231)	29% (8)
		Lepidic predominant	2% (16)	0% (0)
		Micropapillary predom.	0% (1)	0% (0)
		Mucinous	1% (6)	0% (0)
		Papillary predominant	1% (4)	0% (0)
		Solid predominant	63% (447)	71% (20)
Age at diagnosis	887			<0.001
		(Q1, median, Q3)	56.9 63.9 70.5	45.8 55.6 61.7

Supplemental Figure 5

METamp, positive (%)

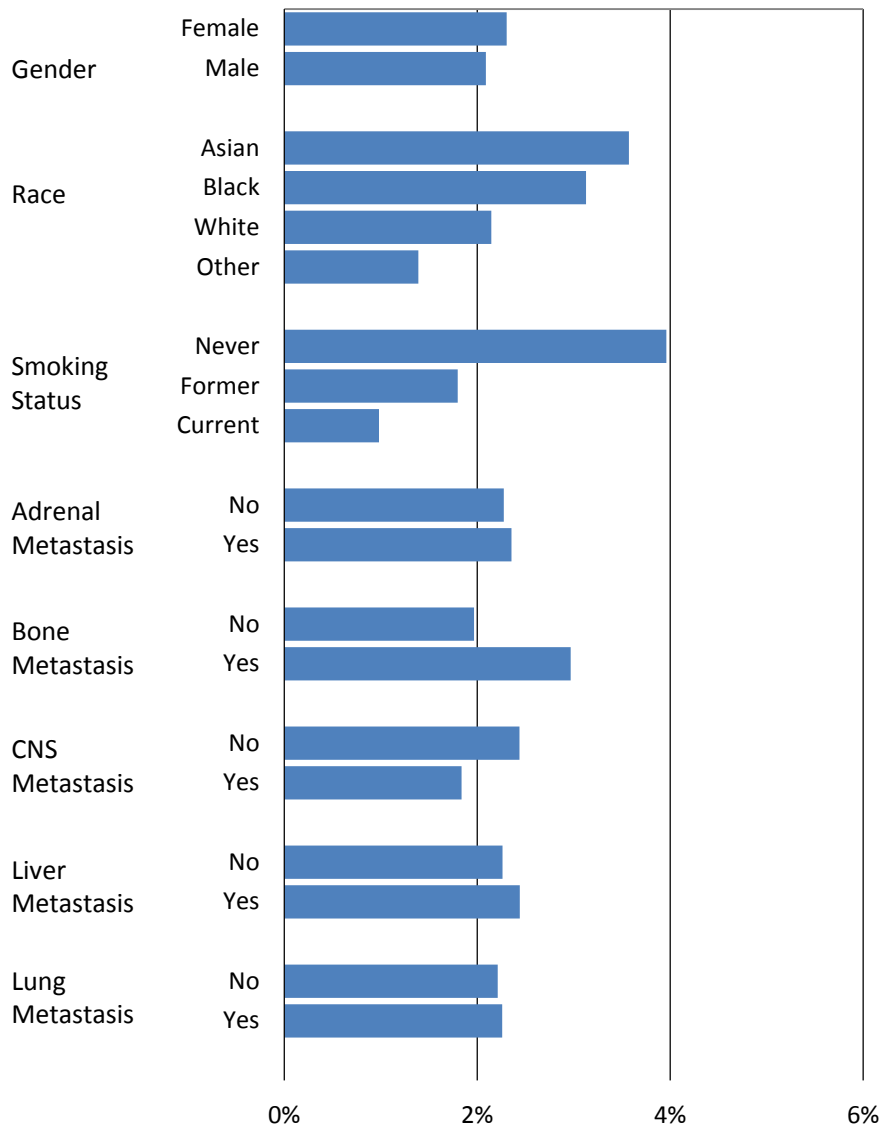


Gene: MET (amplification) (n=689)

Variable	N	WT (n=656)	Mut (n=33)	P
Sex	904			1
		Female	52% (339)	52% (17)
		Male	48% (317)	48% (16)
Race	904			0.11
		Amer Indian/Alaska Native	0% (2)	0% (0)
		Asian	3% (19)	3% (1)
		Black/African American	8% (51)	21% (7)
		Native Hawaii/Pacific Isl.	0% (0)	0% (0)
		White	81% (530)	70% (23)
		Not disclosed	8% (54)	6% (2)
Smoker	890			0.49
		Current	13% (83)	16% (5)
		Former	62% (402)	69% (22)
		Never	25% (160)	16% (5)
Stage at diagnosis	887			0.81
		I	8% (49)	9% (3)
		II	7% (45)	6% (2)
		III	11% (72)	6% (2)
		IV	74% (479)	79% (26)
Adrenal metastasis	863			0.04
		No	90% (565)	77% (24)
		Yes	10% (66)	23% (7)
Bone metastasis	871			0.81
		No	69% (440)	71% (22)
		Yes	31% (198)	29% (9)
Brain metastasis	869			0.81
		No	73% (464)	71% (22)
		Yes	27% (172)	29% (9)
Liver metastasis	864			0.19
		No	90% (566)	97% (30)
		Yes	10% (66)	3% (1)
Lung metastasis	882			0.88
		No	45% (289)	43% (13)
		Yes	55% (357)	57% (17)
Histopathologic subtype	788			0.55
		Acinar predominant	33% (185)	17% (4)
		Lepidic predominant	2% (9)	0% (0)
		Micropapillary predom.	0% (1)	0% (0)
		Mucinous	1% (4)	0% (0)
		Papillary predominant	1% (4)	0% (0)
		Solid predominant	64% (359)	83% (20)
Age at diagnosis	887			0.46
		(Q1, median, Q3)	56.8 63.6 70.2	55.7 62.2 69.9

Supplemental Figure 5

RETr, Positive (%)

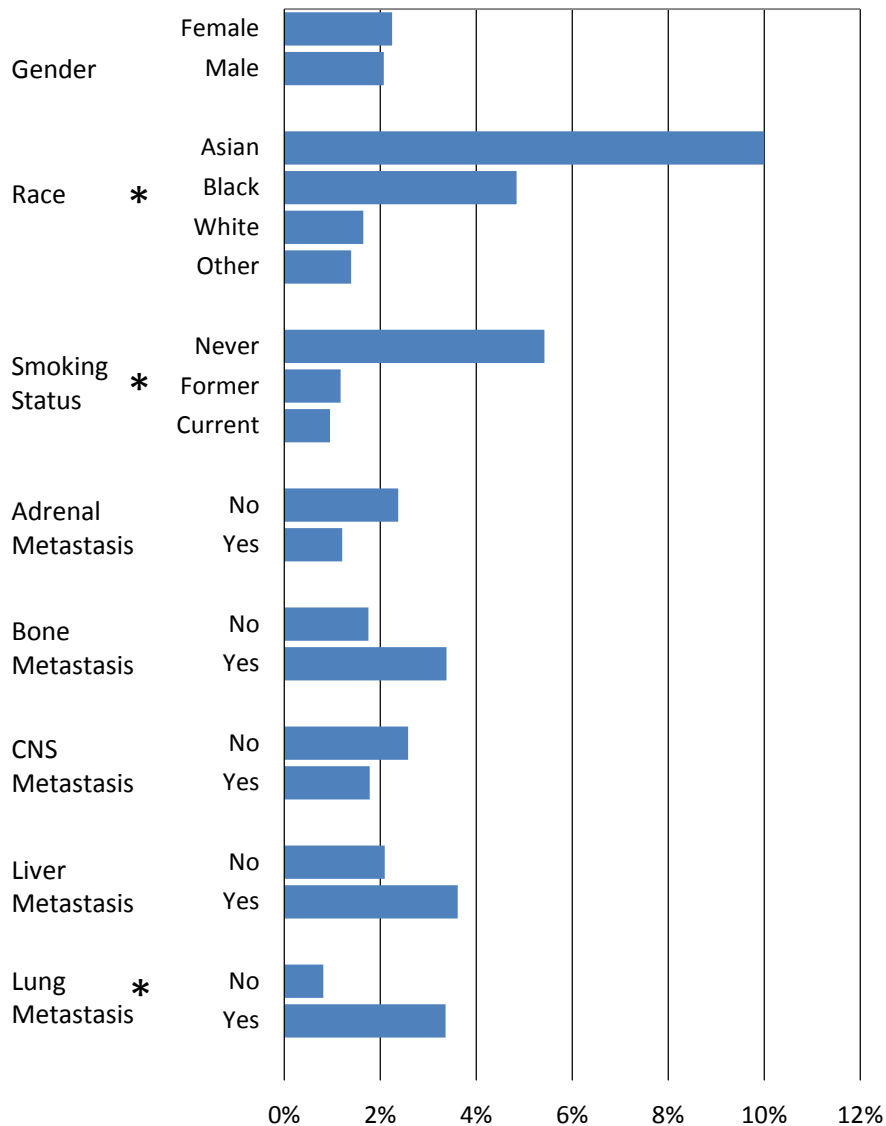


Gene: *RET* (rearrangement) (n=817)

Variable	N	WT (n=799)	Mut (n=18)	P
Sex	904			0.83
Female		53% (424)	56% (10)	
Male		47% (375)	44% (8)	
Race	904			0.98
Amer Indian/Alaska Native		0% (2)	0% (0)	
Asian		3% (27)	6% (1)	
Black/African American		8% (62)	11% (2)	
Native Hawaii/Pacific Isl.		0% (1)	0% (0)	
White		80% (639)	78% (14)	
Not disclosed		9% (68)	6% (1)	
Smoker	890			0.14
Current		13% (101)	6% (1)	
Former		63% (492)	50% (9)	
Never		25% (194)	44% (8)	
Stage at diagnosis	887			0.43
I		10% (77)	11% (2)	
II		7% (52)	11% (2)	
III		12% (92)	0% (0)	
IV		72% (568)	78% (14)	
Adrenal metastasis	863			0.96
No		89% (688)	89% (16)	
Yes		11% (83)	11% (2)	
Bone metastasis	871			0.39
No		71% (549)	61% (11)	
Yes		29% (229)	39% (7)	
Brain metastasis	869			0.61
No		72% (561)	78% (14)	
Yes		28% (214)	22% (4)	
Liver metastasis	864			0.92
No		90% (692)	89% (16)	
Yes		10% (80)	11% (2)	
Lung metastasis	882			0.96
No		45% (354)	44% (8)	
Yes		55% (433)	56% (10)	
Histopathologic subtype	788			0.90
Acinar predominant		33% (233)	21% (3)	
Lepidic predominant		2% (15)	0% (0)	
Micropapillary predom.		0% (1)	0% (0)	
Mucinous		1% (5)	0% (0)	
Papillary predominant		0% (3)	0% (0)	
Solid predominant		63% (439)	79% (11)	
Age at diagnosis	887			0.49
(Q1, median, Q3)		56.7 63.6 70.4	58.7 65.1 73.2	

Supplemental Figure 5

ROS1r, Positive (%)

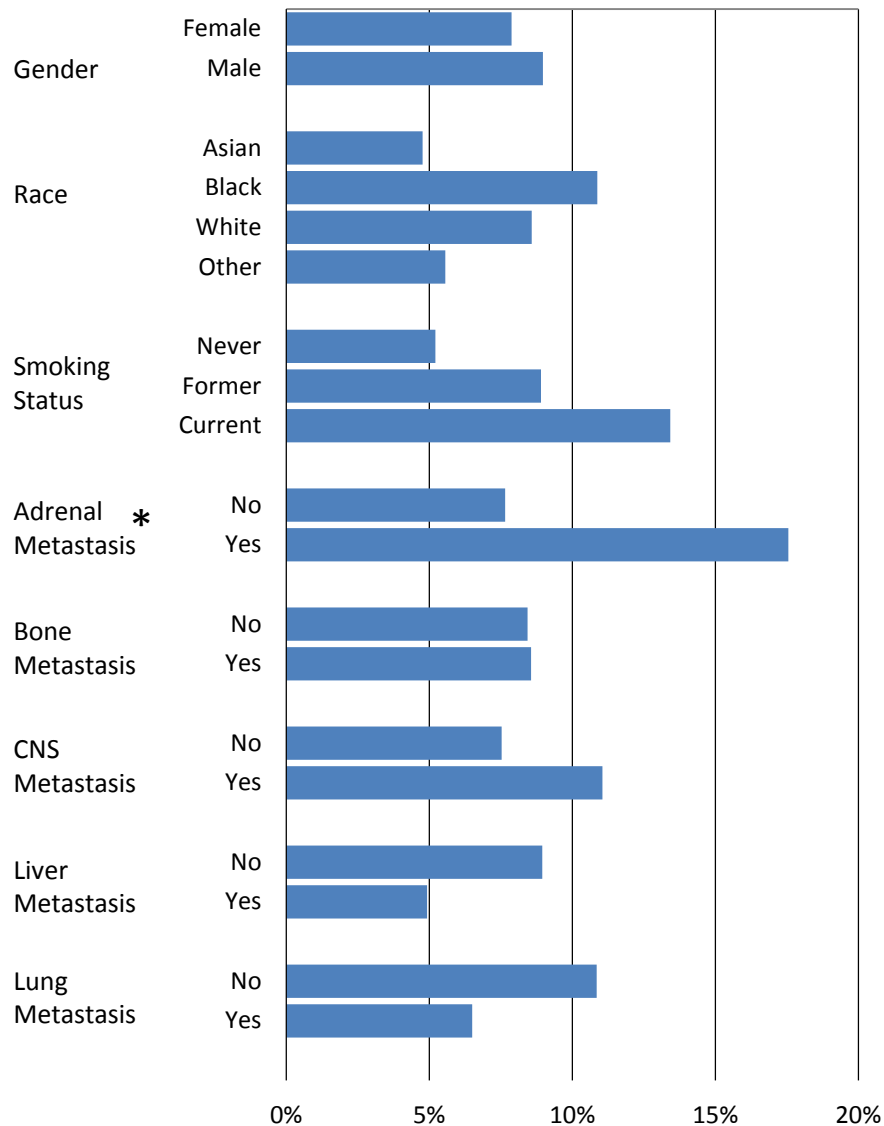


Gene: ROS1 (rearrangement) (n=832)

Variable	N	WT (n=814)	Mut (n=18)	P
Sex	904			0.87
	Female	54% (436)	56% (10)	
	Male	46% (378)	44% (8)	
Race	904			0.04
	Amer Indian/Alaska Native	0% (2)	0% (0)	
	Asian	3% (27)	17% (3)	
	Black/African American	7% (59)	17% (3)	
	Native Hawaii/Pacific Isl.	0% (1)	0% (0)	
	White	81% (657)	61% (11)	
	Not disclosed	8% (68)	6% (1)	
Smoker	890			0.001
	Current	13% (104)	6% (1)	
	Former	63% (506)	33% (6)	
	Never	24% (192)	61% (11)	
Stage at diagnosis	887			0.59
	I	10% (79)	0% (0)	
	II	7% (53)	6% (1)	
	III	12% (93)	12% (2)	
	IV	72% (579)	82% (14)	
Adrenal metastasis	863			0.5
	No	90% (700)	94% (17)	
	Yes	10% (82)	6% (1)	
Bone metastasis	871			0.15
	No	71% (561)	56% (10)	
	Yes	29% (229)	44% (8)	
Brain metastasis	869			0.59
	No	72% (567)	78% (14)	
	Yes	28% (221)	22% (4)	
Liver metastasis	864			0.38
	No	90% (703)	83% (15)	
	Yes	10% (80)	17% (3)	
Lung metastasis	882			0.01
	No	46% (366)	17% (3)	
	Yes	54% (432)	83% (15)	
Histopathologic subtype	788			<0.001
	Acinar predominant	33% (230)	35% (6)	
	Lepidic predominant	2% (15)	0% (0)	
	Micropapillary predom.	0% (0)	6% (1)	
	Mucinous	1% (5)	0% (0)	
	Papillary predominant	1% (4)	0% (0)	
	Solid predominant	64% (450)	59% (10)	
Age at diagnosis	887			0.03
	(Q1, median, Q3)	56.7 63.8 70.5	51.0 57.8 67.4	

Supplemental Figure 5

PTEN loss by IHC, Positive(%)

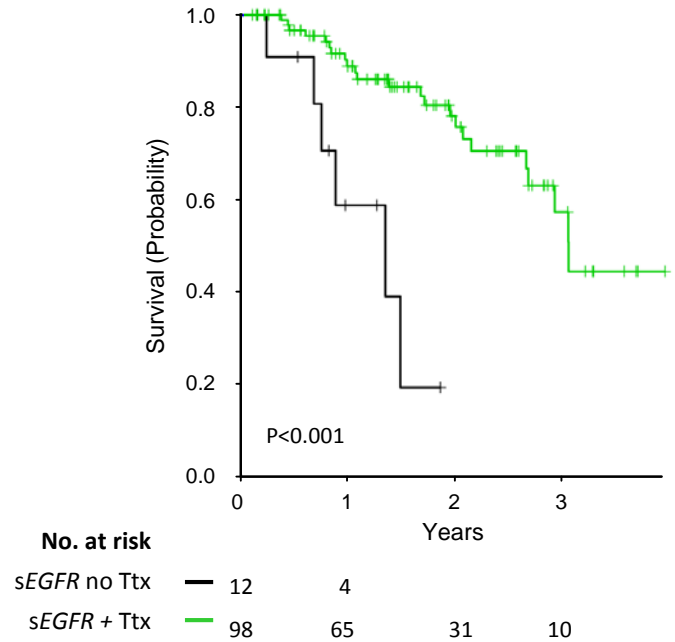


Gene: PTEN (IHC loss) (n=646)

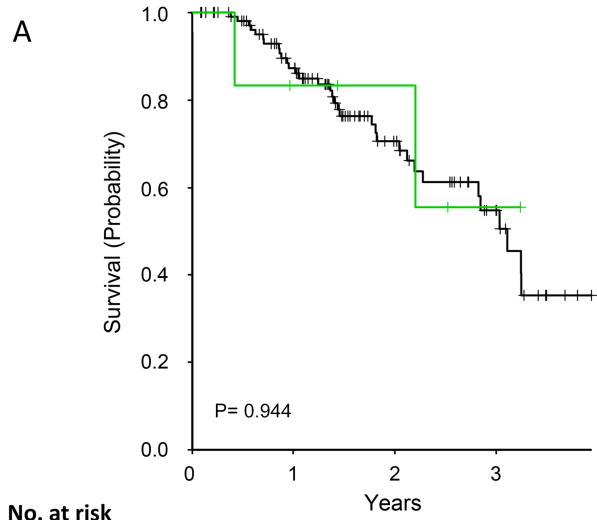
Variable	N	WT (n=592)	Mut (n=54)	P
Sex	904			0.62
		Female	55% (328)	52% (28)
		Male	45% (264)	48% (26)
Race	904			0.92
		Amer Indian/Alaska Native	0% (2)	0% (0)
		Asian	3% (20)	2% (1)
		Black/African American	7% (41)	9% (5)
		Native Hawaii/Pacific Isl.	0% (1)	0% (0)
		White	81% (480)	83% (45)
		Not disclosed	8% (48)	6% (3)
Smoker	890			0.08
		Current	12% (71)	20% (11)
		Former	60% (348)	63% (34)
		Never	28% (164)	17% (9)
Stage at diagnosis	887			0.03
		I	11% (65)	2% (1)
		II	8% (44)	4% (2)
		III	11% (63)	20% (11)
		IV	71% (413)	74% (40)
Adrenal metastasis	863			0.01
		No	92% (519)	81% (43)
		Yes	8% (47)	19% (10)
Bone metastasis	871			0.96
		No	70% (402)	70% (37)
		Yes	30% (171)	30% (16)
Brain metastasis	869			0.16
		No	73% (418)	64% (34)
		Yes	27% (153)	36% (19)
Liver metastasis	864			0.29
		No	90% (509)	94% (50)
		Yes	10% (58)	6% (3)
Lung metastasis	882			0.05
		No	45% (263)	59% (32)
		Yes	55% (317)	41% (22)
Histopathologic subtype	788			0.11
		Acinar predominant	33% (175)	15% (7)
		Lepidic predominant	2% (12)	0% (0)
		Micropapillary predom.	0% (1)	0% (0)
		Mucinous	1% (3)	0% (0)
		Papillary predominant	1% (3)	0% (0)
		Solid predominant	63% (336)	85% (39)
Age at diagnosis	887			0.92
		(Q1, median, Q3)	56.2 63.5 70.7	57.3 64.7 68.9

Supplemental Figure 6

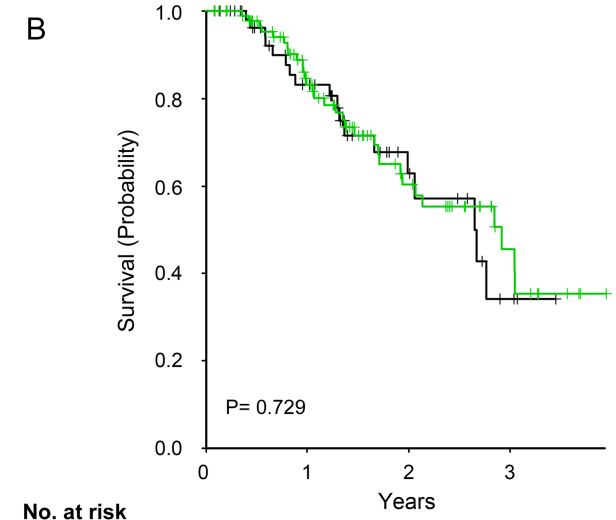
sEGFR survival comparison according to targeted treatment (Ttx) or not.



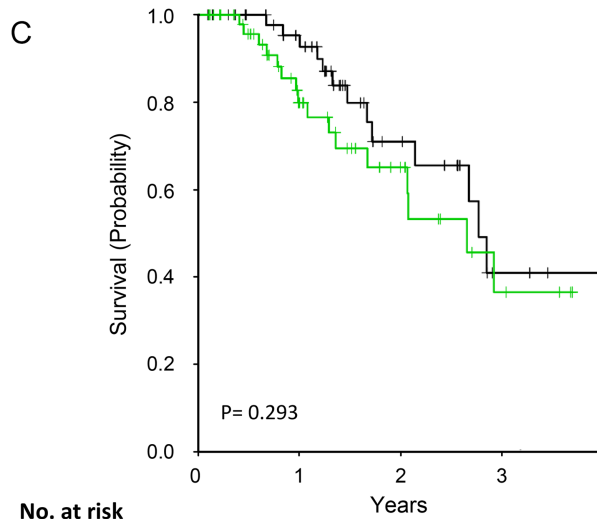
Supplemental Figure 7



No. at risk		0	1	2	3
PTEN expression by IHC	—	110	70	28	9
No PTEN expression by IHC	—	6	4	3	1



No. at risk		0	1	2	3
MET expression by IHC	—	57	36	12	3
No MET expression by IHC	—	93	57	25	9

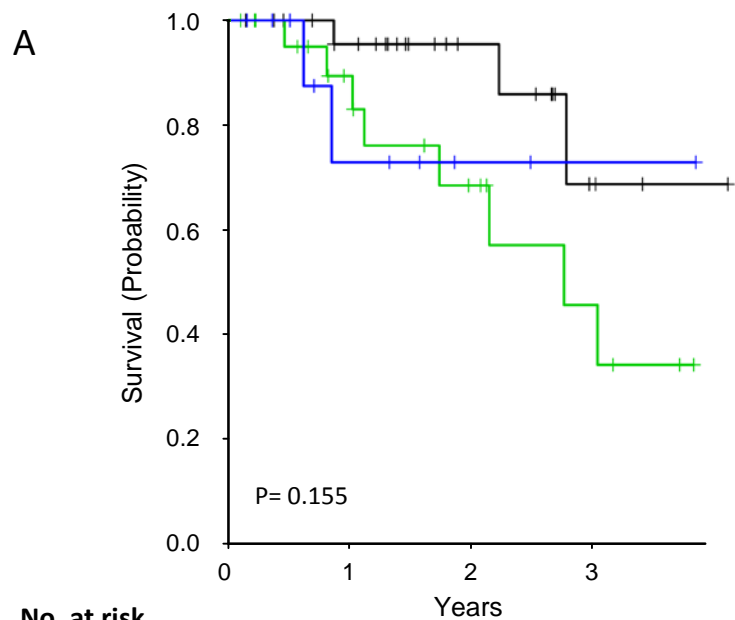


No. at risk		0	1	2	3
No <i>PTEN/PIK3CA/TP53</i>	—	51	36	14	3
+ <i>PTEN/PIK3CA/TP53</i>	—	52	26	12	4

Comparison of survival in the targeted therapy cohort according to presence or absence of several molecular features

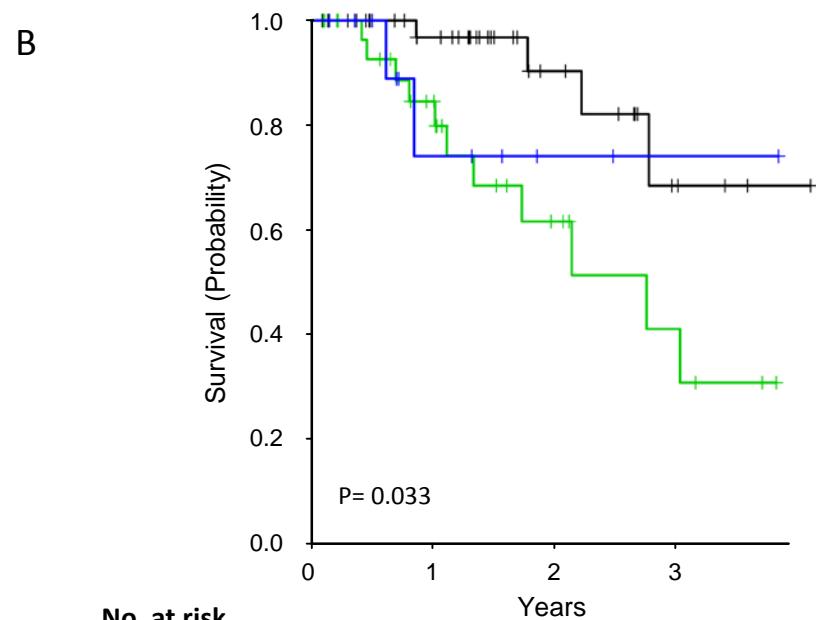
Supplemental Figure 8

Survival comparisons according to presence or absence of TP53 mutation.



No. at risk

<i>sEGFR</i> Ttx No <i>TP53</i>	26	20	10	2
<i>sEGFR</i> Ttx <i>TP53</i> disruptive	23	12	7	3
<i>sEGFR</i> Ttx <i>TP53</i> nondisruptive	11	5	2	1



No. at risk

Driver Ttx No <i>TP53</i>	40	28	12	3
Driver Ttx <i>TP53</i> disruptive	31	15	7	3
Driver Ttx <i>TP53</i> nondisruptive	12	5	2	1