

# The Future of Combined Modality Therapy



# Considerations for Adjuvant Therapy

## PRO

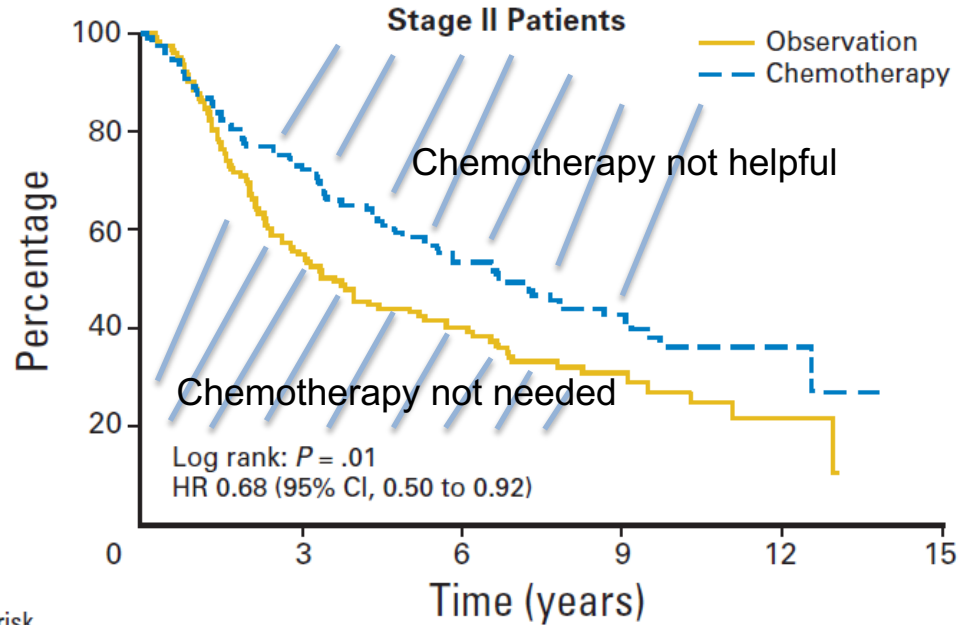
- Adjuvant is evidence-based standard of care for resected stage IB and II disease
- Tumor biomarkers can guide therapeutic decisions
- No delay of surgery
- No hilar or mediastinal fibrosis
- No risk of disease progression resulting in missed opportunity for curative surgery

## CON

- Poor tolerance and compliance with adjuvant protocols
- Longer treatment (4 cycles or much longer if TKI/IO)
- No intermediate endpoints
- Long follow up required for DFS or OS

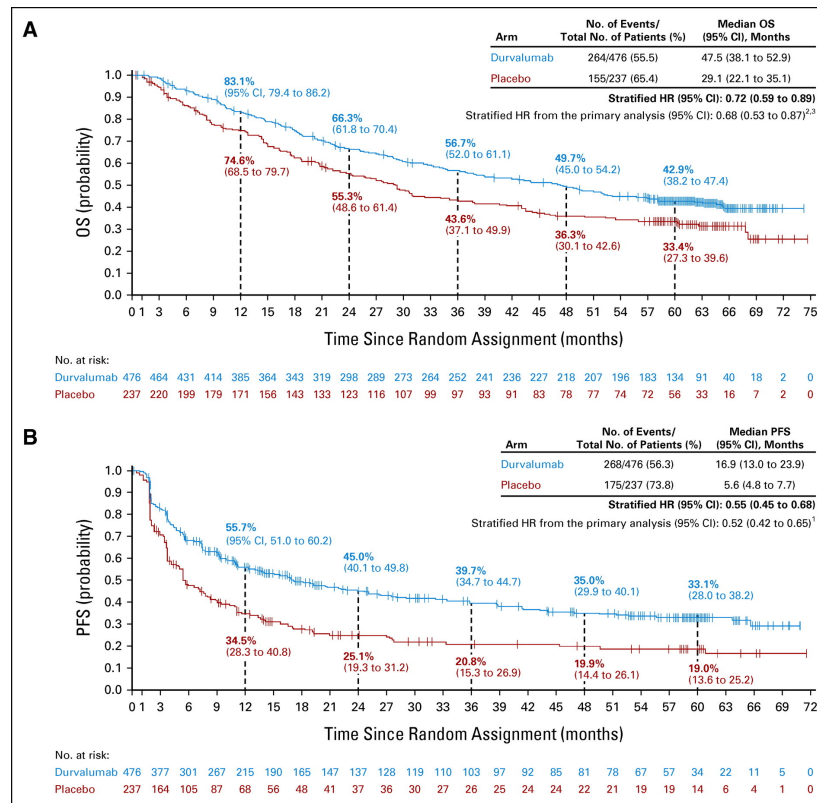
# JBR.10 – Overall Survival

**B**



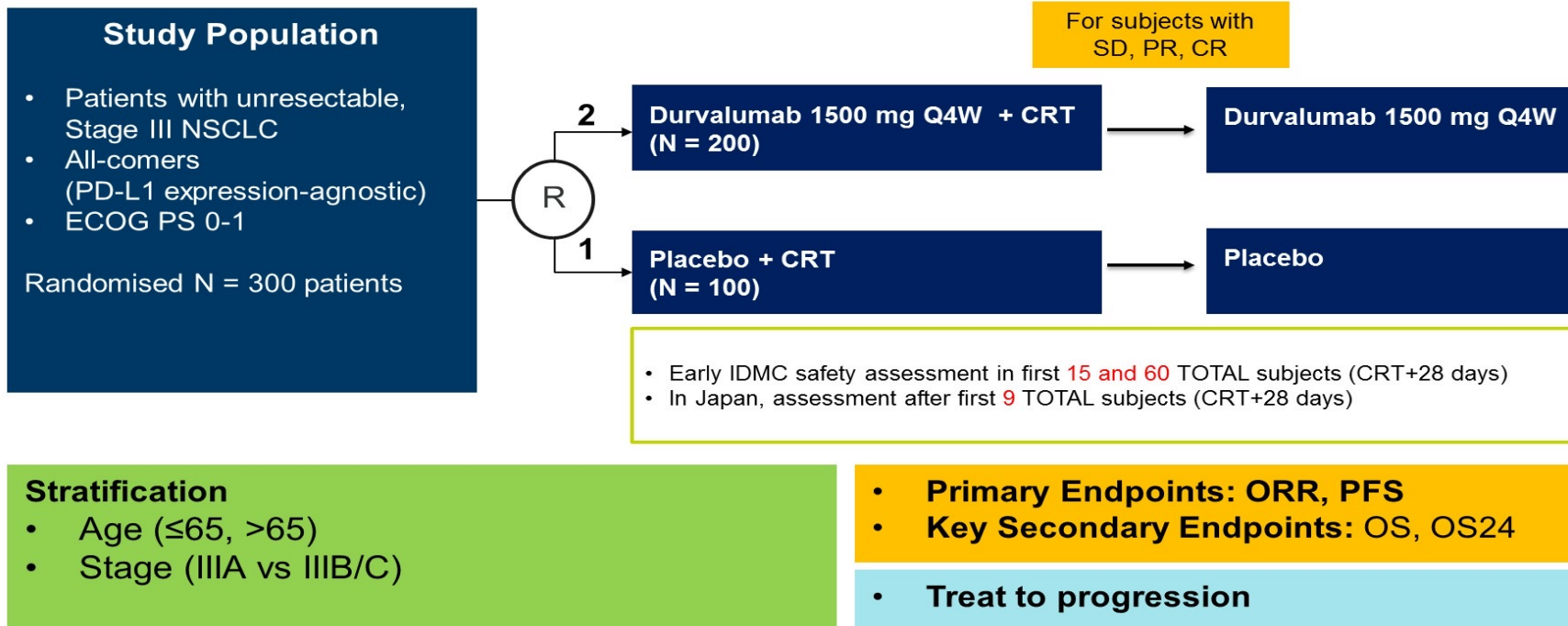
No. at risk						
Observation	132	72	48	18	4	0
Chemotherapy	131	95	65	31	6	0

# Updated OS and PFS in the intent-to-treat population





# PACIFIC-2: Study Design



Study Chair: Bradley



# Central Challenge to Curative Therapies

- Does current staging provide accurate enough information
- How do we know when enough is enough
- Can we achieve cure with less toxicity (duration, intensity, cost of treatment)
- Can biomarkers provide information in real time
- Can we design De-escalation Trials

# Curative Therapies need to be personalized

## Surrogate Endpoints:

- DFS (CTs, PET)
- PRO's
- (mPR), pCR
- Systemic Markers of Minimal Residual Disease

# Biomarkers in (Neo)adjuvant Setting

## Specific Tumor Biomarker (mutation):

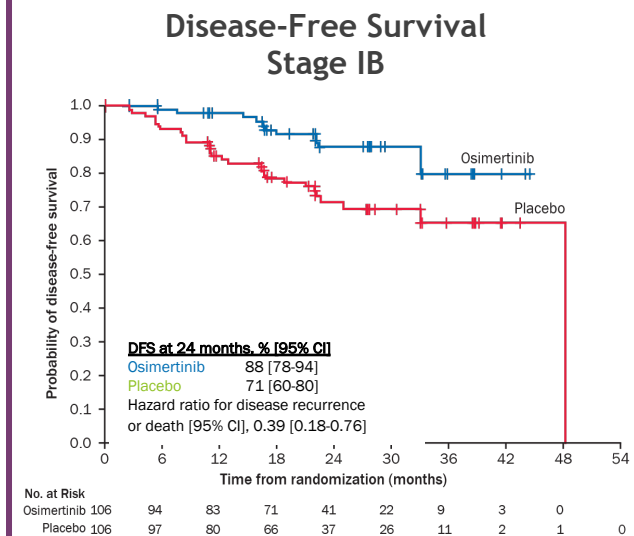
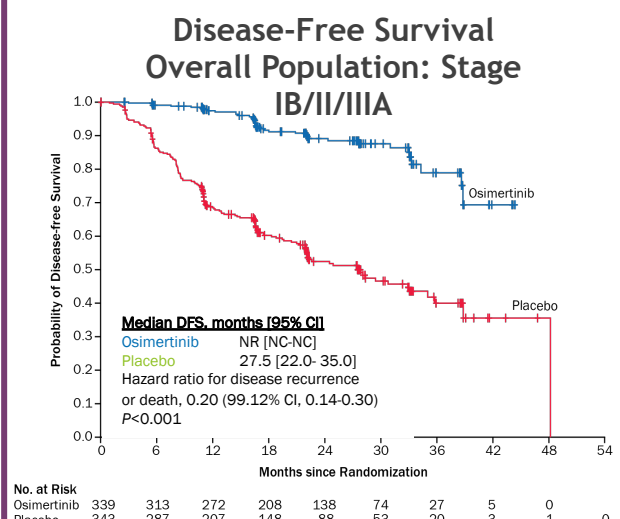
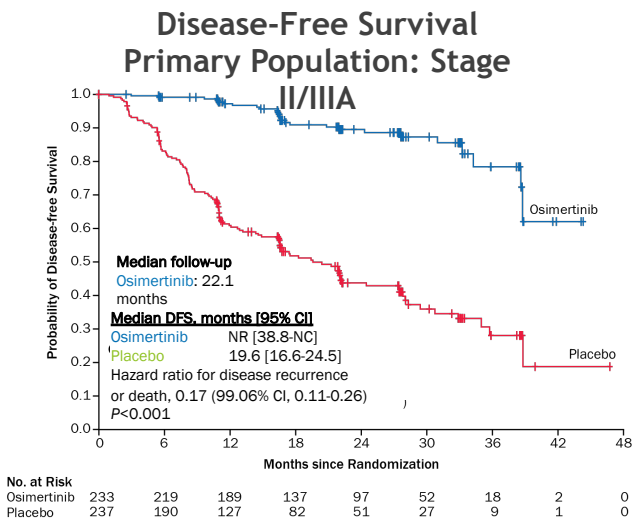
- Validates choice of drug
- Can be used to measure efficacy and relapse (Guardant)
- Refines optimal patient population
- Limits cost and toxicity

# Role of Biomarkers

## Related to Disease:

- Allow to measure treatment efficacy
- ctDNA
- Imaging technologies

# ADAURA: Efficacy

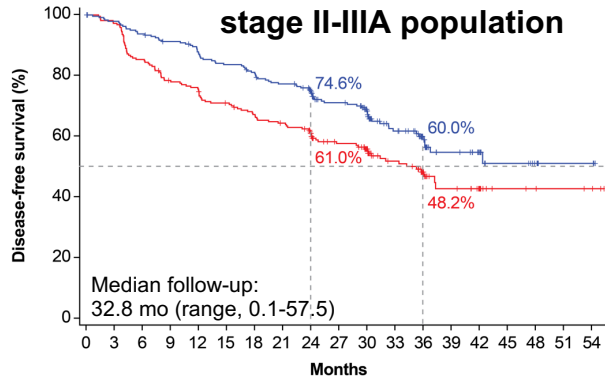


	Stage IB	Stage II	Stage IIIA
<b>2 year DFS rate, % (95% CI)</b>			
– Osimertinib	87 (77, 93)	91 (82, 95)	88 (79, 94)
– Placebo	73 (62, 81)	56 (45, 65)	32 (23, 42)
<b>Overall HR (95% CI)</b>	<b>0.39 (0.18, 0.76)</b>	<b>0.17 (0.08, 0.31)</b>	<b>0.12 (0.07, 0.20)</b>



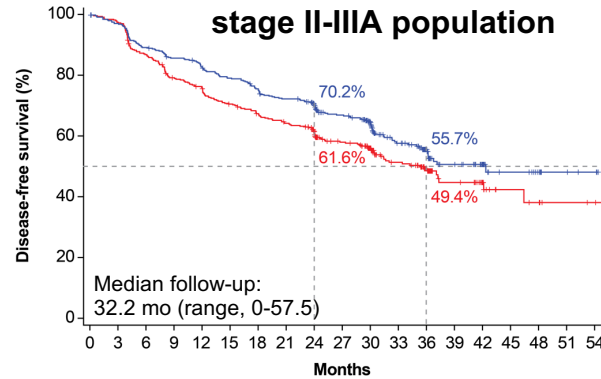
# DFS in the PD-L1 TC $\geq 1\%$ Stage II-III A, All-Randomized Stage II-III A and ITT Populations (primary endpoint)

**PD-L1 TC  $\geq 1\%$   
stage II-III A population**



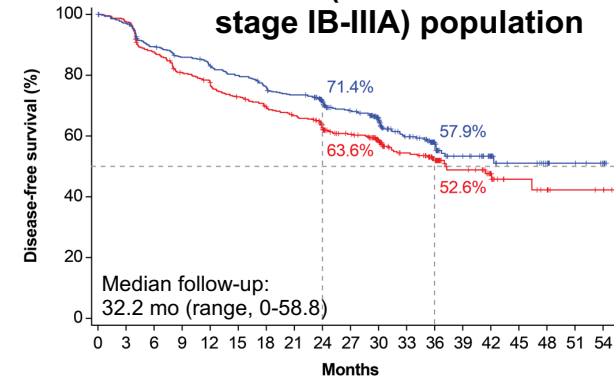
No. at risk																			
Atezolizumab	248	235	225	217	206	198	190	181	159	134	111	76	54	31	22	12	8	3	
BSC	228	212	186	169	160	151	142	135	117	97	80	59	38	21	14	7	6	4	3

**All-randomized  
stage II-III A population**



No. at risk																			
Atezolizumab	442	418	384	367	352	337	319	305	269	225	185	120	84	48	34	16	11	5	3
BSC	440	412	366	331	314	292	277	263	230	182	146	102	71	35	22	10	8	4	3

**ITT (randomized  
stage IB-III A) population**



No. at risk																			
Atezolizumab	507	478	437	418	403	387	367	353	306	257	212	139	97	53	38	19	14	8	4
BSC	498	467	418	383	365	342	324	309	269	219	173	122	90	46	30	13	10	5	4

	Atezolizumab (n=248)	BSC (n=228)
Median DFS (95% CI), mo	NE (36.1, NE)	35.3 (29.0, NE)
Stratified HR (95% CI)	0.66 (0.50, 0.88)	
P value <sup>b</sup>	0.004 <sup>c</sup>	

	Atezolizumab (n=442)	BSC (n=440)
Median DFS (95% CI), mo	42.3 (36.0, NE)	35.3 (30.4, 46.4)
Stratified HR (95% CI)	0.79 (0.64, 0.96)	
P value <sup>b</sup>	0.02 <sup>c</sup>	

	Atezolizumab (n=507)	BSC (n=498)
Median DFS (95% CI), mo	NE (36.1, NE)	37.2 (31.6, NE)
Stratified HR (95% CI)	0.81 (0.67, 0.99)	
P value <sup>b</sup>	0.04 <sup>d</sup>	

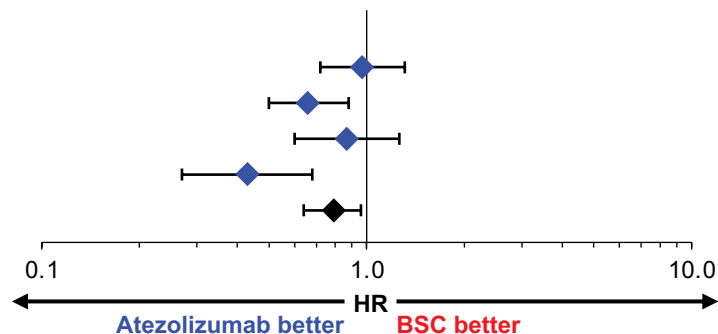
Clinical cutoff: 21 January 2021. <sup>a</sup> Per SP263 assay. <sup>b</sup> Stratified log-rank. <sup>c</sup> Crossed the significance boundary for DFS.

<sup>d</sup> The statistical significance boundary for DFS was not crossed. 1. Wakelee H, et al. J Clin Oncol. 2021;39(suppl 15):8500.

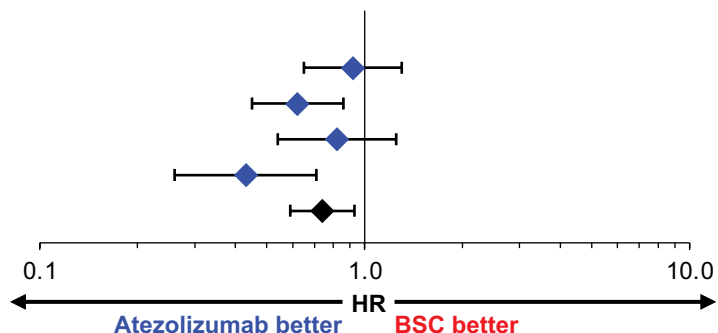
# DFS by PD-L1 Status<sup>a</sup>

*All-randomized stage II-IIIa population ( +/-EGFR/ALK+ disease)*

<u>Subgroup (including EGFR/ALK+)</u>	<u>n</u>	<u>HR (95% CI)<sup>b,c</sup></u>
<b>PD-L1 status by SP263</b>		
TC <1%	383	0.97 (0.72, 1.31)
TC ≥1%	476	0.66 (0.50, 0.88)
TC 1-49%	247	0.87 (0.60, 1.26)
TC ≥50%	229	0.43 (0.27, 0.68)
<b>All patients<sup>d</sup></b>	882	0.79 (0.64, 0.96)



<u>Subgroup (excluding EGFR/ALK+)<sup>e</sup></u>	<u>n</u>	<u>HR (95% CI)<sup>f,g</sup></u>
<b>PD-L1 status by SP263</b>		
TC <1%	312	0.92 (0.65, 1.30)
TC ≥1%	410	0.62 (0.45, 0.86)
TC 1-49%	201	0.82 (0.54, 1.25)
TC ≥50%	209	0.43 (0.26, 0.71)
<b>All patients<sup>h</sup></b>	743	0.74 (0.59, 0.93)



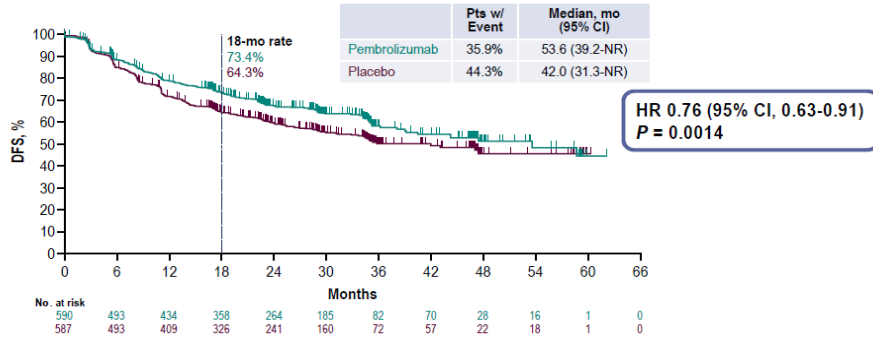
Clinical cutoff: 21 January 2021. <sup>a</sup> Per SP263 assay.

<sup>b</sup> Stratified for all patients and PD-L1 TC ≥1%; unstratified for all other subgroups. <sup>c</sup> DFS analyses in the PD-L1 TC <1% and TC 1-49% subgroups were exploratory. <sup>d</sup> 23 patients had unknown PD-L1 status as assessed by SP263. <sup>e</sup> Excluding patients with known EGFR/ALK+ NSCLC. <sup>f</sup> Unstratified for all subgroups. <sup>g</sup> EGFR/ALK+ exclusion analyses were post hoc. <sup>h</sup> 21 patients had unknown PD-L1 status as assessed by SP263.

# PEARLS/KEYNOTE-091

## Randomized, Triple-Blind, Phase 3 Trial

### DFS, Overall Population

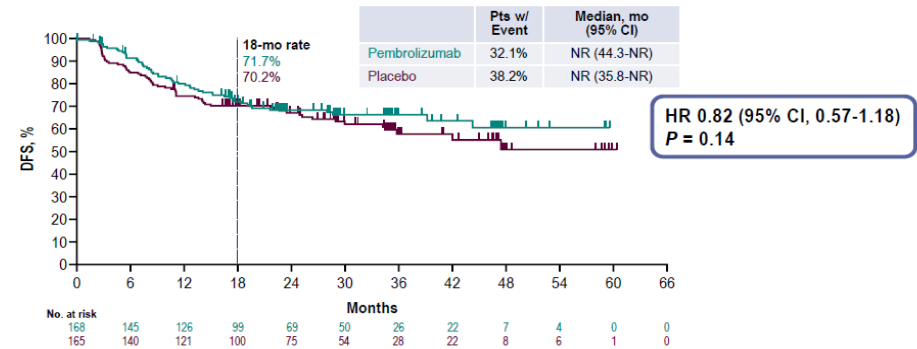


ESMO VIRTUAL PLenary

Response assessed per RECIST v1.1 by investigator review.  
Data cutoff date: September 20, 2021

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### DFS, PD-L1 TPS ≥50% Population

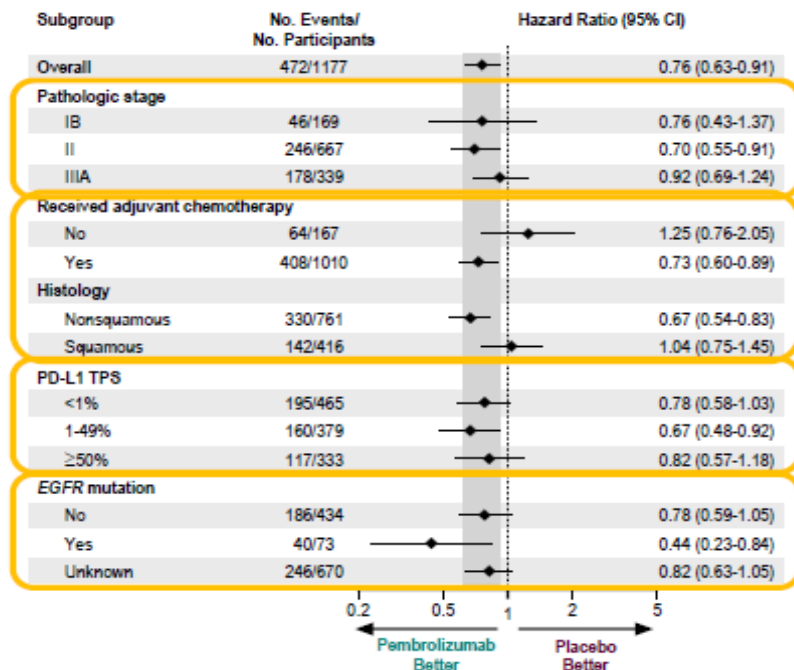
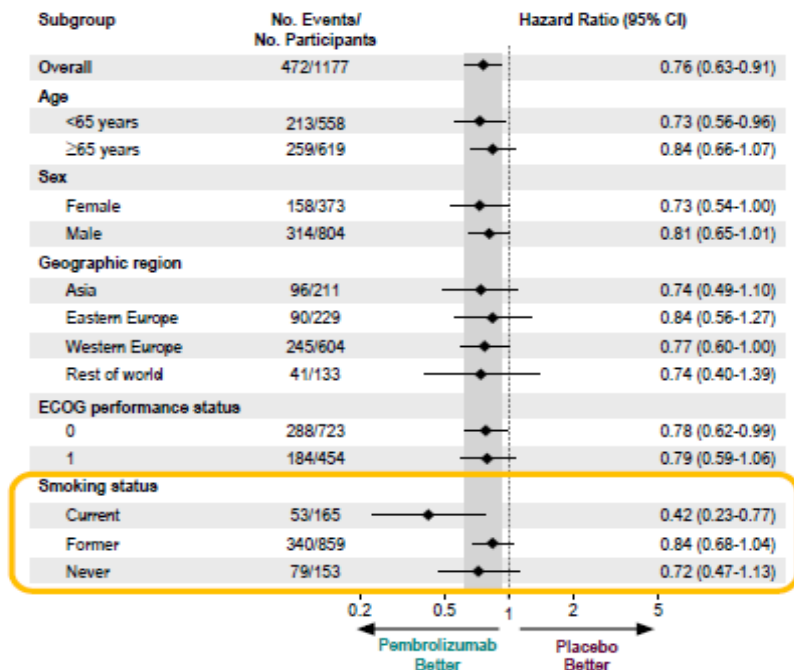


ESMO VIRTUAL PLenary

Response assessed per RECIST v1.1 by investigator review.  
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# DFS in Key Subgroups, Overall Population



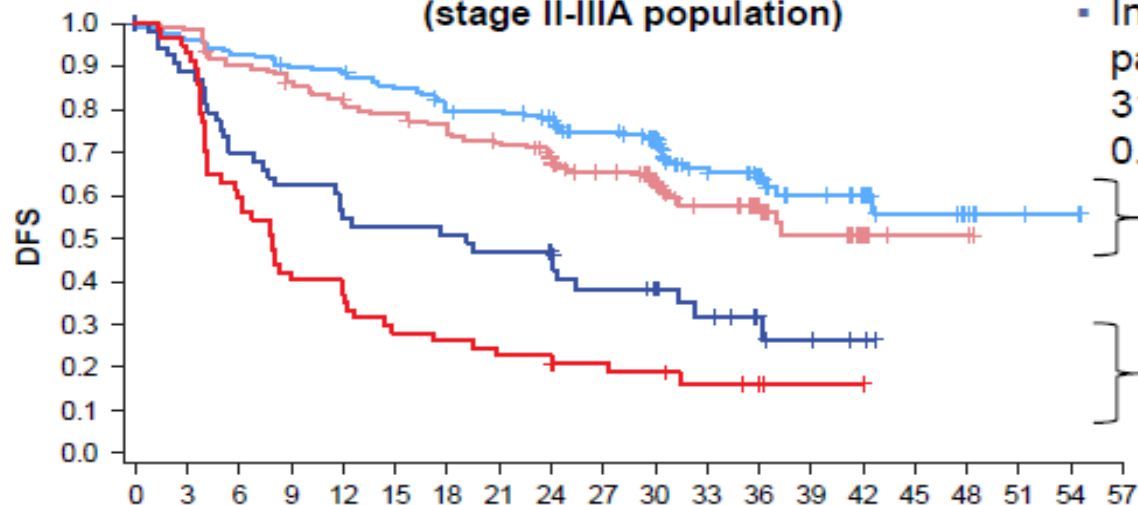
ESMO VIRTUAL PLENARY

Response assessed per RECIST v1.1 by investigator review.  
Data cutoff date: September 20, 2021

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# Impower 010 – Exploratory results for ctDNA

DFS in ctDNA-defined subgroups  
(stage II-IIIa population)



- In all ctDNA-evaluable stage II-IIIa patients, mDFS was NR (atezo) vs 31.4 months (BSC), with an HR of 0.69 (95% CI: 0.53, 0.89)

ctDNA-	Atezo (n=218)	BSC (n=204)
mDFS, mo	NR	NR
HR (95% CI)	0.72 (0.52, 1.00)	

ctDNA+	Atezo (n=53)	BSC (n=59)
mDFS, mo	19.1	7.9
HR (95% CI)	0.61 (0.39, 0.94)	

No. at risk	Months																			
	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	54	57
Atezo, ctDNA-	218	206	199	192	189	180	170	166	151	131	112	73	58	33	24	12	8	3	2	0
Atezo, ctDNA+	53	47	37	33	29	28	27	25	23	17	14	10	6	3	2	0	0	0	0	0
BSC, ctDNA-	204	193	176	167	158	152	143	137	124	106	88	62	44	19	9	3	3	0	0	0
BSC, ctDNA+	59	53	34	24	21	16	15	13	13	9	8	6	4	1	1	0	0	0	0	0

Clinical cutoff: 21 January 2021. Unstratified HRs are shown.

Zhou et al. Impower010 biomarkers. <https://bit.ly/3F2Kr1O>  
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# Immune Checkpoint Inhibitors for Resectable NSCLC

## Neoadjuvant setting



# Considerations for Neoadjuvant I/O or Targeted Therapy

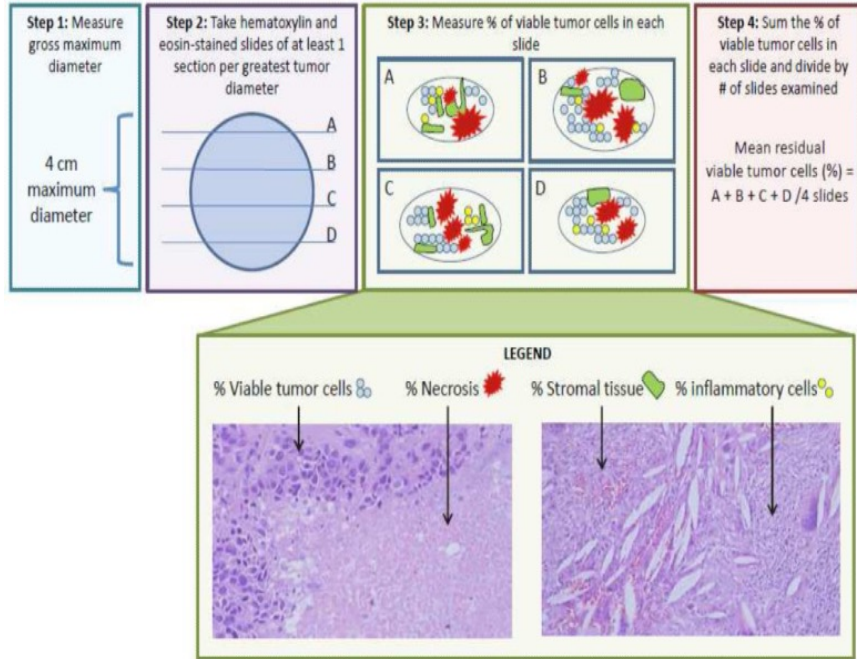
## PRO

- Early eradication of micrometastatic disease
- Improved tolerance of toxicities
- Improved compliance and higher drug exposure
- Pre- and post-treatment tissue to assess biomarkers or adjust treatment
- Guide for need of adjuvant therapy
- Early trial endpoints and shorter trial duration
- Presence of whole tumour allows activation of broader & more diverse immune response

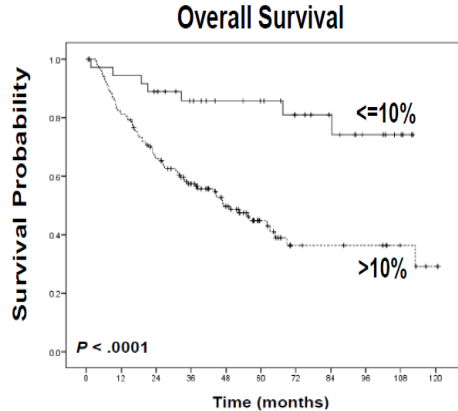
## CON

- Delays of surgery (treatment-related toxicity)
- Increased surgical complications or fewer minimally invasive resection
- Risk of disease progression resulting in missed opportunity for curative surgery
- However, phase 2 neoadjuvant immunotherapy data show approximately **90% of patients underwent surgery**, similar to studies with adjuvant chemotherapy

# Major Pathological Response (<10% viable tumor cells) after Neoadjuvant Chemotherapy as Surrogate Endpoint

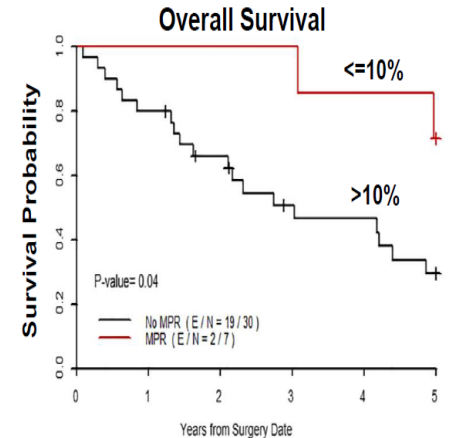


Histopathologic Response Criteria Predict Survival of NSCLC Patients Treated with Neoadjuvant Chemotherapy



Pataer A et al., *J Thorac Oncol*, 2012

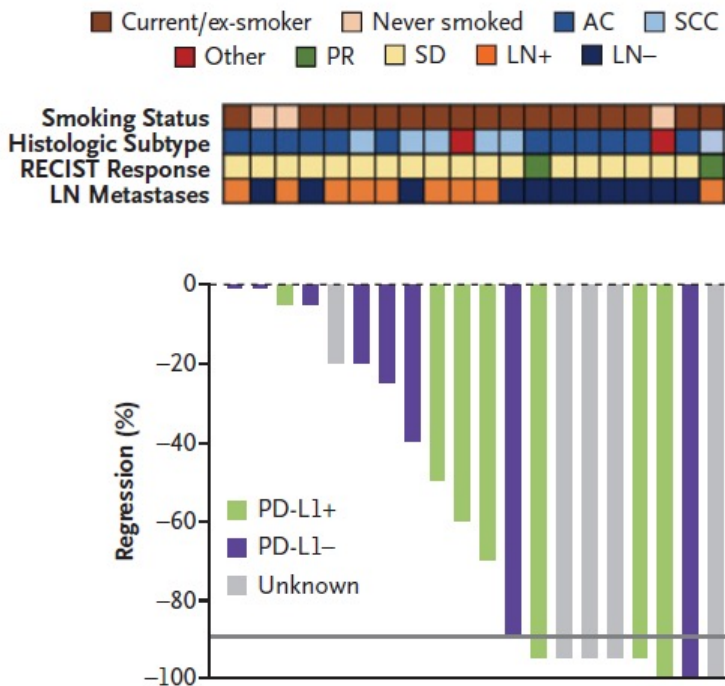
Neoadjuvant Cisplatin Docetaxel Followed by Surgery and Erlotinib in NSCLC Patients



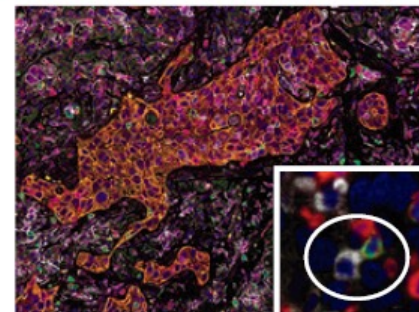
Cascione T et al., *Ann Thorac Surg*, 2017

# Pathological Assessment of Response to Neoadjuvant Blockade of Programmed Death 1

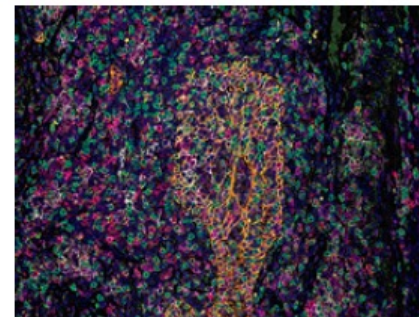
**A Percentage of Pathological Regression, According to Subgroup**



**B Biopsy Sample before Nivolumab**



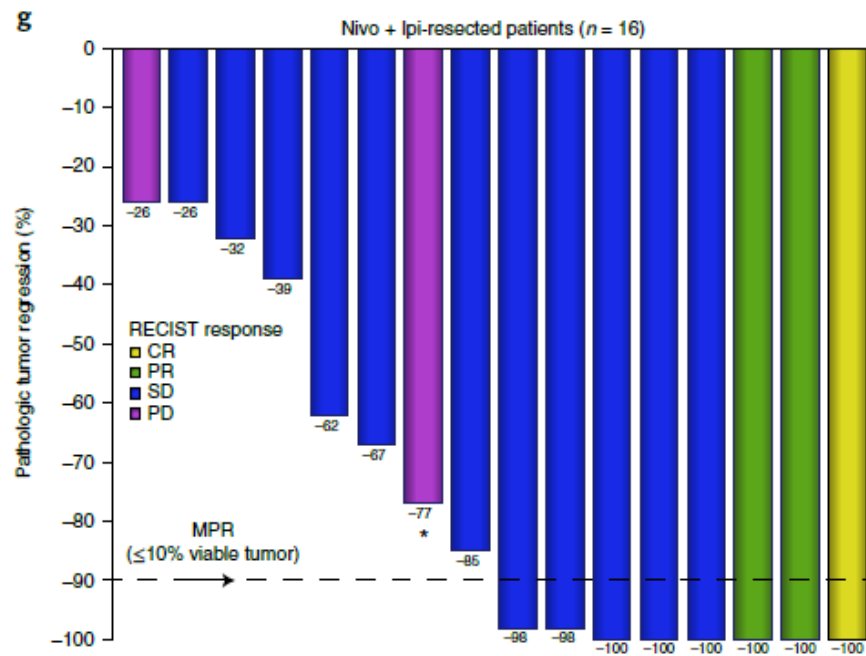
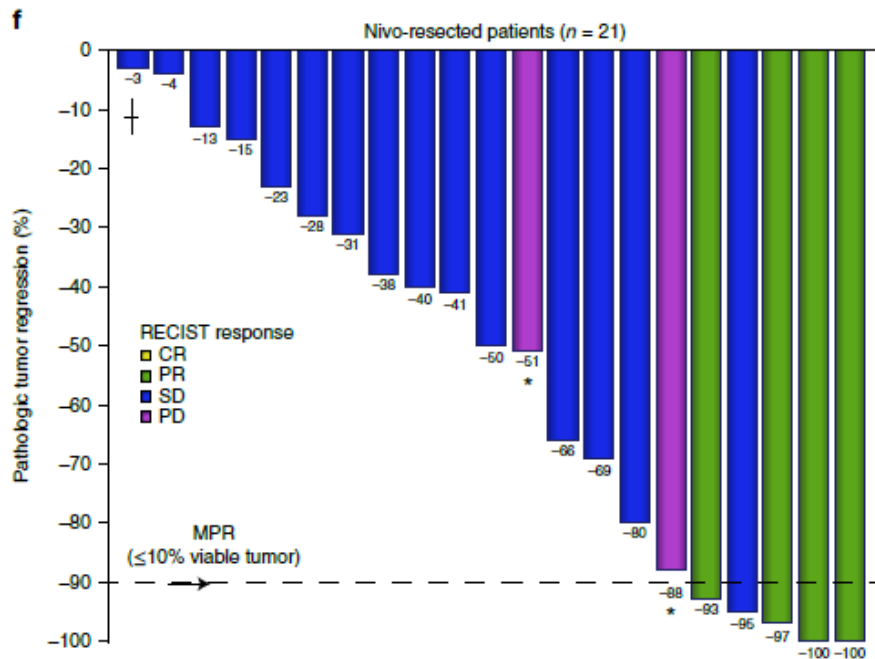
**C Biopsy Sample after Nivolumab**



# Neoadjuvant Nivolumab or Nivolumab Plus Ipilimumab in Operable NSCLC: The Phase 2 Randomized NEOSTAR Trial

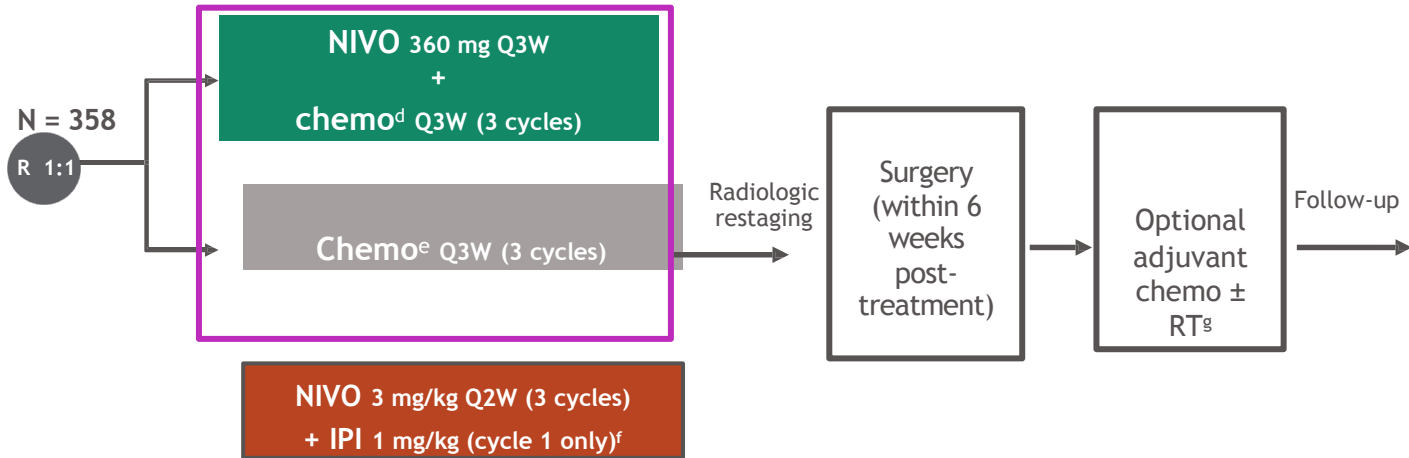
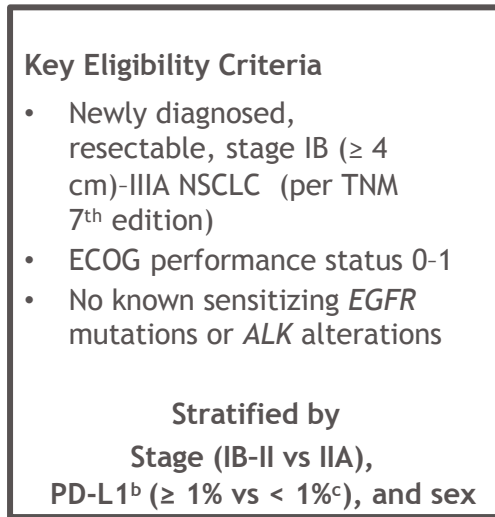
Endpoints: MPR (vs historical control with chemotherapy)  
Goal of 28% (6/21) vs 15%

# Pathologic Responses to Neoadjuvant Nivolumab and Nivolumab + Ipilimumab in Resected Patients



# CheckMate 816 Study Design<sup>a</sup>

## Primary analysis population



### Primary endpoints

- pCR by BIPR
- EFS by BICR

### Secondary endpoints

- MPR by BIPR
- OS
- Time to death or distant metastases

### Exploratory endpoints

- ORR by BICR
- Predictive biomarkers (PD-L1, TMB, ctDNA<sup>h</sup>)

Database lock: September 16, 2020; minimum follow-up: 7.6 months for NIVO + chemo and chemo arms.

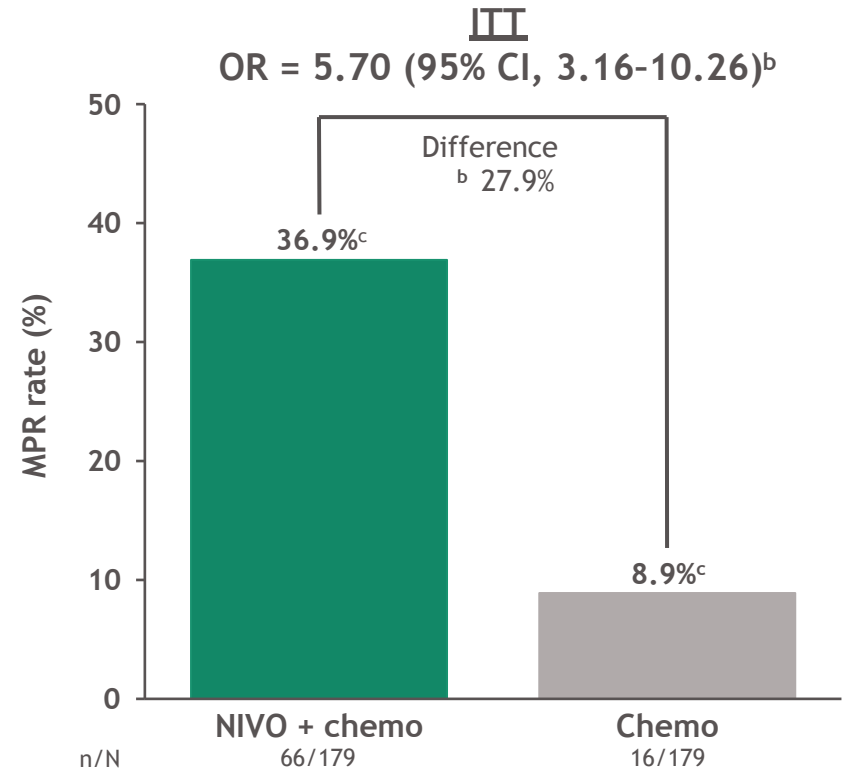
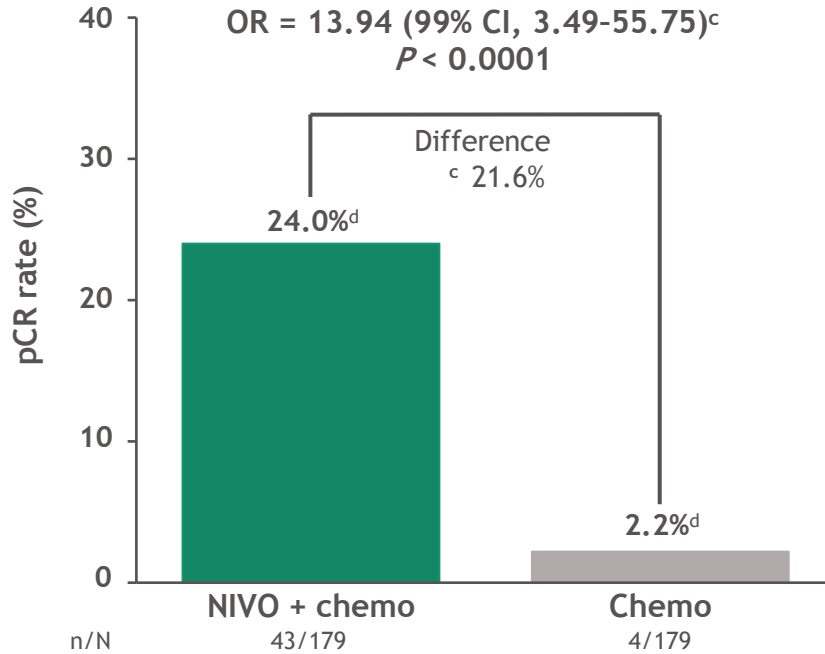
<sup>a</sup>NCT02998528; <sup>b</sup>Determined by the PD-L1 IHC 28-8 pharmDx assay (Dako); <sup>c</sup>Included patients with PD-L1 expression status not evaluable and indeterminate; <sup>d</sup>NSQ: pemetrexed + cisplatin or paclitaxel + carboplatin; SQ: gemcitabine + cisplatin or paclitaxel + carboplatin; <sup>e</sup>Vinorelbine + cisplatin, docetaxel + cisplatin, gemcitabine + cisplatin (SQ only), pemetrexed + cisplatin (NSQ only), or paclitaxel + carboplatin; <sup>f</sup>Randomized exploratory arm (enrollment closed early); <sup>g</sup>Per healthcare professional choice; <sup>h</sup>Performed using tumor-guided personalized ctDNA panel (ArcherDX Personalized Cancer Monitoring).



# CM 816: Primary Endpoint: pCR rate

# mPR rate

## Primary endpoint: ITT (ypT0N0)<sup>b</sup>

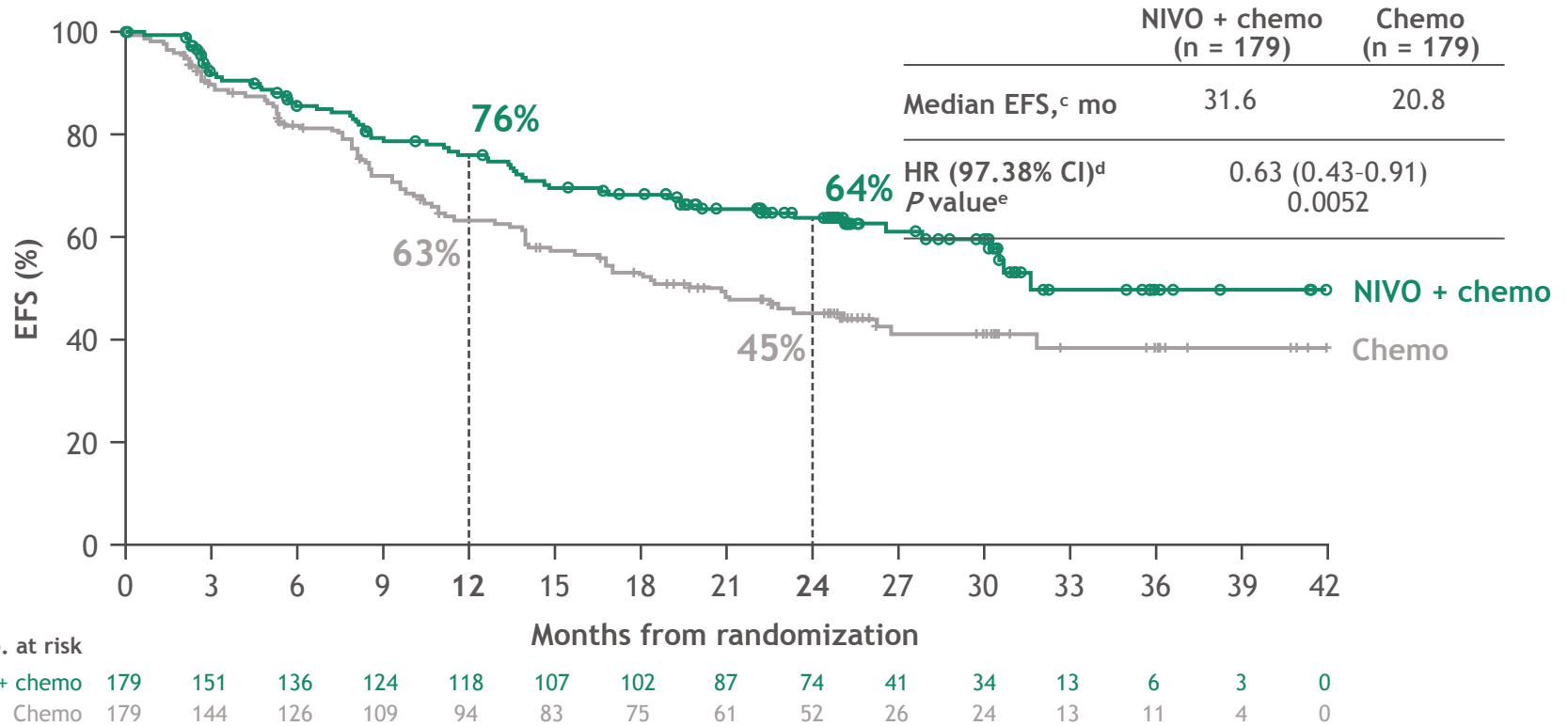


- pCR rate in the exploratory NIVO + IPI arm (ITT) was 20.4% (95% CI, 13.4-29.0)

<sup>a</sup>Per BIPR; pCR: 0% residual viable tumor cells in both primary tumor (lung) and sampled lymph nodes; <sup>b</sup>ITT principle: patients who did not undergo surgery counted as non-responders for primary analysis;

<sup>c</sup>Calculated by stratified Cochran-Mantel-Haenszel method; <sup>d</sup>pCR rates 95% CI: NIVO + chemo, 18.0-31.0; chemo, 0.6-5.6; <sup>e</sup>Patients who underwent definitive surgery with an evaluable pathology sample for BIPR.

# Primary endpoint: EFS<sup>a,b</sup> with neoadjuvant NIVO + chemo vs chemo

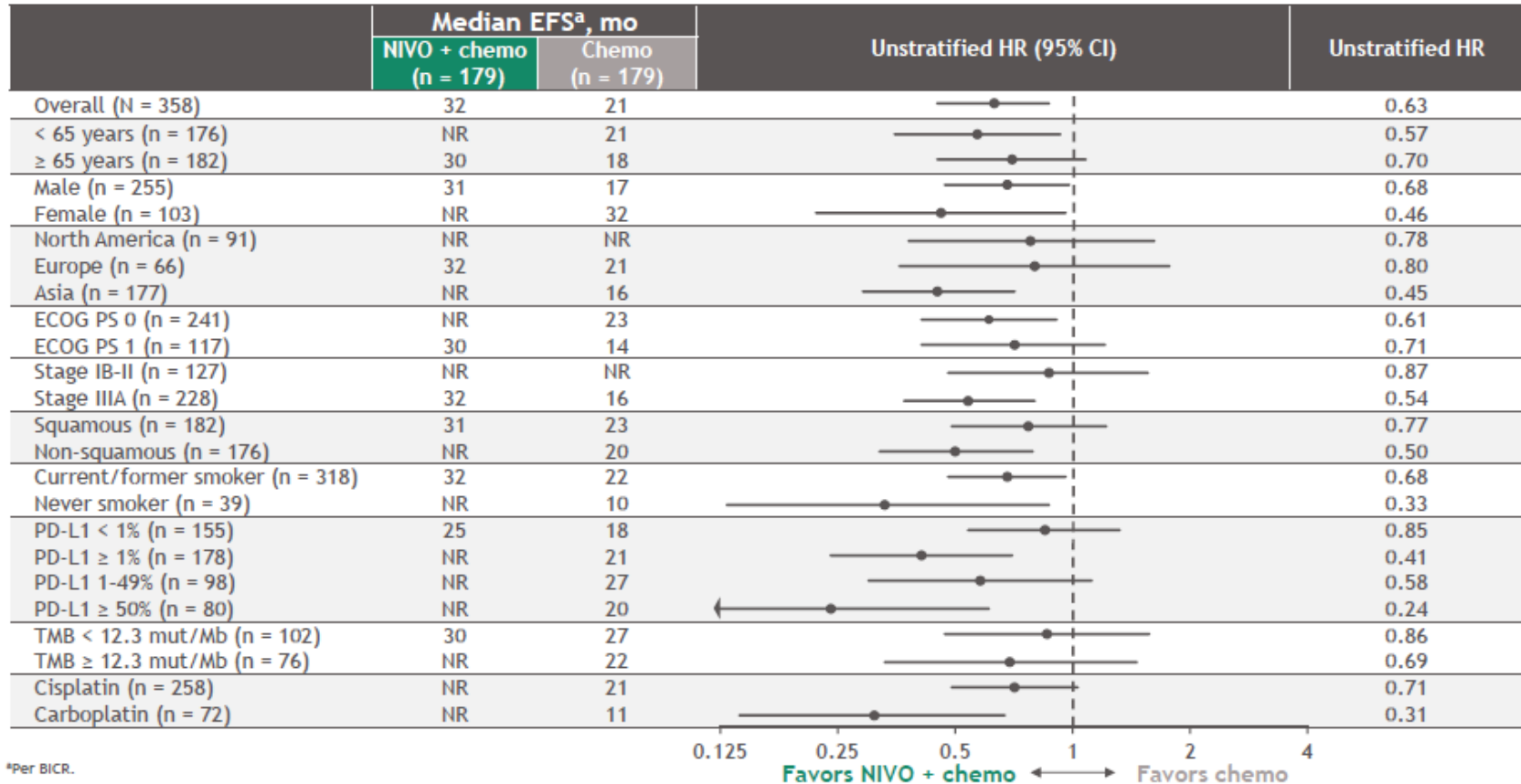


Minimum follow-up: 21 months; median follow-up, 29.5 months.

<sup>a</sup>Per BICR; <sup>b</sup>EFS defined as the time from randomization to any progression of disease precluding surgery, progression or recurrence of disease after surgery, progression for patients without surgery, or death due to any cause; patients with subsequent therapy were censored at the last evaluable tumor assessment on or prior to the date of subsequent therapy; <sup>c</sup>95% CI = 30.2-NR (NIVO + chemo) and 14.0-26.7 (chemo);

<sup>d</sup>95% CI = 0.45-0.87; <sup>e</sup>The significance boundary at this interim analysis was 0.0262.

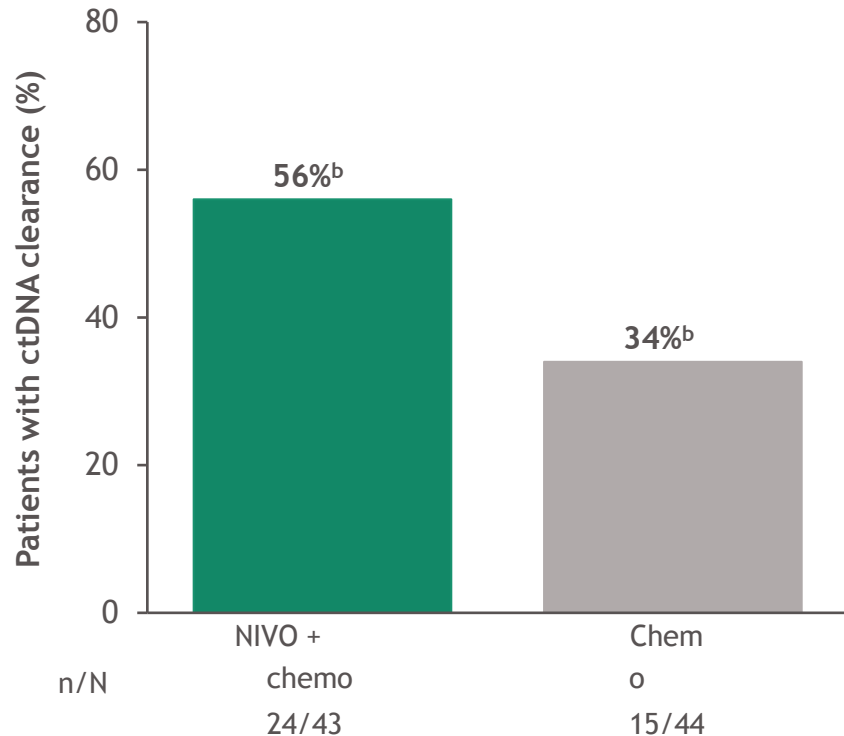
# EFS subgroup analysis



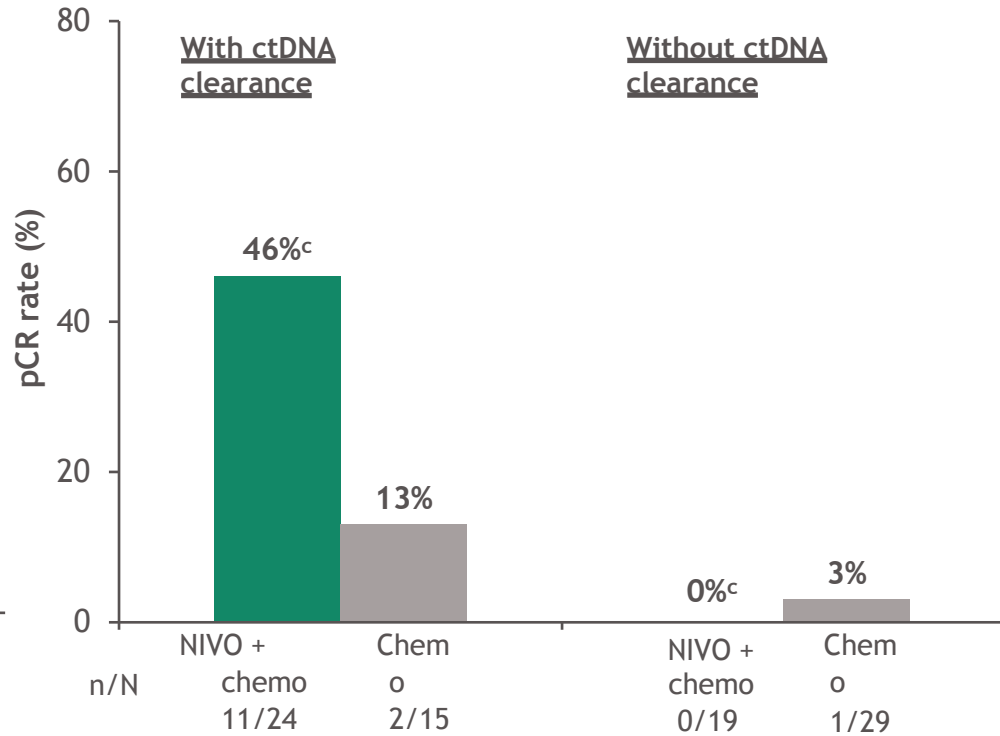
<sup>a</sup>Per BICR.

# ctDNA Clearance and Association With Pathological Response

## ctDNA clearance and mPR rates

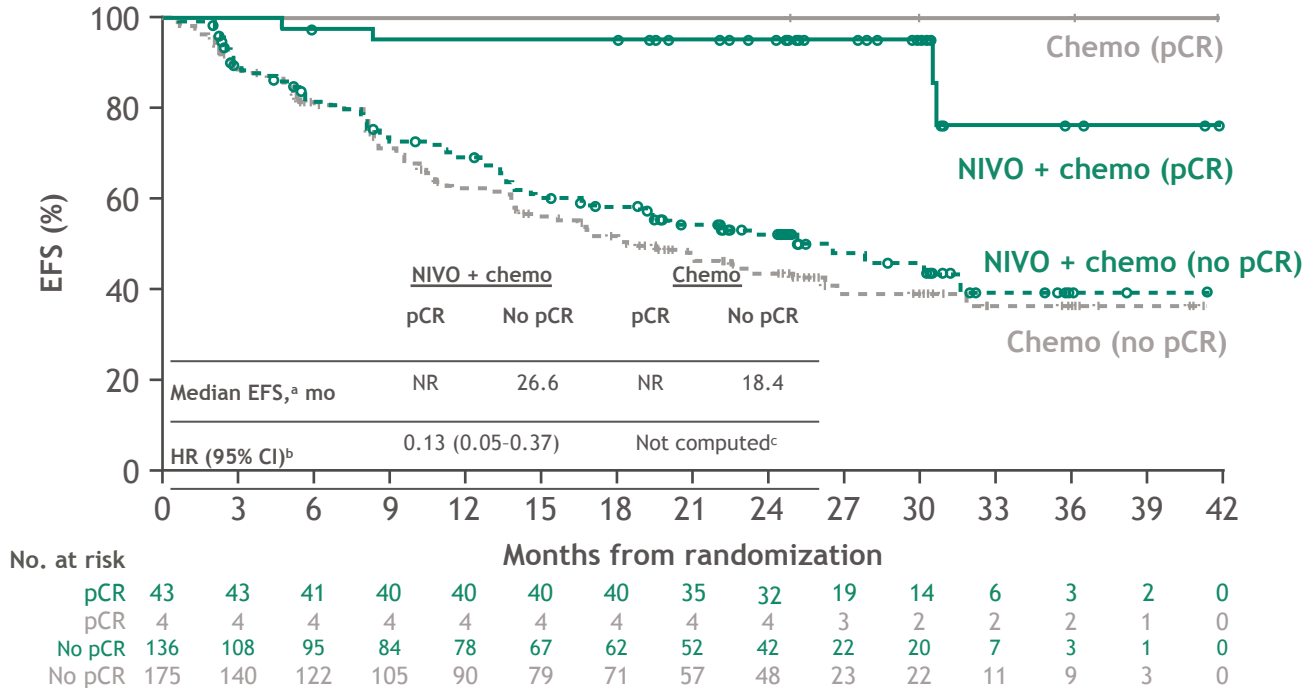


## ctDNA clearance and pCR rates



<sup>a</sup>Performed using tumor-guided personalized ctDNA panel (ArcherDX Personalized Cancer Monitoring); 90 patients were ctDNA evaluable and 87 had detectable ctDNA at C1D1; main reason for sample attrition were lack of tissue for WES and lack of quality control pass for tissue and plasma; <sup>b</sup>ctDNA clearance 95% CI: NIVO + chemo, 40-71; chemo, 20-50; <sup>c</sup>pCR rates 95% CI for NIVO + chemo: with ctDNA clearance, 26-67; without ctDNA clearance, 0-18.

# Exploratory analysis: EFS by pCR status



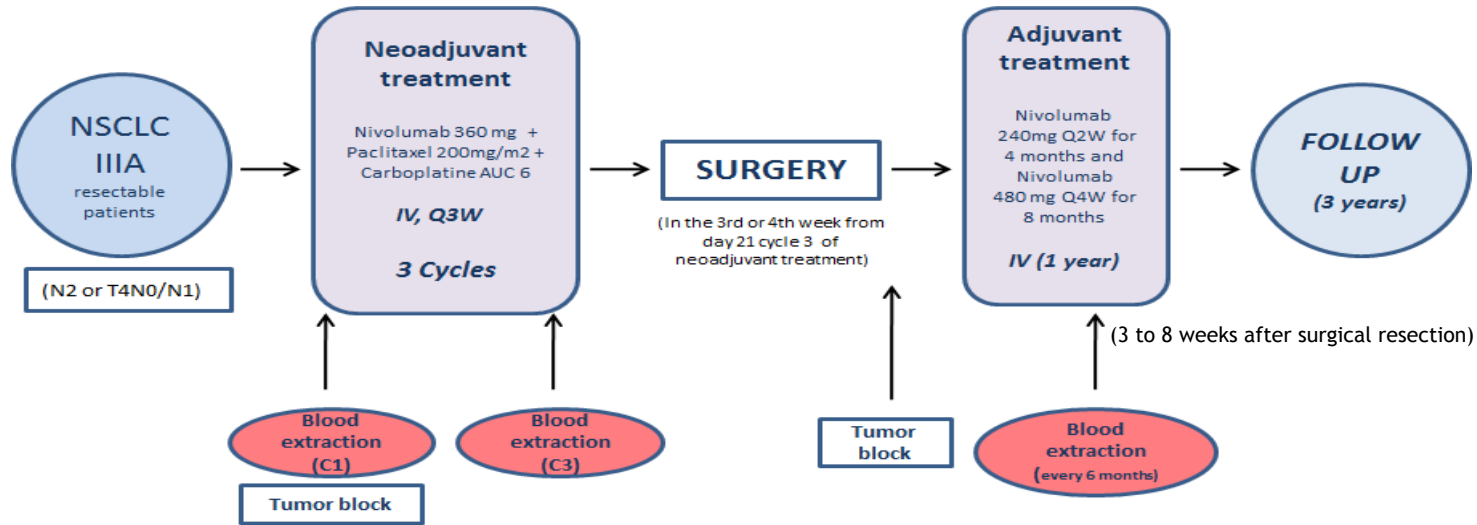
- pCR rates were significantly improved with NIVO + chemo vs chemo (24.0% vs 2.2%)
- In patients without pCR, HR (95% CI) for NIVO + chemo vs chemo was 0.84 (0.61-1.17)

Minimum follow-up: 21 months; median follow-up, 29.5 months.

<sup>a</sup>95% CI = 30.6-NR (NIVO + chemo, pCR), 16.6-NR (NIVO + chemo, no pCR) and NR-NR (chemo, pCR), 13.9-26.2 (chemo, no pCR); <sup>b</sup>In the pooled patient population (NIVO + chemo and chemo arms combined), EFS HR (95% CI) was 0.11 (0.04-0.29) for patients with pCR vs those without pCR; <sup>c</sup>HR was not computed for the chemo arm due to only 4 patients having a pCR.

# Overall Survival and Biomarker Analysis of Neoadjuvant Nivolumab Plus Chemotherapy in Operable Stage IIIA NSCLC (NADIM phase II trial)

Open-label, multicenter (18), single-arm phase 2



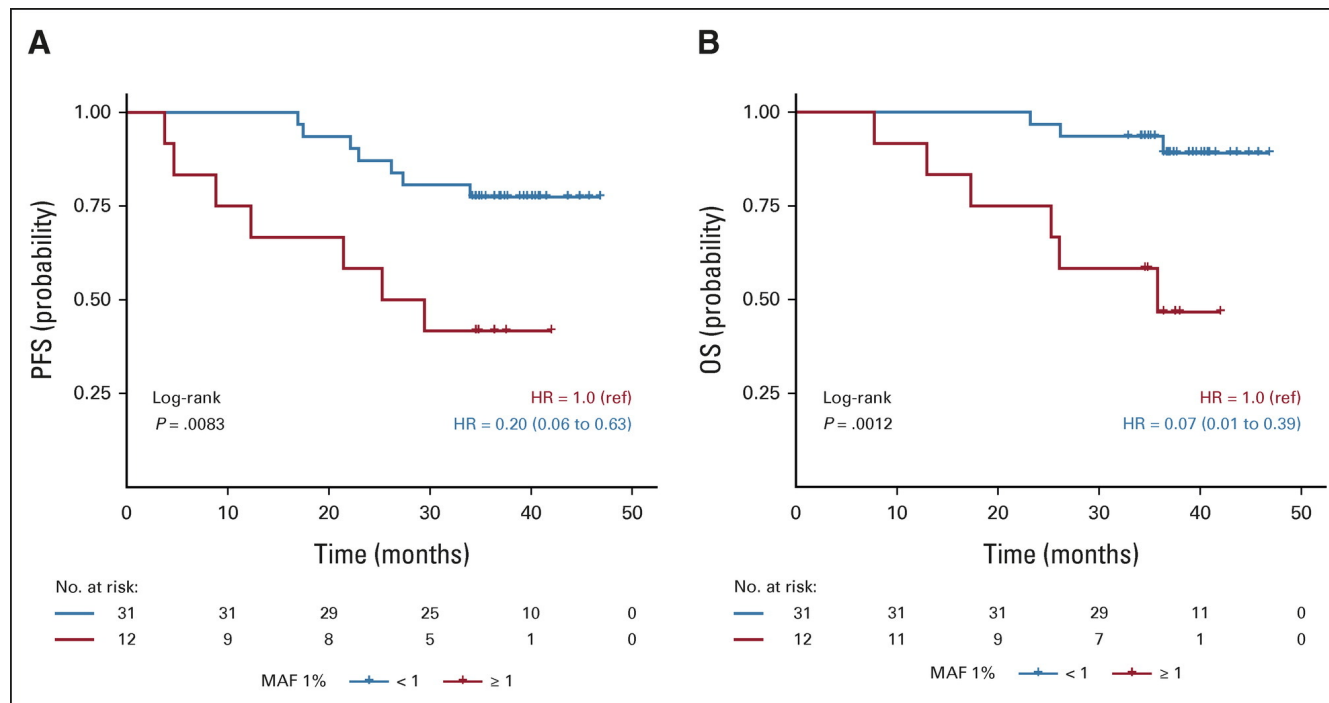
**Sample size: 46**  
**Study start: April 2017**  
**Enrollment completion: August 2018**

**Primary Endpoint:**  
PFS at 24 months

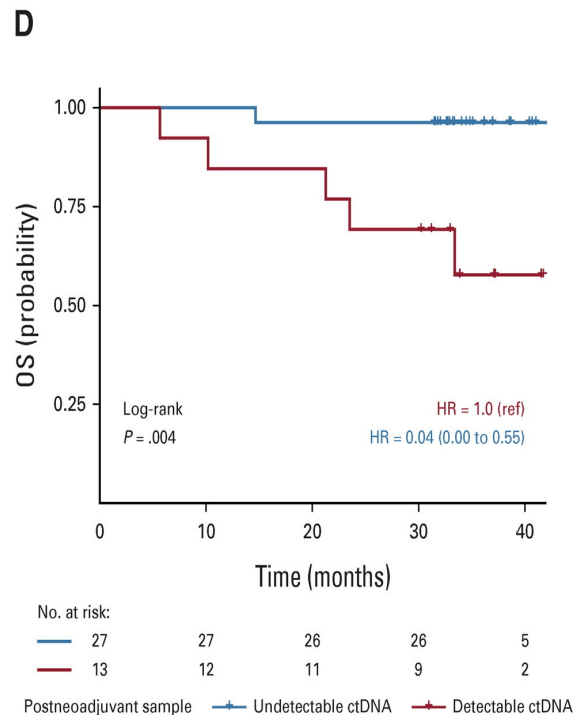
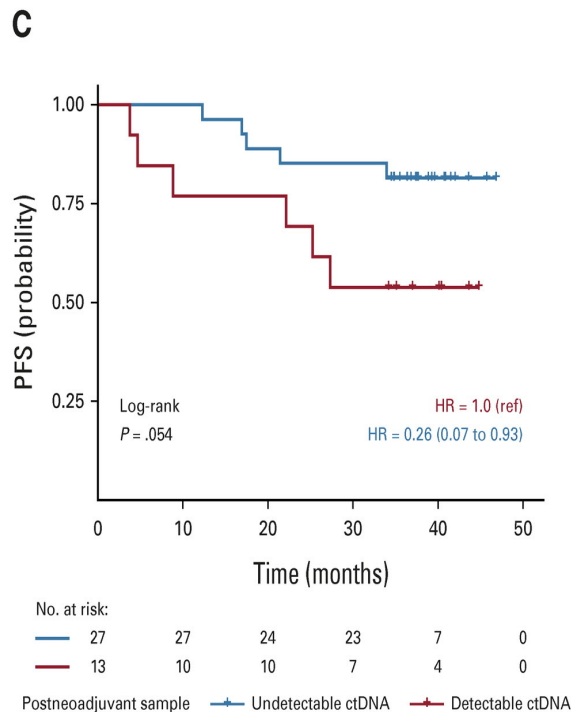
**Secondary Endpoints:** Down-staging rate, complete resection rate, ORR, safety, TTP, OS at 3 years



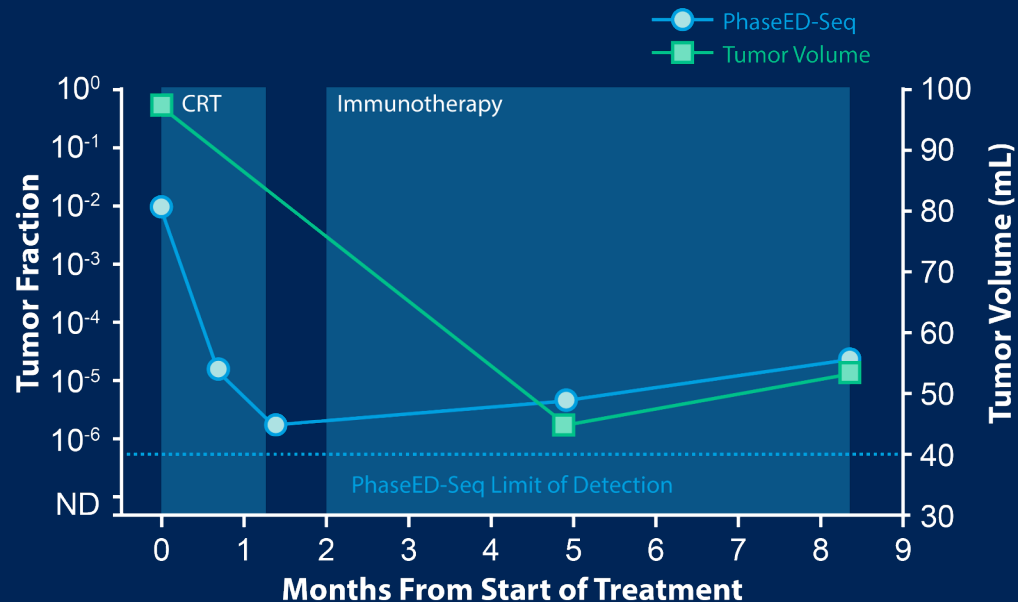
# PFS and OS by ctDNA levels at baseline, using a cutoff of <1% MAF



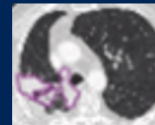
# PFS and OS according to ctDNA detection after neoadjuvant treatment



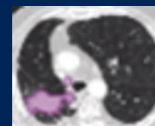
# “Liquid Biopsy” — an Innovation in Diagnostic Testing



Pretreatment Mass



Treatment Response with Residual Fibrosis, Tumor DNA Detected in Plasma



Biopsy confirmed recurrent disease

# Impact of Neoadjuvant IO in Early-Stage NSCLC

- **Ideal primary endpoint: a surrogate for EFS, PFS, or OS**
- **Potential surrogate endpoints and predictors of IO response**
  - mPR (based on neoadjuvant chemotherapy)
  - pCR (supported by CM-816)
  - ctDNA (also to monitor recurrence)
  
  - TMB
  - Tumor microenvironment (ex. Immune cellular infiltrates, cytokines, PD-L1)
  - Microbiome

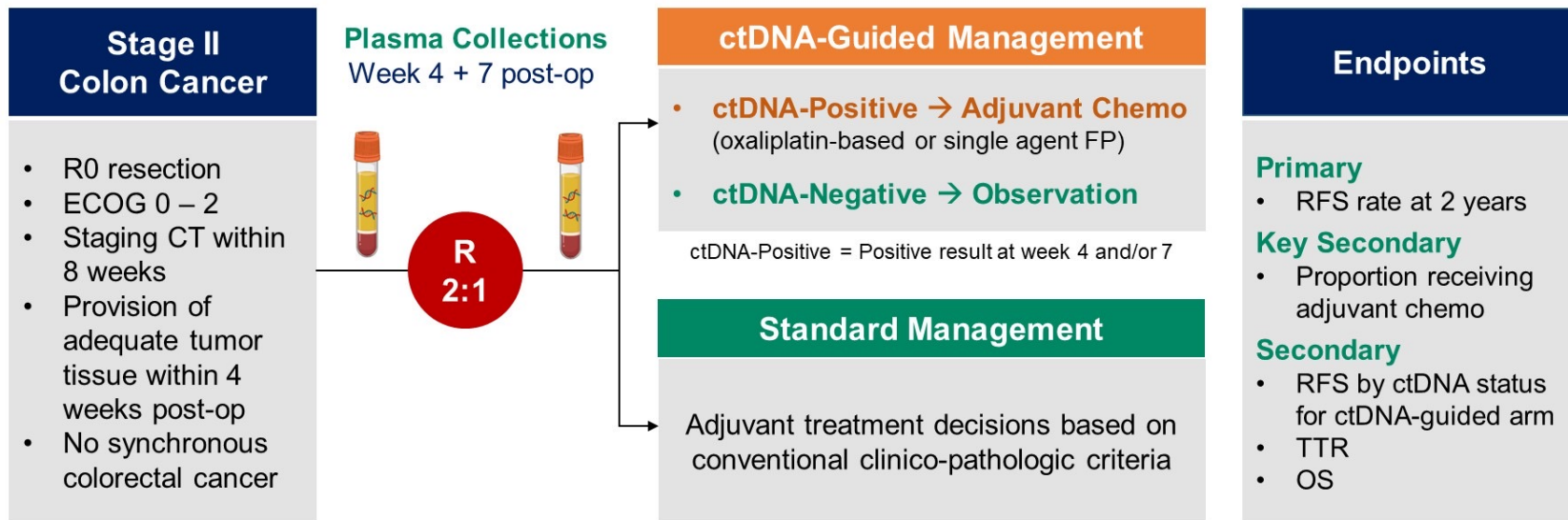
# Biomarker-Based De-escalated curative-intent Therapy

## Examples:

- Dynamic Trial in Colorectal Cancer
- Dostarlimab De-escalation Trial in Rectal Cancer as example for HPV-related Head and Neck Cancer

# DYNAMIC Study Design

ACTRN12615000381583



## Stratification Factors

- T stage (T3 vs T4)
- Type of participating center (metropolitan vs regional)

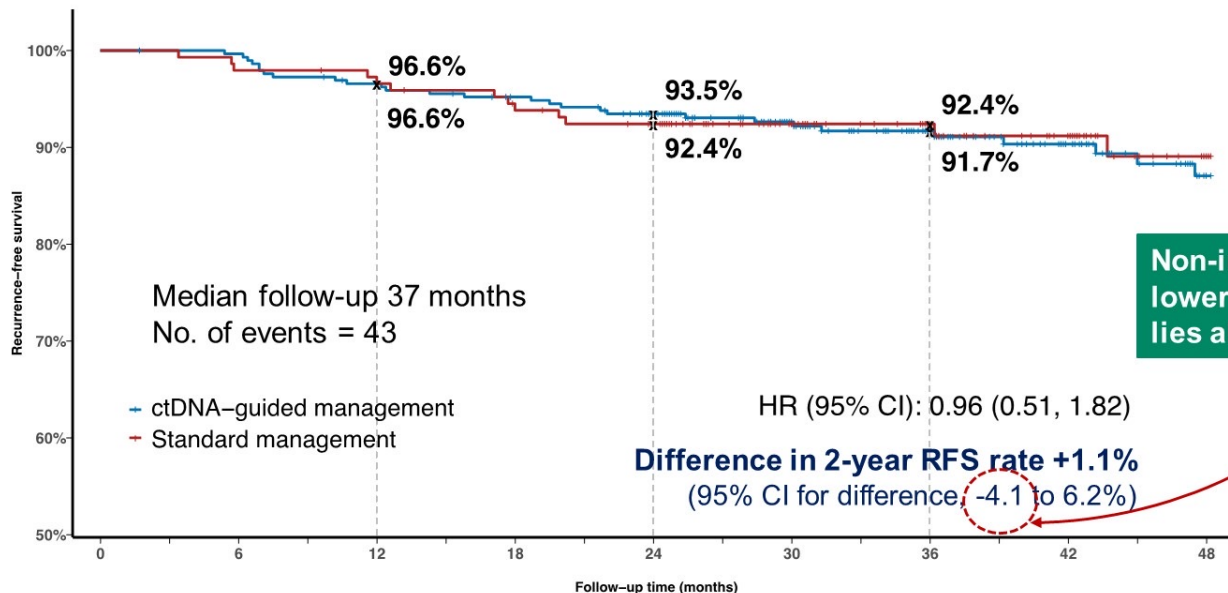
## Surveillance:

- CEA → 3-monthly for 24M, then 6-monthly for 36M
- CT C/A/P → 6-monthly for 24M, then at 36M

# Adjuvant Treatment Delivery

Treatment Information	ctDNA-Guided N = 294	Standard Management N = 147	P-value
Adjuvant Chemotherapy received, n	45 (15%)	41 (28%)	0.0017
Chemotherapy regimen received, n			
Oxaliplatin-based doublet	28/45 (62%)	4/41 (10%)	<.0001
Single agent fluoropyrimidine	17/45 (38%)	37/41 (90%)	
Time from surgery to commencing chemotherapy, median (IQR), days	83 (76, 89)	53 (49, 61)	<.0001
Treatment duration, median (IQR), weeks	24 (19, 24)	24 (21, 24)	0.9318
Completed planned treatment, n	38 (85%)	32 (78%)	0.7036
Percentage of full dose delivered, median (IQR)	78 (56, 100)	84 (64, 100)	0.6194

# Recurrence-Free Survival



	Numbers at risk								
	0	6	12	18	24	30	36	42	48
ctDNA-guided	294	292	281	273	259	207	155	109	64
Standard	147	144	142	136	128	97	78	57	33



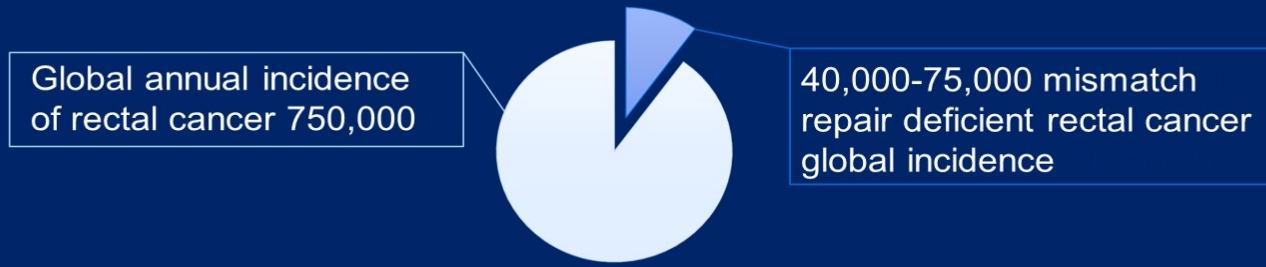
# Summary

- **For patients with stage II colon cancer, a ctDNA-guided approach (treating only patients with a positive ctDNA after surgery) compared to standard-of-care**
  - Substantially reduced the proportion receiving adjuvant chemotherapy (28% → 15%)
  - Did not compromise recurrence-free survival (2-year RFS: 93.5% vs 92.4%)
- **Patients with a positive ctDNA after surgery may derive RFS benefit from adjuvant chemotherapy**
  - Favorable 3-year RFS in patients treated with adjuvant chemotherapy (86.4%) versus low RFS in historical series (< 20%) if untreated
  - Ongoing trials (e.g., COBRA, CIRCULATE, CIRCULATE-PRODIGE) will provide further guidance regarding the optimal use of ctDNA-informed management
- **ctDNA-negative patients have a low recurrence risk without adjuvant chemotherapy**
  - 3-year RFS 92.5% (clinical low risk: 96.7%; T3: 94.2%)

# Dynamic Trial as Example for NSCLC

- Continue to use stage to determine need for neoadjuvant therapy at diagnosis but obtain ctDNA
- pCR as optimal goal and primary endpoint
- Use ctDNA to determine number of neoadjuvant chemo-IO cycles and
- To determine the need to give additional adjuvant therapy (NADIM II)

# Mismatch repair deficient rectal cancer



- Approximately 5-10% of rectal cancers are mismatch repair deficient
- Relatively resistant to chemotherapy
- Checkpoint blockade is highly effective in metastatic mismatch repair deficient cancers with a complete response rate ~10%

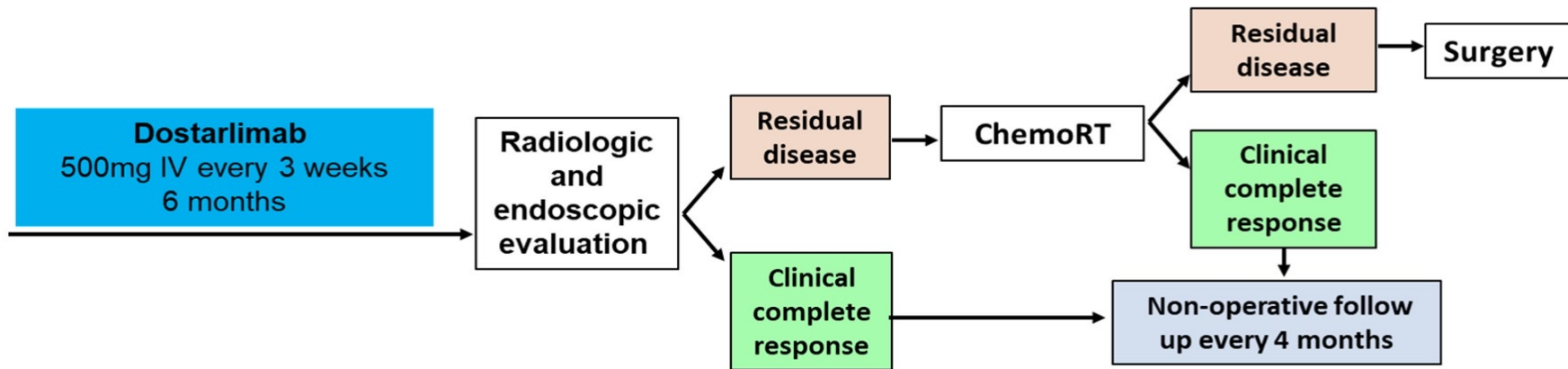
Cercek A, Clin Cancer Res 2020  
Andre T, N Engl J Med 2020  
Le DT, N Engl J Med 2015



# Hypothesis:

In mismatch repair deficient rectal cancer, PD-1 blockade may be able to either:

- a) replace chemotherapy
- b) replace chemo *and* radiation therapy
- c) replace chemo *and* radiation, *and* surgery



**Patient population:** Stage II and III mismatch repair deficient rectal cancer

**Target Enrollment:** 30 subjects

**Study Design:** Simon's two stage minimax design

NCT04165772



# Individual responses to PD-1 blockade with dostarlimab

Patients who completed 6-months of dostarlimab

ID	Age	Stage T	Stage N	FU (months)	Digital rectal exam response	Endoscopic best response	Rectal MRI best response	Overall response <b>100%</b>
1	38	T4	N+	23.8	CR	CR	CR	cCR
2	30	T3	N+	20.5	CR	CR	CR	cCR
3	61	T1/2	N+	20.6	CR	CR	CR	cCR
4	28	T4	N+	20.5	CR	CR	CR	cCR
5	53	T1/2	N+	9.1	CR	CR	CR	cCR
6	77	T1/2	N+	11.0	CR	CR	CR	cCR
7	77	T1/2	N+	8.7	CR	CR	CR	cCR
8	55	T3	N+	5.0	CR	CR	CR	cCR
9	68	T3	N+	4.9	CR	CR	CR	cCR
10	78	T3	N-	1.7	CR	CR	CR	cCR
11	55	T3	N+	4.7	CR	CR	CR	cCR
12	27	T3	N+	4.4	CR	CR	CR	cCR
13	26	T3	N+	0.8	CR	CR	CR	cCR
14	43	T3	N+	0.7	CR	CR	CR	cCR

# Potential Impact on Patients is Huge

- Unprecedented 100% clinical complete response rate
- Possibility of decreased morbidity from elimination of pelvic radiation and surgery
  - Bowel dysfunction
  - Urinary dysfunction
  - Sexual dysfunction
  - Infertility
  - Permanent ostomy
- Particularly relevant as incidence of rectal cancer is increasing steadily in young people

American Cancer Society. *Cancer Facts & Figures 2022*. Atlanta: American Cancer Society; 2022.

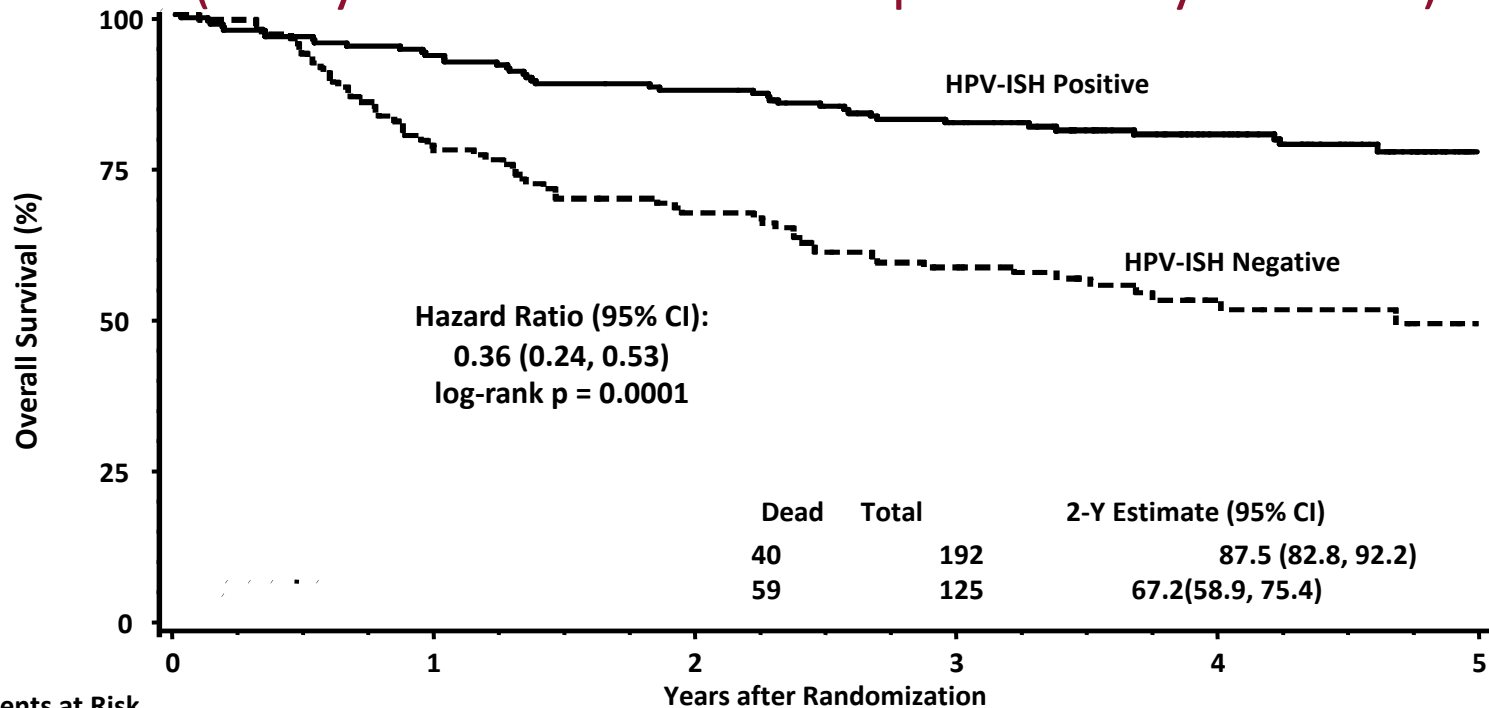
# Remaining Questions

- Can tumor regrowth and recurrence be successfully salvaged?
- Are there biomarkers (e.g., ctDNA, PET scans) that can better predict pCR?
- What is the optimal duration of neoadjuvant immunotherapy?
- Is there a role for combination with anti-CTLA4 antibodies, chemotherapy, and/or radiation?
- Can neoadjuvant checkpoint blockade alone prevent additional Lynch-related CRCs and other Lynch-associated cancers?



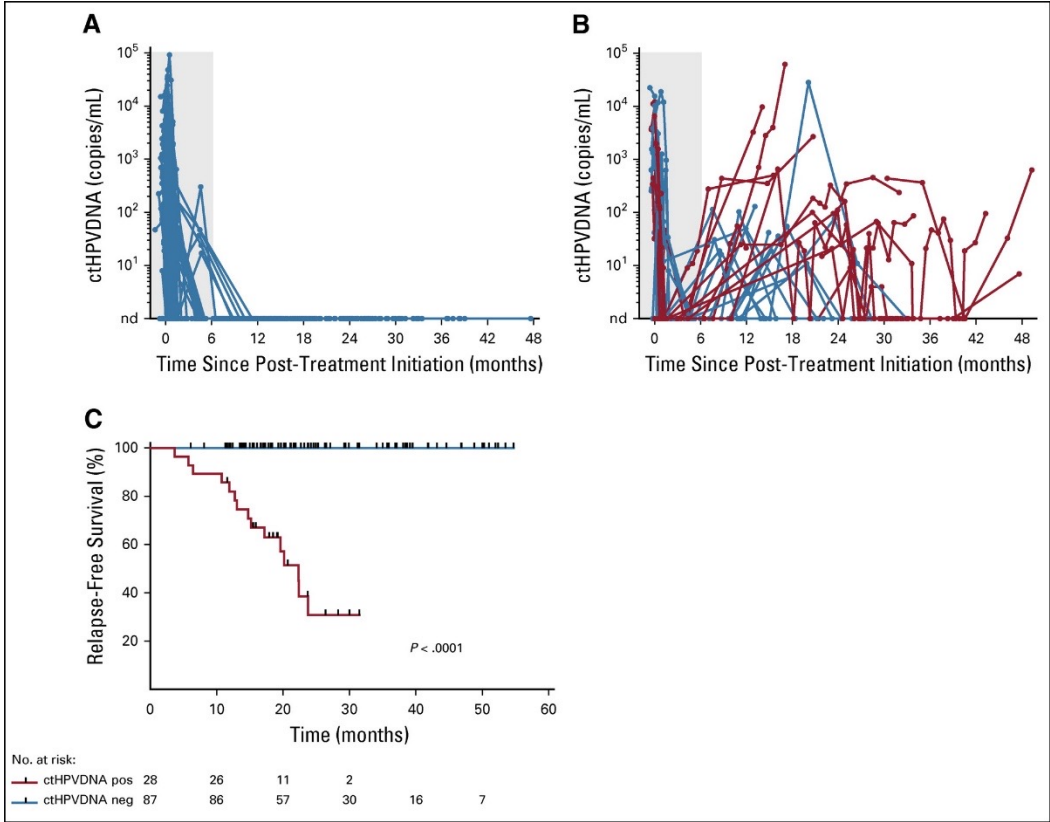
# Role of HPV in Locoregionally Advanced HNC

## Phase III Trial RTOG 0129 – Survival by HPV Status (70 Gy and concomitant cisplatin every 3 weeks)

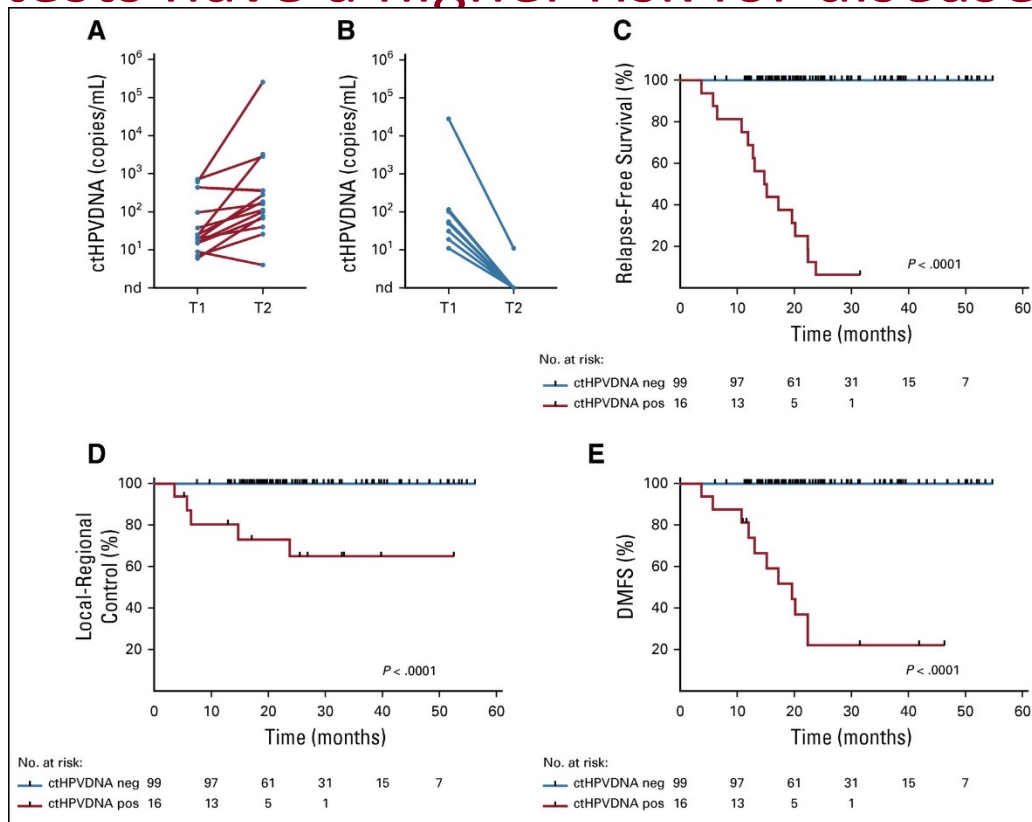


Patients at Risk	0	1	2	3	4	5
HPV-ISH Positive	192	179	167	148	104	19
HPV-ISH Negative	125	97	83	69	35	6

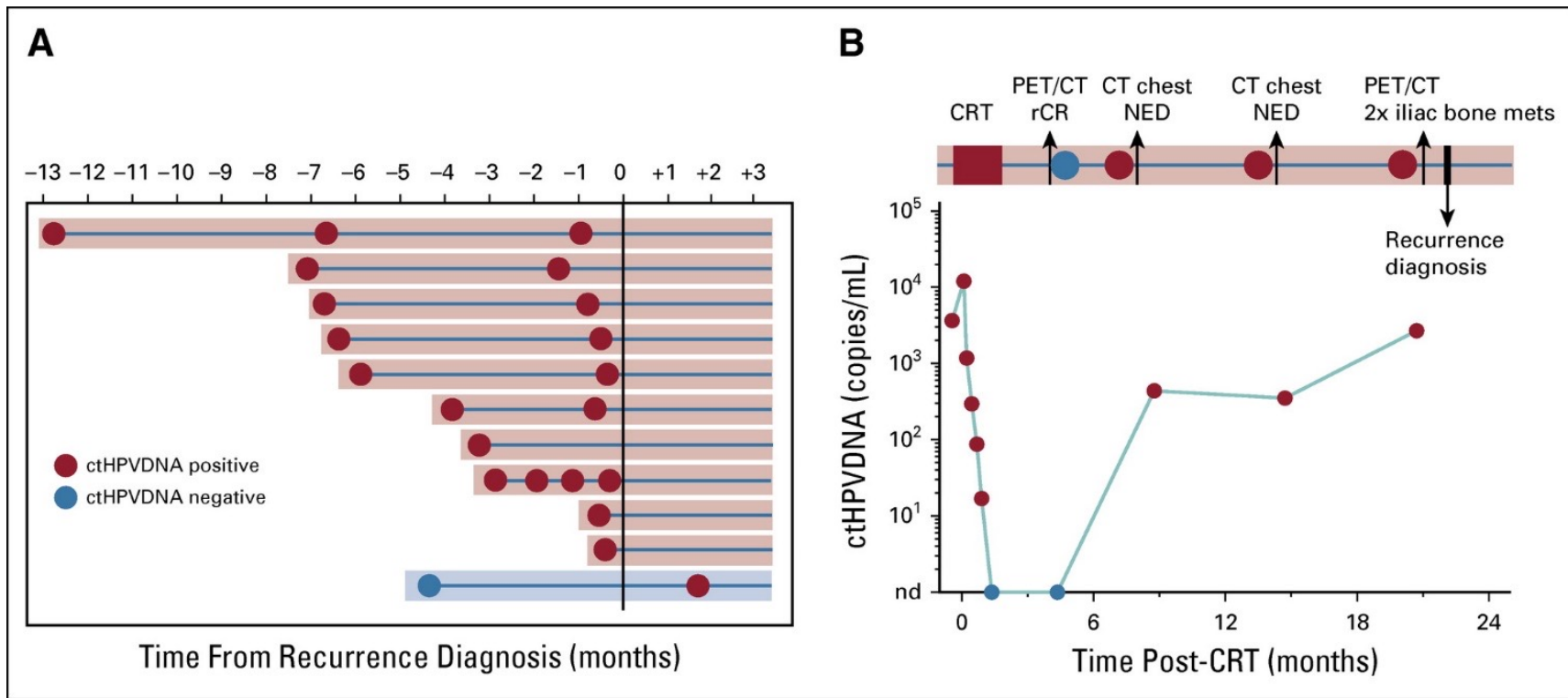
# Longitudinal ctHPVDNA surveillance identifies patients at high risk of disease recurrence



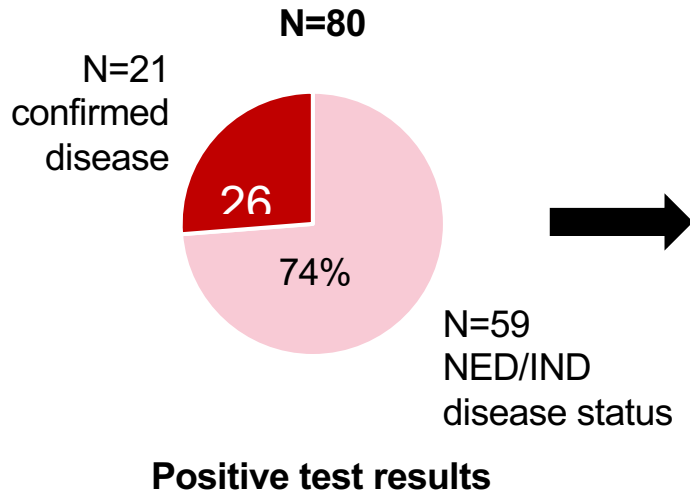
# Patients with two consecutively abnormal ctHPVDNA surveillance tests have a higher risk for disease recurrence



# ctHPVDNA surveillance facilitates early detection of disease recurrence



# Detection of Occult Recurrence



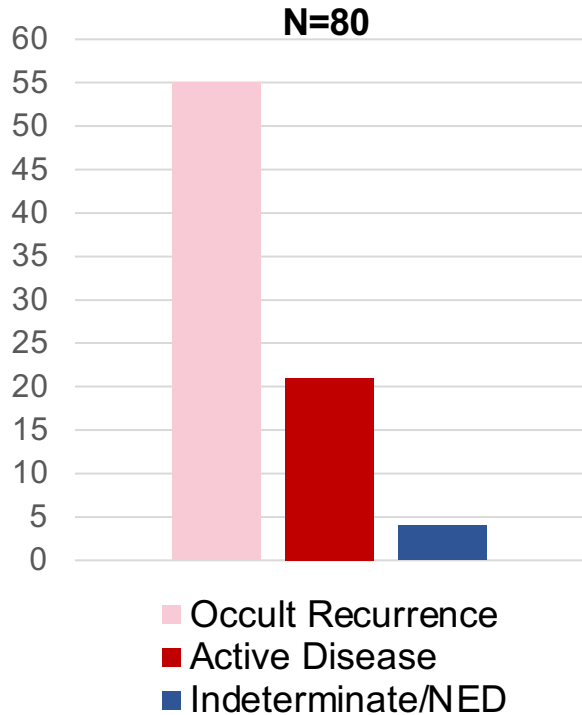
55/59 (93%) later had **proven recurrent, metastatic disease** on imaging and/or biopsy

Of the remaining 4/59 (7%), **2** have **clinically suspicious lesions** (tongue base, pulmonary nodule), and **2** are **clinically NED**

All 4 have TTMV-HPV DNA values ranging from **16-79 frg/mL**



# PPV of TTMV-HPV DNA to Detect Recurrence



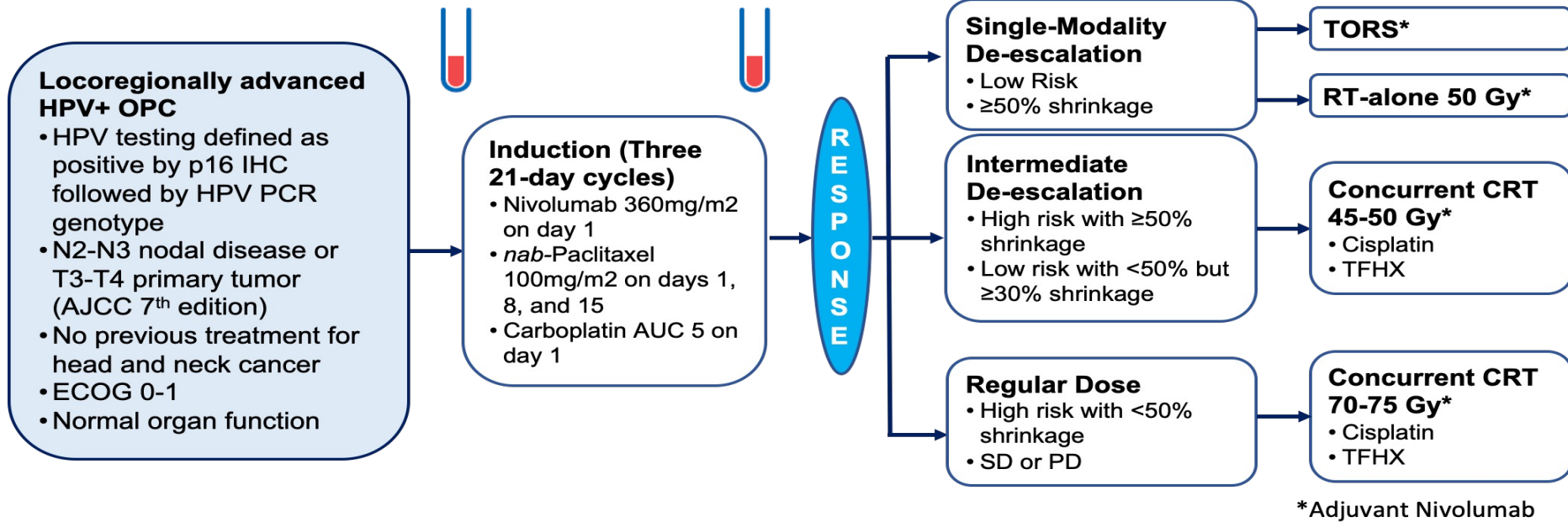
55/59 (93%) later had **proven recurrent, metastatic disease** on imaging and/or biopsy

	Disease	No disease	Total
Positive	76	4	80
Negative		996	996
Total	76	1000	1076

$$55 + 21 = 76/80 = \mathbf{95\% \text{ PPV}}$$

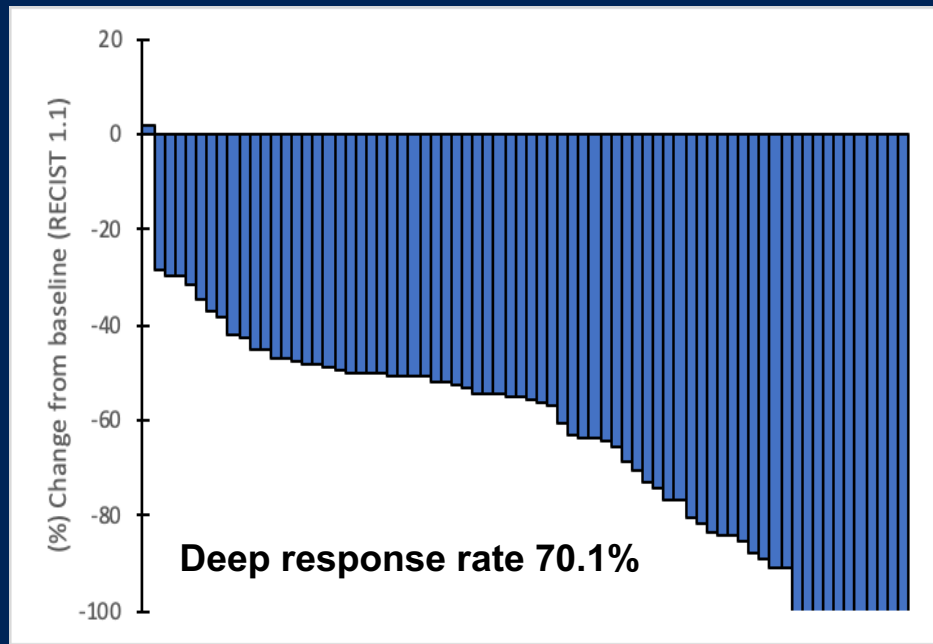


# Optima 2 study design



# Response Following Induction

## Radiographic Response

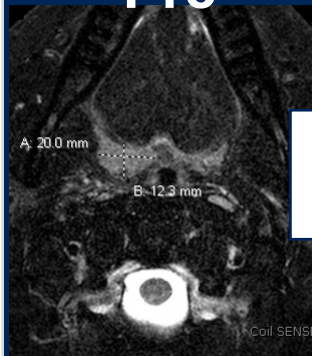


## Pathologic Response

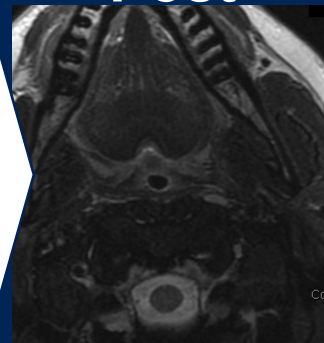
Pathologic complete response rate among TORS patients was 67% (6/9)

### Patient #1

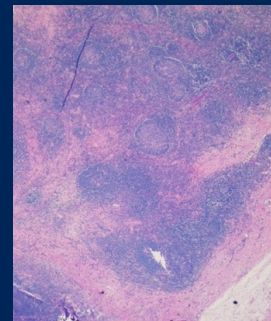
Pre



Post



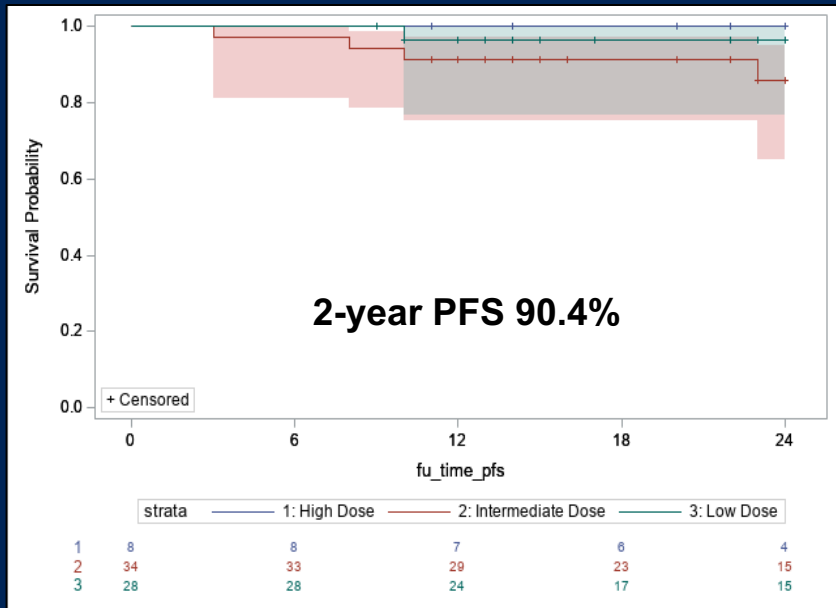
TORS



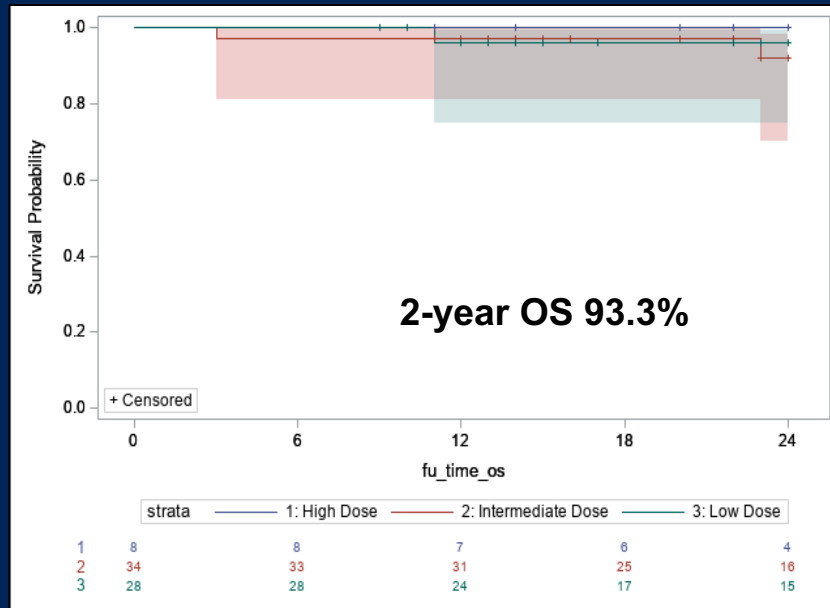


# Survival Outcomes

## Progression-Free Survival

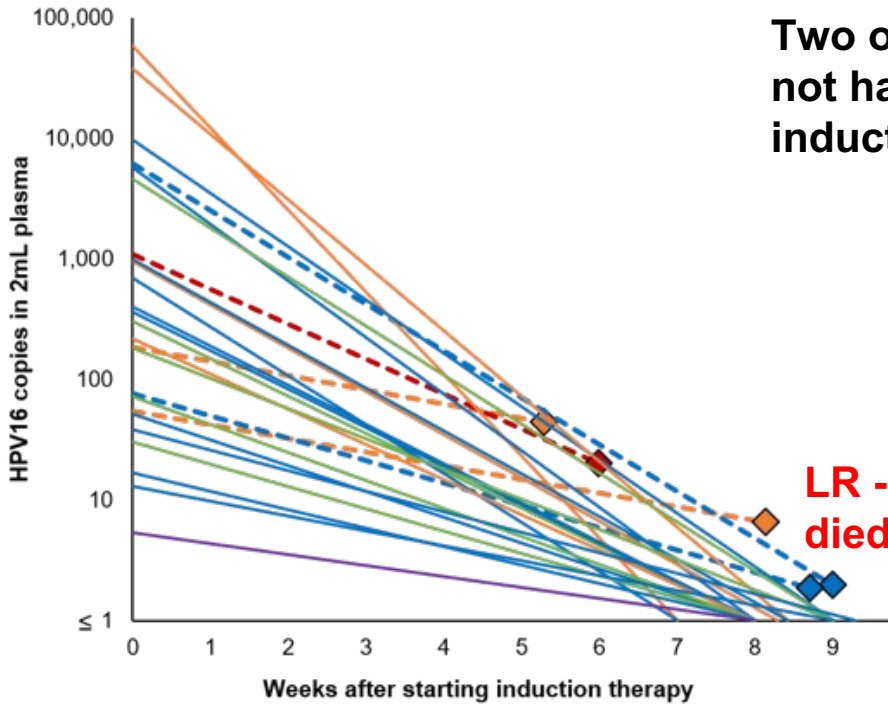


## Overall Survival



Median f/u 23.1 months (IQR 13.7,31.2)

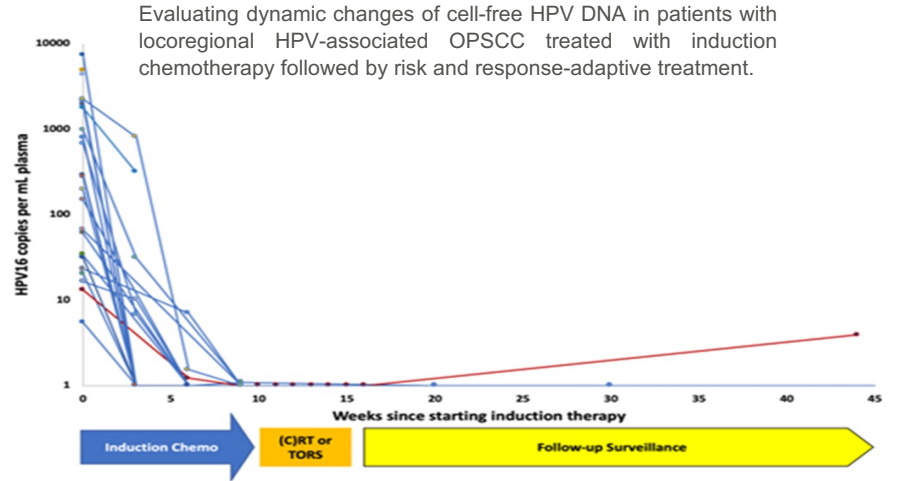
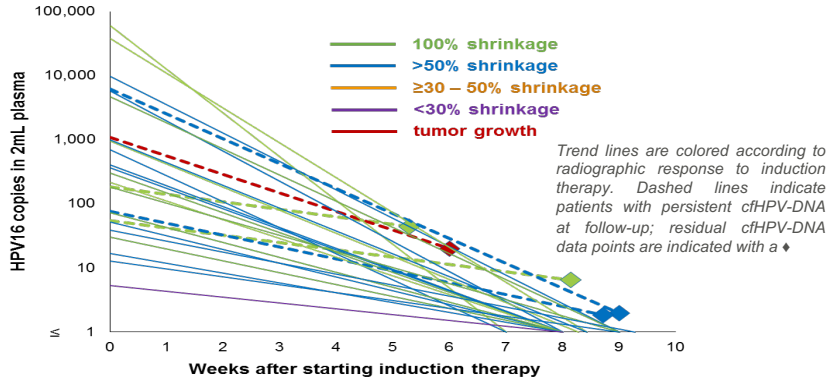
# Clearance of cfDNA with induction may predict treatment failure



Two other recurrences did not have ctDNA post-induction collected.

LR -> RT50 -> recurred -> died

# Preliminary data

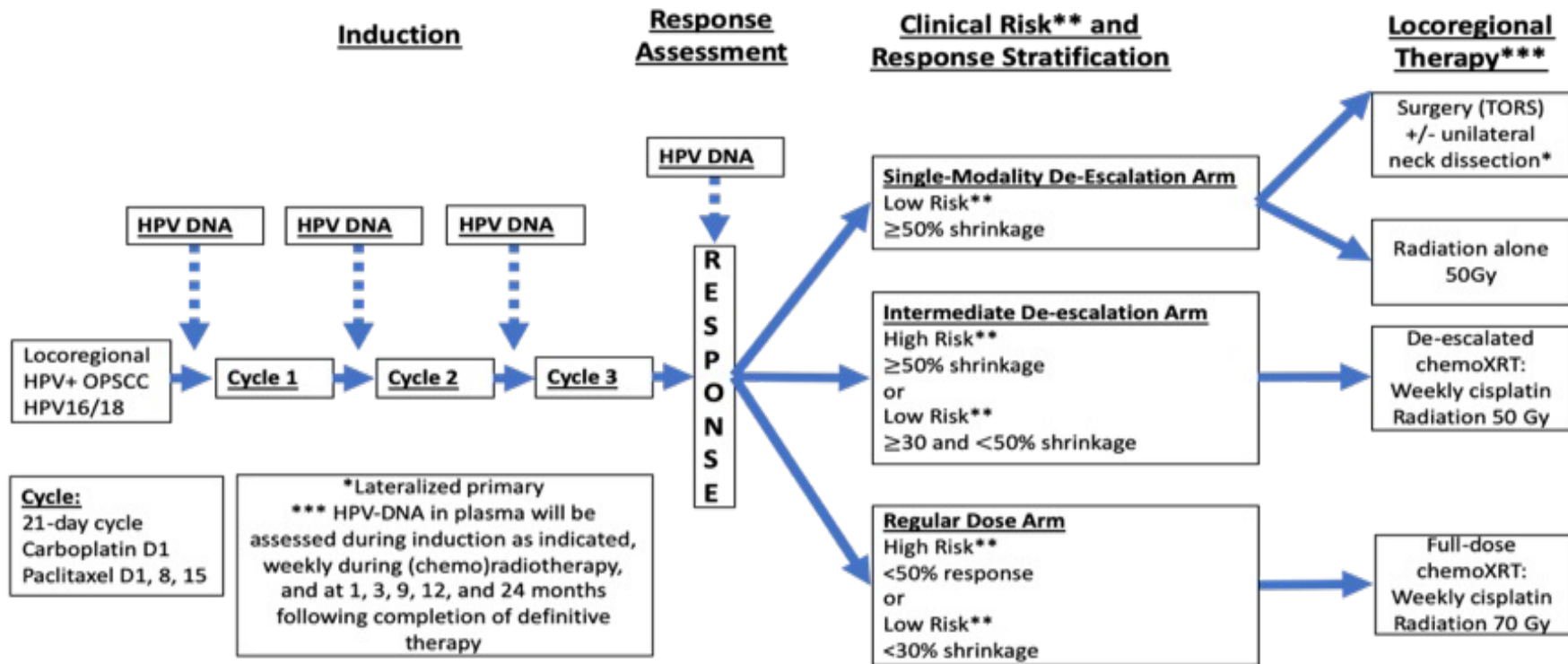


Evaluating dynamic changes of cell-free HPV DNA in patients with locoregional HPV-associated OPSCC treated with induction chemotherapy followed by risk and response-adaptive treatment.

**Figure 1. Serial quantitative cfHPV DNA among first 25 patients on study. Quantitative change in ctDNA during induction chemotherapy, response-adaptive locoregional therapy including deescalated RT +/- chemotherapy or transoral robotic surgery (TORS), and during follow-up.**

- Longitudinal ctHPV-DNA analysis in patients receiving chemotherapy with immunotherapy. Trend lines are colored according to radiographic response to chemotherapy with immunotherapy. Dashed lines indicate patients with persistent ctHPV-DNA at follow-up; residual ctHPV-DNA data points are indicated with a ♦.
- All 25 patients with baseline cfHPV-DNA showed a decrease in cfHPV-DNA level at follow-up, with complete clearance observed in 21/25 of patients, consistent with tumor response (shrinkage) to induction therapy.
- Of 4 patients with persistent cfHPV-DNA: 2 patients progressed on induction therapy, 1 patient demonstrated subsequent recurrence and death, 1 patient demonstrated concern for distant metastasis followed by death.

# Prospective study of dynamic changes of cell-free HPV DNA in HPV+ OPC treated with risk and response-adaptive treatment



# Conclusion

- Early results of low-risk de-escalated arm of OPTIMA 2 suggest that induction chemoimmunotherapy with carboplatin/nab-paclitaxel/nivolumab followed by de-escalated locoregional therapy leads to excellent oncologic outcomes with low rates of acute toxicity.
- High pCR rate to induction chemoimmunotherapy in TORS arm suggests that RT or surgery may be selectively omitted.
- cfHPV-DNA may assist in more precise selection of patients for treatment de-intensification