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TRIAL OF STAGE IB to IIIA LUNG CANCER	Neoadjuvant chemotherapy alone	Chemotherapy PLUS NIVOLUMAB
Event-free survival	20.8 months	31.6 months
Pathological complete response	2.2%	24%
Grade 3 or 4 adverse events	36.9%	33.5%
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CHECKMATE 816: BIG CHANGES!!
Patients with unresectable and/or locally advanced lung cancer can now be resected after neoadjuvant chemoimmunotherapy
Some of these cancers have complete pathologic responses
Response to therapy can be measured by pathologists (with major work by pathologists' assistants)
Patients are doing better! Some may be cured
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	NEOADJUVANT CHEMOIMMUNOTHERAPY NEED IEED TO PROVIDE LOTS OF INFO ON THESE CASES
COMPLETE PATHOLOGIC RESPONSE (CPR) = NO VIABLE TUMOR CELLS MAJOR PATHOLOGIC RESPONSE (MPR) = LESS THAN OR EQUAL TO 10% VIABLE TUMOR	Diagnosis Comment [4] Type of neoaduvant therapy: chemoimmunotherapy (checkmate 816 protocol) Assessment of primary tumor/tumor bed (post neoadjuvant therapy): Percentage of viable tumor: 80% Percentage of neorosis: 10% Percentage of stroma (inflammation/fibrosis): 10% Grade of inflammation: moderate Method: Correlation with gross photograph: Yes Evaluation aided by tumor mapping to match gross photograph to histologic sections: Yes Evaluation aided by correlation of pathologic findings with imaging: Yes Overall assessment: Complete pathologic response (CPR) present (no viable tumor cells): No Major pathologic response (CPR) present (no viable tumor cells): No Major pathologic response (MPR) present (less than or equal to 10% viable tumor): No Treatment effect in lymph node metastases: Total number of lymph nodes examined: 8 Total number of lymph nodes examined: 31 Carcinoma present? No Total number of lymph nodes with metastatic carcinoma: 0 Lymph nodes stations involved by tumor with treatment-related changes: 0 Lymph nodes stations involved by treatment-related changes without viable tumor: 1 Largest tumor focus: NA Extracapsular invasion: N/A



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ro				s? A stud			Case	Histological assessment	Original report	Molecular analysis	Driver mutation in tumour 1	Driver mutation in tumour 2
	secte	d lung no	dules in 3	2 natient	s using a	routine	1	R	R	UR	Negative	KRAS c.35G>T
						routine	2	R	R	18	KRAS ¢ 34G>T	KRAS c.34G>T
ne	ext-a	eneration	sequenci	ng panel f	or		3	UR	UR	UR	EGFR c.2240T>C	KRAS c.34G>T
	-			.g paner i			4	UR	UR	E	KRAS c.34G>T	KRAS c.34G>T
dr	iver I	mutations					5	R	R	UR	BRAF c.1406G>T	KRAS c.35G>T
	i ci i	natations					6	UR	UR	UR	Negative	KRAS c.35G>T
Erid	O V Day	ehl 😑 ,1 Erika E	Doxtador 1 V.	Mai Chang 1 D	niel U Farler	Carol Eanuar ²	7	R	R	R	MET c.3029C>T	MET c.3029C>T
FI10	IO K BIU	eni 🔍, Erika E	Doxiader, Yu-	wer cheng, "Di	aniel n Farkas,	Caror rarver,-	8	B	R	R	EGFR c.2573T>G	EGFR c.2573T>G
San	ijay Mul	khopadhyay 💿					9	UR	UR	UR	KRAS c.34G>T	Negative
							10	UR	R	1	KRAS c.35G>T	KRAS c.35G>T
							11	R	R	R	ERBB2 c.2264T>C	ERBB2 c.2264T>C
							12	8	8	R	EGFR c.2240_2257del	EGFR c.2240 2257del
							13	B	R	B	KRAS c.57G>T	KRAS c.57G>T
							14	R	R	T.	Negative (only EGFR tested)	Negative
							15	R	R	1	Negative	Negative
							16	8	R	R	BRAF NS815	BRAF N5815
hle 3	Cases they	unht to be related by histol	any best proven to be since	lated by molecular analysi	6		17	R			KRAS C 183A>T	KRAS C 34G>T
ble 3	Cases tho	ught to be related by histol	ogy but proven to be unre	elated by molecular analysi	6	Orininal turnour	17		UR	UR	KRAS c.183A>T FGFR exon 19 n Glu746, Ala750del	KRAS c.34G>T FGFR c 2303_2304insTGTGGCC
ble 3 se	Cases thou Location	ught to be related by histol	ogy but proven to be unre Driver gene variant	elated by molecular analys Tumour histology	is Tumour size (mm)	Original tumour stage	17 18 19	R UR UR	UR	UR UR UR	EGFR exon 19 p.Glu746_Ala750del	EGFR c.2303_2304insTGTGGCC
	Location RML		Driver gene variant	Tumour histology MP80, P10, A10, NM	Turnour size (mm) 30	stage T4N0	18	UR		UR	EGFR exon 19 p.Glu746_Ala750del KRAS c.34G>T	EGFR c.2303_2304insTGTGGCC Negative
	Location RML RUL	Histological assessment R	Driver gene variant NVD KRAS c.35G>T	Tumour histology MP80, P10, A10, NM MP80, A20, MX	Tumour size (mm) 30 25	stage T4N0 T4N0	18 19	UR	UR UR	UR	EGFR exon 19 p.Glu746_Ala750del	EGFR c.2303_2304insTGTGGCC Negative Negative (ERBB2 not tested)
	Location RML RUL RUL	Histological assessment	Driver gene variant NVD KRAS c.35G>T BRAF c.1406G>T	Tumour histology MP80, P10, A10, NM MP80, A20, MX P70, A30, NM	Tumour size (mm) 30 25 14	stage T4N0 T4N0 T4N0 T4N0	18 19 20 21	UR UR R UR	UR UR R UR	UR UR I UR	EGFR exon 19 p.Glu746_Ala750del KRAS c.34G>T ER882 p.A775_G776insYVMA KRAS c.35G>T	EGFR c.2303_2304insTGTGGCC Negative Negative (ERBB2 not tested) Negative
	Location RML RUL	Histological assessment R	Driver gene variant NVD KRAS c.35G>T	Tumour histology MP80, P10, A10, NM MP80, A20, MX	Tumour size (mm) 30 25	stage T4N0 T4N0	18 19 20 21 22	UR UR R UR UR	UR UR R UR UR	UR UR I UR UR	EGFR exon 19 p.Glu746_Ala750del KRAS c.34G>T ERB82 p.A775_G776insYVMA KRAS c.35G>T KRAS c183A>C	EGFR c.2303_2304insTGTGGCC Negative Negative (ERB82 not tested) Negative MET c.3028+2T>C
se	Location RML RUL RUL RLL LUL RLL	Histological assessment R	Driver gene variant NVD KRAS c.35G>T BRAF c.1406G>T KRAS c.35G>T KRAS c.35G>T KRAS c.34G>T	Tumour histology MP80, P10, A10, NM MP80, A20, MX P70, A30, NM P70, A20, MP10, NM	Tumour size (mm) 30 25 14 23 20 18	stage T4N0 T4N0 T4N0 T4N0 T4N0 T2aN0 T2aN0	18 19 20 21 22 23	UR UR R UR	UR UR R UR UR UR	UR UR I UR UR UR	EGFR exon 19 p.Glu746_Ala750del KRA5 c.34G>T ERB82 p.A775_G776insYVMA KRA5 c.35G>T KRA5 c.183A>C KRA5 c.35G>T	EGFR c.2303_2304insTGTGGCC Negative Negative (ERBB2 not tested) Negative MET c.3028+2T>C KRAS c.34G>T
se	Location RML RUL RUL RLL LUL RLL RUL	Histological assessment R	Driver gene variant NVD KRAS c.35Gs-T BRAF c.1406Gs-T KRAS c.35Gs-T KRAS c.35Gs-T KRAS c.183As-T KRAS c.38Gs-T KRAS c.38Gs-T KRAS c.38Gs-T KRAS c.38Gs-T KRAS c.38Gs-T	Tumour histology MP80, P10, A10, NM MP80, A20, MX P70, A30, NM P70, A30, MM P70, A30, MM A60, P30, MP10, NM A50, P10, MP10, NM S50, MP30, P20, NM	Turnour size (mm) 30 25 14 23 20 18 18	stage T4N0 T4N0 T4N0 T4N0 T4N0 T4N0 T2AN0 T1DN0	18 19 20 21 22	UR UR R UR UR UR	UR UR R UR UR	UR UR I UR UR	EGFR exon 19 p.Glu746_Ala750del KRAS c.34G>T ERB82 p.A775_G776insYVMA KRAS c.35G>T KRAS c183A>C	EGFR c.2303_2304insTGTGGCC Negative Negative (ERBB2 not tested) Negative MET c.3028+2T>C KRAS c.34G>T KRAS c.34G>A
se	Location RML RUL RUL RUL LUL RUL RUL RUL	Histological assessment R R R R	Driver gene variant NVO KRAS (35G:57 KRAS (35G:57 KRAS (35G:57 KRAS (35G:57 KRAS (35G:57 KRAS (35G:54 KRAS (35G:54 KRAS (34G:54	Tumour histology MP80, P10, A10, NM MP80, A20, MX P70, A30, NM P70, A20, MP10, NM A60, P30, MP10, NM A60, P30, MP10, NM S50, MP30, P20, NM S50, MP30, P20, NM	Tumour size (mm) 30 25 14 23 20 18 18 17	stage T4N0 T4N0 T4N0 T4N0 T4N0 T2aN0 T2aN0	18 19 20 21 22 23 24 25	UR UR R UR UR UR UR	UR UR R UR UR UR UR UR	UR UR I UR UR UR UR UR	EGFR exon 19 p.Glu746_Ala/250del KRAS c.34Gs-T ER882 p.A/75_G776insYVMA KRAS c.35Gs-T KRAS c.183A>C KRAS c.35Gs-T KRAS c.35Gs-T KRAS c.35Gs-T KRAS c.37Gs-T	EGFR c.2303_2304(instGTGGCC Negative Negative (ERB82 not tested) Negative MET c.3028+27>C KRAS c.34G>T KRAS c.34G>T KRAS c.34G>T
se	Location RML RUL RUL RUL LUL RUL RUL RUL	Histological assessment R R R	Driver gene variant NVO KRAS (35G:57 KRAS (35G:57 KRAS (35G:57 KRAS (35G:57 KRAS (35G:57 KRAS (35G:54 KRAS (35G:54 KRAS (34G:54	Tumour histology MP80, P10, A10, NM MP80, A20, MX P70, A30, NM P70, A20, MP10, NM A60, P30, MP10, NM A60, P30, MP10, NM S50, MP30, P20, NM S50, MP30, P20, NM	Tumour size (mm) 30 25 14 23 20 18 18 17	stage T4N0 T4N0 T4N0 T4N0 T4N0 T4N0 T2AN0 T1DN0	18 19 20 21 22 23 24	UR UR R UR UR UR R	UR UR R UR UR UR UR	UR UR I UR UR UR UR	EGFR exon 19 p.Glu746_Ala750del KRAS c.34G5-T ER882_DA775_G776insYVMA KRAS c.35G5-T KRAS c.33G5-T KRAS c.33G5-T KRAS c.33G5-T KRAS c.33G5-T KRAS c.33G5-T Negative	EGFR c.2303_2304ins1GTGGCC Negative Negative (ER882 not tested) Negative MET c.3028+2T>C KRAS c.34G>T KRAS c.34G>T KRAS c.34G>T
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