

The role of co-mutations in immunotherapy response in NSCLC

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Associate Professor

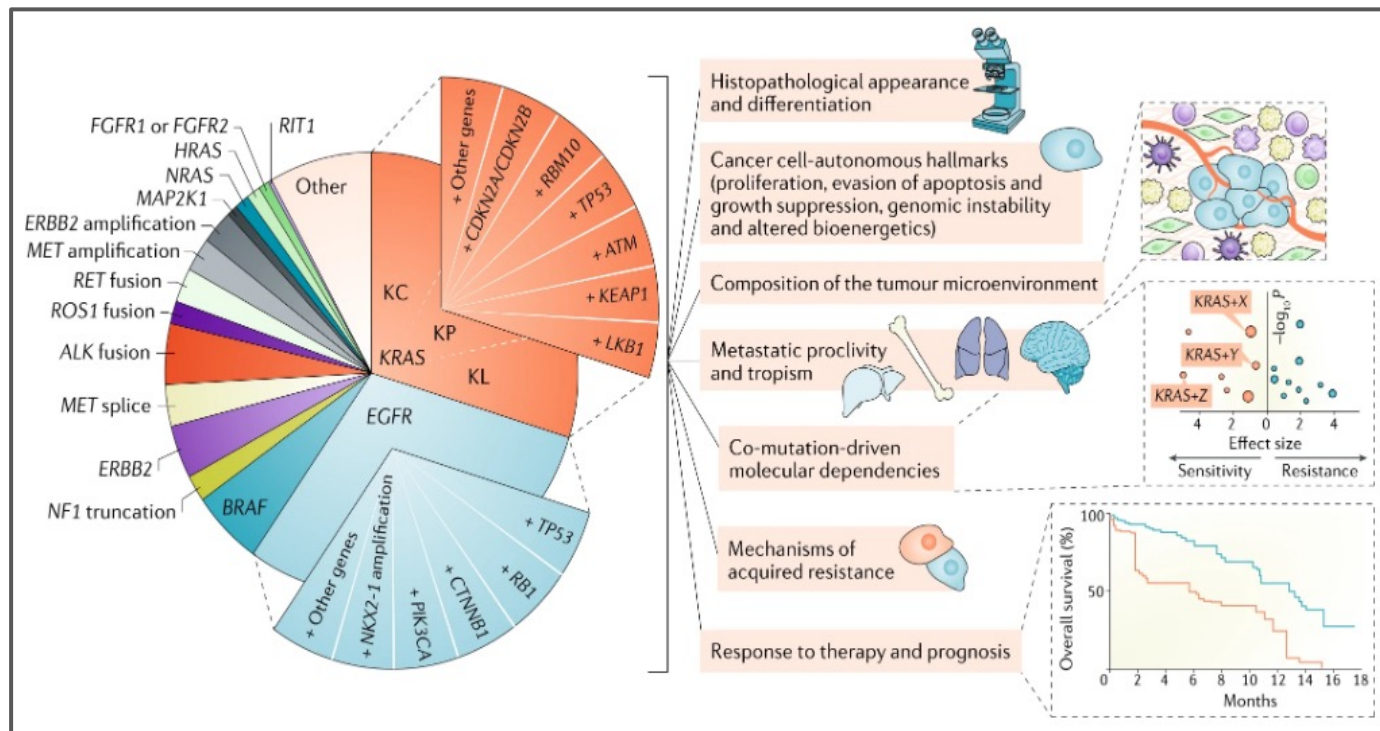
Department of Thoracic/Head and Neck Medical Oncology

Masters in Thoracic Oncology Summit (MaTOS™), Albuquerque, New Mexico

November 18, 2023

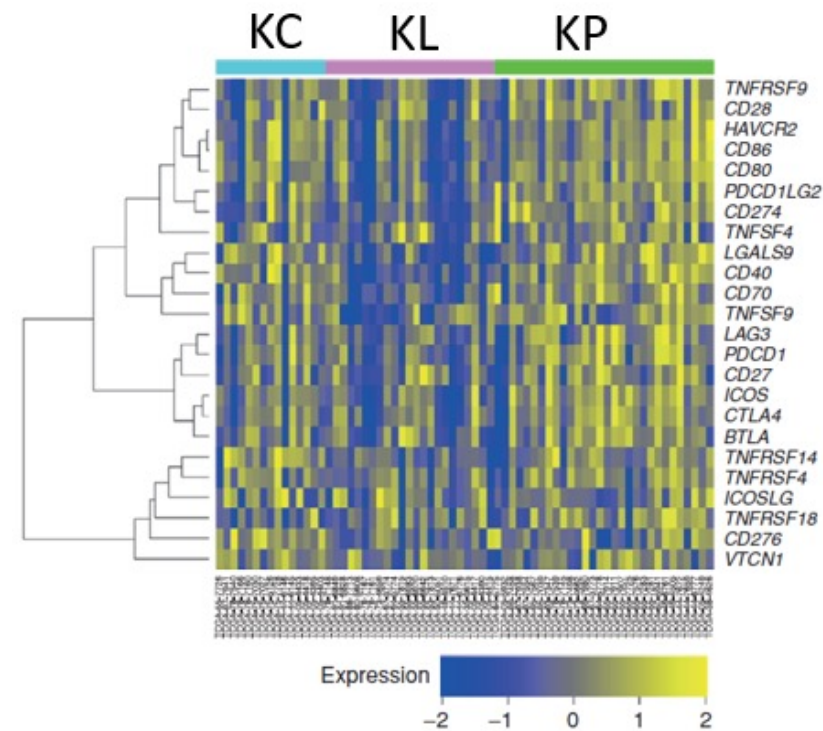
Oncogenic drivers and co-alterations collectively impact both tumor cell-intrinsic and non-tumor cell autonomous cancer hallmark traits

A.



Skoulidis F and Heymach JV, *Nat Rev Cancer*, 2019

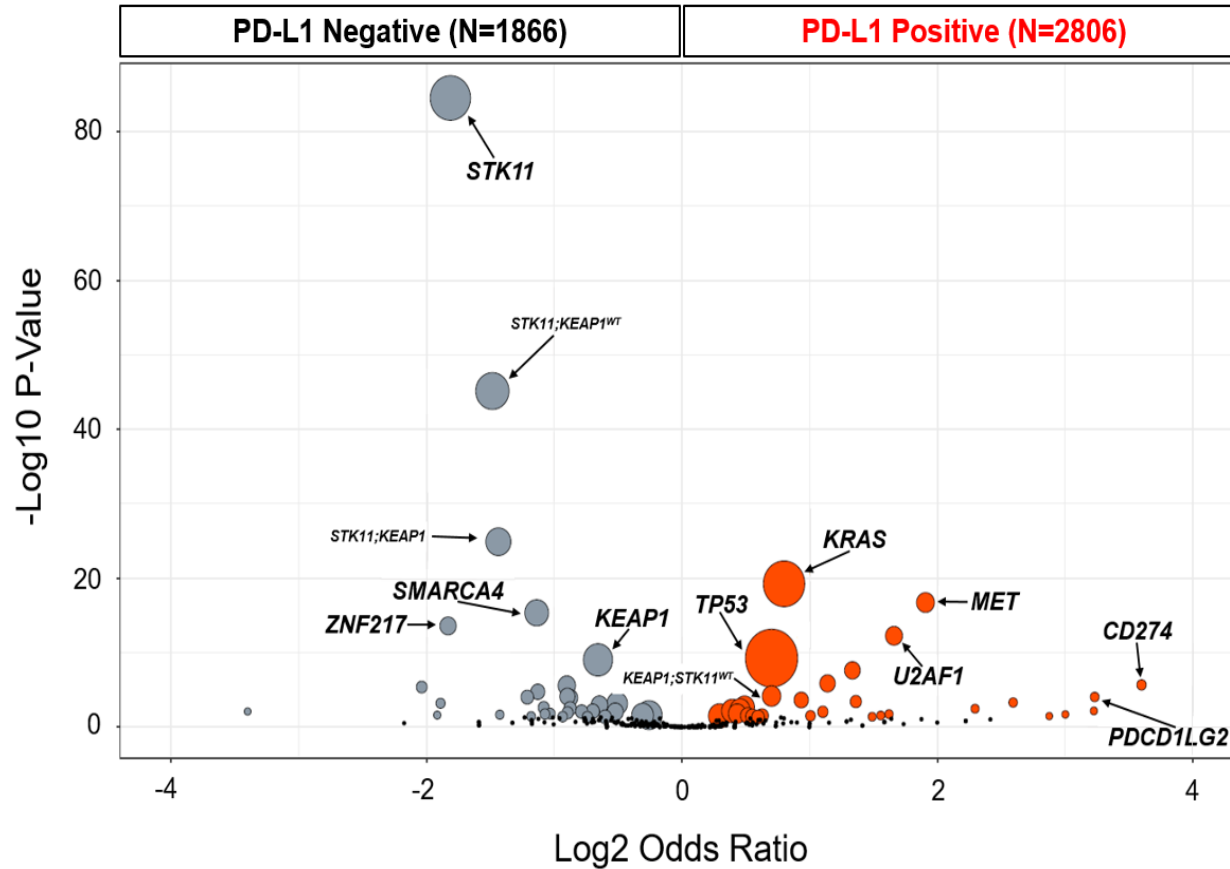
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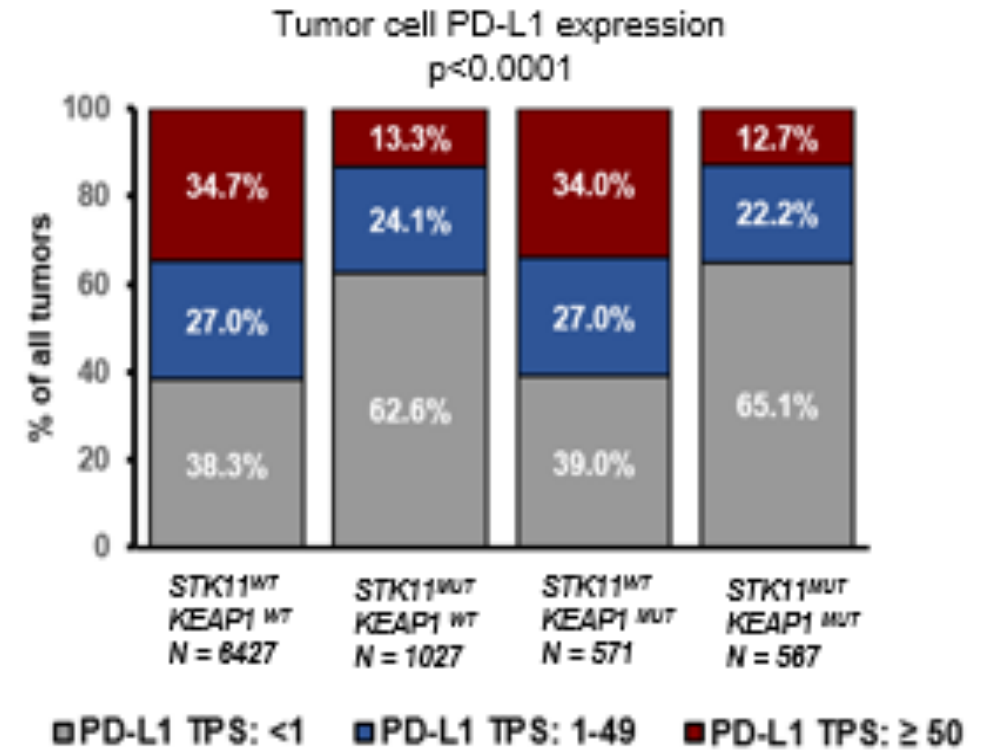
Skoulidis F et al., *Cancer Discovery*, 2015

Oncogenotype and tumor cell PD-L1 expression in lung adenocarcinoma

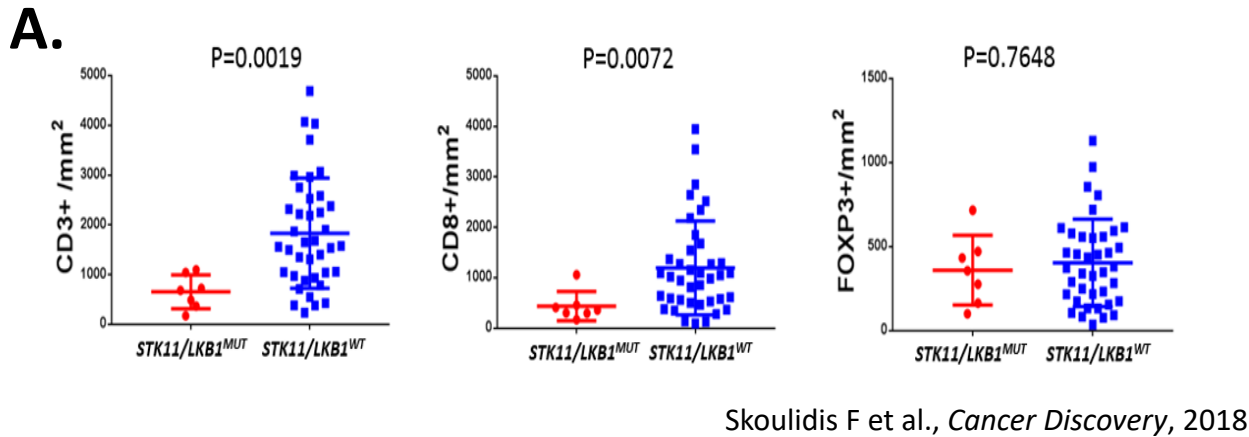
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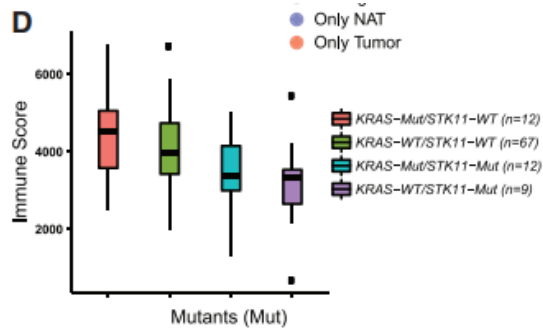
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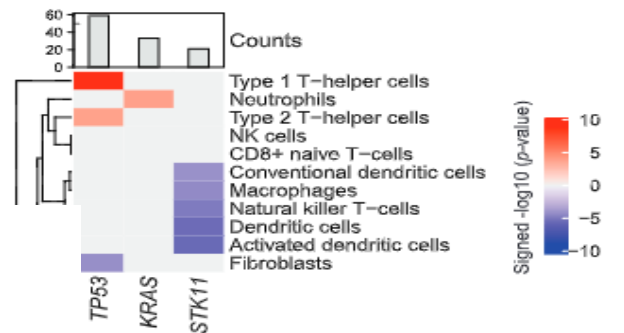
Somatic mutations in *STK11/LKB1* (KL) promote establishment of a T-cell depleted and suppressive myeloid cell –enriched TIME in NSCLC



B.

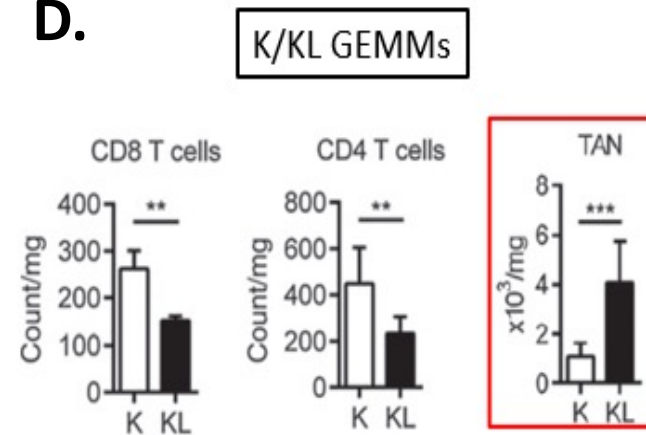


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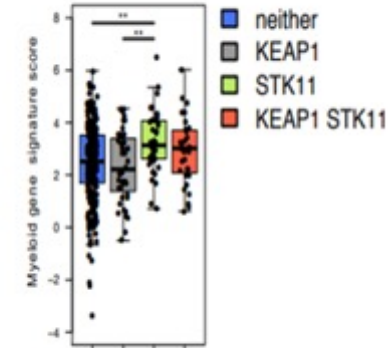
Gillette MA et al., *Cell*, 2020

D.



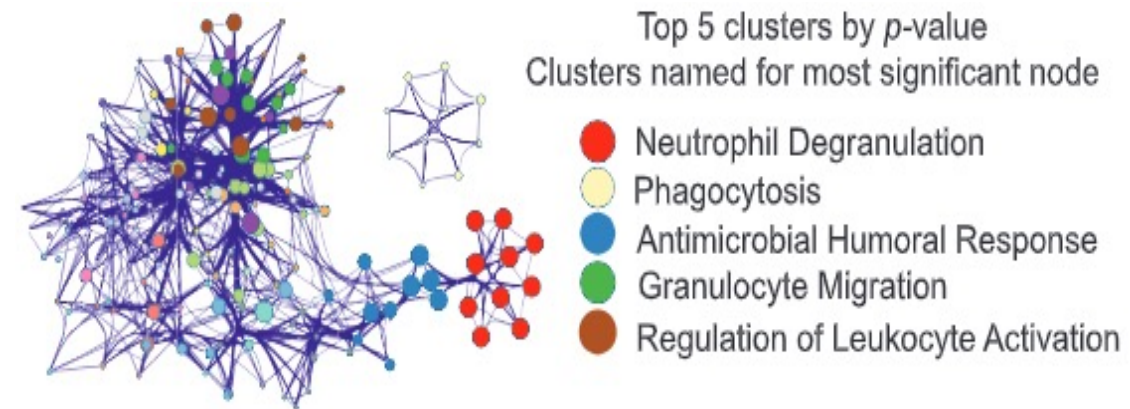
Koyama S et al, *Cancer Research*, 2016

OAK study, GEP



Singh A et al., *Clin Cancer Res*, 2020

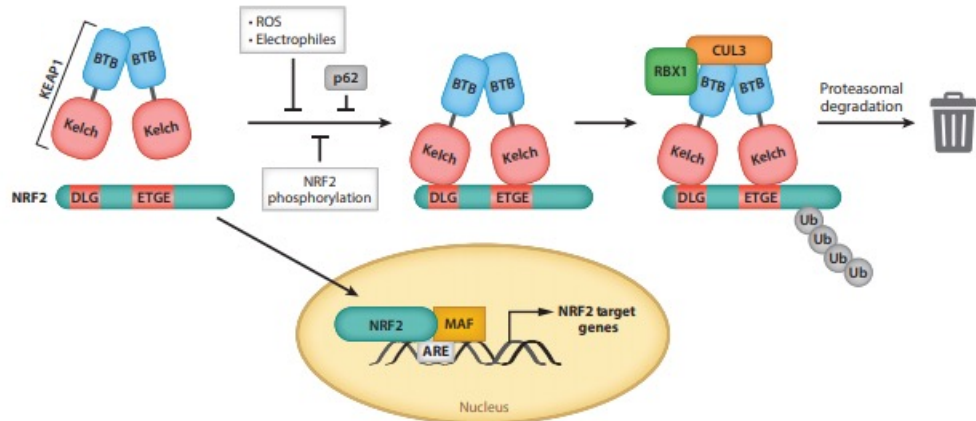
E.



Gillette MA et al., *Cell*, 2020

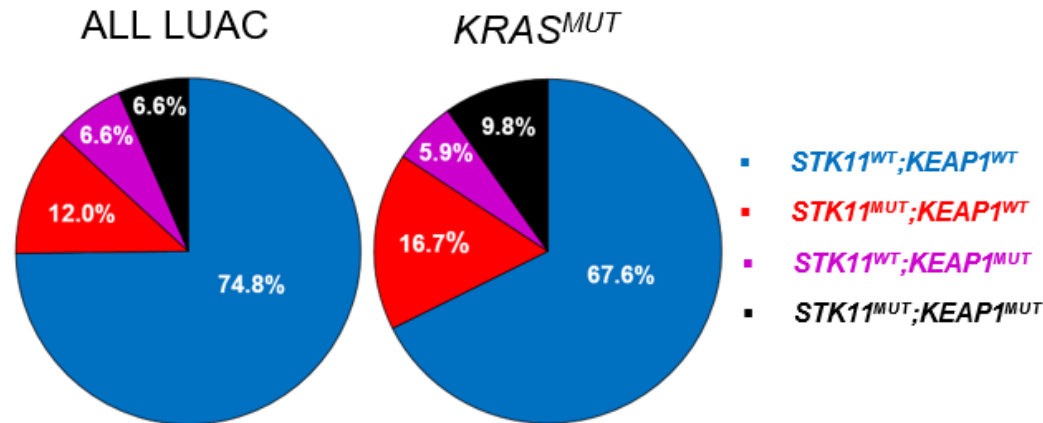
KEAP1 inactivation is also associated with an altered NSCLC TIME

A.



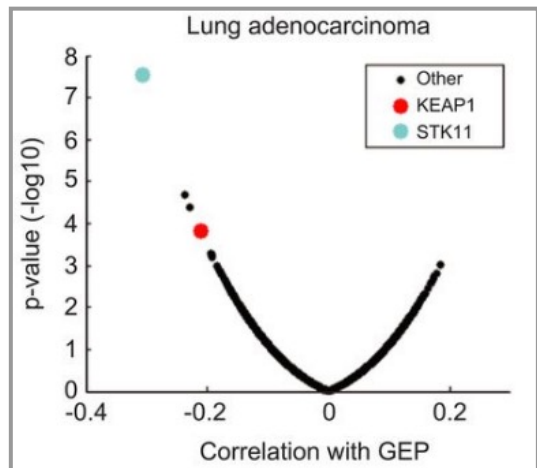
Wu WL, Papagiannakopoulos T, 2021

B.



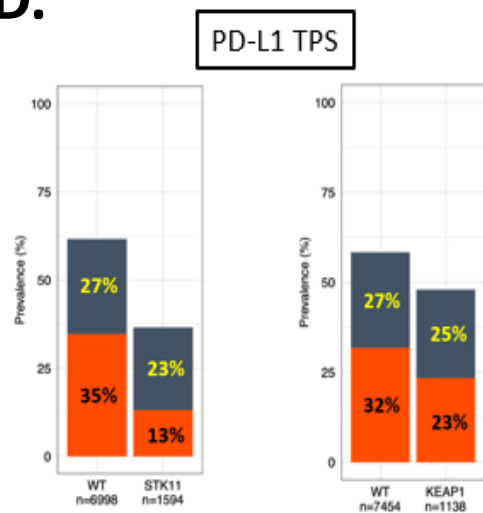
Collaboration with Lee Albacker, FMI

C.



Cristescu R et al, *Science*, 2018

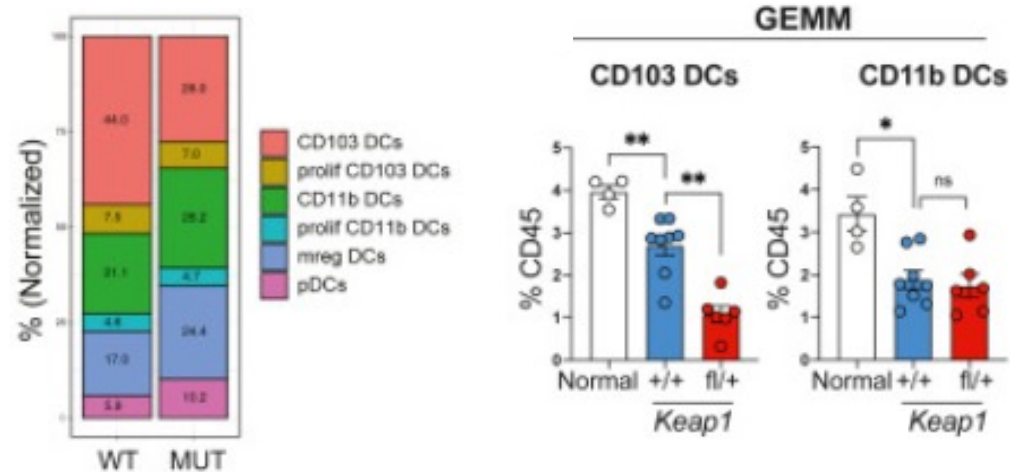
D.



Skoulidis F et al, EON meeting, 2020
Collaboration with Dr Albacker, Foundation Medicine

E.

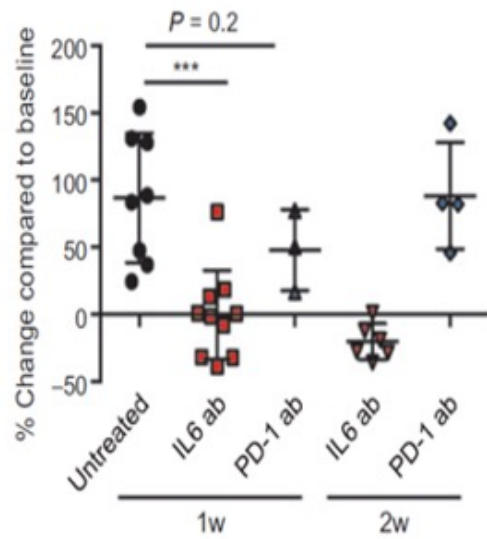
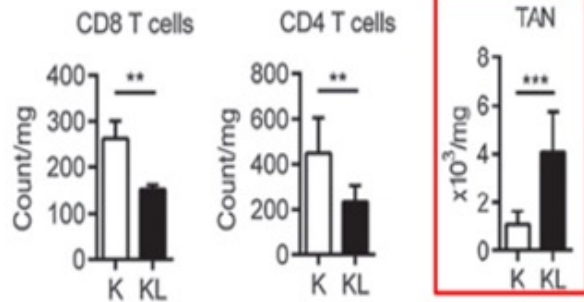
Depletion of CD103+ DCs



Zavitsanou AM et al., *Cell Reports*, 2023

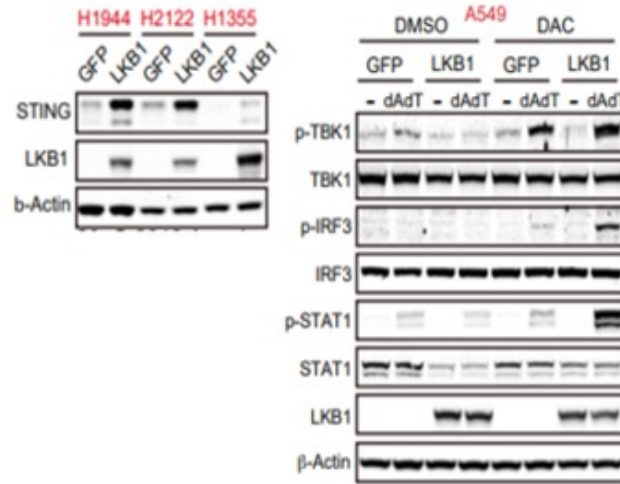
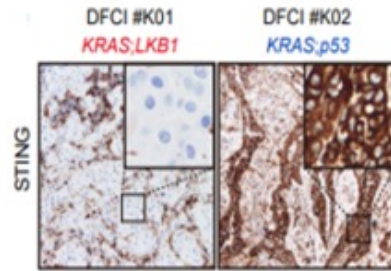
Mechanisms of *STK11* and *KEAP1* loss-mediated immune escape

Recruitment of MDSCs



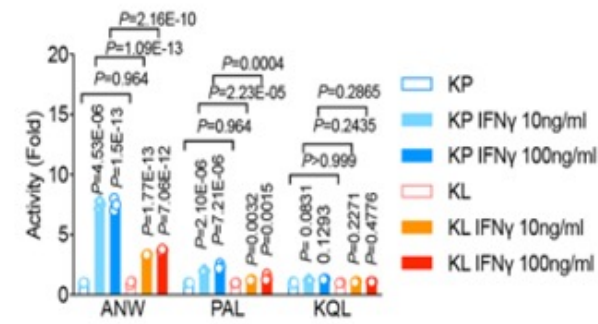
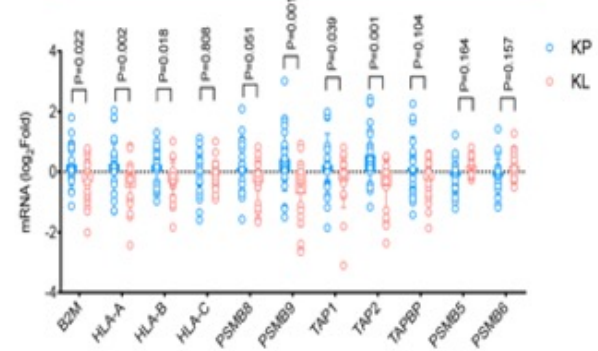
Koyama S et al., *Cancer Research*, 2016

Repression of STING



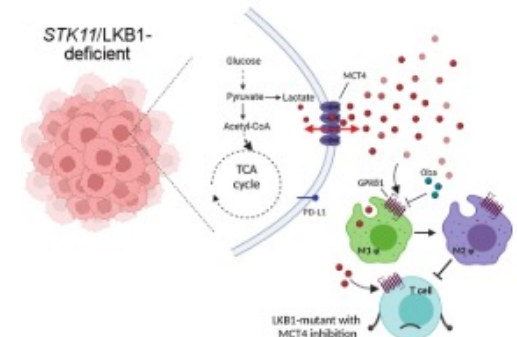
Kitajima S et al., *Cancer Discovery*, 2018

Impaired antigen presentation

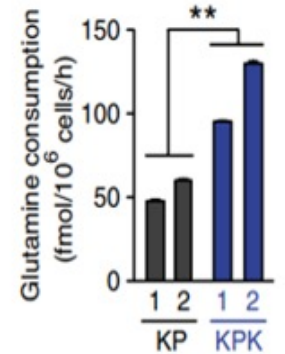


Deng J et al., *Nat Cancer*, 2021

Metabolically adverse TME

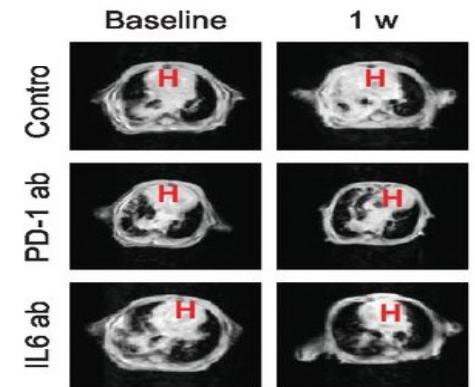
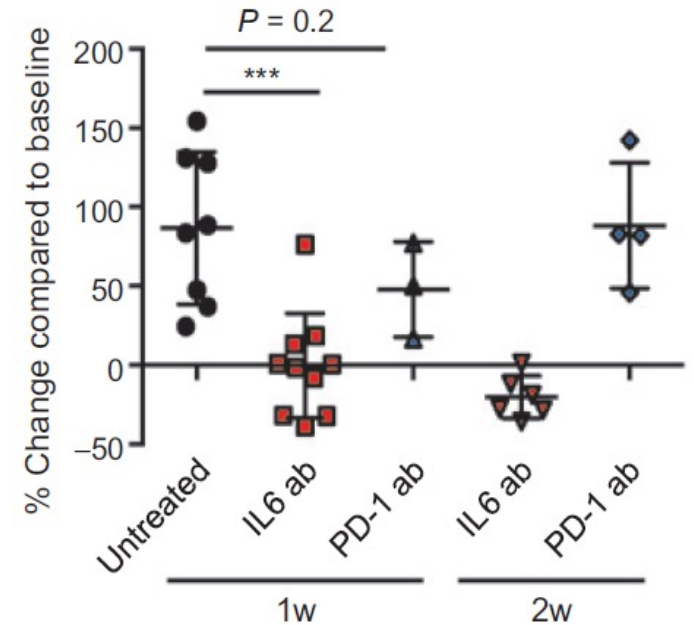
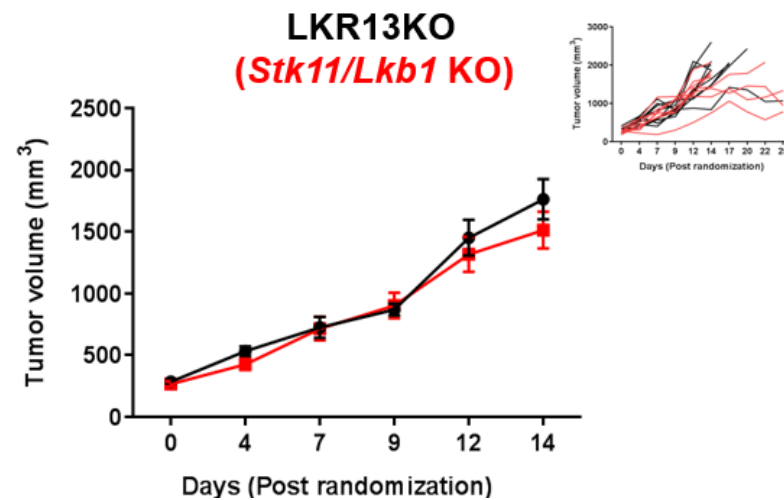
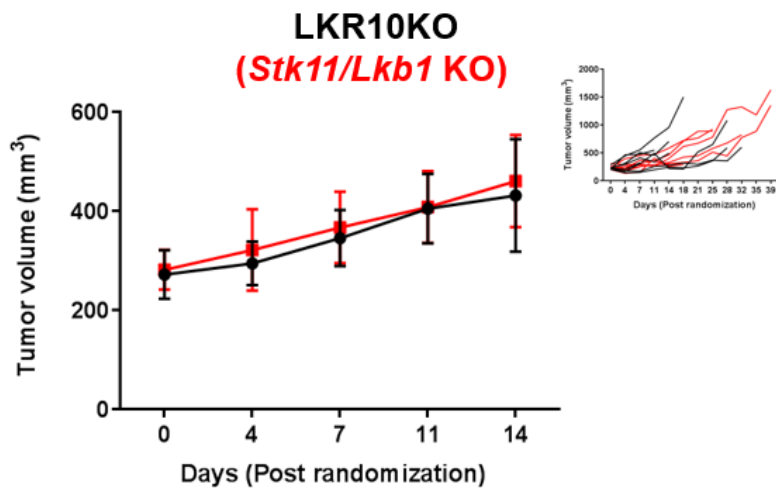
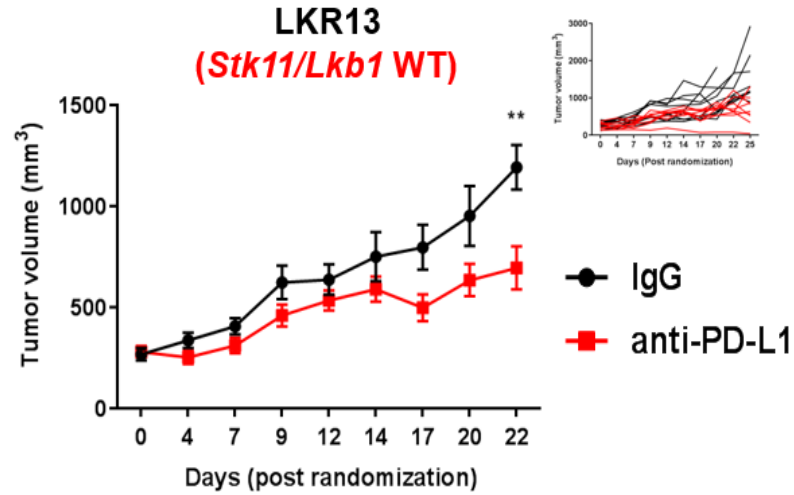
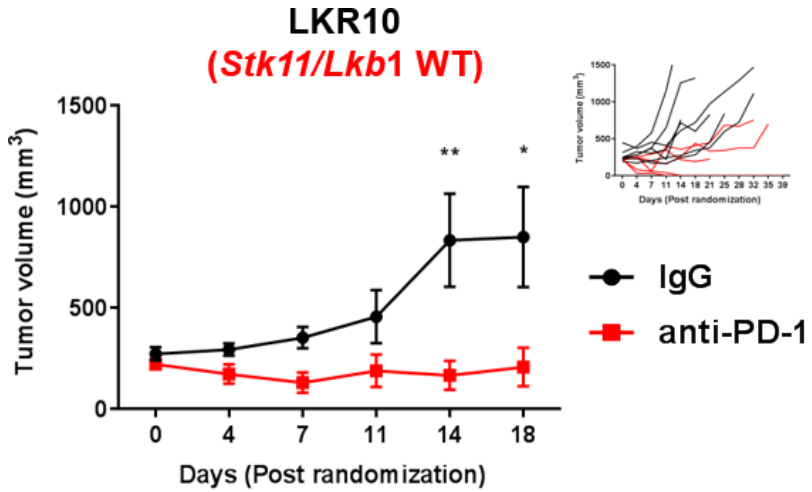


Qian et al., *Cancer Cell*, 2023



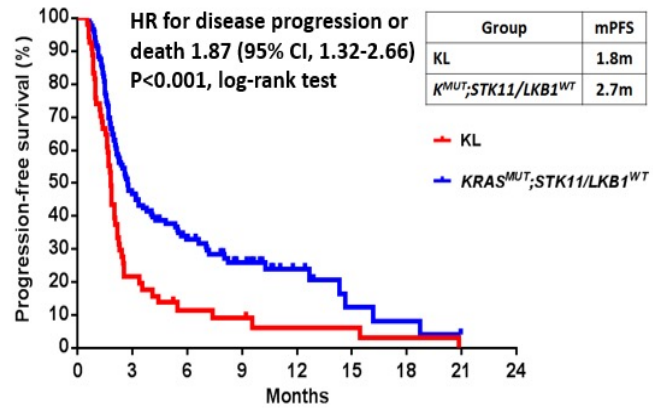
Romero et al., *Nat Med*, 2017

Stk11/Lkb1 loss promotes primary resistance to PD-1/PD-L1 blockade in immune-competent models of *Kras*-mutant LUAC

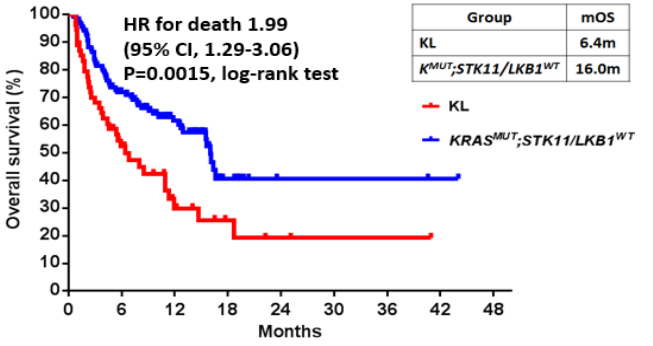


STK11 and KEAP1 alterations drive inferior clinical outcomes with PD-1 axis inhibitor monotherapy in KRAS-mutant NSCLC

Skoulidis F et al., Cancer Discov, 2018

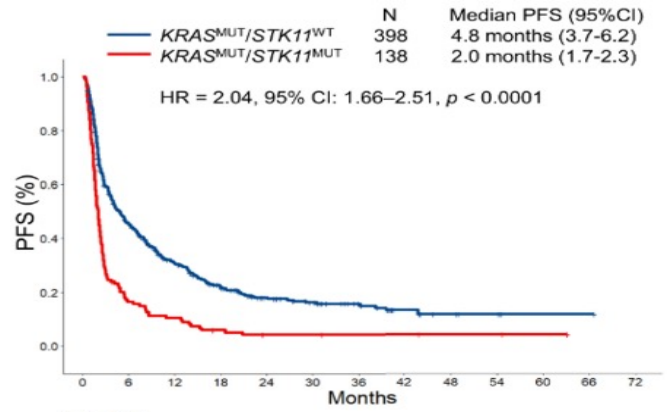


KL	54(0)	11(2)	5(3)	4(3)	2(4)	2(4)	1(4)	1(4)	0(4)
$K^{MUT};STK11^{WT}$	120(0)	55(3)	34(9)	18(18)	8(27)	3(29)	2(29)	1(29)	0(30)

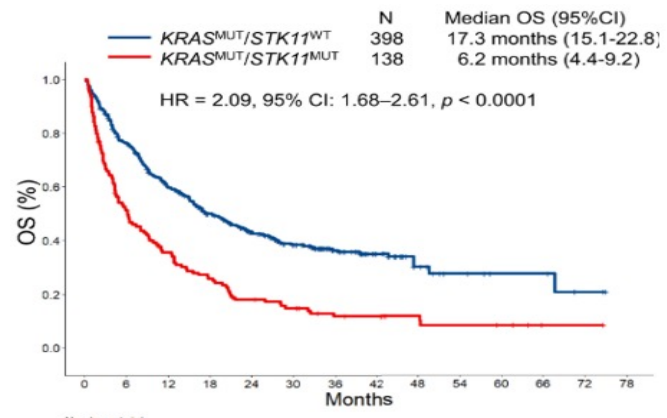


KL	54(0)	25(5)	10(12)	4(16)	2(17)	1(18)	1(18)	0(19)	0(19)
$K^{MUT};STK11^{WT}$	120(0)	81(6)	46(32)	8(60)	2(66)	2(66)	2(66)	1(67)	0(68)

Ricciuti B et al., JTO, 2021

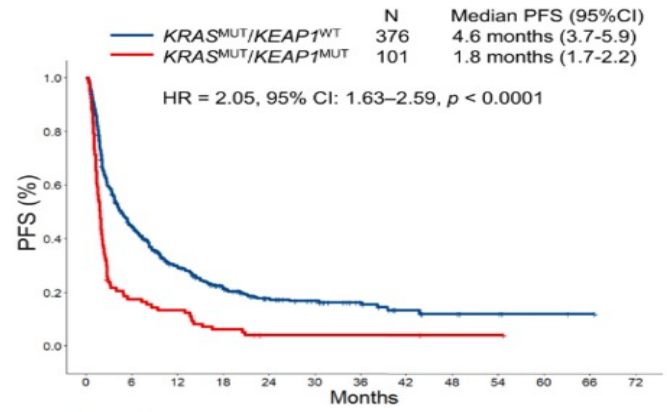


Group	0	6	12	18	24	30	36	42	48	54	60	66	72
$KRAS^{MUT};STK11^{WT}$	398	172	108	89	44	34	21	9	5	3	1	1	0
$KRAS^{MUT};STK11^{MUT}$	138	22	14	7	4	4	3	3	2	2	1	0	0

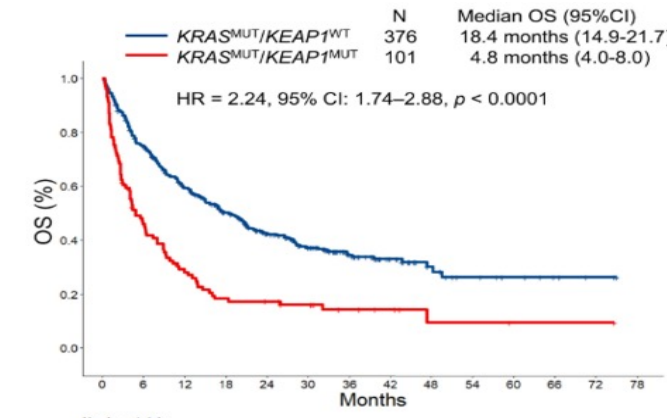


Group	0	6	12	18	24	30	36	42	48	54	60	66	72	78
$KRAS^{MUT};STK11^{WT}$	398	296	223	170	129	92	59	34	14	9	5	5	2	0
$KRAS^{MUT};STK11^{MUT}$	138	69	47	34	23	18	11	10	7	5	4	1	1	0

Ricciuti B et al., JTO, 2021



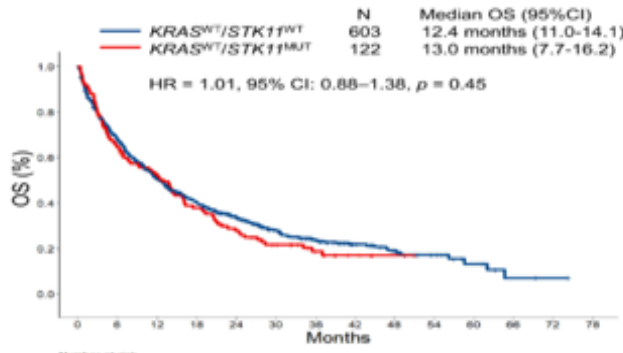
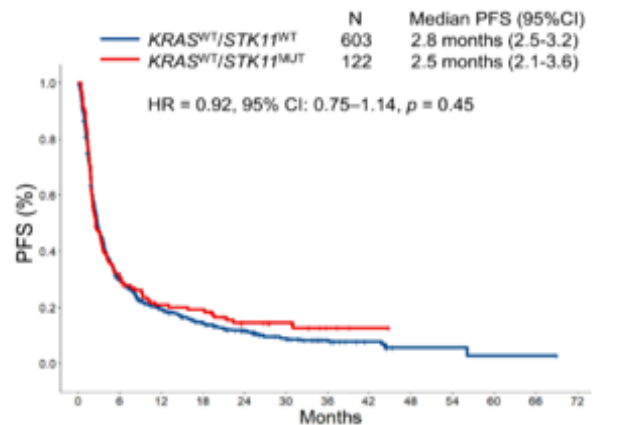
Group	0	6	12	18	24	30	36	42	48	54	60	66	72
$KRAS^{MUT};KEAP1^{WT}$	376	100	68	53	40	32	19	9	6	4	2	1	0
$KRAS^{MUT};KEAP1^{MUT}$	101	17	13	6	2	2	2	2	1	1	0	0	0



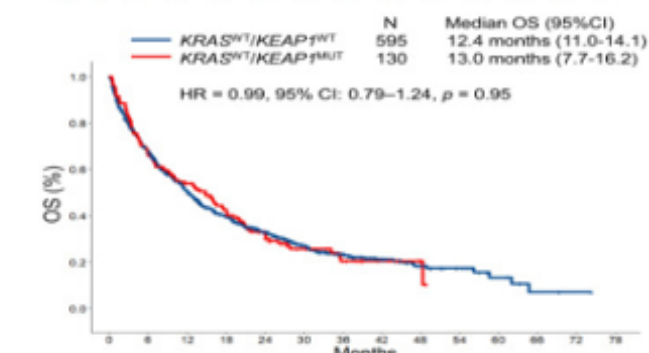
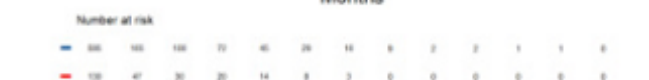
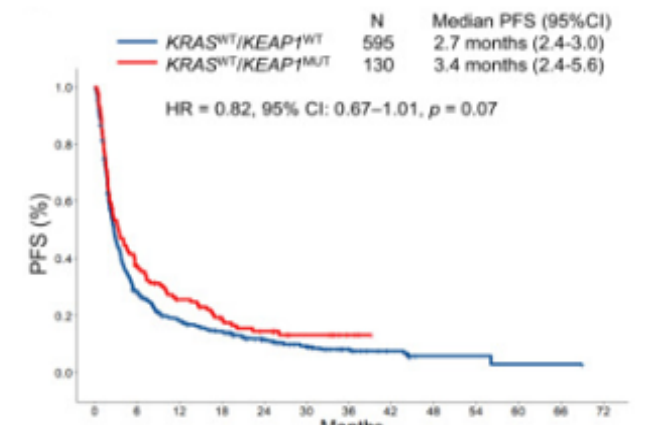
Group	0	6	12	18	24	30	36	42	48	54	60	66	72	78
$KRAS^{MUT};KEAP1^{WT}$	376	276	210	162	116	83	52	32	18	10	7	4	2	0
$KRAS^{MUT};KEAP1^{MUT}$	101	44	28	17	15	12	7	5	2	2	1	1	1	0

Potential modifiers of the impact of *STK11* and *KEAP1* alterations on IO outcomes : *KRAS* status

***KRAS*^{WT}/*STK11*^{WT} vs *KRAS*^{WT}/*STK11*^{MUT}**



***KRAS*^{WT}/*KEAP1*^{WT} vs *KRAS*^{WT}/*KEAP1*^{MUT}**

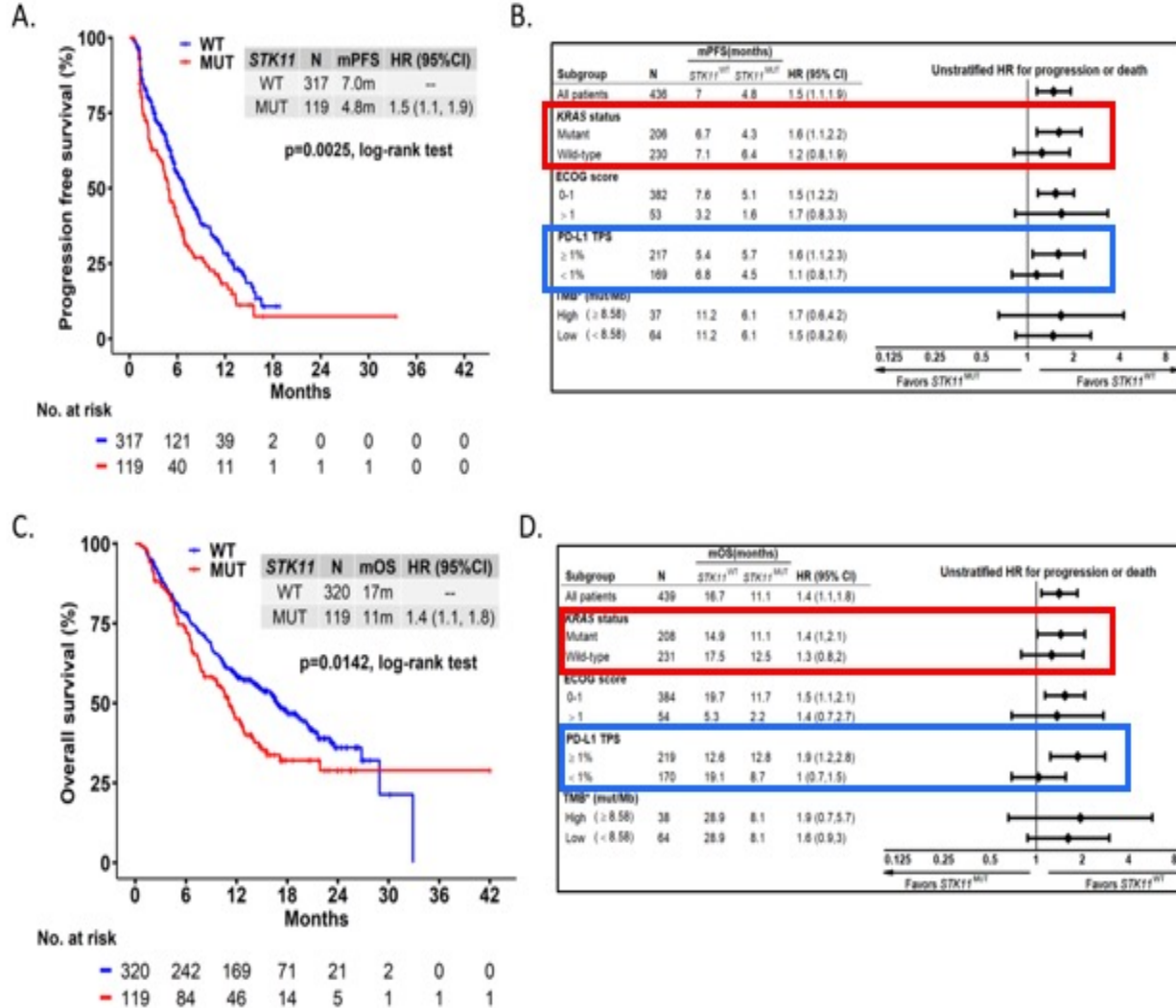


Onco-genotype	mPFS	mOS
<i>KRAS</i> ^{MUT} ; <i>STK11</i> ^{WT}	4.8m	17.3m
<i>KRAS</i> ^{MUT} ; <i>STK11</i> ^{MUT}	2.0m	6.2m
<i>KRAS</i> ^{WT} ; <i>STK11</i> ^{WT}	2.8m	12.4m
<i>KRAS</i> ^{WT} ; <i>STK11</i> ^{MUT}	2.5m	13.0m
<i>KRAS</i> ^{MUT} ; <i>KEAP1</i> ^{WT}	4.6m	18.4m
<i>KRAS</i> ^{MUT} ; <i>KEAP1</i> ^{MUT}	1.8m	4.8m
<i>KRAS</i> ^{WT} ; <i>KEAP1</i> ^{WT}	2.7m	12.4m
<i>KRAS</i> ^{WT} ; <i>KEAP1</i> ^{MUT}	3.4m	13.0m

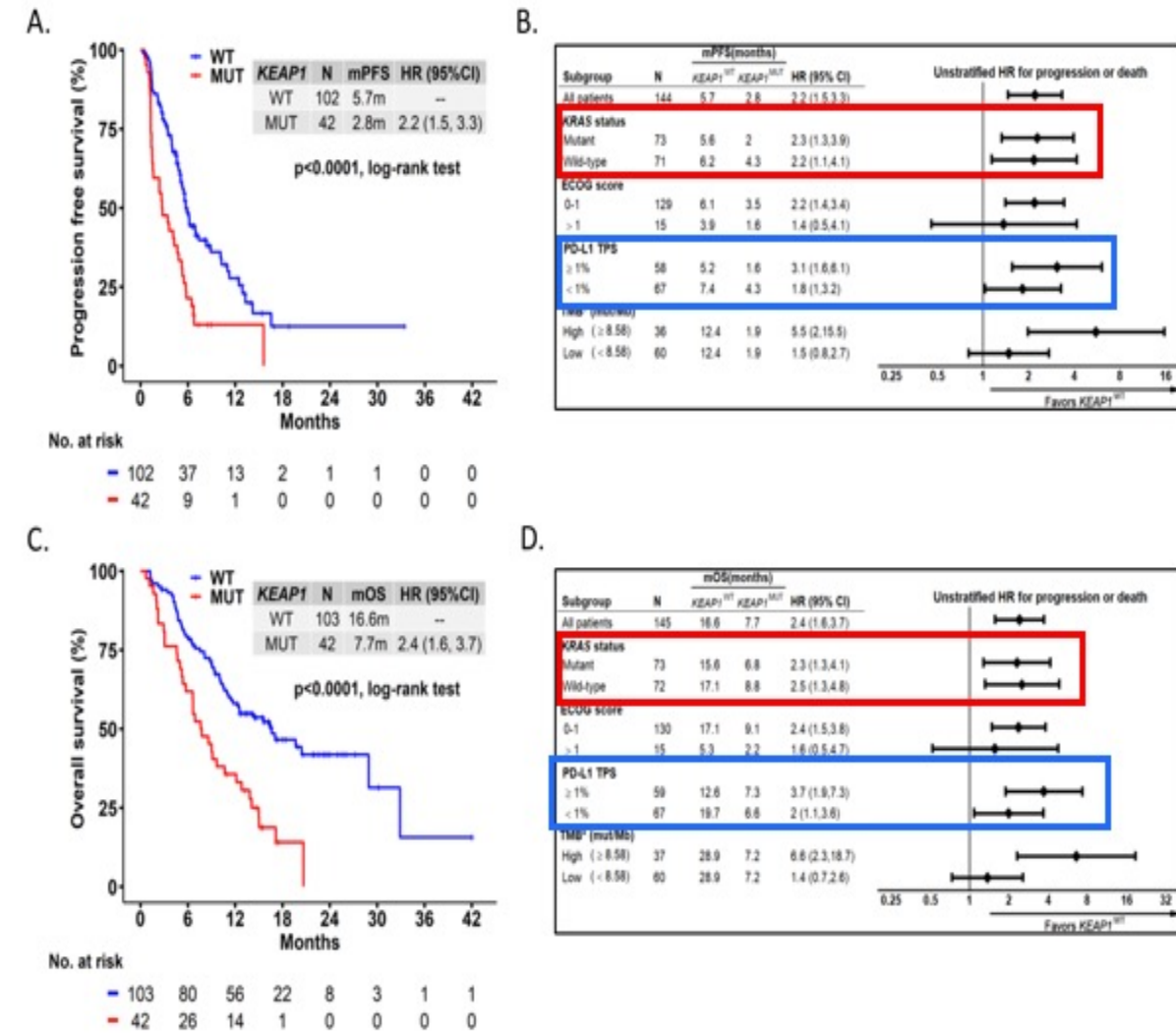
Ricciuti B et al., *JTO*, 2021

STK11 and KEAP1 alterations and clinical outcomes with first-line PCP chemoIO (platinum, pemetrexed, pembrolizumab)

STK11



KEAP1



STK11 and KEAP1 alterations and clinical outcomes with 1st line chemolO

Forest-plot for progression-free survival (PFS)

Mutation status	N	HR (95% CI) PFS	P-value
KRAS ^{WT}	560	reference	
KRAS ^{MUT}	351	1.10 (0.95-1.28)	0.21
TP53 ^{WT}	370	reference	
TP53 ^{MUT}	412	0.83 (0.71-0.97)	0.020
STK11 ^{WT}	516	reference	
STK11 ^{MUT}	191	1.46 (1.22-1.76)	<0.001
KEAP1 ^{WT}	518	reference	
KEAP1 ^{MUT}	189	1.53 (1.28-1.84)	<0.001
SMARCA4 ^{WT}	593	reference	
SMARCA4 ^{MUT}	114	1.62 (1.30-2.02)	<0.001
KRAS ^{WT} TP53 ^{WT}	182	reference	
KRAS ^{WT} TP53 ^{MUT}	287	0.88 (0.71-1.08)	0.21
KRAS ^{WT} STK11 ^{WT}	322	reference	
KRAS ^{WT} STK11 ^{MUT}	99	1.19 (0.93-1.54)	0.16
KRAS ^{WT} KEAP1 ^{WT}	327	reference	
KRAS ^{WT} KEAP1 ^{MUT}	104	1.37 (1.07-1.75)	0.01
KRAS ^{WT} SMARCA4 ^{WT}	361	reference	
KRAS ^{WT} SMARCA4 ^{MUT}	70	1.36 (1.03-1.81)	<0.001
KRAS ^{MUT} TP53 ^{WT}	188	reference	
KRAS ^{MUT} TP53 ^{MUT}	125	0.75 (0.58-0.97)	0.027
KRAS ^{MUT} STK11 ^{WT}	184	reference	
KRAS ^{MUT} STK11 ^{MUT}	92	1.92 (1.46-2.53)	<0.001
KRAS ^{MUT} KEAP1 ^{WT}	191	reference	
KRAS ^{MUT} KEAP1 ^{MUT}	85	1.82 (1.38-2.41)	<0.001
KRAS ^{MUT} SMARCA4 ^{WT}	232	reference	
KRAS ^{MUT} SMARCA4 ^{MUT}	44	2.39 (1.67-3.42)	<0.001

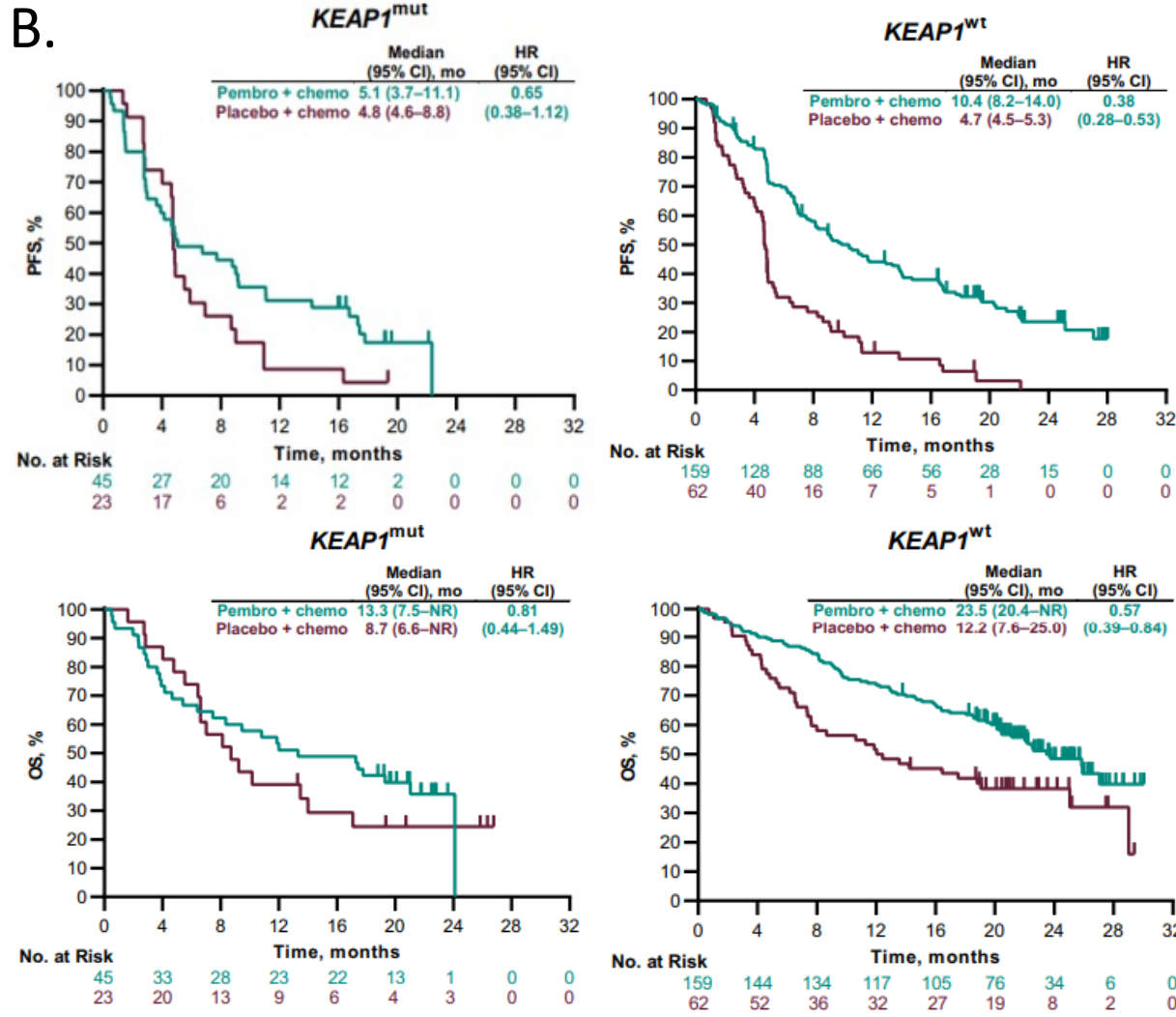
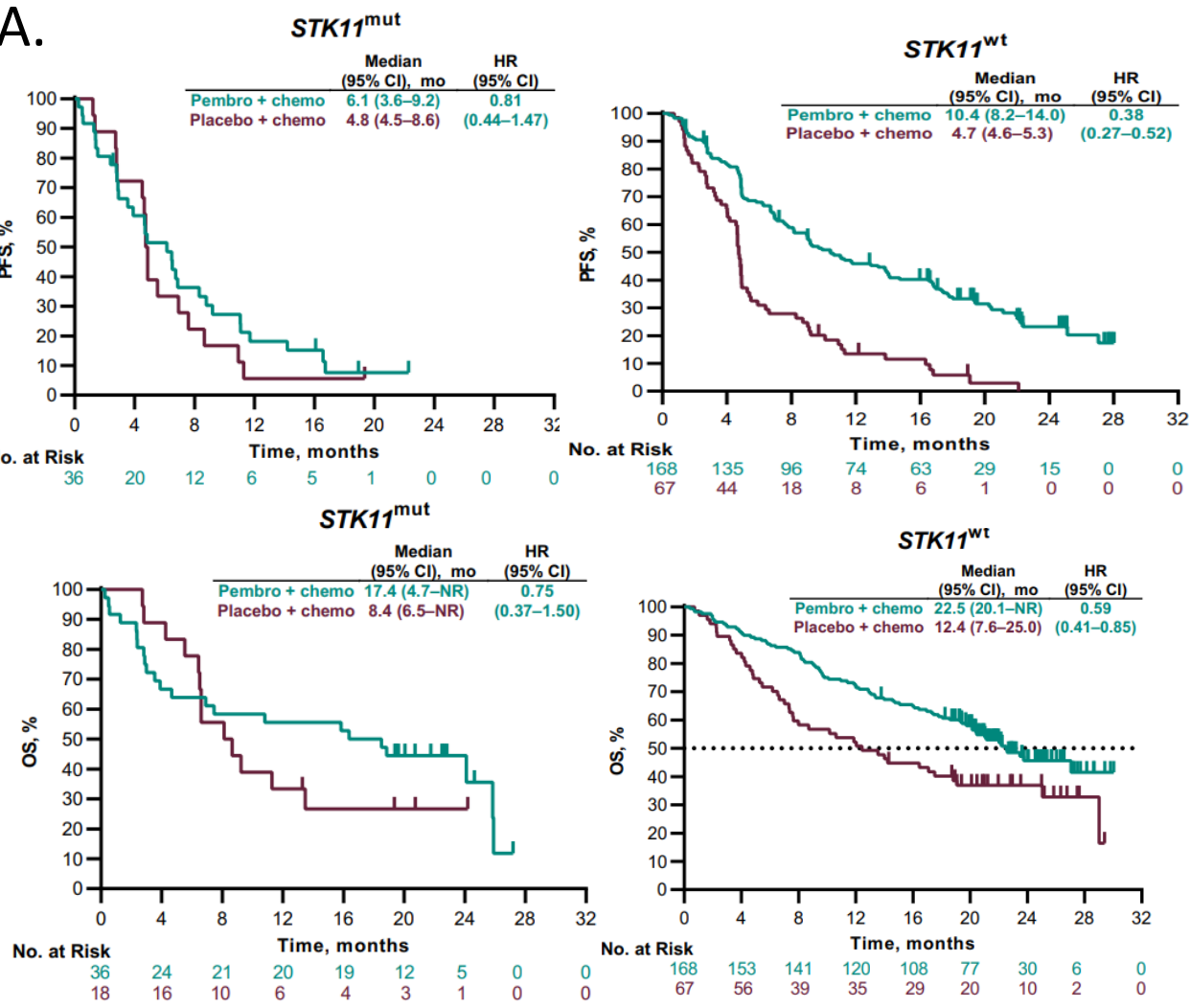
0.25 0.50 1.0 1.5 2.0 2.5
 ← Better PFS Worse PFS →

Forest-plot for overall survival (OS)

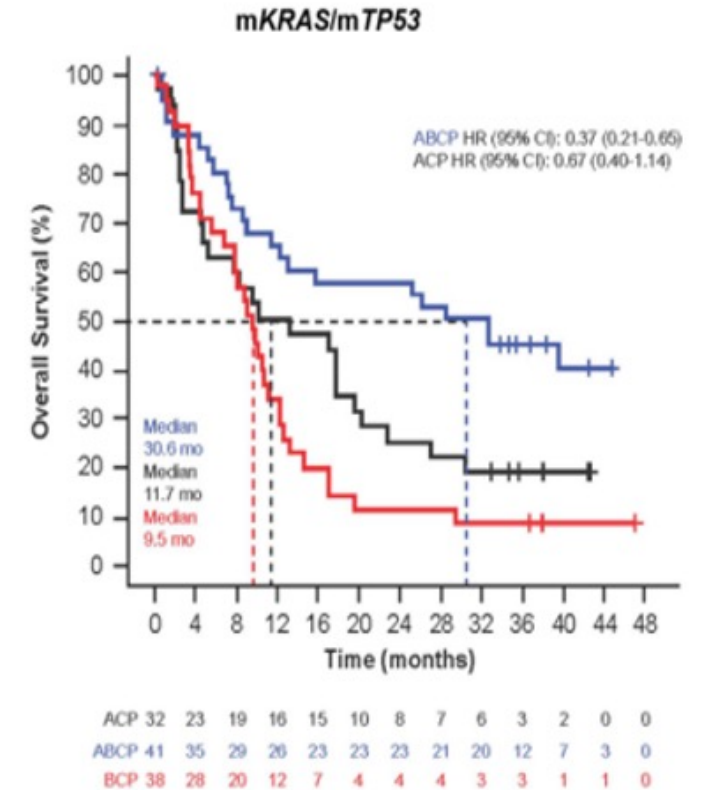
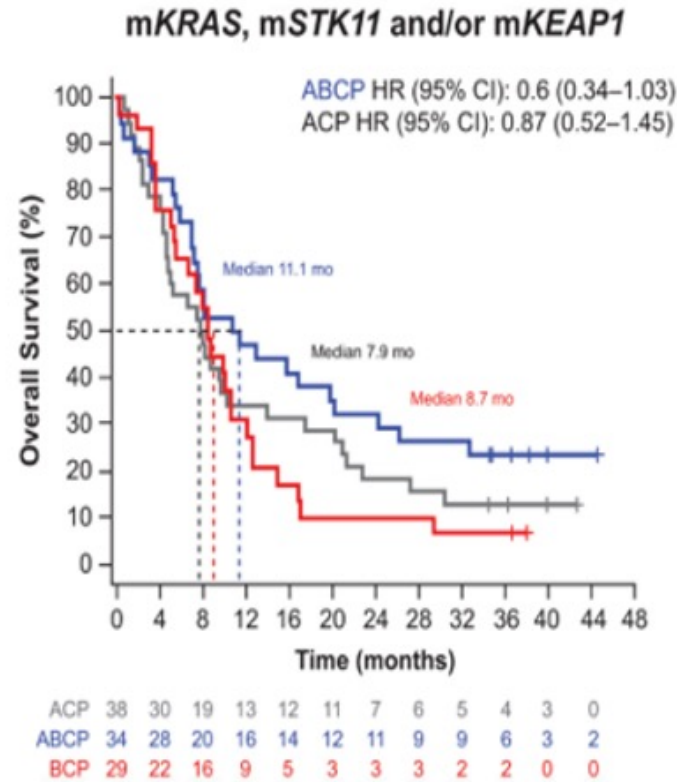
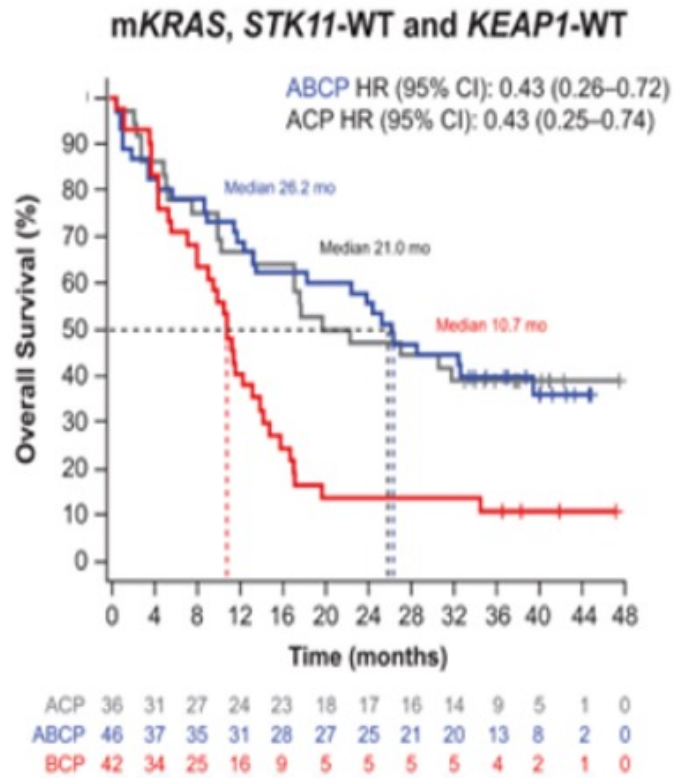
Mutation status	N	HR (95% CI) OS	P-value
KRAS ^{WT}	560	reference	
KRAS ^{MUT}	351	1.07 (0.93-1.28)	0.42
TP53 ^{WT}	370	reference	
TP53 ^{MUT}	412	0.87 (0.73-1.04)	0.13
STK11 ^{WT}	516	reference	
STK11 ^{MUT}	191	1.36 (1.10-1.67)	0.004
KEAP1 ^{WT}	518	reference	
KEAP1 ^{MUT}	189	1.71 (1.40-2.10)	<0.001
SMARCA4 ^{WT}	593	reference	
SMARCA4 ^{MUT}	114	1.70 (1.33-2.17)	<0.001
KRAS ^{WT} TP53 ^{WT}	182	reference	
KRAS ^{WT} TP53 ^{MUT}	287	0.89 (0.70-1.13)	0.35
KRAS ^{WT} STK11 ^{WT}	322	reference	
KRAS ^{WT} STK11 ^{MUT}	99	1.14 (0.85-1.53)	0.38
KRAS ^{WT} KEAP1 ^{WT}	327	reference	
KRAS ^{WT} KEAP1 ^{MUT}	104	1.64 (1.25-2.17)	<0.001
KRAS ^{WT} SMARCA4 ^{WT}	361	reference	
KRAS ^{WT} SMARCA4 ^{MUT}	70	1.36 (0.98-1.88)	0.06
KRAS ^{MUT} TP53 ^{WT}	188	reference	
KRAS ^{MUT} TP53 ^{MUT}	125	0.85 (0.63-1.13)	0.27
KRAS ^{MUT} STK11 ^{WT}	184	reference	
KRAS ^{MUT} STK11 ^{MUT}	92	1.66 (1.22-2.26)	0.001
KRAS ^{MUT} KEAP1 ^{WT}	191	reference	
KRAS ^{MUT} KEAP1 ^{MUT}	85	1.78 (1.31-2.43)	<0.001
KRAS ^{MUT} SMARCA4 ^{WT}	232	reference	
KRAS ^{MUT} SMARCA4 ^{MUT}	44	2.52 (1.72-3.68)	<0.001

0.25 0.50 1.0 1.5 2.0 2.5
 ← Better OS Worse OS →

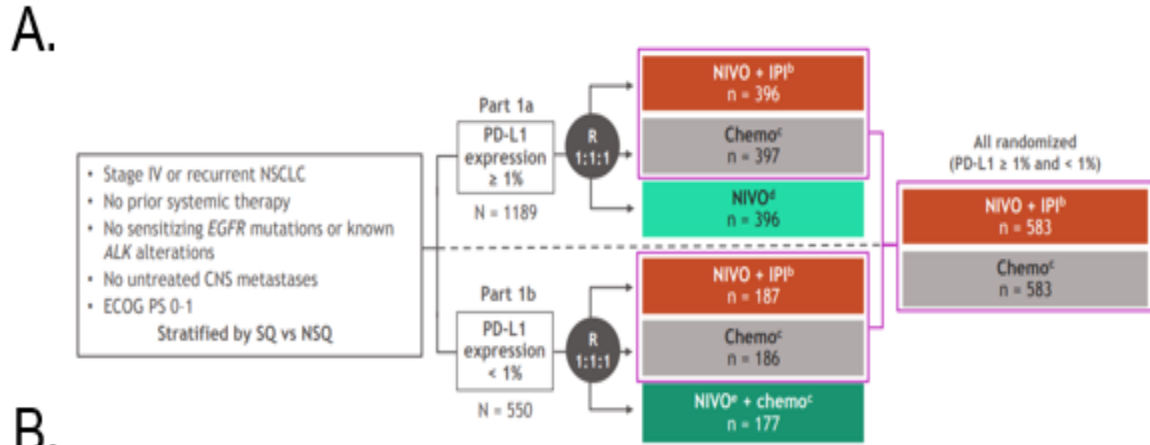
Reduced benefit from the addition of pembrolizumab to platinum doublet chemotherapy in patients with *STK11* and *KEAP1*-mutant NSCLC in KEYNOTE-189



Clinical outcomes in *KRAS* co-mutational subgroups in IMpower150



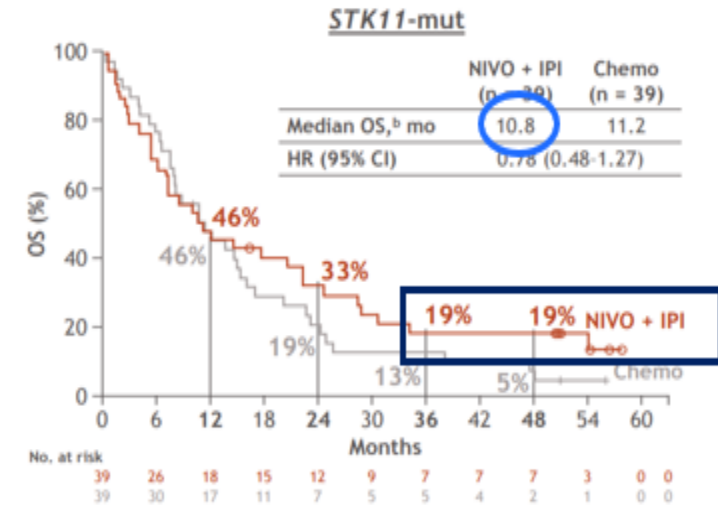
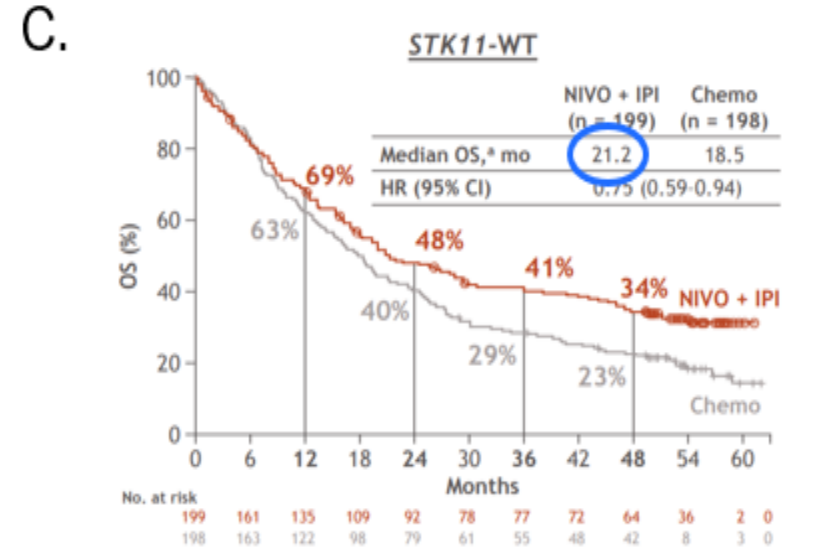
STK11 and KEAP1 alterations and clinical outcomes with ipi/nivo in Part 1 of CheckMate 227



B.

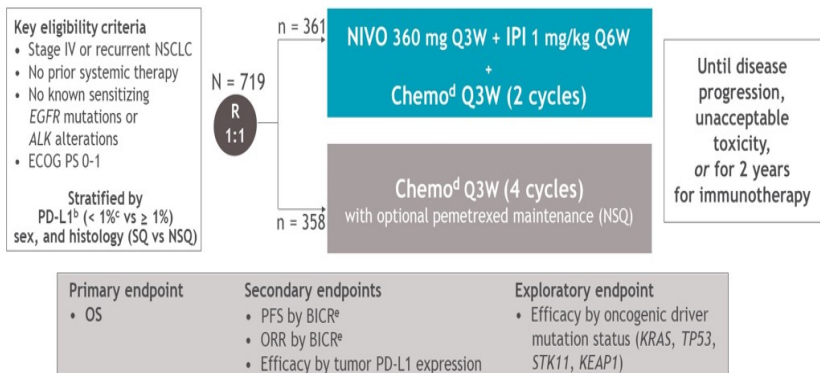
Subgroup, n ^b	4-y PFS rate, %		Median PFS, mo		Unstratified HR	Unstratified HR (95% CI)
	NIVO + IPI	Chemo	NIVO + IPI	Chemo		
NSQ (n = 419, 419)	14	3	5.2	5.6	0.82	
Mut-eval (n = 238, 237)	14	3	5.6	5.6	0.76	
KRAS-WT (n = 150, 162)	19	6	5.6	5.6	0.75	
KRAS-mut (n = 88, 75)	17	2	5.4	5.8	0.78	
TP53-WT (n = 111, 106)	10	5	5.4	5.6	0.88	
TP53-mut (n = 127, 131)	24	7	5.8	6.6	0.69	
STK11-WT (n = 199, 198)	19	6	8.1	6.1	0.72	
STK11-mut (n = 39, 39)	13	0	2.8	4.3	1.04	
KEAP1-WT (n = 218, 219)	16	6	5.5	5.8	0.83	
KEAP1-mut (n = 20, 18)	41	0	11.1	2.9	0.25	

KEAP1^{MUT} (N=38)
Ipi/Nivo: mOS 24.4m
Chemo: mOS 8.9m



Clinical outcomes with the CheckMate 9LA regimen in *STK11*-mutant NSCLC

A.



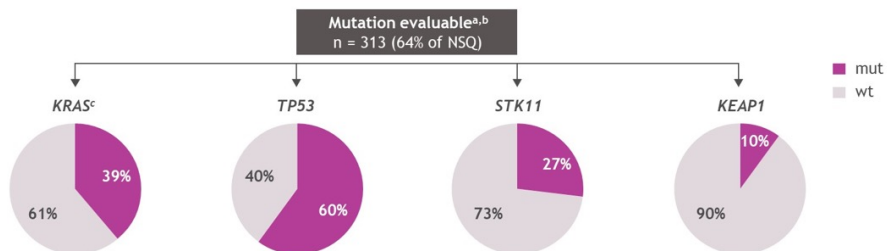
Database lock: February 15, 2022; minimum/median follow-up for OS: 36.1/42.6 months.

Reprinted from *Lancet Oncology*, 22, Paz-Ares L, et al, First-line nivolumab plus ipilimumab combined with two cycles of chemotherapy in patients with non-small-cell lung cancer (CheckMate 9LA): an international, randomised, open-label, phase 3 trial, 190-211, Copyright 2021, with permission from Elsevier.

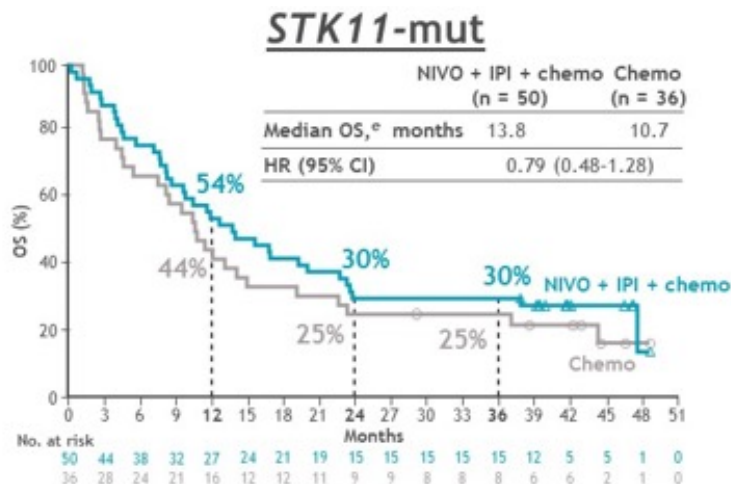
^aIC103215706; ^bDetermined by the PD-L1 IHC 28-8 pharmDx assay (Dako); ^cPatients unevaluable for PD-L1 were stratified to PD-L1 < 1% and capped to 10% of all randomized patients; ^dNSQ: pemetrexed + cisplatin or carboplatin; SQ: paclitaxel + carboplatin; ^eHierarchically statistically tested.

1. Paz-Ares L, et al. *Lancet Oncol* 2021;22:190-211; 2. Reck M, et al. *ESMO Open* 2021;6:100273.

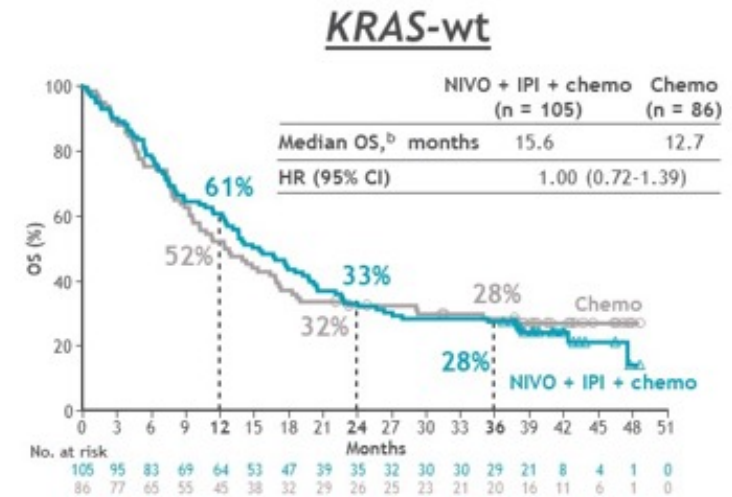
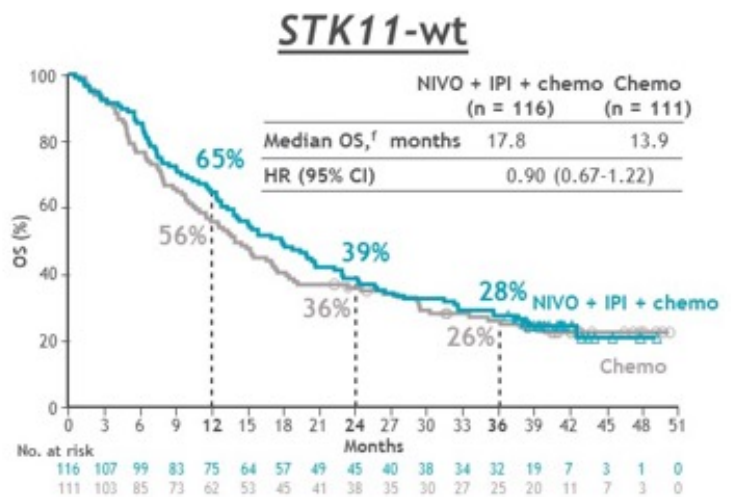
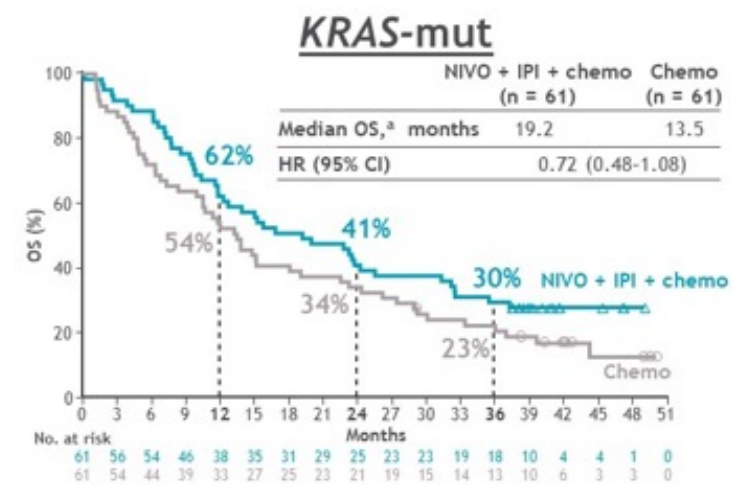
B.



C.



D.



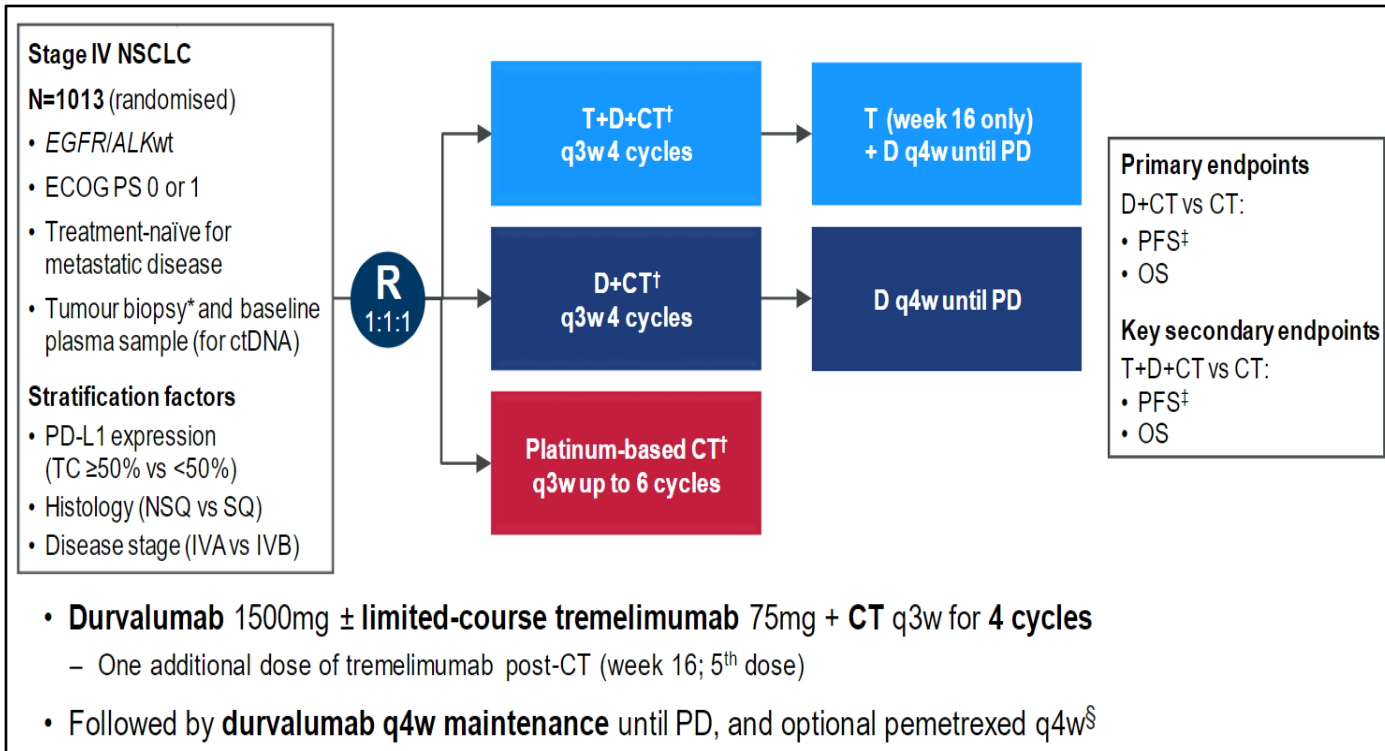
• Similar trend of OS benefit was seen with NIVO + IPI + chemo vs chemo in *KRAS* G12C-mut (n = 50) and *KEAP1*-mut (n = 32) subgroups

Database lock: February 15, 2022; minimum follow-up: 36.1 months.

^a95% CI, 11.9-25.5 (NIVO + IPI + chemo) and 10.0-19.1 (chemo); ^b95% CI, 12.3-19.9 (NIVO + IPI + chemo) and 9.5-17.0 (chemo); ^c95% CI, 12.6-22.7 (NIVO + IPI + chemo) and 9.5-15.4 (chemo); ^d95% CI, 10.4-22.9 (NIVO + IPI + chemo) and 9.5-23.3 (chemo); ^e95% CI, 8.6-22.7 (NIVO + IPI + chemo) and 5.4-14.9 (chemo); ^f95% CI, 13.2-22.8 (NIVO + IPI + chemo) and 10.6-17.4 (chemo).

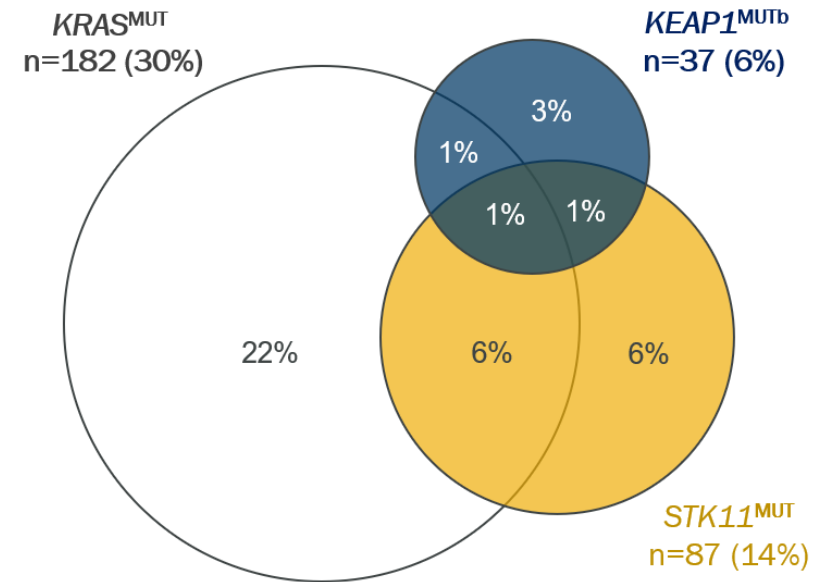
POSEIDON Study of Durvalumab+-Tremelimumab+Chemo for the 1st line Treatment of Metastatic NSCLC

A.



B.

Mutation-Evaluable Population^a
(n=612; 96% of randomized patients with NSQ histology)

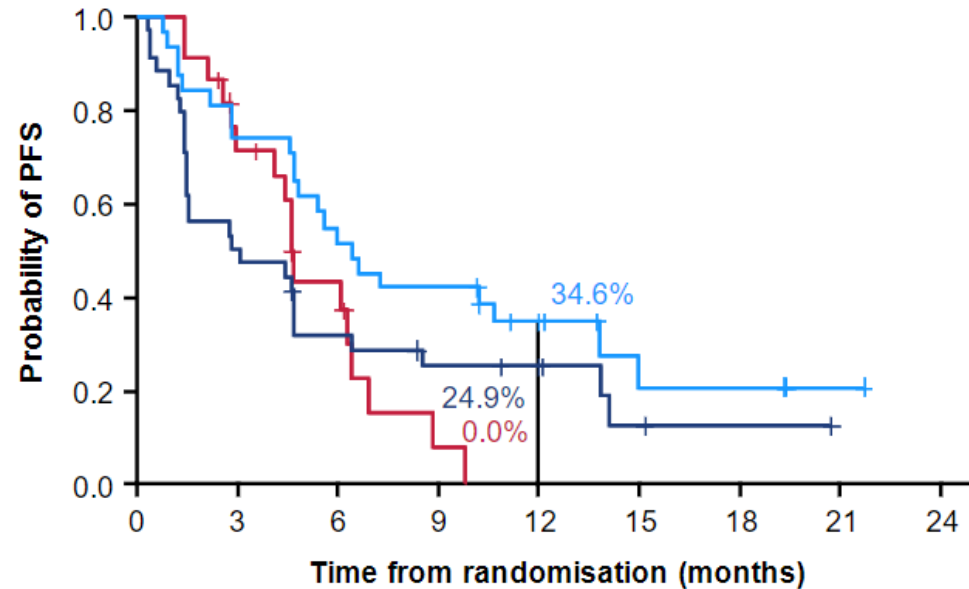


PFS and ORR with the POSEIDON regimen (D+T+chemo) in *STK11*-mutant NSCLC

A.

PFS

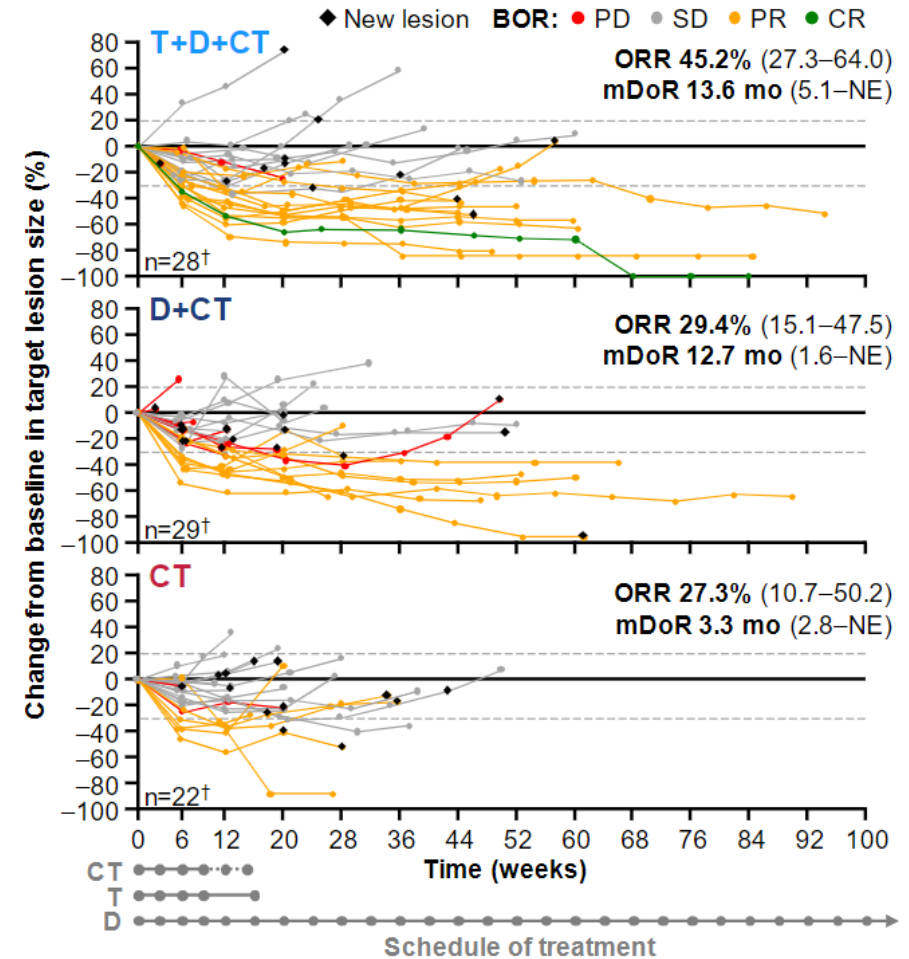
	T+D+CT	D+CT	CT
Events, n/N	22/31	27/34	17/22
mPFS, mo (95% CI)	6.4 (4.7–13.8)	2.9 (1.4–4.7)	4.6 (2.9–6.4)
HR* (95% CI)	0.47 (0.23–0.93)	1.02 (0.55–1.93)	–



No. at risk

	0	3	6	9	12	15	18	21	24
T+D+CT	31	23	16	13	7	3	3	1	0
D+CT	34	17	10	7	5	2	1	0	0
CT	22	14	7	1	0	0	0	0	0

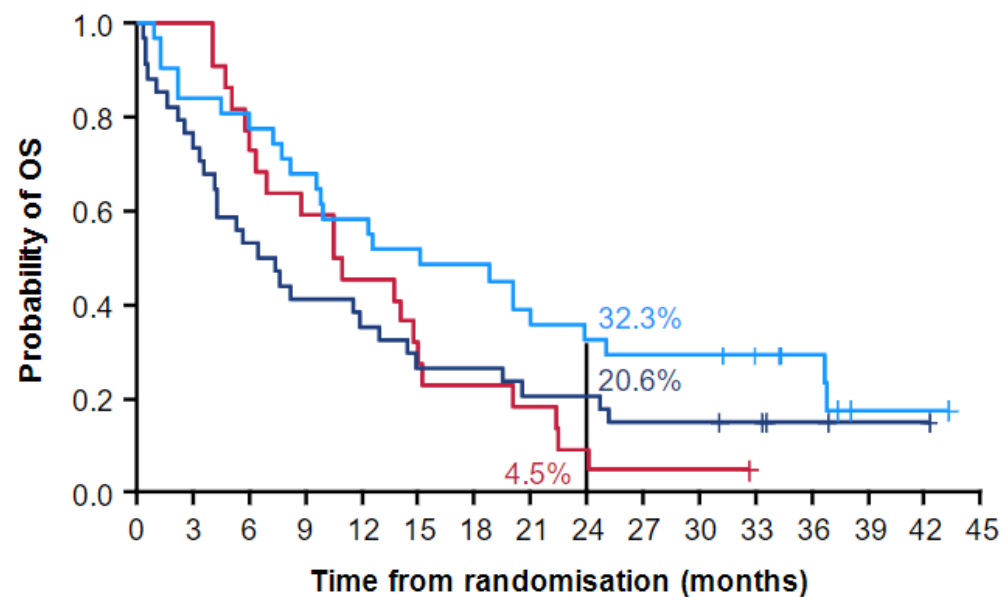
B.



OS with the POSEIDON regimen (D+T+chemo) in *STK11*-mutant NSCLC

*STK11*m

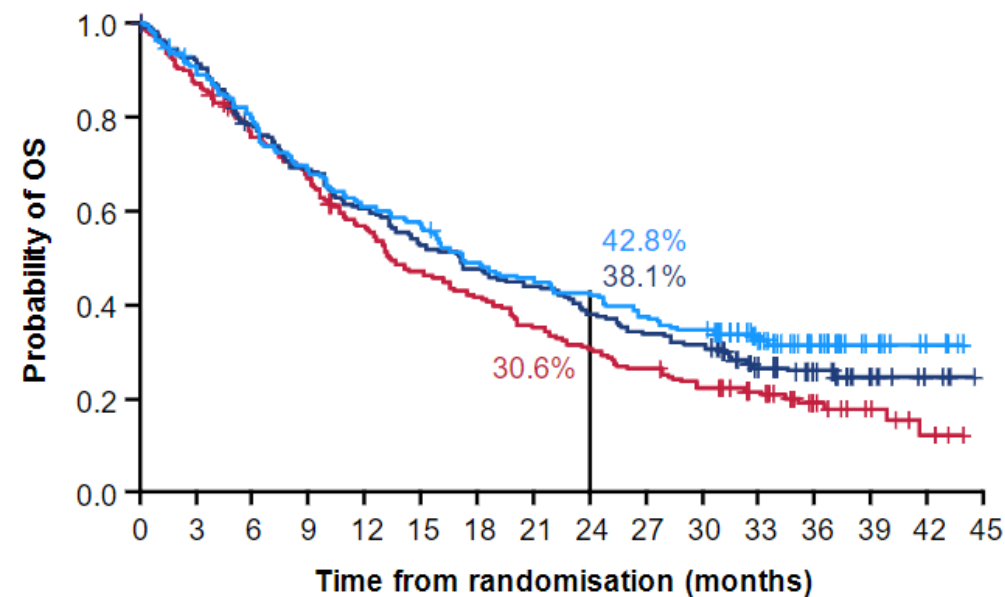
	T+D+CT	D+CT	CT
Events, n/N	24/31	29/34	21/22
mOS, mo (95% CI)	15.0 (8.2–23.8)	6.9 (3.6–12.9)	10.7 (6.0–14.9)
HR* (95% CI)	0.56 (0.30–1.03)	1.03 (0.59–1.84)	–



No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45
T+D+CT	31	26	24	21	18	15	15	11	10	9	9	7	5	1	1	0
D+CT	34	26	18	14	12	9	9	7	7	5	5	4	2	1	0	0
CT	22	22	16	13	10	6	5	4	1	1	1	0	0	0	0	0

*STK11*wt

	T+D+CT	D+CT	CT
Events, n/N	118/177	123/169	141/179
mOS, mo (95% CI)	17.2 (14.9–22.1)	17.1 (13.3–22.3)	13.4 (11.5–17.5)
HR* (95% CI)	0.73 (0.57–0.93)	0.81 (0.64–1.04)	–



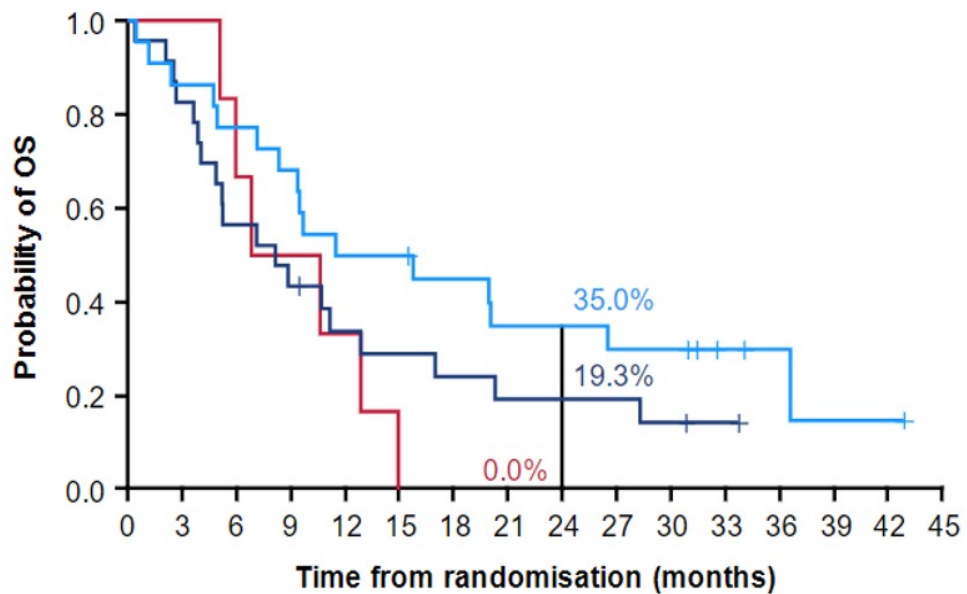
No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45
T+D+CT	177	159	140	120	107	100	85	79	74	65	60	40	25	11	5	0
D+CT	169	155	130	114	100	87	79	73	63	56	52	33	23	10	4	0
CT	179	154	131	116	97	80	71	60	52	45	37	29	15	8	4	0

OS and ORR with the POSEIDON regimen (D+T+chemo) in *KEAP1*-mutant NSCLC

A.

*KEAP1*m

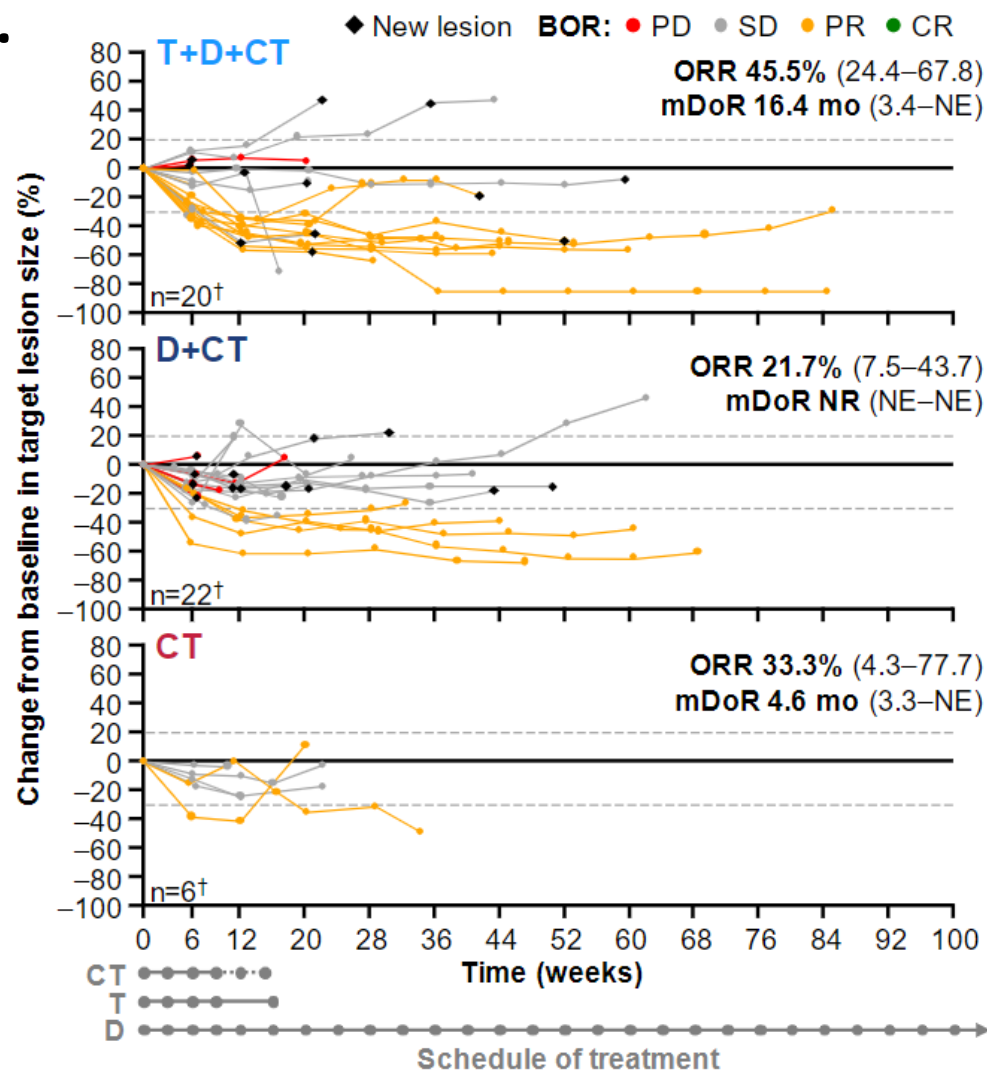
	T+D+CT	D+CT	CT
Events, n/N	16/22	19/23	6/6
mOS, mo (95% CI)	13.7 (7.2–26.5)	8.1 (4.0–12.9)	8.7 (5.1–NE)
HR* (95% CI)	0.43 (0.16–1.25)	0.77 (0.31–2.15)	–



No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45
T+D+CT	22	19	17	15	11	11	9	7	7	6	6	3	2	1	1	0
D+CT	23	19	13	10	7	6	5	4	4	4	3	1	0	0	0	0
CT	6	6	4	3	2	0	0	0	0	0	0	0	0	0	0	0

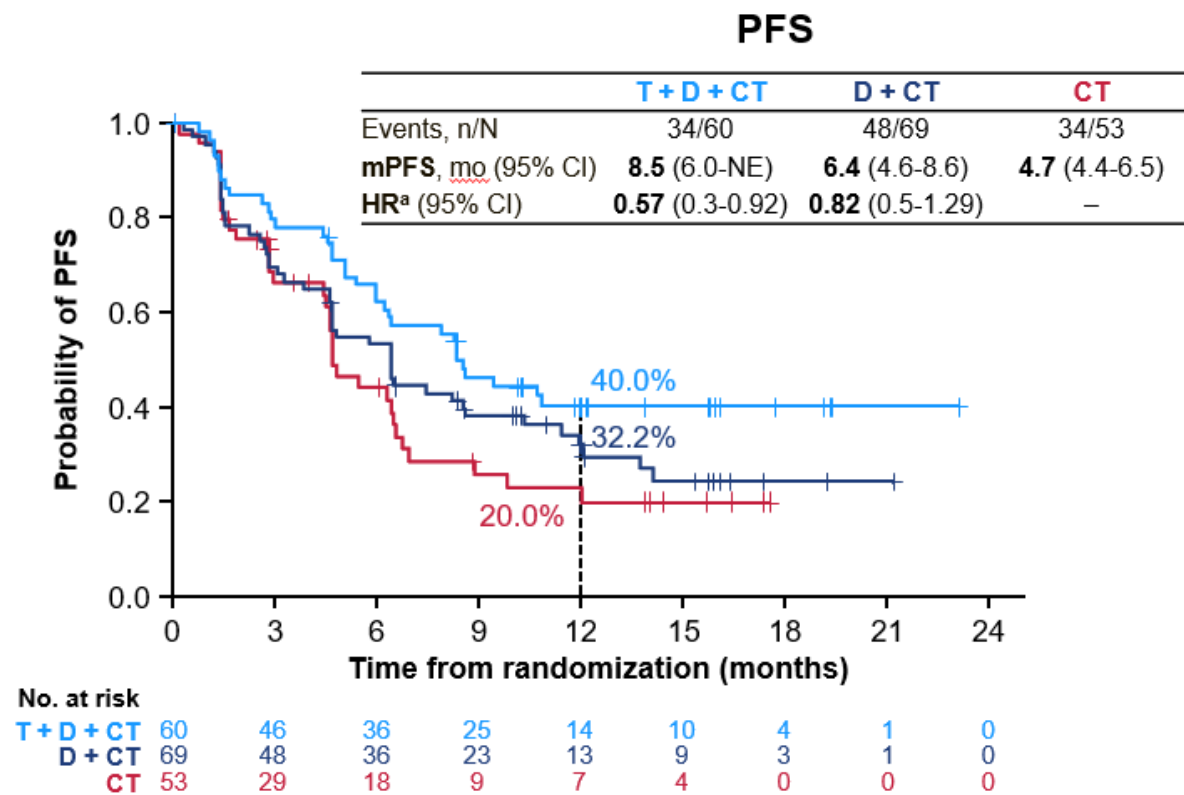
HR (95% CI) vs CT in NSQ *KEAP1*m was 0.33 (0.10–1.15) with T+D+CT and 0.67 (0.23–2.17) with D+CT

B.

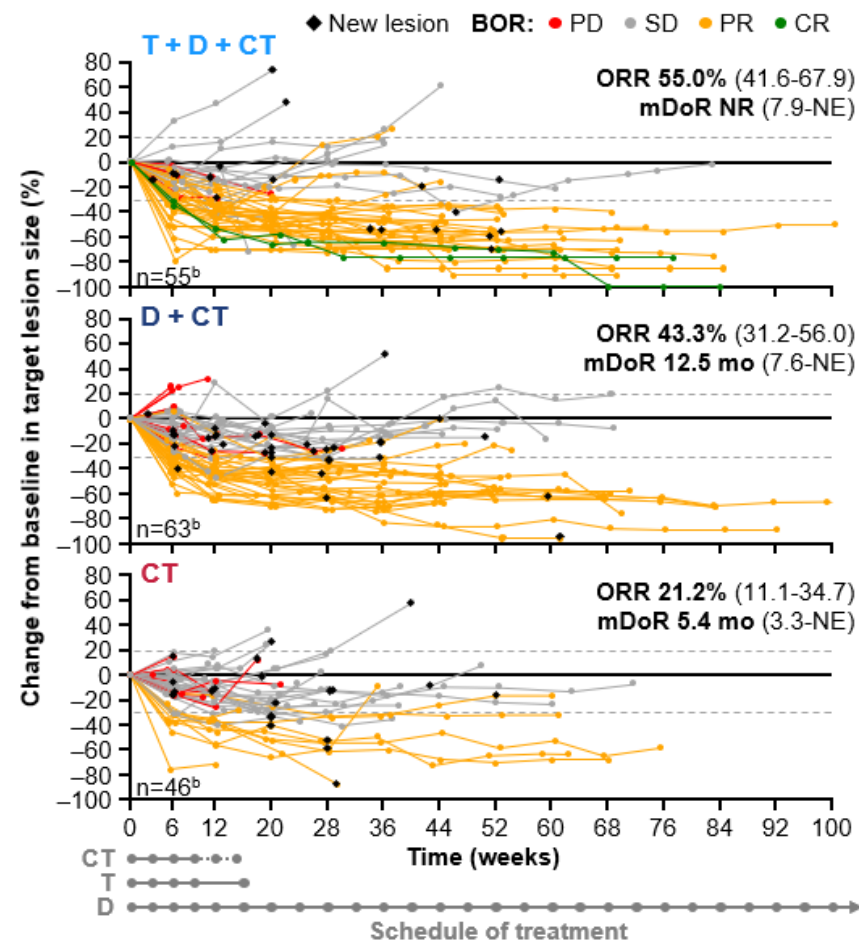


PFS and ORR with the POSEIDON regimen (D+T+chemo) in *KRAS*-mutant NSCLC (4-year update)

A.



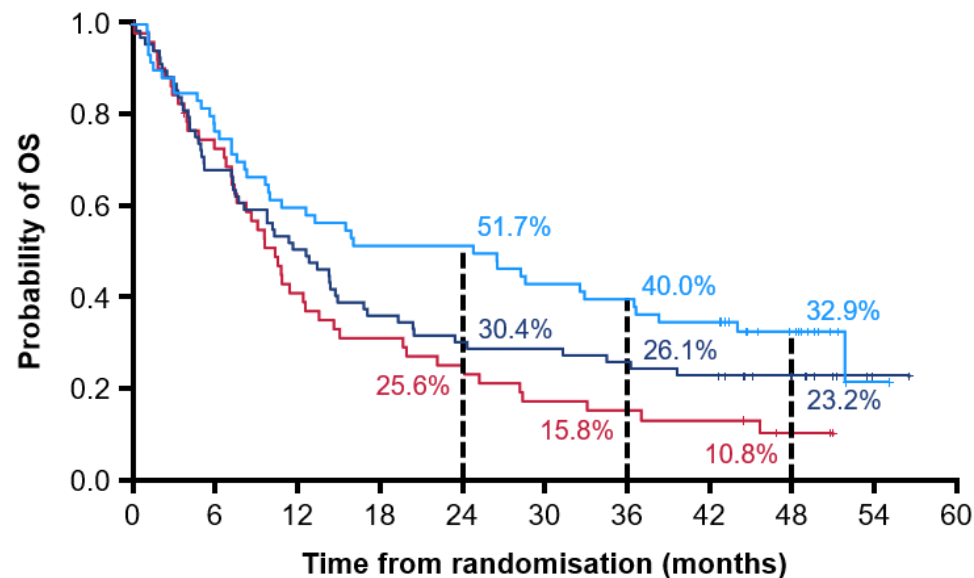
B.



OS with the POSEIDON regimen (D+T+chemo) in *KRAS*-mutant and wild-type NSCLC (4-year update)

*KRAS*_m

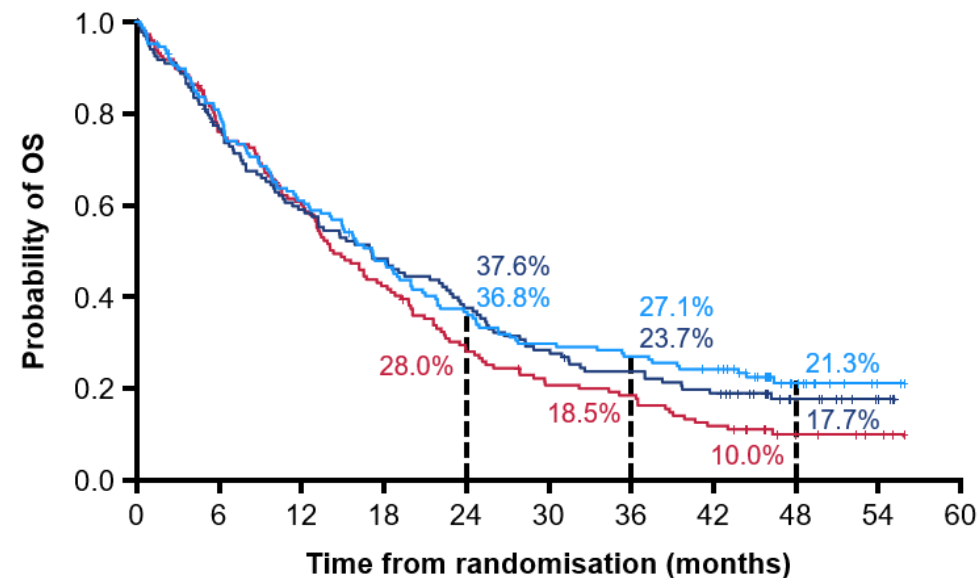
	T+D+CT	D+CT	CT
Events, n/N	41/60	53/69	45/53
mOS, months (95% CI)	25.7 (9.9–36.7)	12.6 (7.5–16.9)	10.4 (7.5–13.6)
HR* (95% CI)	0.55 (0.36–0.85)	0.78 (0.52–1.16)	–



No. at risk	0	6	12	18	24	30	36	42	48	54	60
T+D+CT	60	46	36	31	31	26	24	21	12	1	0
D+CT	69	47	35	25	21	20	18	16	9	1	0
CT	53	37	21	16	13	9	7	6	2	0	0

*KRAS*_{wt}

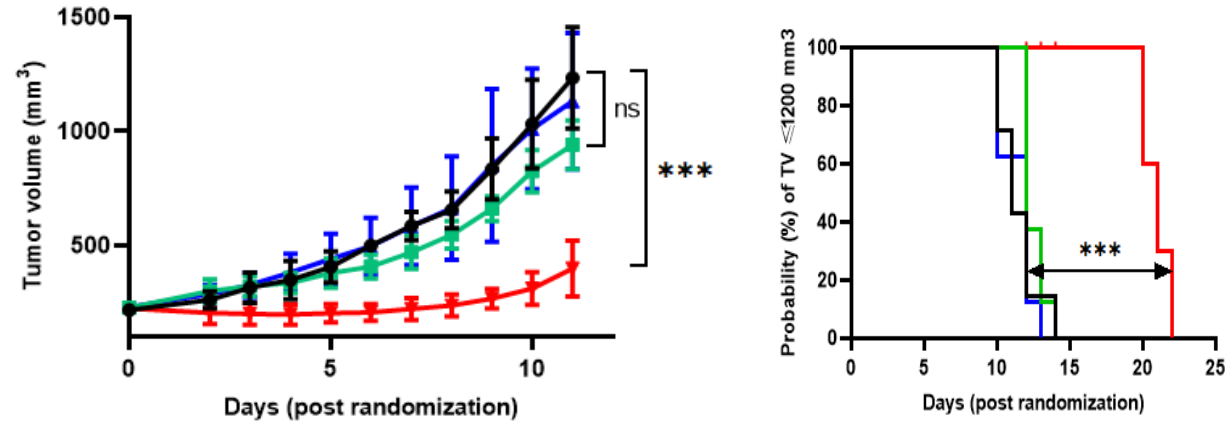
	T+D+CT	D+CT	CT
Events, n/N	113/148	107/134	127/148
mOS, months (95% CI)	17.1 (13.4–20.1)	17.1 (12.3–22.6)	14.4 (12.6–18.3)
HR* (95% CI)	0.78 (0.60–1.00)	0.83 (0.64–1.08)	–



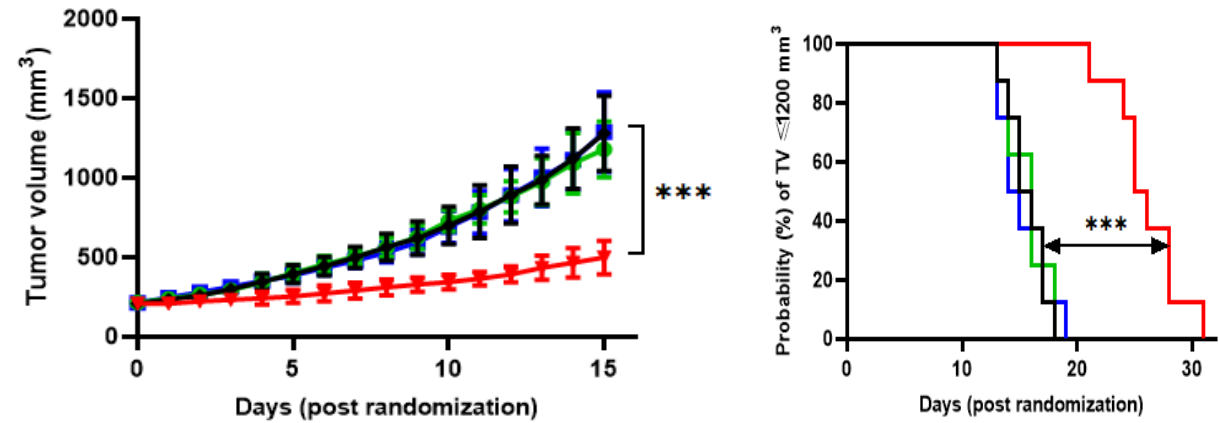
No. at risk	0	6	12	18	24	30	36	42	48	54	60
T+D+CT	148	118	89	69	53	43	39	34	12	2	0
D+CT	134	101	77	63	49	37	30	24	11	3	0
CT	148	110	86	60	39	28	25	16	7	4	0

Sensitivity of $Kras^{MUT};Stk11^{-/-}$ lung adenocarcinomas to dual anti-PD-1/anti-CTLA-4 ICB is recapitulated in syngeneic models

KL2 ($Kras^{G12C};Stk11^{-/-}$)



KL5 ($Kras^{G12C};Stk11^{-/-}$)



— IgG control — αPD-1 — αCTLA-4 — αPD-1 + αCTLA-4

Conclusions

- Co-mutations in key tumor suppressor genes – most prominently *STK11* and *KEAP1* shape the immune contexture of ns-NSCLC.
- *STK11* and *KEAP1* alterations frequently co-occur (and are both enriched in *KRAS*-mutant NSCLC) and loss of both TSGs promotes lung oncogenesis. However, their inactivation imparts both overlapping as well as distinct effects on the TIME.
- Only *STK11* alterations are associated with lack of/low PD-L1 expression on tumor cells.
- Somatic mutations in *KEAP1* and/or *STK11* identify difficult to treat subgroups of patients with mNSCLC that exhibit poor clinical outcomes with PD-(L)1 inhibitor – based chemo-immunotherapy (such as the KEYNOTE-189 regimen) or PD-(L)1 monotherapy, especially in patients harboring *KRAS*-mutant NSCLC.
- Loss of *KEAP1* and/or *STK11* may impart selective sensitivity to dual immune checkpoint blockade with anti-PD-(L)1+ anti-CTLA-4.
- Chemo-IO regimens that incorporate anti-CTLA-4 in addition to anti-PD-(L)1 (such as 9LA and POSEIDON) may represent a preferred approach in *STK11* and/or *KEAP1*-mutated NSCLC with good PS. Data from POSEIDON appear the most robust to date in this patient population.
- A randomized controlled clinical trial (TRITON) (POSEIDON regimen vs KEYNOTE 189) in patients with previously untreated metastatic NSCLC with *STK11*, *KEAP1* or *KRAS* alterations is under development to confirm findings from POSEIDON.
- *STK11*, *KEAP1* represent emerging biomarkers for selection of first-line regimens in advanced NSCLC.
- A number of novel therapeutic strategies to induce/re-invigorate effective anti-tumor immunity in *STK11* and/or *KEAP1*-mutant NSCLC are currently in development.

Thank you !