

# Prise en charge thérapeutique des CBNPC

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# Traitement

## But :

1. Guérir le malade, / obtenir une rémission la plus longue possible.
2. Augmenter la survie.
3. Améliorer et garder une bonne QOL

## Armes :

1. Chirurgie
2. Radiothérapie
3. Traitement médicamenteux
4. Traitement symptomatique

## Stratégie thérapeutique :

1. Traitement de maintenance
2. Traitement de deuxième ligne ,
3. traitement de 3<sup>eme</sup> ligne et au delà.

## Eléments d'évaluation :

1. Réponse au traitement
2. Survie globale ,
3. Survie sans progression
4. Qualité de vie

# Pré requis au traitement

1

## RCP

- Age / Comorbidités.
- PS
- L'extension de la maladie (TNM)
- Le type histologique
- Les drivers moléculaires

- âge, [ $< 70$  ans,  $70 - 89$  ans]
- Comorbidités tel que insuffisance rénale, cardiopathies, neuropathies importantes....
- PS = 0, 1 / PS = 2 / PS = 3, 4.

Choix du protocole thérapeutique médicale

- Cisplatine / carboplatine
- Monothérapie / bithérapie
- Chimiothérapie / Thérapie ciblée (TKI / antiangiogénique) / Immunothérapie.

# Pré requis au traitement

2

## RCP

- Age / Comorbidités.
- PS
- L'extension de la maladie (TNM)
- Le type histologique
- Les drivers moléculaires

On aura 3 grands groupes :

- Stades localisés opérables
- Stades localement avancés non opérables
- Stades métastatiques

# Classification par stade

		Tis	NO	MO
<b>I</b>	IA1	T1mi, T1a	NO	MO
	IA2	T1b	NO	MO
	IA3	T1c	NO	MO
	IB	T2a	NO	MO
<b>II</b>	IIA	T2b	NO	MO
	IIB	T1a, T1b, T1c, T2a, T2b T3	N1 NO	MO MO
<b>III</b>	IIIA	T1a, T1b, T1c, T2a, T2b T3	N2 N1	MO MO
	IIIA	T4	NO, N1	MO
	IIIB	T1a, T1b, T1c, T2a, T2b T3, T4	N3 N2	MO MO
	IIIC	T3, T4	N3	
<b>IV</b>	IVA	Quel que soit T	Quel que soit N	M1a, M1b
	IVB	Quel que soit T	Quel que soit N	M1c

### Stades opérables

<b>I</b>	IA1	T1mi, T1a	N0
	IA2	T1b	N0
	IA3	T1c	N0
	IB	T2a	N0
<b>II</b>	IIA	T2b	N0
	IIB	T1a, T1b, T1c, T2a, T2b T3	N1 N0
<b>III</b>	IIIA	T1a, T1b, T1c, T2a, T2b T3	N2 N1

**BUT du traitement**

Guérir le patient  
Eviter les récurrences

### Stades localement avancés

<b>III</b>	IIIA	T4	N0, N1	M0
	IIIB	T1a, T1b, T1c, T2a, T2b	N3	M0
		T3, T4	N2	M0
IIIC	T3, T4	N3	M0	

Obtenir une rémission  
Eviter les récurrences

### Stades métastatiques

<b>IV</b>	IVA	Quel que soit T	Quel que soit N	M1a, M1b
	IVB	Quel que soit T	Quel que soit N	M1c

Obtenir une rémission  
Prolonger la SSP et SG  
Améliorer et maintenir QOL

## Stades opérables

<b>I</b>	IA1	T1mi, T1a	N0
	IA2	T1b	N0
	IA3	T1c	N0
	IB	T2a	N0
<b>II</b>	IIA	T2b	N0
	IIB	T1a, T1b, T1c, T2a, T2b T3	N1 N0
<b>III</b>	IIIA	T1a, T1b, T1c, T2a, T2b T3	N2 N1

## Armes thérapeutique

Chirurgie  
+ Chimiothérapie  
+/- Radiothérapie

## Stades localement avancés

<b>III</b>	IIIA	T4	N0, N1	M0
	IIIB	T1a, T1b, T1c, T2a, T2b	N3	M0
		T3, T4	N2	M0
IIIC	T3, T4	N3	M0	

CT/RT concomitante ou  
séquentielle

## Stades métastatiques

<b>IV</b>	IVA	Quel que soit T	Quel que soit N	M1a, M1b
	IVB	Quel que soit T	Quel que soit N	M1c

Chimiothérapie (1970)  
Thérapie ciblée (2004)  
Immunothérapie (2016)  
Traitement symptomatique

# Pré requis au traitement

3

## RCP

- Age / Comorbidités.
- PS
- L'extension de la maladie (TNM)
- Le type histologique
- Les drivers moléculaires



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## NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

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CLINICAL PRESENTATION	HISTOLOGIC SUBTYPE <sup>a</sup>	BIOMARKER TESTING <sup>mm</sup>
Advanced or metastatic disease	<b>Adénocarcinome</b> Carcinome grandes cellules CBNPC NOS	<ul style="list-style-type: none"> <li>• Molecular testing, including:                             <ul style="list-style-type: none"> <li>▸ EGFR mutation (category 1), ALK (category 1), KRAS, ROS1, BRAF, NTRK1/2/3, METex14 skipping, RET</li> <li>▸ Testing should be conducted as part of broad molecular profiling<sup>nn</sup></li> </ul> </li> <li>• PD-L1 testing (category 1)</li> </ul>
	Squamous cell carcinoma	<ul style="list-style-type: none"> <li>• Consider molecular testing, including:<sup>oo</sup> <ul style="list-style-type: none"> <li>▸ EGFR mutation, ALK, KRAS, ROS1, BRAF, NTRK1/2/3, MET exon 14 skipping, RET</li> <li>▸ Testing should be conducted as part of broad molecular profiling<sup>nn</sup></li> </ul> </li> <li>• PD-L1 testing (category 1)</li> </ul>

Establish histologic subtype<sup>a</sup> with adequate tissue for molecular testing (consider rebiopsy<sup>ll</sup> if appropriate)

Smoking cessation counseling

Integrate palliative care<sup>c</sup> (See [NCCN Guidelines for](#)



# NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

## PRINCIPLES OF PATHOLOGIC REVIEW

### Classification

The types of NSCLC are: adenocarcinoma, squamous cell carcinoma, adenosquamous carcinoma, large cell carcinoma, and sarcomatoid carcinoma.

▶ **Squamous cell carcinoma:** A malignant epithelial tumor that either shows keratinization and/or intercellular bridges, or a morphologically undifferentiated NSCC that expresses immunohistochemical markers of squamous cell differentiation.

▶ **Adenocarcinoma:**

- ◊ For small (<3 cm), resected lesions, determining extent of invasion is critical.
  - Adenocarcinoma in situ (AIS; formerly BAC): A small ( $\leq 3$  cm) localized nodule with lepidic growth, mostly non-mucinous, although mucinous types can occur. Multiple synchronous AIS tumors can also occur.
  - Minimally invasive adenocarcinoma (MIA): A small ( $\leq 3$  cm) solitary adenocarcinoma with a predominantly lepidic pattern and  $\leq 5$  mm invasion in greatest dimension. MIA is usually non-mucinous, but rarely may be mucinous. MIA is, by definition, solitary and discrete.
  - Invasive adenocarcinoma: A malignant epithelial tumor with glandular differentiation, mucin production, or pneumocyte marker expression. The tumors show an acinar, papillary, micropapillary, lepidic, or solid growth pattern, with either mucin or pneumocyte marker expression. The invasive adenocarcinoma component should be present in at least one focus measuring  $>5$  mm in greatest dimension.
  - Invasive adenocarcinoma variants: invasive mucinous adenocarcinoma, solid papillary adenocarcinoma, and micropapillary adenocarcinoma.
  - Refer to College of American Pathologists [Protocols](#) for additional information.

▶ **Adenosquamous carcinoma:** A carcinoma showing components of both adenocarcinoma and squamous cell carcinoma. A squamous component constituting at least 10% of the tumor. Definitive diagnosis is based on findings in small biopsies, cytology, or excisional biopsies. For large resections, otherwise squamous should trigger molecular testing.

▶ **Large cell carcinoma:** Undifferentiated NSCC that lacks the cytologic features of adenocarcinoma, or squamous cell carcinoma. The diagnosis requires immunohistochemistry on resection or cytology specimens.

▶ **Sarcomatoid carcinoma** is a general term that includes pleomorphic carcinoma, carcinosarcoma, and pulmonary blastoma. It is best to use the specific term for these entities whenever possible.

- ◊ Pleomorphic carcinoma is a poorly differentiated NSCC that consists of a mixture of spindle and giant cells. Spindle cell carcinoma consists of spindle cells, and giant cell carcinoma consists almost entirely of tumor giant cells.
- ◊ Carcinosarcoma is a malignant tumor that consists of a mixture of carcinoma and sarcoma (e.g., rhabdomyosarcoma, chondrosarcoma, osteosarcoma).
- ◊ Pulmonary blastoma is a biphasic tumor that consists of fetal-type squamous cell carcinoma and embryonal rhabdomyosarcoma.

### Classification histologique OMS 2015

Carcinome épidermoïde	Pont d'union +/- Kératine
Adénocarcinome	Différenciation glandulaire Production mucine (BA +)
Carcinome adéno-squameux	Double composante épidermoïde et adénocarcinome avec pour chaque composante au moins 10 % de la Tumeur
Carcinome à grandes cellules	Pas de différenciation Diagnostic sur pièce opératoire
Carcinome	Déliomorphe / Carcino-sarcome

## PRINCIPLES OF PATHOLOGIC REVIEW

### Immunohistochemistry

- Judicious use of IHC is strongly recommended to preserve tissue for molecular testing, most notably in small specimens. When adenocarcinoma or squamous cell carcinomas are poorly differentiated, the defining morphologic criteria that would allow for specific diagnosis may be inconspicuous or absent. In this case, IHC or mucin staining may be necessary to determine a specific diagnosis.
- In small specimens, a limited number of immunostains with one lung adenocarcinoma marker (TTF1, napsin A) and one squamous carcinoma marker (p40, p63) should suffice for most diagnostic problems. Virtually all tumors that lack squamous cell morphology and show co-expression of p63 and TTF1 are preferably classified as adenocarcinoma. A simple panel of TTF1 and p40 may be sufficient to classify most NSCC-NOS cases.
- Testing for NUT expression by IHC should be considered in all poorly differentiated carcinomas that lack glandular differentiation or specific etiology, particularly in non-smokers or in patients presenting at a young age, for consideration of a pulmonary NUT carcinoma.
- IHC should be used to differentiate primary lung adenocarcinoma from squamous cell carcinoma, large cell carcinoma, metastatic carcinoma, and primary pleural mesothelioma (particularly for pleural specimens).
- Primary pulmonary adenocarcinoma:
  - ▶ In patients for whom the primary origin of the carcinoma is uncertain, an appropriate panel of immunohistochemical stains is recommended to assess for metastatic carcinoma to the lung.

	C épidermoïde	Adénocarcinome
Morphologie	- Pont d'union - +/- Kératine	Morphologie glandulaire Mucine (BA +)
IHC	P63 ou p40	TTF1 ou Napsin A

- ▶ TTF1 is a homeodomain-containing nuclear transcription protein of the embryonal and mature lung and thyroid. TTF1 immunoreactivity is seen in non-mucinous adenocarcinoma subtypes. Metastatic adenocarcinoma to the thyroid malignancies, in which case thyroglobulin and PAX8 are also used. Metastatic adenocarcinoma to the gynecologic tract, pancreatobiliary) have been noted, and may be of value in establishing correlation with clinical and radiologic features.
- ▶ Napsin A—an aspartic proteinase expressed in normal type II pneumocytes—is expressed in >80% of lung adenocarcinomas and may be a useful adjunct to TTF1.
- ▶ The panel of TTF1 (or alternatively napsin A) and p40 (or alternatively p63) may be useful in refining the diagnosis to either adenocarcinoma or squamous cell carcinoma in small biopsy specimens previously classified as NSCC NOS.

# Pré requis au traitement

4

## RCP

- Age / Comorbidités.
- PS
- L'extension de la maladie (TNM)
- Le type histologique
- Les drivers moléculaires

Stade Avancé / métastatique

Tous stades



## NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

### CLINICAL PRESENTATION

### HISTOLOGIC SUBTYPE<sup>a</sup>

### BIOMARKER TESTING<sup>mm</sup>

- EGFR
- ALK
- ROS-1
- BRAF
- NTRK 1/2/3
- MET exon 14
- RET
- KRAS
- PDL-1

- Adenocarcinoma
- Large cell
- NSCLC not otherwise specified (NOS)

Squamous cell carcinoma

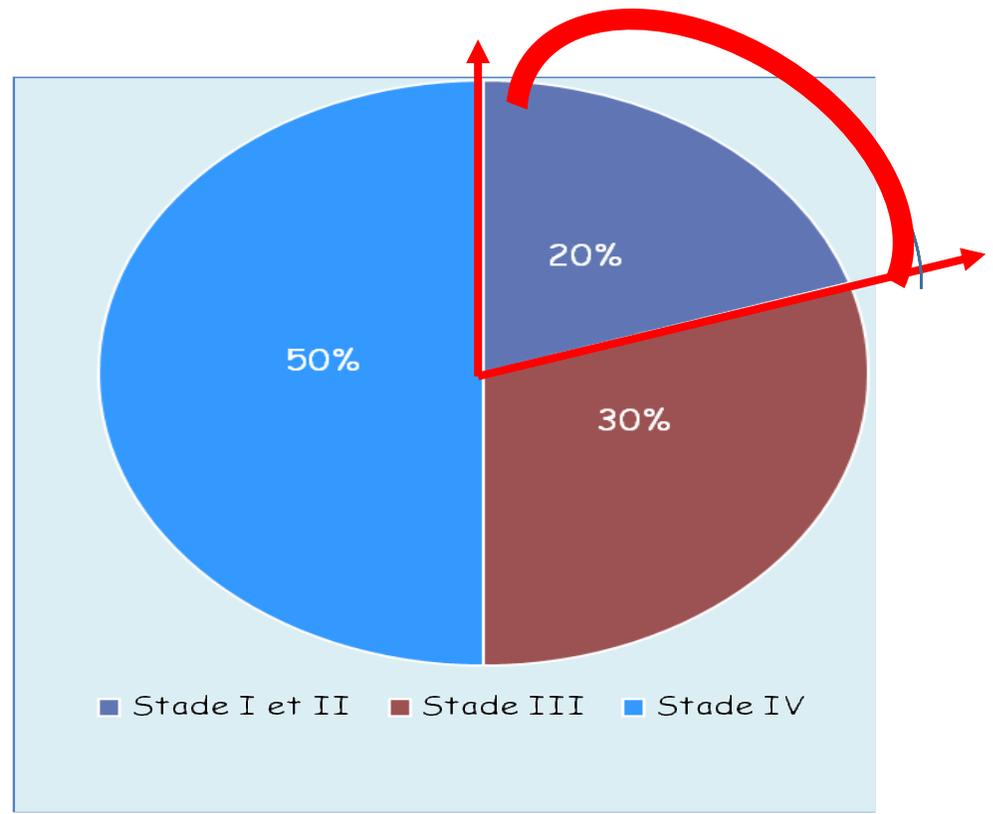
- Molecular testing, including:
  - EGFR mutation (category 1), ALK (category 1), KRAS, ROS1, BRAF, NTRK1/2/3, METex14 skipping, RET
  - Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)

- Consider molecular testing, including:<sup>oo</sup>
  - EGFR mutation, ALK, KRAS, ROS1, BRAF, NTRK1/2/3, MET exon 14 skipping, RET
  - Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)

**Stades localisés**

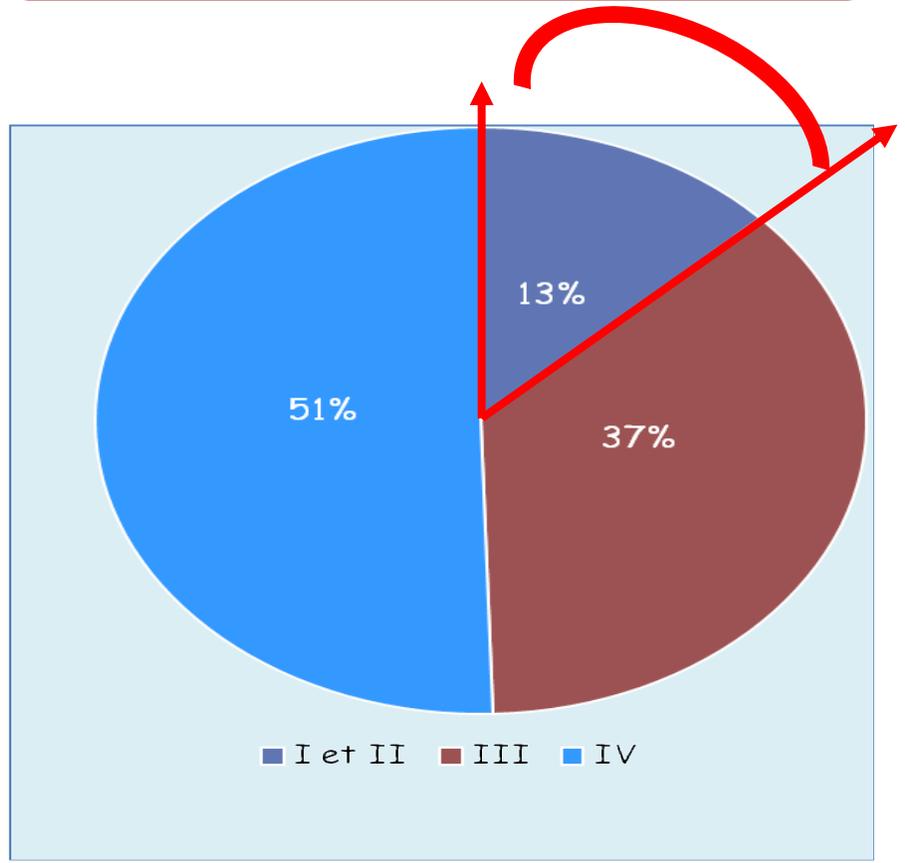
**Les indispensables**

**Dans le monde**



Fry WA, et al. Cancer. 1996;77:1949–1995

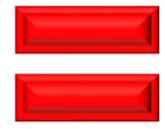
**Algérie**



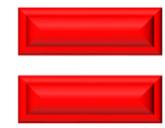
CPMC, HDJ Oncologie médicale. 2000

**Les indispensables**

Stades localisés			
<b>I</b>	IA1	T1mi, T1a	N0
	IA2	T1b	N0
	IA3	T1c	N0
	IB	T2a	N0
<b>II</b>	IIA	T2b	N0
	IIB	T1a, T1b, T1c, T2a, T2b T3	N1 N0
<b>III</b>	IIIA	T1a, T1b, T1c, T2a, T2b	N2
		T3	N1



**Stade opérable**

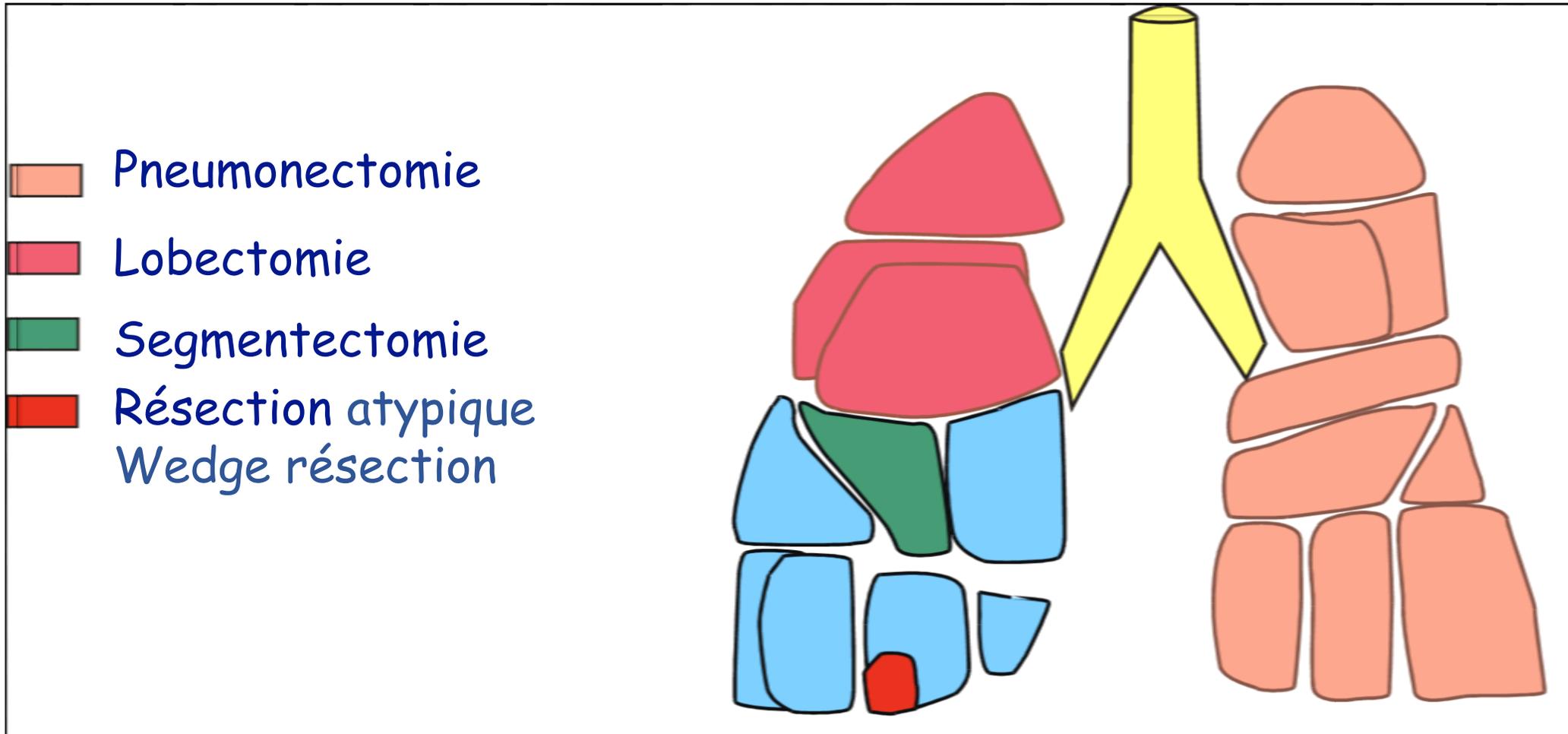


**Chirurgie  
+/- Chimiothérapie  
+/- Radiothérapie**

1

Chirurgie = 1- Résection Tumorale  
2- Curage ganglionnaire

Chirurgie  
=  
Seul traitement curateur



# NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

**Résection tumorale**

## PRINCIPLES OF SURGICAL TH

- Determination of resectability, surgical staging, and *pulmonary resection should be a prominent part of their practice.*
- CT and PET/CT used for staging should be within 60 days before proceeding with surgery.
- For medically operable disease, resection is the preferred local treatment modality (including radiofrequency ablation, and cryotherapy). Thoracic surgical oncology consultation should be considered for curative local therapy. In cases where SABR is considered for high-stage disease, a radiation oncologist is recommended.
- The overall plan of treatment as well as needed imaging studies should be determined by the multidisciplinary team.
- Thoracic surgeons should actively participate in multidisciplinary discussions and decisions (including multidisciplinary clinic and/or tumor board).
- Patients who are active smokers should be provided counseling and smoking cessation. While active smokers have a mildly increased incidence of postoperative pulmonary complications, this does not constitute a prohibitive risk for surgery. Surgeons should not deny surgery to patients solely because of active smoking. Smoking cessation should be a predominant therapy for patients with early-stage lung cancer.

### Resection

- Anatomic pulmonary resection is preferred for the majority of patients with NSCLC.
- Sublobar resection - Segmentectomy and wedge resection should achieve parenchymal margins.
- Sublobar resection should also sample appropriate N1 and N2 lymph node stations, increasing the surgical risk.
- Segmentectomy (preferred) or wedge resection is appropriate in selected patients:
  - ▶ Poor pulmonary reserve or other major comorbidity that contraindicates lobectomy
  - ▶ Peripheral nodule<sup>1</sup> ≤ 2 cm with at least one of the following:
    - ◊ Pure AIS histology
    - ◊ Nodule has ≥50% ground-glass appearance on CT
    - ◊ Radiologic surveillance confirms a long doubling time (≥400 days)
- VATS or minimally invasive surgery (including robotic-assisted approaches) should be considered in patients without significant surgical contraindications, as long as there is no compromise of standard oncologic outcomes.
- In high-volume centers with significant VATS experience, VATS lobectomy in selected patients is associated with decreased pain, reduced hospital length of stay, more rapid return to function, fewer complications, and improved quality of life outcomes.
- Lung-sparing anatomic resection (sleeve lobectomy) is preferred over pneumonectomy when complete resection is achieved.
- T3 (invasion) and T4 local extension tumors require en-bloc resection of the involved structures. If complete resection is uncertain, consider obtaining an additional surgical margin.

Margins and Nodal Assessment (see [NSCL-B 2 of 4](#))

<sup>1</sup>Peripheral is defined as the outer one third of the lung parenchyma.

## Principes généraux pour une bonne chirurgie cancer bronchique

- 1- Chirurgien expert
- 2- TDM et Pet-scan ≤ 60 jours
- 3- Résection anatomique
- 4- Si résection sub-lobaire, segmentaire ou wedge → marge de résection ≥ 2 cm ou ≥ taille du nodule.
- 5- Curage gg stations N1 et N2
- 6- Segmentectomie ou wedge :
  - Si comorbidités et Fonction respiratoire CI lobectomie
  - Si nodule périphérique ≤ 2 cm avec au moins un des éléments suivant :
    - histo = ADK in situ
    - nodule ≥ 50 % composante en verre dépoli à la TDM.
    - si surveillance radiologique confirme un temps de dédoublement ≥ 400 jours.

1

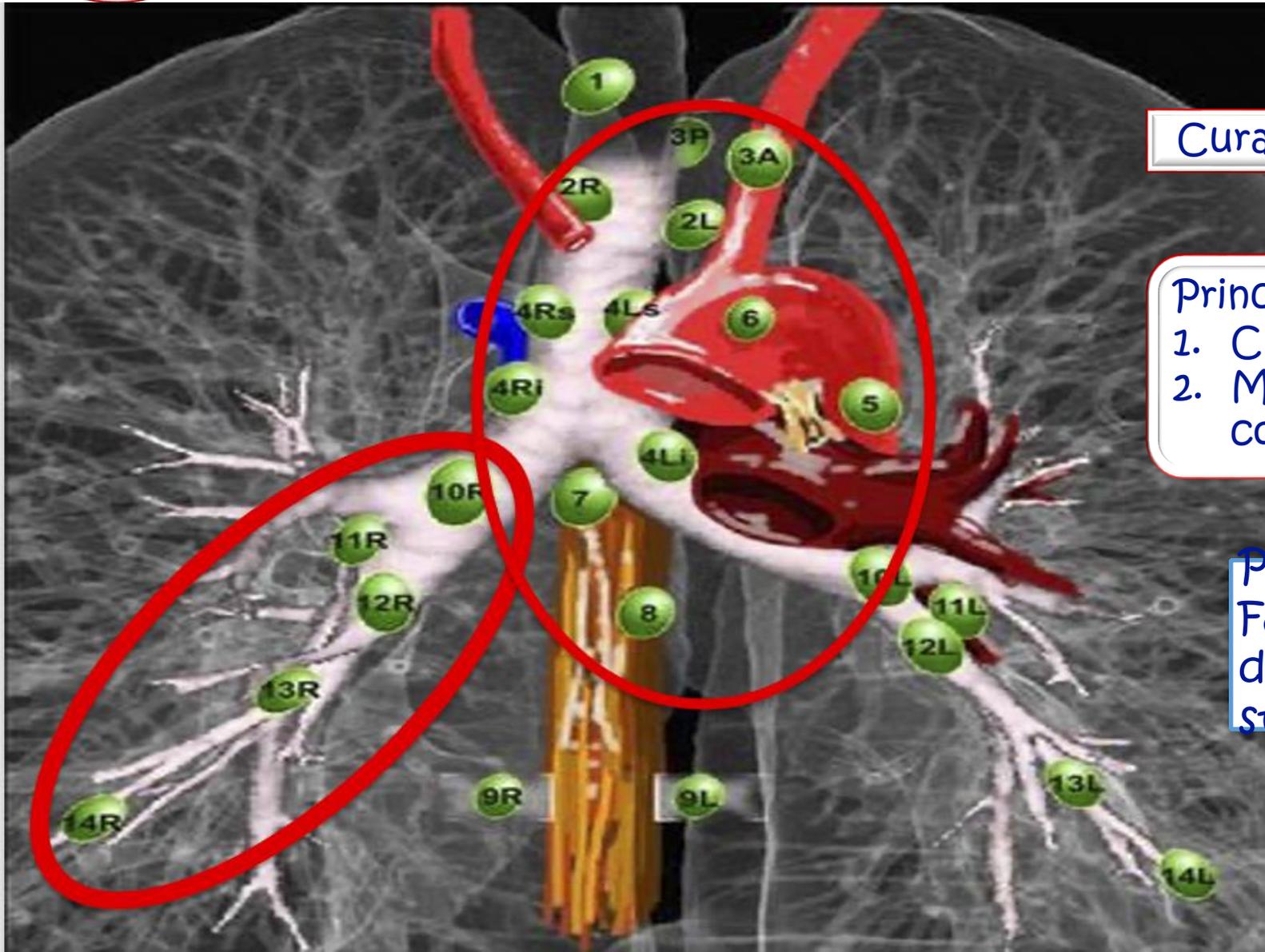
# Chirurgie = 1- Résection Tumorale 2- Curage ganglionnaire

Curage ganglionnaire doit être systématique

Principe d'un bon curage gg

1. Curage N1 et N2 / routine
2. Minimum de 3 stations N2, curage gg complet.

Pour les patients No ou N1 =  
Faire un prélèvement systématique  
de 1 à 2 ganglions de chaque  
station gg



## Curage ganglionnaire

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mediastinal lymph node dissection did not improve survival.<sup>348,349</sup> Thus, systematic lymph node sampling is appropriate during pulmonary resection; one or more nodes should be sampled from all mediastinal stations. For right-sided cancers, an adequate mediastinal lymphadenectomy should include stations 2R, 4R, 7, 8, and 9. For left-sided cancers, stations 4L, 5, 6, 7, 8, and 9 should be sampled.<sup>348</sup> Patients should have N1 and N2 node resection and mapping (American Thoracic Society map) with a minimum of 3 N2 stations sampled or a complete lymph node dissection.<sup>135</sup> The lymph node map from the IASLC may be useful.<sup>350</sup> Formal ipsilateral mediastinal lymph node dissection is indicated for patients undergoing resection for stage IIIA (N2) disease. For patients undergoing sublobular resection, the appropriate N1 and N2 lymph node stations should be sampled unless not technically feasible because sampling would substantially increase the surgical risk.

### A gauche :

- 4L para trachéaux inférieur
- 5 sous aortique
- 6 para aortique
- 7 sous Carénaire
- 8 para œsophagien
- 9 ligament pulmonaire

### A droite :

- 2R para trachéaux supérieur
- 4R para trachéaux inférieur
- 7 sous Carénaire
- 8 para œsophagien
- 9 ligament pulmonaire

## Pourquoi traitement adjuvant ?

La survie a long terme des patients ayant bénéficié d'une chirurgie est décevante du fait de **rechutes soit locale ou à distance**

TNM	Survie a 5 ans % CTNM	Survie a 5 ans % pTNM	Récidive locale %	Récidive a distance %
IA = T1a, b No Mo	50	73	10	15
IB = T2a No Mo	43	58	10	30
IIA = T1a, b N1 Mo T2 N1 Mo	36	46		
IIB = T2b N1 Mo T3 No Mo	25 22	36 38	12	40
IIIA = T1, 2 N2 Mo T3 N1 N2 Mo	19	24 25	15	60

Pour améliorer ces résultats et éviter les récurrences locale et à distance, l'idée est née d'associer un traitement adjuvant par CT et/ou RT.

# Stades localisés

But du traitement adjuvant

Eviter les récurrences à distance



Chimiothérapie

Eviter les récurrences locales



Radiothérapie

Essais	n	Stades	Médiane de survie		Survie a 5 ans		p
			Contrôle	Experimental	Contrôle	Experimental	
Intergroupe 0115	488	II, IIIA	37,9	38,8	33	39	0.56
Big Lung Trial	381	I, II, IIIA	24,7	27	-	-	NS
ALPI-EORTC	1209	I, II, IIIA	48	55,2	-	-	0,58
<b>IALT</b>	<b>1867</b>	<b>I, II, IIIA</b>	<b>44,4</b>	<b>50,8</b>	<b>40,4</b>	<b>44,5</b>	<b>&lt; 0,03</b>
UFT	999	I					
<b>Intergroupe JBR10</b>	<b>482</b>	<b>IB, II</b>	<b>73</b>	<b>94</b>	<b>54</b>	<b>69</b>	<b>0,009</b>
CALBG 9633	344	IB	78	95	57	59	0,37
<b>ANITA</b>	<b>840</b>	<b>IB, II, IIIA</b>	<b>43,7</b>	<b>65,7</b>	<b>42,6</b>	<b>51,2</b>	<b>0,017</b>
Dosell et al	110	IB	41,6	44,8	30	28	0,003

- Pour les tumeurs stade IA = aucun bénéfice n'est démontré
  - Pour les tumeurs stade IB =
    - Deux études de sous groupes (ANITA et Intergroupe JBR 10) ne démontrent pas de bénéfice.
    - La méta-analyse LACE montre une tendance au bénéfice mais qui est non significative.
    - L'étude CALGB 9633 montre un bénéfice pour les tumeurs de 4 cm
- Ces résultats ne sont pas suffisant pour faire une CT adjuvante dans les tumeurs stade IB.
- Mais il semble être raisonnable de la proposer pour les tumeurs de plus de 4 cm

CT adjuvante  
Stade II IIIA

# Stades localisés

Plusieurs essais sont franchement positive et montrent un bénéfice absolu en terme de survie à 5 ans Variant entre 4 à 15 % avec les CT à base de platine

- Etude IALT ( I II et III, CT avec CDDP)  
Bénéfice de survie à 5 ans de 4,1 %
- Etude NCIC JBR 10 ( IB et II, CDDP + VNR)  
Gain de survie à 5 ans de 15 %
- Etude CALGB 9633 (IB, Carbo + TXL)  
Gain de survie à 4 ans de 12 %
- Essai ANITA  
SM 43,7 mois ----- 65, 7 mois (CDDP VNR)

Deux grandes méta-analyses :

- Medical research Council portant sur 8147 patients , 30 essais
- Lung Adjuvant Cisplatin Evaluation LACE portant sur 4584 patients, 5 essais

Confirme ce bénéfice en survie entre de 4 à 5,3 % en faveur du bras CT adjuvante

Mais ce gain de survie, concerne essentiellement les stades qui s'accompagnent d'une extension ganglionnaire stade II et IIIA

- Le risque de **récidive locorégionale** est compris entre 15 à 40 % selon les stades
- D'où l'intérêt de la radiothérapie adjuvante
- De nombreuses études randomisées ont évalué cette option.
- La méta-analyse PORT a montré que la RT adjuvante bien qu'elle diminue le risque de récidive locale de près de 25 %, a un effet défavorable sur la survie globale.
- Cet **effet défavorable** était plus prononcé en cas de tumeur de stade I II et NO N 1.

- Pas d'indication de radiothérapie adjuvante (post-opératoire) pour les N0 et N1.
- RT adjuvante post-opératoire pour les stades IIIA N2 :
  - Diminue le risque de rechute locale.
  - mais pas d'amélioration de la survie.

Mais...

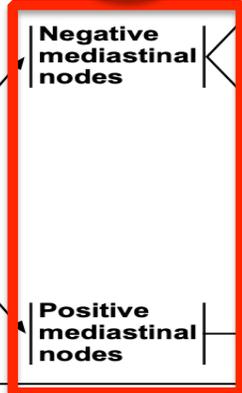
# Stades localisés

CLINICAL ASSESSMENT PRETREATMENT EVALUATION<sup>9</sup>

Stage IA (peripheral T1abc, N0)

- Pulmonary function tests (PFTs) (if not previously done)
- Bronchoscopy (intraoperative preferred)
- Consider pathologic mediastinal lymph node evaluation<sup>h,i</sup>
- FDG PET/CT scan<sup>j</sup> (if not previously done)

1



Patient opérable

Patient inopérable

Résection tumorale + curage gg

Radiothérapie

**Patients stades localisés:**

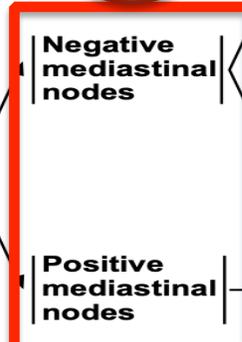
- 1- No ou N1 ?
- 2- Opérable / non opérable ?
- 3- Marge de résection saines ou infiltrées ?

CLINICAL ASSESSMENT PRETREATMENT EVALUATION<sup>9</sup>

Stage IB (peripheral T2a, N0)  
 Stage I (central T1abc-T2a, N0)  
 Stage II (T1abc-2ab, N1; T2b, N0)  
 Stage IIB (T3, N0)<sup>e</sup>  
 Stage IIIA (T3, N1)

- PFTs (if not previously done)
- Bronchoscopy
- Pathologic mediastinal lymph node evaluation<sup>h</sup>
- FDG PET/CT scan<sup>j</sup> (if not previously done)
- Brain MRI with contrast<sup>o</sup> (Stage II, IIIA) (Stage IB [optional])

1



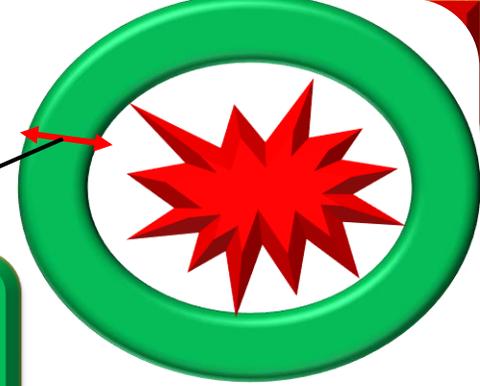
Patient opérable

Patient inopérable

Résection tumorale + curage gg  
 Suivi de CT adjuvante

No = RT suivi CT adj  
 Haut risque Ib-IIB  
 N1 = RTCT suivi  
 Durvalumab

# Stades localisés



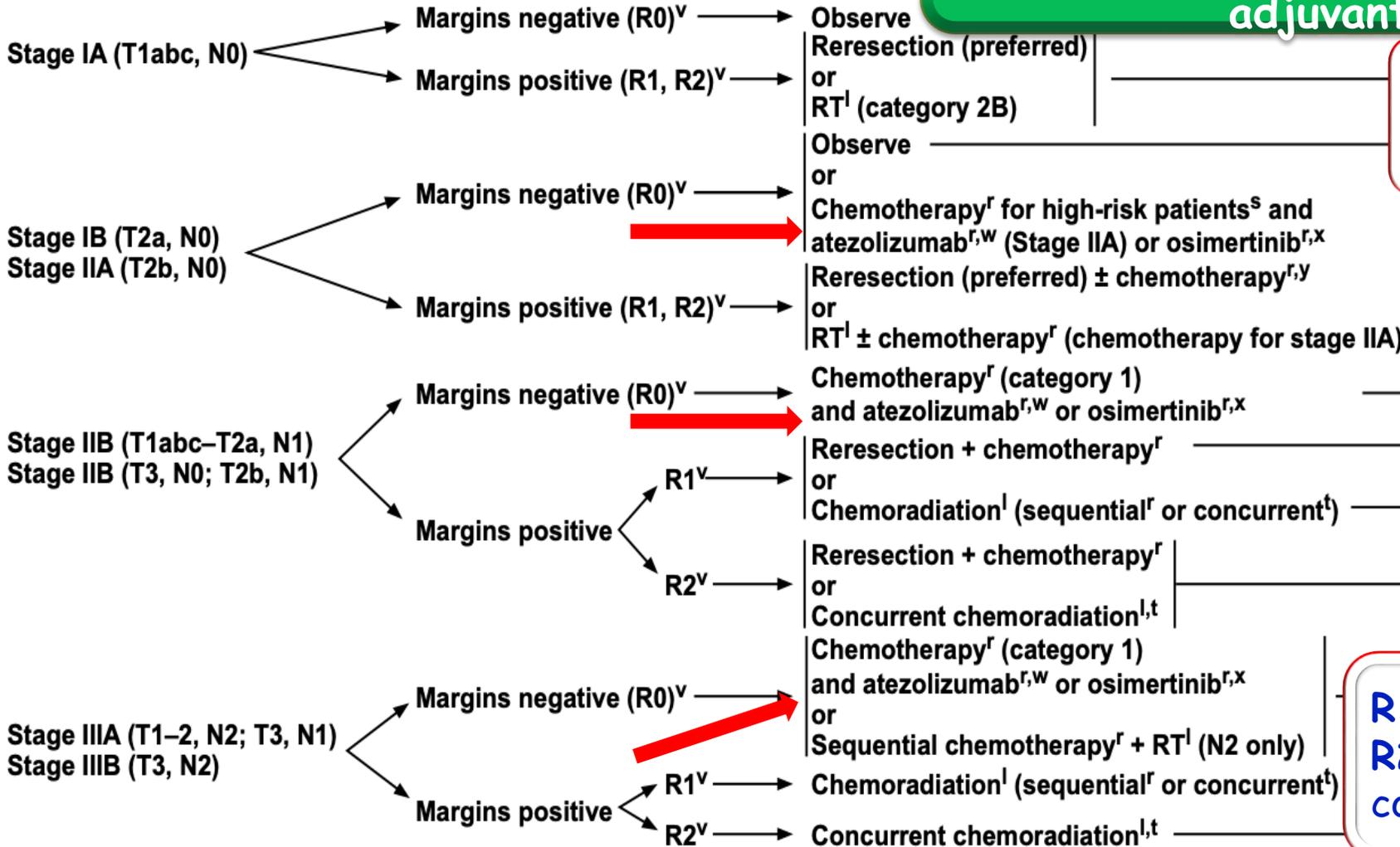
**Marges négative =  
Surveillance ou traitement  
adjuvant**

R0 = Pas de résidu tumoral  
R1 = Résidu tumoral microscopique  
R1 = Résidu tumoral macroscopique

R1 = Reprise Xie +/- CT  
R2 = Reprise chirurgicale ou RT-CT  
concomitante / séquentielle

**FINDINGS AT SURGERY**

**ADJUVANT**



# Stades localisés

- T > 4 cm
- T peu différenciée
- Invasión vasculaire
- Wedge resection
- Atteinte plèvre viscérale
- Statut ganglionnaire Nx

## FINDINGS AT SURGERY

## ADJUVANT TREATMENT

Stage IA (T1abc, N0)  
 T1a ≤ 1 cm  
 T1b > 1 ≤ 2 cm  
 T1b > 2 ≤ 3 cm

Margins negative (R0)<sup>v</sup>  
 Margins positive (R1, R2)<sup>v</sup>

Observe  
 Reresection (preferred)  
 or  
 RT<sup>l</sup> (category 2B)

Surveillance  
 Xie ou RT

Stage IB (T2a, N0)  
 Stage IIA (T2b, N0)  
 T2a > 3 ≤ 4 cm  
 T1b > 4 ≤ 5 cm

Margins negative (R0)<sup>v</sup>  
 Margins positive (R1, R2)<sup>v</sup>

Observe  
 or  
 Chemotherapy<sup>r</sup> for high-risk patients<sup>s</sup> and atezolizumab<sup>r,w</sup> (Stage IIA) or osimertinib<sup>r,x</sup>  
 Reresection (preferred) ± chemotherapy<sup>r,y</sup>  
 or  
 RT<sup>l</sup> ± chemotherapy<sup>r</sup> (chemotherapy for stage IIA)

Surveillance  
 CT si F risque + Atezolizumab ou Osimertinib  
 Xie +/-CT  
 RT +/-CT

Stage IIB (T1abc-T2a, N1)  
 Stage IIB (T3, N0; T2b, N1)  
 T3 > 5 ≤ 7 cm

Margins negative (R0)<sup>v</sup>  
 Margins positive  
 R1<sup>v</sup>  
 R2<sup>v</sup>

Chemotherapy<sup>r</sup> (category 1) and atezolizumab<sup>r,w</sup> or osimertinib<sup>r,x</sup>  
 Reresection + chemotherapy<sup>r</sup>  
 or  
 Chemoradiation<sup>l</sup> (sequential<sup>r</sup> or concurrent<sup>t</sup>)  
 Reresection + chemotherapy<sup>r</sup>  
 or  
 Concurrent chemoradiation<sup>l,t</sup>

CT + Atezo ou Osimertinib  
 RT + CT  
 RT-CT séquentielle ou concomitante  
 Xie + CT  
 RT-CT concomitante

# Essai ADAURA

## Phase III en adjuvant chez les patients atteints de CBNPC EGFR muté

Patients atteints d'un CBNPC réséqué de stade IB, II, IIIA, ayant reçu ou non une chimiothérapie adjuvante

- Âge  $\geq$  18 ans (Japon/Taïwan :  $\geq$  20 ans)
- Performance OMS status : 0/1
- CBNPC non épidermoïde avec del exon 19/L858R
- Imagerie cérébrale, si non réalisée en préopératoire
- Résection complète avec marges négatives
- Délai maximal entre la chirurgie et la randomisation
  - 10 semaines sans chimiothérapie adjuvante
  - 26 semaines si chimiothérapie adjuvante

### Schéma de l'étude

**Stratification :**  
stade (IB vs II vs IIIA)  
EGFRm (del exon 19 vs L858R)  
Ethnie : Asiatiques vs non-Asiatiques

Osimertinib  
80 mg/j

Randomisation  
1:1  
(n = 682)

Placebo  
1/j

- Plan de durée de traitement : 3 ans
- Traitement poursuivi jusqu'à :
  - rechute de la maladie
  - fin du traitement
  - arrêt selon critères de l'étude
- **Suivi :**
  - jusqu'à rechute : S12 et S24, puis toutes les 24 semaines pendant 5 ans, puis tous les ans
  - après la rechute : toutes les 24 semaines pendant 5 ans, puis tous les ans

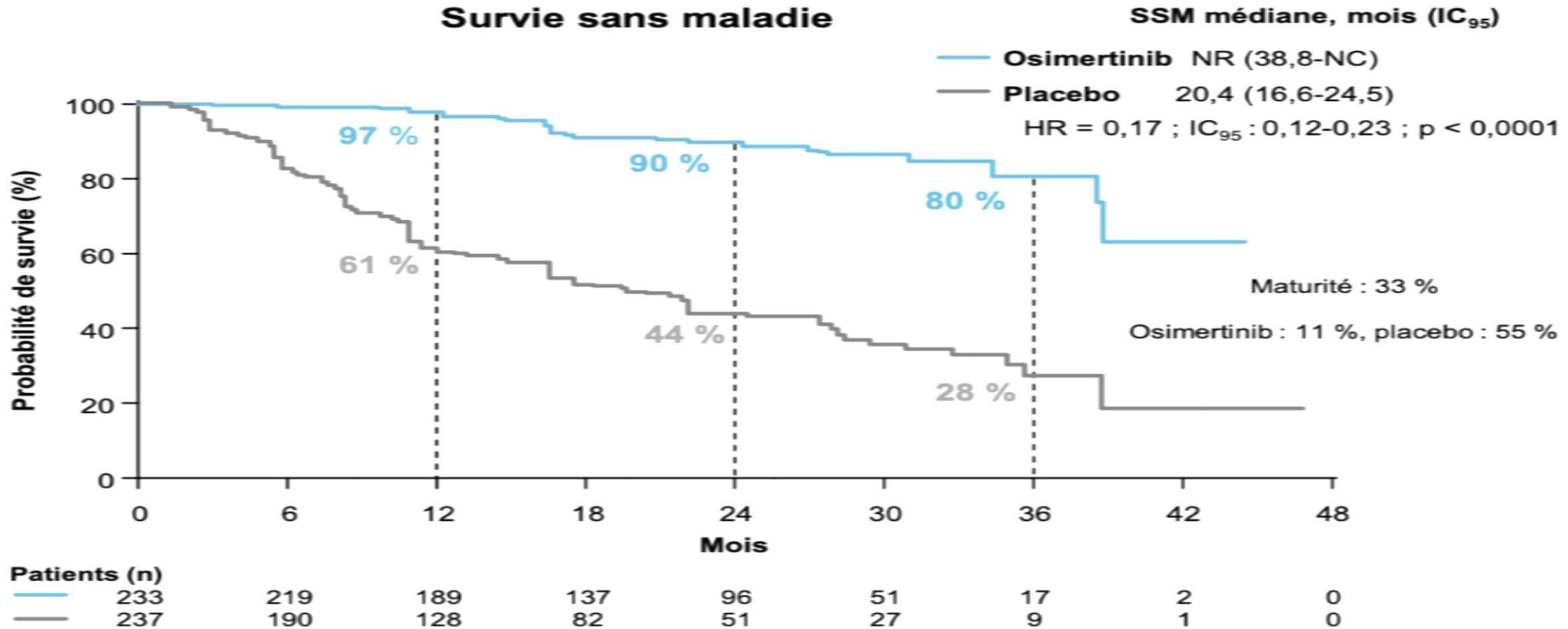
### • Critères d'évaluation

- Principal : survie sans maladie (SSM), évaluée par les investigateurs, chez les patients de stade II-III A
- Secondaires : SSM dans la population globale, taux de SSM à 2, 3, 4 et 5 ans, SG, tolérance, qualité de vie
- **Levée de l'aveugle du fait de l'efficacité observée, réalisée après les recommandations de l'IDMC ; analyse non planifiée**
- **Au moment de la levée d'aveugle, tous les patients avaient été inclus et suivis pendant au moins 1 an**

L'essai randomisé de phase III ADAURA a randomisé 682 patients opérés d'un CBNPC de stade IB à IIIA, avec mutation activatrice de l'EGFR L858R ou délétion dans l'exon 19.

L'étude comparait un groupe traité par un placebo et un traitement de 3 ans maximum par osimertinib (80 mg/j).

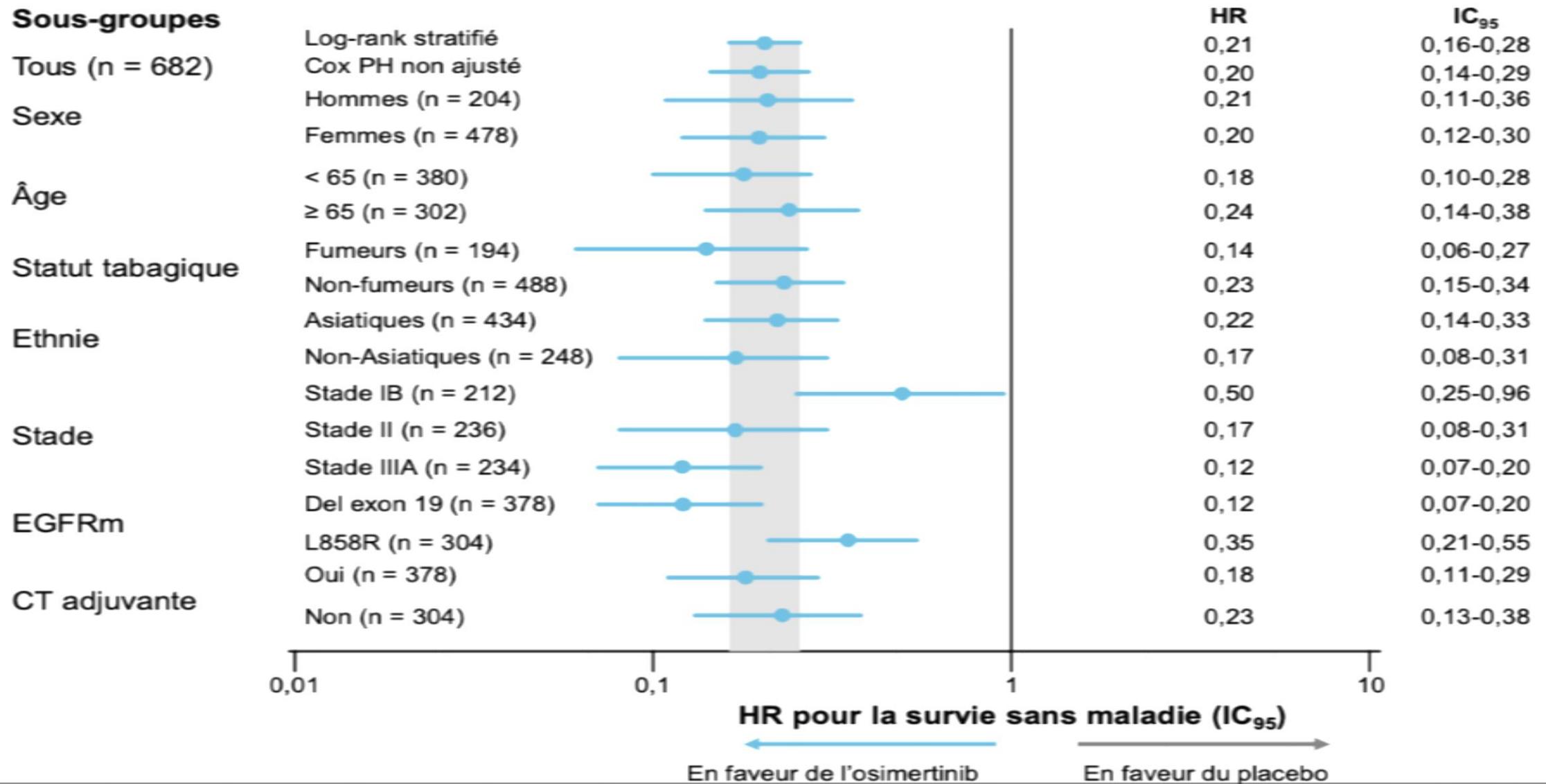
# Essai ADAURA



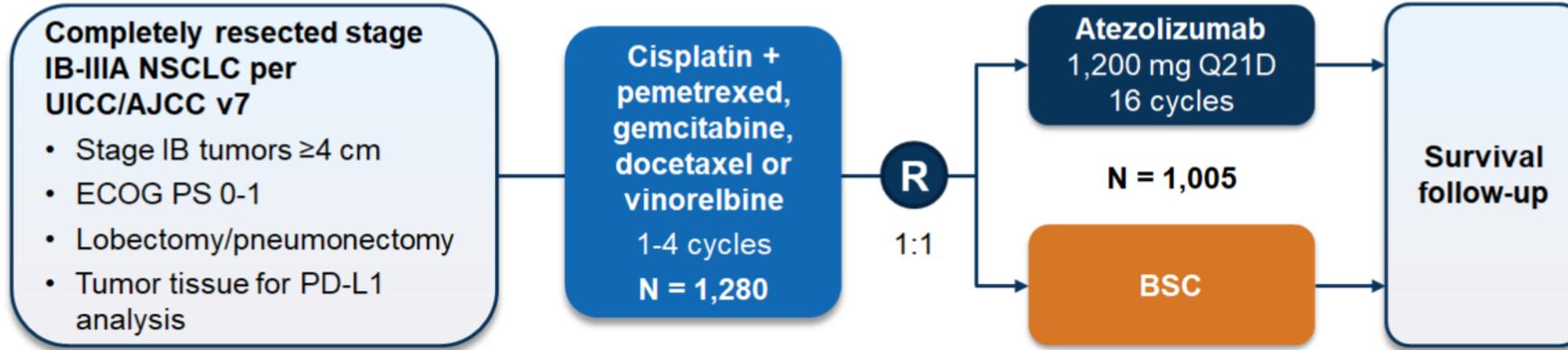
Les résultats montrent un bénéfice majeur de l'osimertinib, avec une réduction du risque de récurrence pour les tumeurs de stades II et IIIA de 83 % par rapport au placebo (HR = 0,17 ; IC95 : 0,12-0,23 ; p < 0,0001).

Ce niveau de bénéfice est l'un des plus élevés rapportés en oncologie thoracique ; la survie sans récurrence à 2 ans était de 90 % avec l'osimertinib et de 44 % avec le placebo.

# Essai ADAURA



# Phase 3 IMpower010 Study: Schema<sup>1</sup>



## Stratification factors

- Sex
- Stage (IB vs II vs IIIA)
- Histology
- PD-L1 tumor expression status<sup>a</sup>: TC2/3 and any IC vs TC0/1 and IC2/3 vs TC0/1 and IC0/1

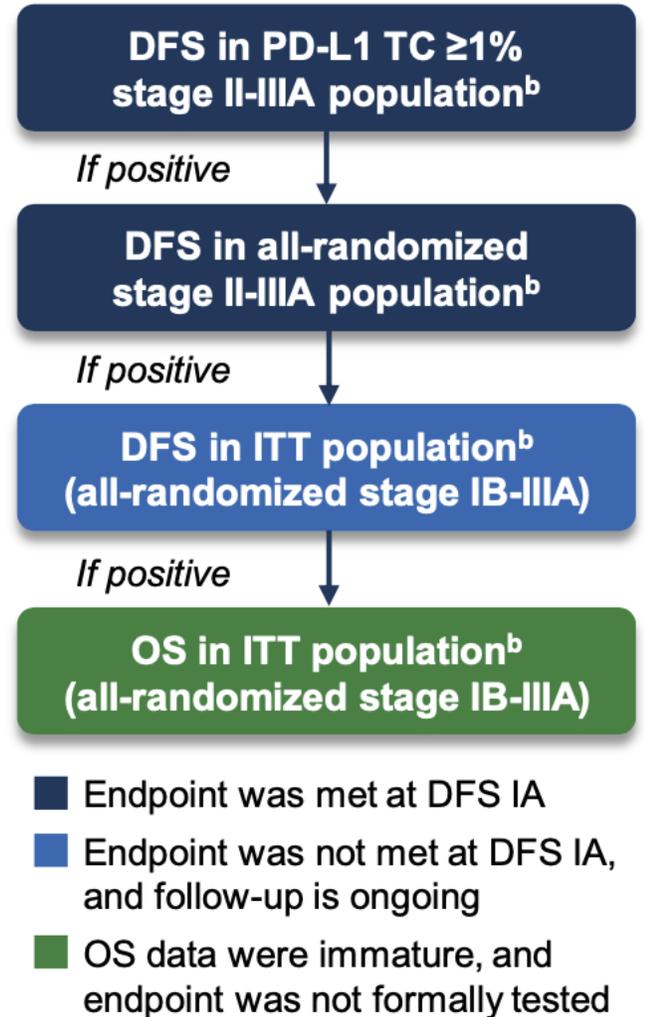
## Primary endpoints

- Investigator-assessed DFS tested hierarchically:
  - PD-L1, TC  $\geq 1\%$  (per SP263) stage II-IIIa population
  - All-randomized stage II-IIIa population
  - ITT population (stage IB-IIIa)

## Exploratory endpoints

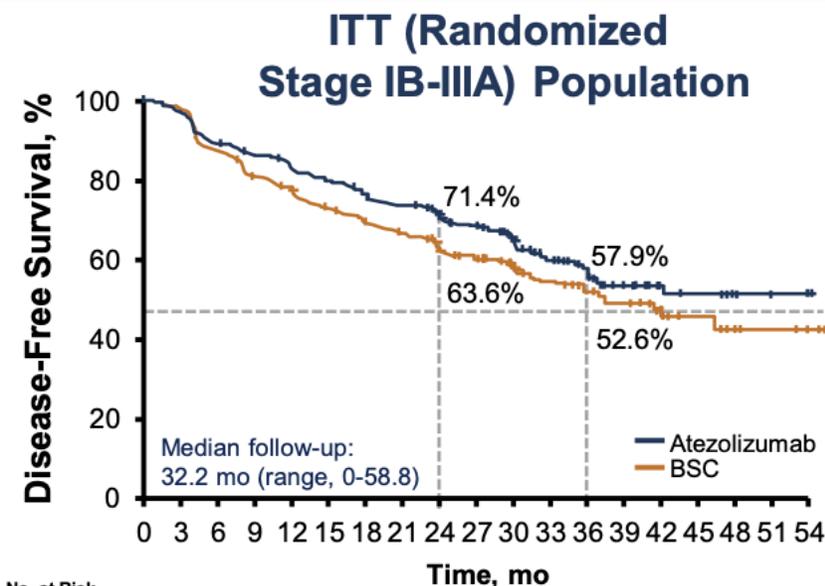
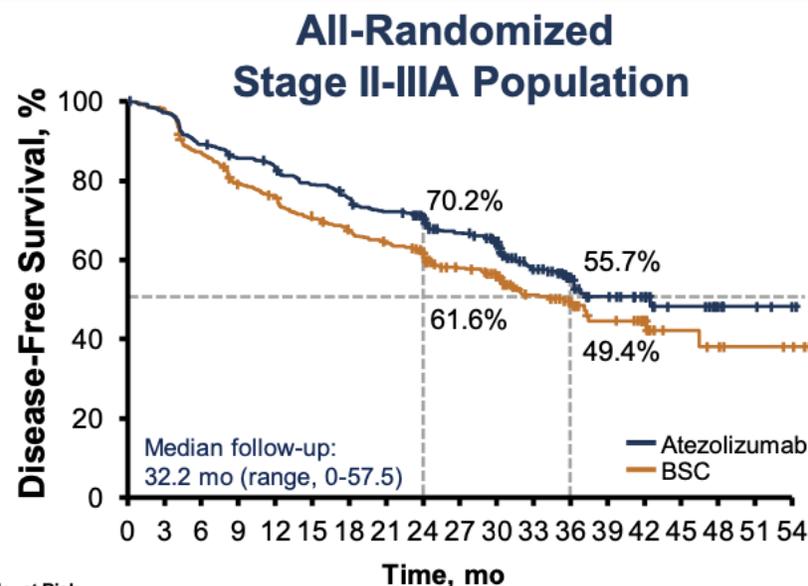
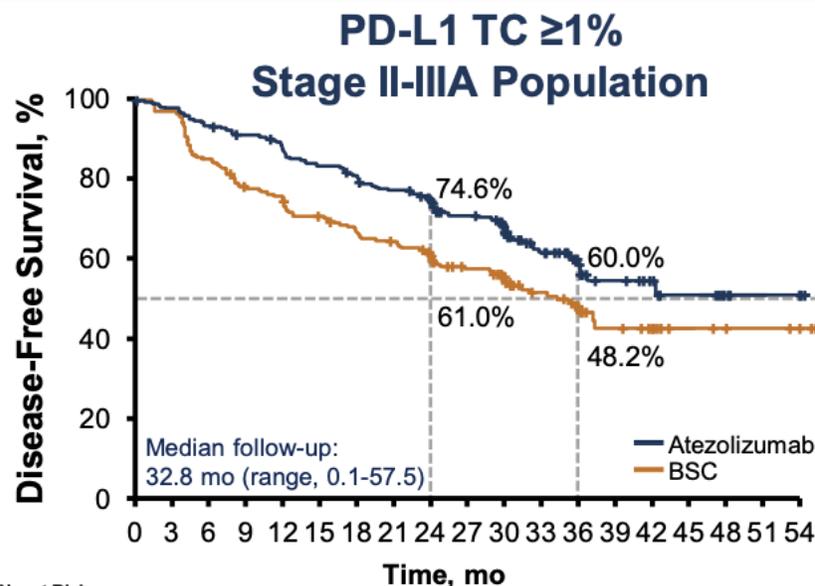
- OS in ITT population
- DFS in PD-L1 TC  $\geq 50\%$  (per SP263) stage II-IIIa population
- 3-year and 5-year DFS in all 3 populations

## Hierarchical statistical testing



Both arms included observation and regular scans for disease recurrence on the same schedule. <sup>a</sup> Per SP142 assay. <sup>b</sup> Two-sided  $\alpha = .05$ .  
1. Wakelee HA et al. ASCO 2021. Abstract 8500.

# IMpower010: DFS in PD-L1 TC ≥1%<sup>a</sup> Stage II-IIIa, All-Randomized, Stage II-IIIa, and ITT (Randomized Stage IB-IIIa) Populations<sup>1</sup>



No. at Risk

Atezolizumab	248	235	225	217	206	198	190	181	159	134	111	76	54	31	22	12	8	3	3
BSC	228	212	186	169	160	151	142	135	117	97	80	59	38	21	14	7	6	4	3

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

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)	
<i>P</i> <sup>b</sup>	.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)	
<i>P</i> <sup>b</sup>	.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)	
<i>P</i> <sup>b</sup>	.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 HA et al. ASCO 2021. Abstract 8500.

# FDA Approval



On October 15, 2021, the FDA approved atezolizumab for adjuvant treatment following resection and platinum-based chemotherapy in patients with stage II to IIIA NSCLC whose tumors have PD-L1 expression on  $\geq 1\%$  of tumor cells, as determined by an FDA-approved test

The FDA also approved the VENTANA PD-L1 (SP263) Assay as a companion diagnostic device to select patients with NSCLC for adjuvant treatment with atezolizumab



### SYSTEMIC THERAPY REGIMENS FOR NEOADJUVANT AND ADJUVANT THERAPY

#### Preferred (nonsquamous)

- Cisplatin 75 mg/m<sup>2</sup> day 1, pemetrexed 500 mg/m<sup>2</sup> day 1 every 21 days for 4 cycles<sup>1</sup>

#### Preferred (squamous)

- Cisplatin 75 mg/m<sup>2</sup> day 1; gemcitabine 1250 mg/m<sup>2</sup> days 1 and 8, every 21 days for 4 cycles<sup>2</sup>
- Cisplatin 75 mg/m<sup>2</sup> day 1; docetaxel 75 mg/m<sup>2</sup> day 1 every 21 days for 4 cycles<sup>3</sup>

#### Other Recommended

- Cisplatin 50 mg/m<sup>2</sup> days 1 and 8; vinorelbine 25 mg/m<sup>2</sup> days 1, 8, 15, and 22, every 28 days for 4 cycles<sup>4</sup>
- Cisplatin 100 mg/m<sup>2</sup> day 1; vinorelbine 30 mg/m<sup>2</sup> days 1, 8, 15, and 22, every 28 days for 4 cycles<sup>5,6</sup>
- Cisplatin 75–80 mg/m<sup>2</sup> day 1; vinorelbine 25–30 mg/m<sup>2</sup> days 1 and 8, every 21 days for 4 cycles
- Cisplatin 100 mg/m<sup>2</sup> day 1; etoposide 100 mg/m<sup>2</sup> days 1–3, every 28 days for 4 cycles<sup>5</sup>

#### Useful in Certain Circumstances

##### Chemotherapy Regimens for Patients with Comorbidities or Patients Not Able to Tolerate Cisplatin

- Carboplatin AUC 6 day 1, paclitaxel 200 mg/m<sup>2</sup> day 1, every 21 days for 4 cycles<sup>7</sup>
- Carboplatin AUC 5 day 1, gemcitabine 1000 mg/m<sup>2</sup> days 1 and 8, every 21 days for 4 cycles<sup>8</sup>
- Carboplatin AUC 5 day 1, pemetrexed 500 mg/m<sup>2</sup> day 1 for nonsquamous every 21 days for 4 cycles<sup>9</sup>

All chemotherapy regimens listed above can be used for sequential chemotherapy/RT.

Protocole de chimiothérapie :  
PS 0, 1 = platine + drogue 3eme génération (GMZ, VNB, TXT, PMT) ou VP16  
PS = 2 → Carboplatine + Paclitaxel / GMZ / PMT

Nombre de cure = 4  
Délai entre 4<sup>eme</sup> semaine et 8<sup>eme</sup> semaine post chirurgie

#### Previous Adjuvant Chemotherapy

- Osimertinib 80 mg daily<sup>10</sup>
  - Osimertinib for patients with completely resected stage IIB-III A or high risk stage IB-II A *EGFR* mutation-positive NSCLC who received previous adjuvant chemotherapy or are ineligible to receive platinum-based chemotherapy.
- Atezolizumab 840 mg every 2 weeks, 1200 mg every 3 weeks, or 1680 mg every 4 weeks for up to 1 year<sup>11</sup>
  - Atezolizumab for patients with completely resected stage IIB-III A or high risk stage II A PD-L1 ≥1% NSCLC who received previous adjuvant chemotherapy.

**Au total**

### Stades localisés

<b>I</b>	IA1	T1mi, T1a < 1 cm	N0
	IA2	T1b > 1 cm ≤ 2 cm	N0
	IA3	T1c > 2 cm ≤ 3 cm	N0
	IB	T2a > 3 cm ≤ 4 cm	N0
<b>II</b>	IIA	T2b > 4 cm ≤ 5 cm	N0
	IIB	T1a, T1b, T1c, T2a, T2b T3	N1 N0
<b>III</b>	IIIA	T1a, T1b, T1c, T2a, T2b T3	N2 N1

**Chirurgie**

**Chirurgie +**  
R0 = CT adjuvante si F risque /osimertinib ou Atézo  
R1-R2 = Reresection + CT  
ou RT + CT

**Chirurgie**  
R0 = CT adjuvante /osimertinib ou Atézolizumab  
ou RT-CT séquentielle  
R1 = re resection + CT  
Ou RT-CT séquentielle ou concomitante  
R2 = re resection + CT  
RT + CT concomitante.

**Chirurgie**  
R0 = CT adjuvante /osimertinib ou Atézolizumab  
ou RT-CT séquentielle (N2)  
R1 = RT-CT séquentielle ou concomitante  
R2 = RT + CT concomitante.

- T > 4 cm
- T peu différenciée
- Invasion vasculaire
- Wedge resection
- Atteinte plèvre viscérale
- Statut ganglionnaire Nx

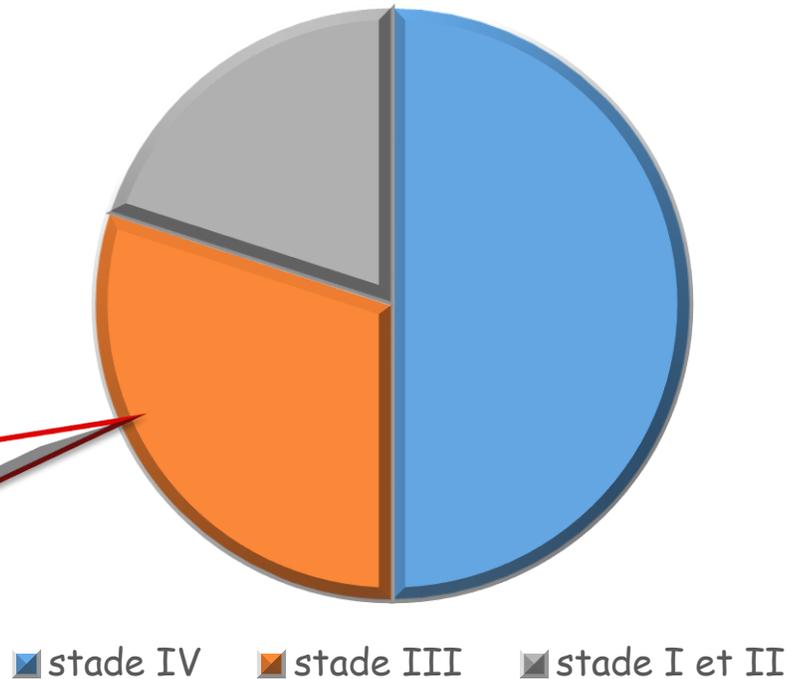
**Stades localement avancés**

Les indispensables

# Stades localement avancés

Groupe de patients  
très hétérogène  
avec un pronostic  
très variable

30 % des patients  
sont au stade  
localement avancé  
IIIA N2, IIIB, IIIC



60 à 70 % des patients  
décèdent de rechute  
métastatique

**Les indispensables**

# Stades localement avancés

## Stades localement avancés

<b>III</b>	IIIA	T4	N0 N1	M0
	IIIB	T1a, T1b, T1c, T2a, T2b T3, T4	N3 N2	M0 M0
	IIIC	T3, T4	N3	M0



**Radio-chimiothérapie  
concomitante**

# Stades localement avancés

## Objectif du traitement

Obtenir le contrôle local de la maladie



**Radiothérapie**

Prévenir l'évolution métastatique en contrôlant la maladie micro métastatique



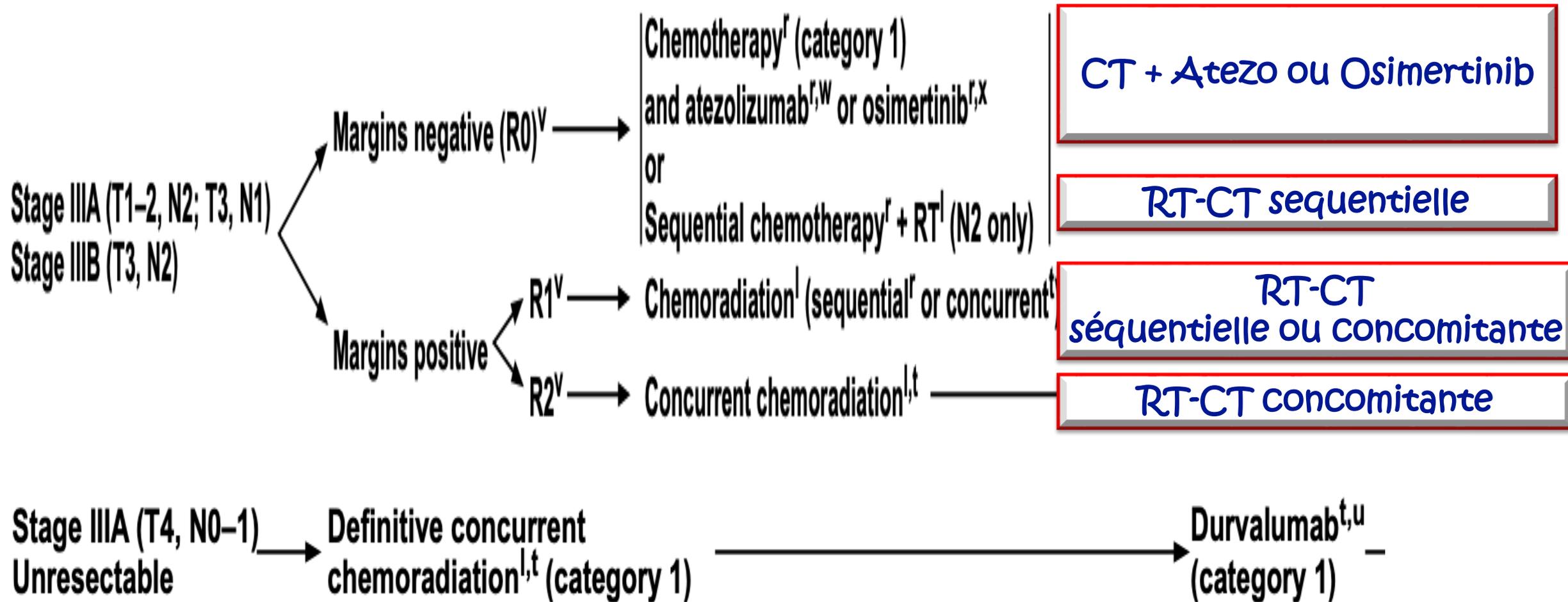
**Chimiothérapie**

# Stades localisés

NCCN

National  
Comprehensive  
Cancer  
Network®

## NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer



Traitement de référence des  
CBNPC localement avancés non  
opérables

Patient PS = 0 ou  
1

Chimio-radiothérapie  
concomitante

Patient PS = 2

Chimio-radiothérapie  
séquentielle

Protocole de chimiothérapie  
: platine + PMT ou  
Paclitaxel ou VP16.

### CONCURRENT CHEMORADIATION REGIMENS

#### Concurrent Chemoradiation Regimens<sup>€</sup>

##### Preferred (nonsquamous)

- Carboplatin AUC 5 on day 1, pemetrexed 500 mg/m<sup>2</sup> on day 1 every 21 days for 4 cycles; concurrent thoracic RT<sup>1,\*†‡</sup>
- Cisplatin 75 mg/m<sup>2</sup> on day 1, pemetrexed 500 mg/m<sup>2</sup> on day 1 every 21 days for 3 cycles; concurrent thoracic RT<sup>2,3,\*†‡</sup>  
± additional 4 cycles of pemetrexed 500 mg/m<sup>2</sup><sup>†,§</sup>
- Paclitaxel 45–50 mg/m<sup>2</sup> weekly; carboplatin AUC 2, concurrent thoracic RT<sup>4,\*†‡</sup> ± additional 2 cycles every 21 days of paclitaxel 200 mg/m<sup>2</sup> and carboplatin AUC 6<sup>†,§</sup>
- Cisplatin 50 mg/m<sup>2</sup> on days 1, 8, 29, and 36; etoposide 50 mg/m<sup>2</sup> days 1–5 and 29–33; concurrent thoracic RT<sup>5,6,\*†‡</sup>

##### Preferred (squamous)

- Paclitaxel 45–50 mg/m<sup>2</sup> weekly; carboplatin AUC 2, concurrent thoracic RT<sup>6,\*†‡</sup> ± additional 2 cycles every 21 days of paclitaxel 200 mg/m<sup>2</sup> and carboplatin AUC 6<sup>†,§</sup>
- Cisplatin 50 mg/m<sup>2</sup> on days 1, 8, 29, and 36; etoposide 50 mg/m<sup>2</sup> days 1–5 and 29–33; concurrent thoracic RT<sup>5,6,\*†‡</sup>

#### Consolidation Immunotherapy for Patients with Unresectable Stage II/III NSCLC, PS 0–1, and No Disease Progression After 2 or More Cycles of Definitive Concurrent Chemoradiation

Durvalumab 10 mg/kg IV every 2 weeks or 1500 mg every 4 weeks for up to 12 months (patients with a body weight of ≥30 kg)<sup>7,8</sup>  
(category 1 for stage III; category 2A for stage II)

# PACIFIC Regimen – Mechanism of Action<sup>1-6</sup>

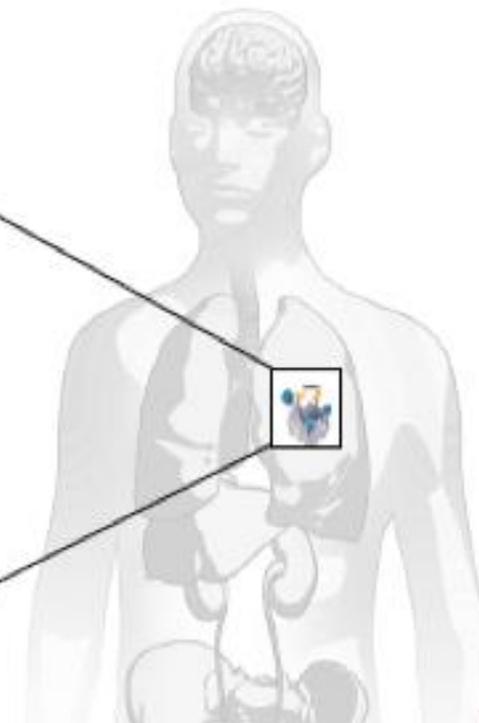
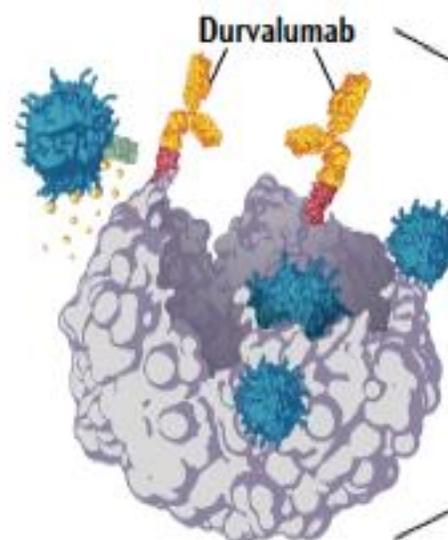
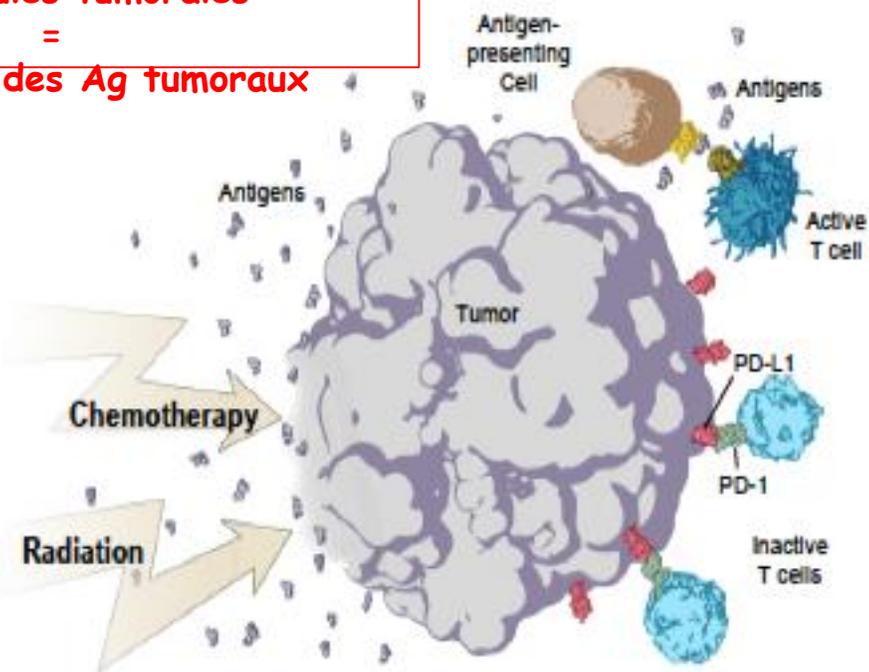
CHEMORADIATION

DURVALUMAB

La RT-CT concomitante entraîne la lyse des cellules tumorales = Relargages des Ag tumoraux

Réaction immunitaire

Annule l'immuno-suppression



## PACIFIC: Study Design

Phase 3, Randomized, Double-blind, Placebo-controlled, Multicenter, International Study<sup>1</sup>

- Unresectable, Stage III NSCLC without progression after definitive platinum-based cCRT ( $\geq 2$  cycles)
- 18 years or older
- WHO PS score 0 or 1
- If available, archived pre-cCRT tumor tissue for PD-L1 testing\*

**All-comers population  
(i.e. irrespective of PD-L1 status)**

**N=713 randomized**

1–42 days  
post-cCRT

**R**

**Durvalumab**  
10 mg/kg q2w for  
up to 12 months  
N=476

2:1 randomization,  
stratified by age, sex, and  
smoking history

**Placebo**  
10 mg/kg q2w for  
up to 12 months  
N=237

### Primary endpoints

- PFS by BICR using RECIST v1.1†
- OS

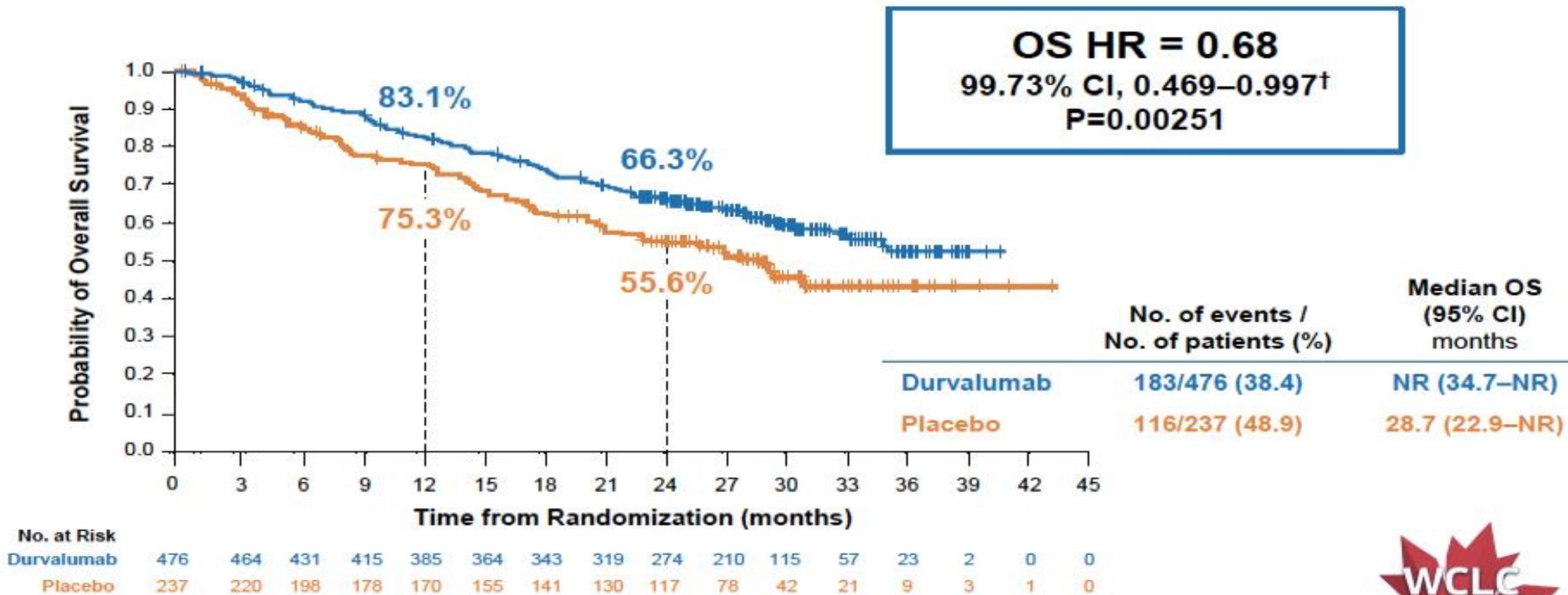
### Key secondary endpoints

- ORR, DoR and TTDM by BICR
- PFS2 by investigator
- Safety
- PROs

\*Using the Ventana SP263 immunohistochemistry assay

†Defined as the time from randomization until the date of objective disease progression or death by any cause in the absence of progression. BICR, blinded independent central review; cCRT, concurrent CRT; PFS2, time to second progression; RECIST, Response Evaluation Criteria in Solid Tumors; TTDM, time to death or distant metastasis. ClinicalTrials.gov number: NCT02125461

## Overall Survival\* (ITT)



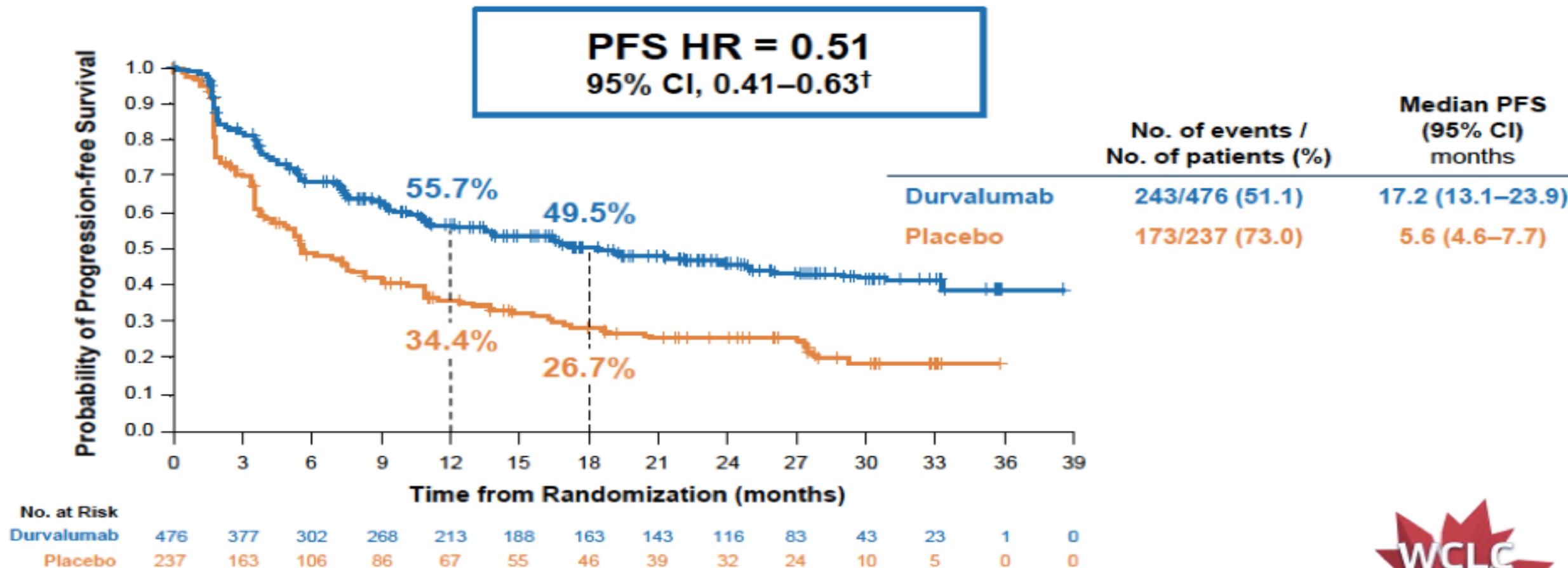
\*Median duration of follow-up for OS was 25.2 months (range 0.2–43.1)

†Adjusted for interim analysis

NR, not reached



## Updated Progression-free Survival by BICR\* (ITT)

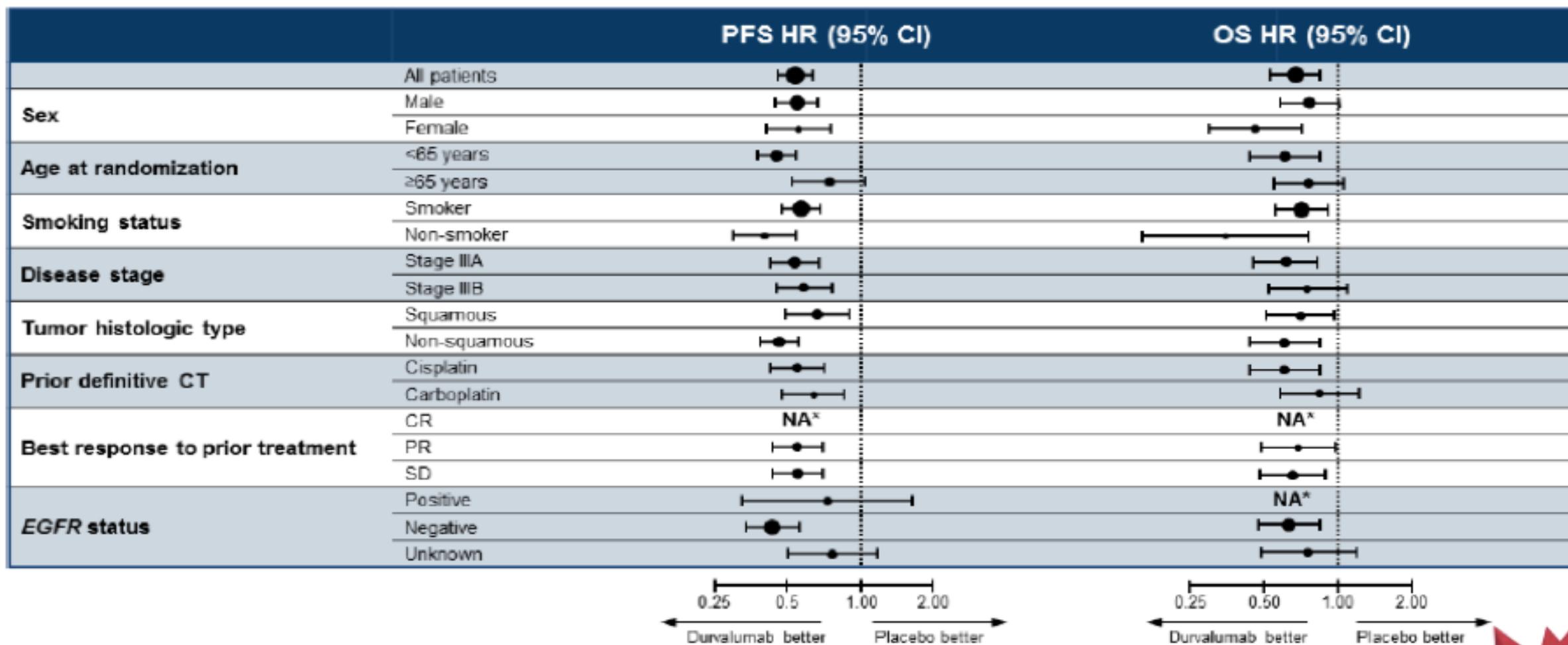


\*Median duration of follow-up was 25.2 months (range 0.2–43.1)

<sup>†</sup>No formal statistical comparison was made because the study had achieved significance for PFS at the first planned IA (data cutoff of Feb 13, 2017)



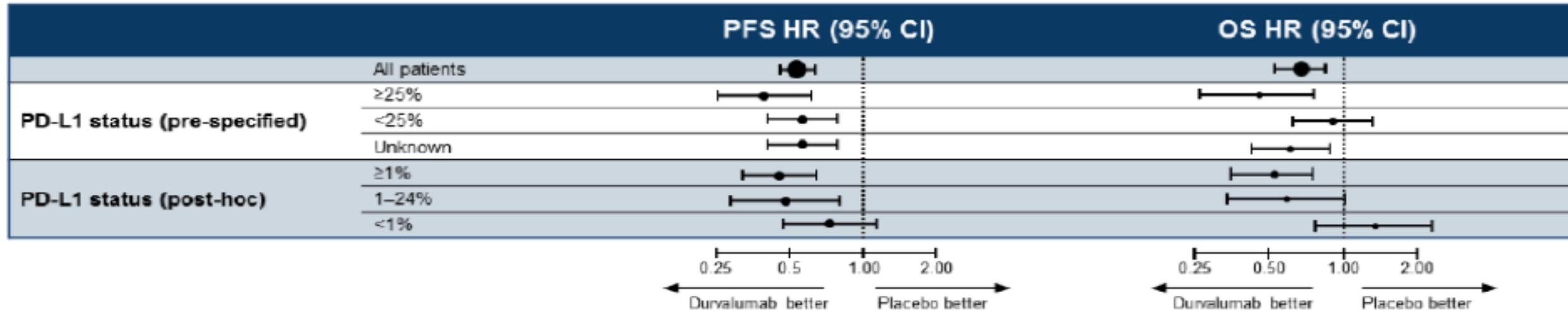
## Progression-free and Overall Survival by Subgroup (ITT)



\*Not calculated if subgroup has <20 events



## Subgroup Analysis by PD-L1 Status



- Important facts regarding PD-L1 status:

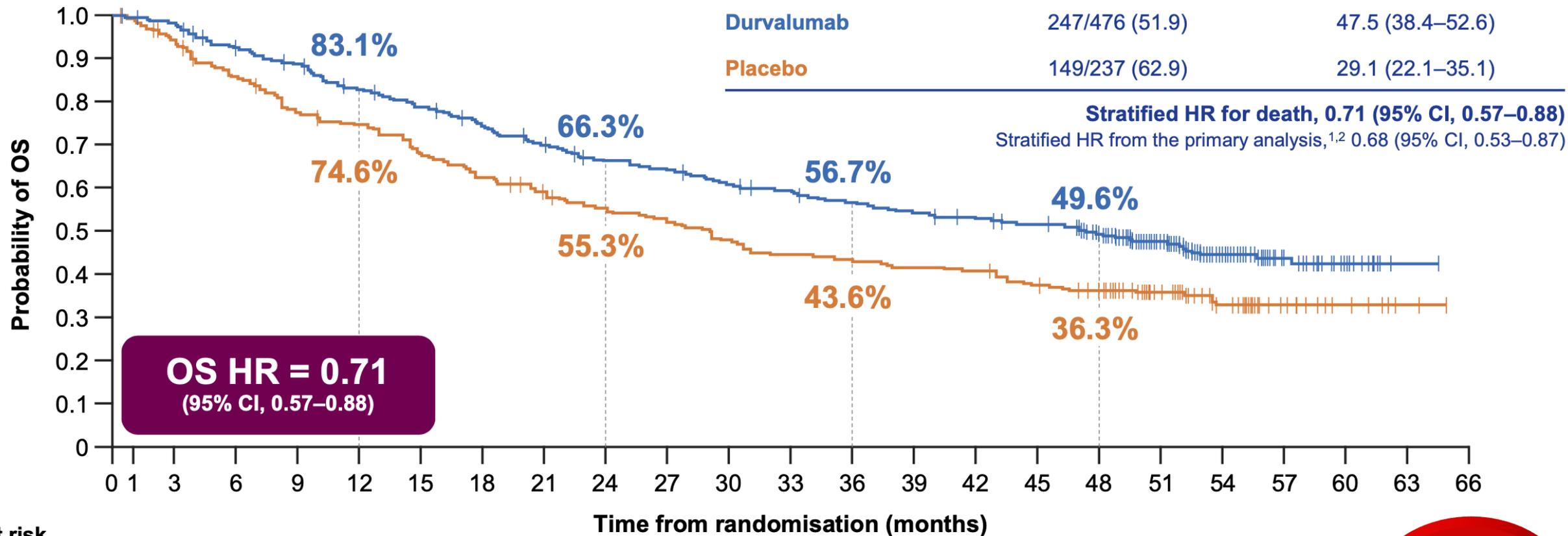
- PD-L1 testing was not required
- 37% of patients with unknown PD-L1 status
- PD-L1 status was obtained pre-CRT (getting a sample post-CRT medically not feasible)
- PD-L1 expression-level cutoff of 1% was part of an unplanned post-hoc analysis requested by a health authority



Survie globale

# PACIFIC: Updated OS

	No. of events/ total no. of patients (%)	Median OS (95% CI), months
<b>Durvalumab</b>	247/476 (51.9)	47.5 (38.4–52.6)
<b>Placebo</b>	149/237 (62.9)	29.1 (22.1–35.1)

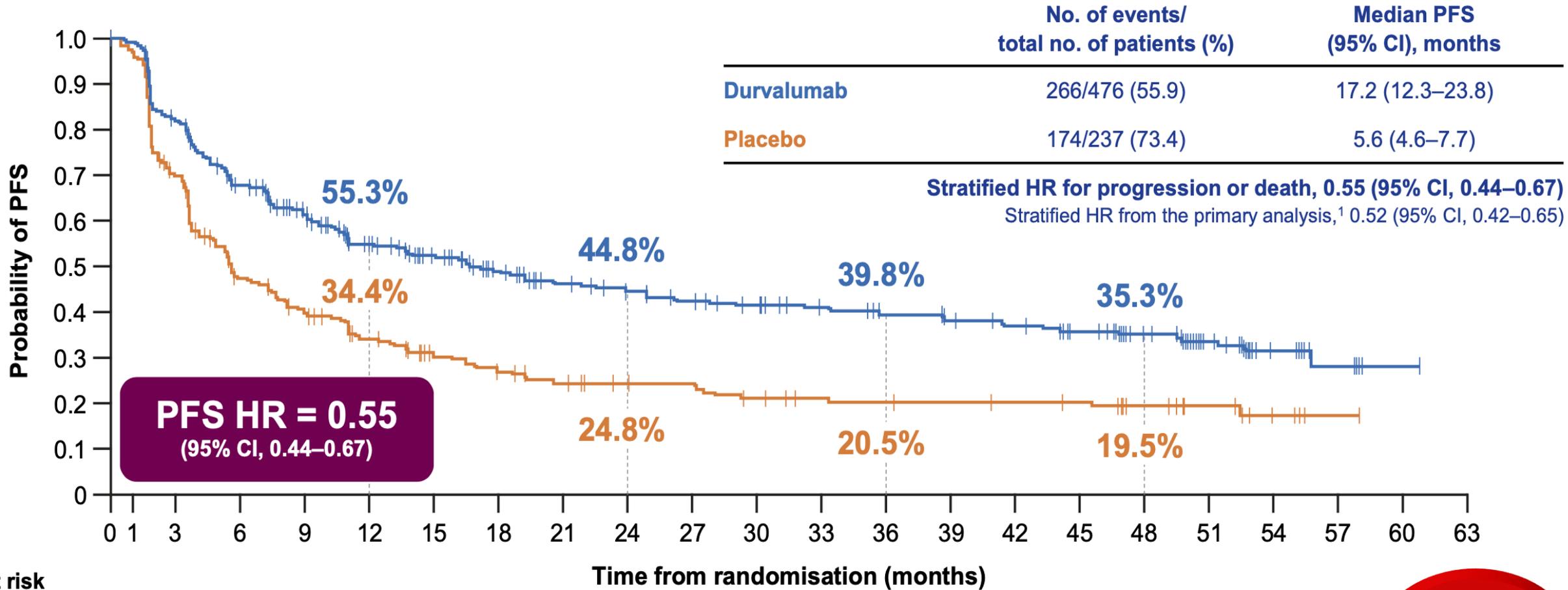


No. at risk	0	1	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	54	57	60	63	66
<b>Durvalumab</b>	476	464	431	414	385	364	343	319	299	290	274	265	252	241	235	225	195	138	75	36	15			
<b>Placebo</b>	237	220	199	179	171	156	143	133	123	116	107	99	97	93	91	83	75	53	29	15	7			



Survie sans progression

# CIFIC: Updated PFS (BIRC)

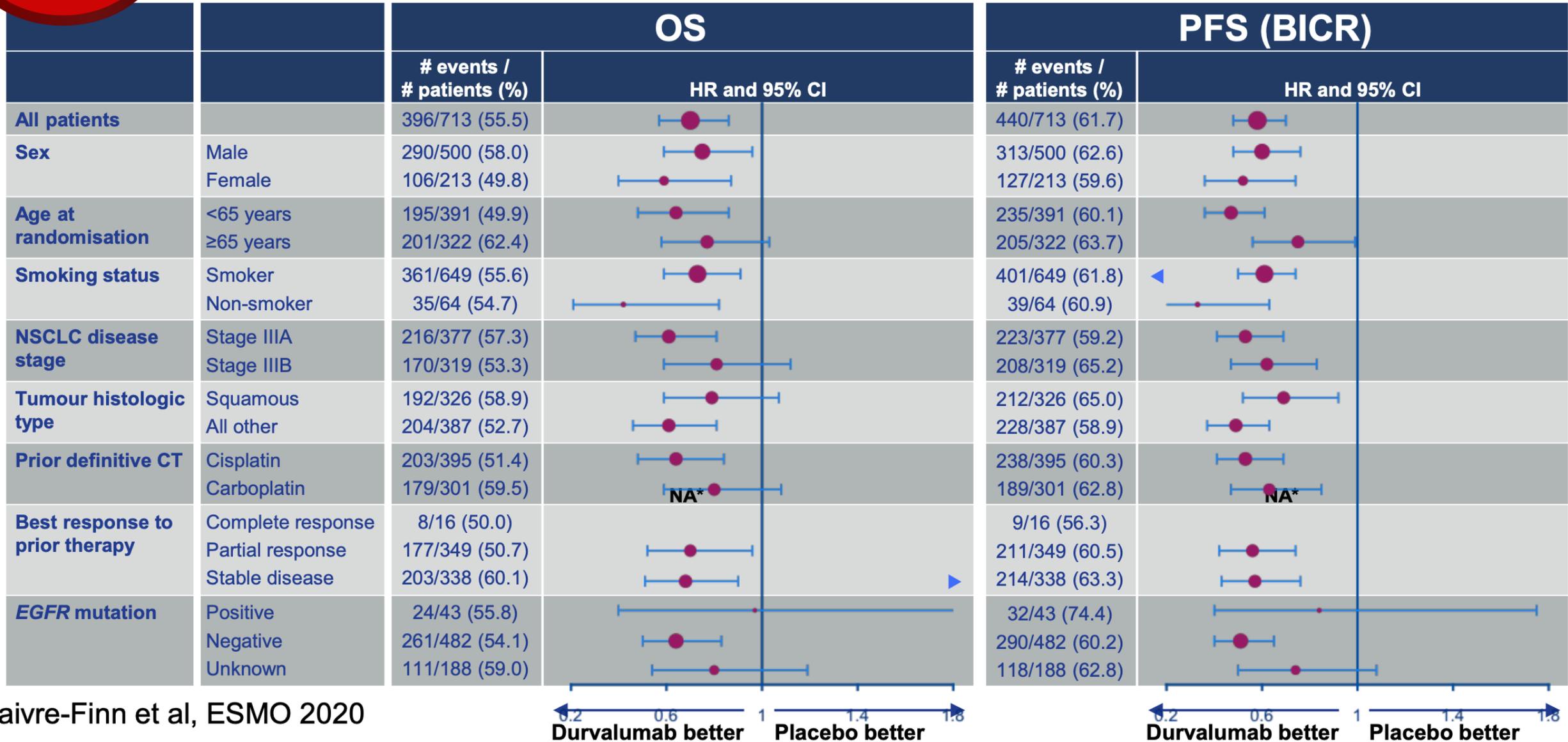


No. at risk	0	1	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	54	57	60	63
<b>Durvalumab</b>	476	377	301	266	213	189	165	146	136	127	119	110	103	97	92	80	59	37	18	8			
<b>Placebo</b>	237	163	105	86	67	55	47	40	36	35	29	26	25	24	23	22	16	11	5	1			



WCLC  
2020

# ACIFIC: Subgroup Analysis



Protocole de chimiothérapie  
: platine + PMT ou  
Paclitaxel ou VP16.

### CONCURRENT CHEMORADIATION REGIMENS

#### Concurrent Chemoradiation Regimens<sup>€</sup>

##### Preferred (nonsquamous)

- Carboplatin AUC 5 on day 1, pemetrexed 500 mg/m<sup>2</sup> on day 1 every 21 days for 4 cycles; concurrent thoracic RT<sup>1,\*†‡</sup>
- Cisplatin 75 mg/m<sup>2</sup> on day 1, pemetrexed 500 mg/m<sup>2</sup> on day 1 every 21 days for 3 cycles; concurrent thoracic RT<sup>2,3,\*†‡</sup>  
± additional 4 cycles of pemetrexed 500 mg/m<sup>2</sup>†,§
- Paclitaxel 45–50 mg/m<sup>2</sup> weekly; carboplatin AUC 2, concurrent thoracic RT<sup>4,\*†‡</sup> ± additional 2 cycles every 21 days of paclitaxel 200 mg/m<sup>2</sup> and carboplatin AUC 6†,§
- Cisplatin 50 mg/m<sup>2</sup> on days 1, 8, 29, and 36; etoposide 50 mg/m<sup>2</sup> days 1–5 and 29–33; concurrent thoracic RT<sup>5,6,\*†‡</sup>

##### Preferred (squamous)

- Paclitaxel 45–50 mg/m<sup>2</sup> weekly; carboplatin AUC 2, concurrent thoracic RT<sup>6,\*†‡</sup> ± additional 2 cycles every 21 days of paclitaxel 200 mg/m<sup>2</sup> and carboplatin AUC 6†,§
- Cisplatin 50 mg/m<sup>2</sup> on days 1, 8, 29, and 36; etoposide 50 mg/m<sup>2</sup> days 1–5 and 29–33; concurrent thoracic RT<sup>5,6,\*†‡</sup>

#### Consolidation Immunotherapy for Patients with Unresectable Stage II/III NSCLC, PS 0–1, and No Disease Progression After 2 or More Cycles of Definitive Concurrent Chemoradiation

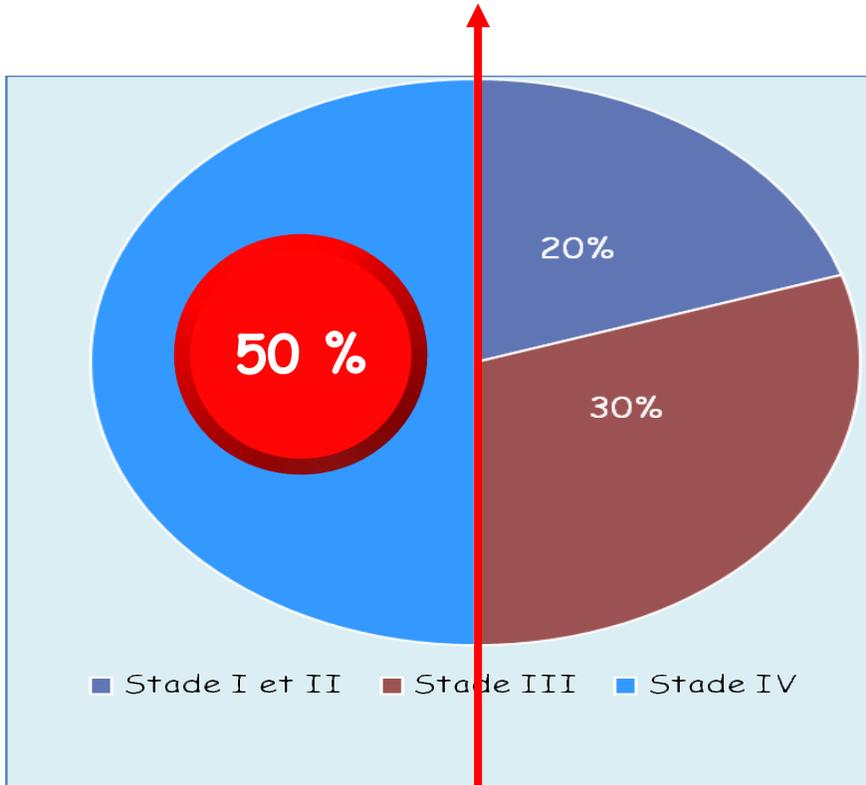
Durvalumab 10 mg/kg IV every 2 weeks or 1500 mg every 4 weeks for up to 12 months (patients with a body weight of ≥30 kg)<sup>7,8</sup>  
(category 1 for stage III; category 2A for stage II)

# Stades métastatiques

Les indispensables

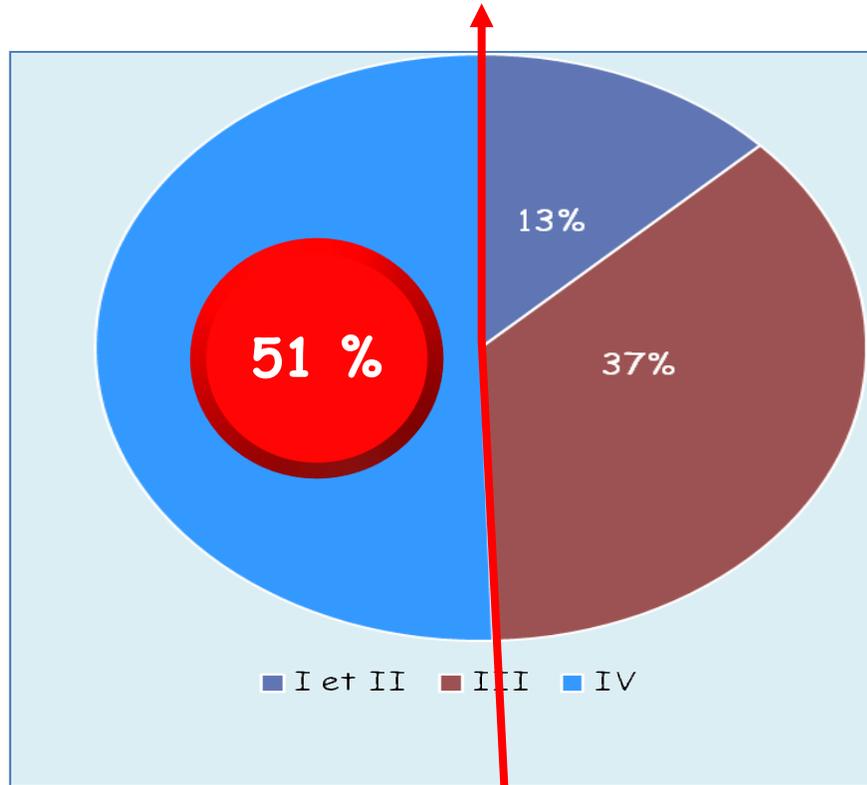
# Stades métastatiques

## Dans le monde



Fry WA, et al. Cancer. 1996;77:1949–1995

## Algérie



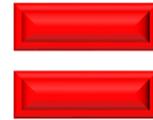
CPMC, HDJ Oncologie médicale. 2000

Les indispensables

# Stades métastatiques

## Stades métastatiques

IV	IVA	Quel que soit T	Quel que soit N	M1a, M1b
	IVB	Quel que soit T	Quel que soit N	M1c



1. Traitement symptomatique
2. La chimiothérapie
3. Les thérapies ciblées
4. L'immunothérapie

1. Prolonger la survie des patients « Survie Globale **SG** » tout en améliorant la qualité de vie.
2. Augmenter le temps jusqu'à progression **SSP**.

# Pré requis au traitement

## RCP

- Age / Comorbidités.
- PS
- L'extension de la maladie (TNM)
- Le type histologique
- Les drivers moléculaires

- âge, [ $< 70$  ans,  $70 - 89$  ans]
- Comorbidités tel que insuffisance rénale, cardiopathies, neuropathies importantes....
- PS = 0, 1 / PS = 2 / PS = 3, 4.

Choix du protocole thérapeutique médicale

- Cisplatine / carboplatine
- Monothérapie / bithérapie
- Chimiothérapie / Thérapie ciblée (TKI / antiangiogénique) / Immunothérapie.

Classification TNM et Stade

### CLINICAL PRESENTATION

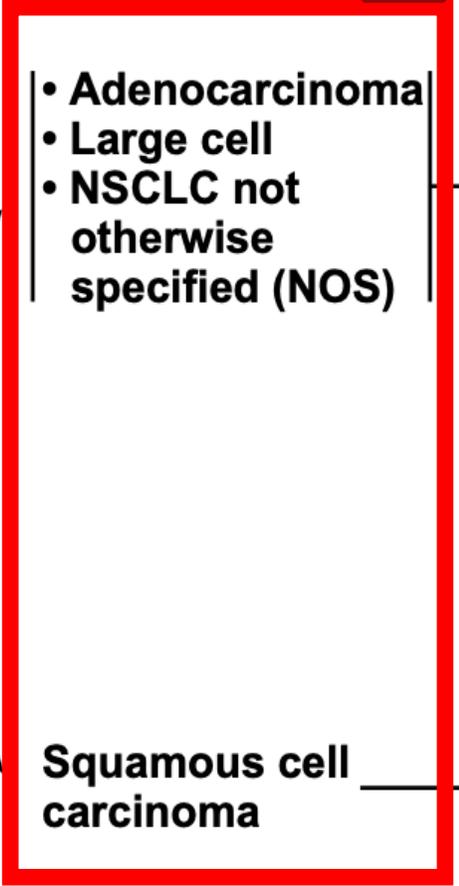
### HISTOLOGIC SUBTYPE<sup>a</sup>



### BIOMARKER TESTING<sup>mm</sup>

Advanced or metastatic disease

- Establish histologic subtype<sup>a</sup> with adequate tissue for molecular testing (consider rebiopsy<sup>ll</sup> if appropriate)
- Smoking cessation counseling
- Integrate palliative care<sup>c</sup> ([See NCCN Guidelines for Palliative Care](#))



- Adenocarcinoma
- Large cell
- NSCLC not otherwise specified (NOS)

Squamous cell carcinoma

- Molecular testing, including:
  - ▶ *EGFR* mutation (category 1), *ALK* (category 1), *KRAS*, *ROS1*, *BRAF*, *NTRK1/2/3*, *MET*ex14 skipping, *RET*
  - ▶ Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)

- Consider molecular testing, including:<sup>oo</sup>
  - ▶ *EGFR* mutation, *ALK*, *KRAS*, *ROS1*, *BRAF*, *NTRK1/2/3*, *MET* exon 14 skipping, *RET*
  - ▶ Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)



# NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

## PRINCIPLES OF PATHOLOGIC REVIEW

### Classification

The types of NSCLC are: adenocarcinoma, squamous cell carcinoma, adenosquamous carcinoma, large cell carcinoma, and sarcomatoid carcinoma.

▶ **Squamous cell carcinoma:** A malignant epithelial tumor that either shows keratinization and/or intercellular bridges, or a morphologically undifferentiated NSCC that expresses immunohistochemical markers of squamous cell differentiation.

▶ **Adenocarcinoma:**

- ◊ For small (<3 cm), resected lesions, determining extent of invasion is critical.
  - Adenocarcinoma in situ (AIS; formerly BAC): A small ( $\leq 3$  cm) localized nodule with lepidic growth, mostly non-mucinous, although mucinous types can occur. Multiple synchronous AIS tumors can also occur.
  - Minimally invasive adenocarcinoma (MIA): A small ( $\leq 3$  cm) solitary adenocarcinoma with a predominantly lepidic pattern and  $\leq 5$  mm invasion in greatest dimension. MIA is usually non-mucinous, but rarely may be mucinous. MIA is, by definition, solitary and discrete.
  - Invasive adenocarcinoma: A malignant epithelial tumor with glandular differentiation, mucin production, or pneumocyte marker expression. The tumors show an acinar, papillary, micropapillary, lepidic, or solid growth pattern, with either mucin or pneumocyte marker expression. The invasive adenocarcinoma component should be present in at least one focus measuring  $>5$  mm in greatest dimension.
  - Invasive adenocarcinoma variants: invasive mucinous adenocarcinoma, solid papillary adenocarcinoma, and micropapillary adenocarcinoma.
  - Refer to College of American Pathologists [Protocols](#) for additional information.

▶ **Adenosquamous carcinoma:** A carcinoma showing components of both adenocarcinoma and squamous cell carcinoma. A squamous component constituting at least 10% of the tumor. Definitive diagnosis is based on findings in small biopsies, cytology, or excisional biopsies. For large resections, otherwise squamous should trigger molecular testing.

▶ **Large cell carcinoma:** Undifferentiated NSCC that lacks the cytologic features of adenocarcinoma, or squamous cell carcinoma. The diagnosis requires immunohistochemistry on resection or cytology specimens.

▶ **Sarcomatoid carcinoma** is a general term that includes pleomorphic carcinoma, carcinosarcoma, and pulmonary blastoma. It is best to use the specific term for these entities whenever possible.

- ◊ Pleomorphic carcinoma is a poorly differentiated NSCC that consists of a mixture of spindle and giant cells. Spindle cell carcinoma consists of spindle cells, and giant cell carcinoma consists almost entirely of tumor giant cells.
- ◊ Carcinosarcoma is a malignant tumor that consists of a mixture of carcinoma and sarcoma (e.g., rhabdomyosarcoma, chondrosarcoma, osteosarcoma).
- ◊ Pulmonary blastoma is a biphasic tumor that consists of fetal

### Classification histologique OMS 2015

Carcinome épidermoïde	Pont d'union +/- Kératine
Adénocarcinome	Différenciation glandulaire Production mucine (BA +)
Carcinome adéno-squameux	Double composante épidermoïde et adénocarcinome avec pour chaque composante au moins 10 % de la Tumeur
Carcinome à grandes cellules	Pas de différenciation Diagnostic sur pièce opératoire
Carcinome	Déliomorphe / Carcino-sarcome

## PRINCIPLES OF PATHOLOGIC REVIEW

### Immunohistochemistry

- Judicious use of IHC is strongly recommended to preserve tissue for molecular testing, most notably in small specimens. When adenocarcinoma or squamous cell carcinomas are poorly differentiated, the defining morphologic criteria that would allow for specific diagnosis may be inconspicuous or absent. In this case, IHC or mucin staining may be necessary to determine a specific diagnosis.
- In small specimens, a limited number of immunostains with one lung adenocarcinoma marker (TTF1, napsin A) and one squamous carcinoma marker (p40, p63) should suffice for most diagnostic problems. Virtually all tumors that lack squamous cell morphology and show co-expression of p63 and TTF1 are preferably classified as adenocarcinoma. A simple panel of TTF1 and p40 may be sufficient to classify most NSCC-NOS cases.
- Testing for NUT expression by IHC should be considered in all poorly differentiated carcinomas that lack glandular differentiation or specific etiology, particularly in non-smokers or in patients presenting at a young age, for consideration of a pulmonary NUT carcinoma.
- IHC should be used to differentiate primary lung adenocarcinoma from squamous cell carcinoma, large cell carcinoma, metastatic carcinoma, and primary pleural mesothelioma (particularly for pleural specimens).
- Primary pulmonary adenocarcinoma:
  - ▶ In patients for whom the primary origin of the carcinoma is uncertain, an appropriate panel of immunohistochemical stains is recommended to assess for metastatic carcinoma to the lung.

	C épidermoïde	Adénocarcinome
Morphologie	- Pont d'union - +/- Kératine	Morphologie glandulaire Mucine (BA +)
IHC	P63 ou p40	TTF1 ou Napsin A

- ▶ TTF1 is a homeodomain-containing nuclear transcription protein of the embryonal and mature lung and thyroid. TTF1 immunoreactivity is seen in non-mucinous adenocarcinoma subtypes. Metastatic adenocarcinoma to the thyroid malignancies, in which case thyroglobulin and PAX8 are also used. Metastatic adenocarcinoma to the gynecologic tract, pancreatobiliary) have been noted, and may be of value in establishing correlation with clinical and radiologic features.
- ▶ Napsin A—an aspartic proteinase expressed in normal type II pneumocytes—is expressed in >80% of lung adenocarcinomas and may be a useful adjunct to TTF1.
- ▶ The panel of TTF1 (or alternatively napsin A) and p40 (or alternatively p63) may be useful in refining the diagnosis to either adenocarcinoma or squamous cell carcinoma in small biopsy specimens previously classified as NSCC NOS.



### CLINICAL PRESENTATION

### HISTOLOGIC SUBTYPE<sup>a</sup>

### BIOMARKER TESTING<sup>mm</sup>

Advanced or metastatic disease

- Establish histologic subtype<sup>a</sup> with adequate tissue for molecular testing (consider rebiopsy<sup>ll</sup> if appropriate)
- Smoking cessation counseling
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- Adenocarcinoma
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Squamous cell carcinoma

- Molecular testing, including:
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  - ▶ Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)
  
- Consider molecular testing, including:<sup>oo</sup>
  - ▶ *EGFR* mutation, *ALK*, *KRAS*, *ROS1*, *BRAF*, *NTRK1/2/3*, *MET* exon 14 skipping, *RET*
  - ▶ Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)

**Marqueurs  
moléculaires**

**PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS**

**Molecular Diagnostic Studies in Non-Small Cell Lung Cancer**

- Numerous gene alterations have been identified that impact therapy selection. Testing of lung cancer specimens for these alterations is essential for the identification of potentially efficacious targeted therapies, as well as avoidance of therapies that are unlikely to be effective.
- Some selection approaches for targeted therapy include predictive immunohistochemical and genomic testing. Genomic testing and other studies utilized to identify tumor type and lineage.
- Major elements of molecular testing that are critical for utilization and interpretation of molecular testing results include:
  - ▶ Use of a laboratory that is properly accredited, with a minimum of CLIA accreditation
  - ▶ Understanding the methodologies that are utilized and the major limitations of those methods
  - ▶ Understanding the spectrum of alterations tested (and those not tested) by a specific assay
  - ▶ Knowledge of whether a tumor sample is subjected to pathologic review and tumor enrichment
  - ▶ The types of samples accepted by the testing laboratory
- Specimen Acquisition and Management:
  - ▶ Although tumor testing has been primarily focused on use of FFPE tissues, increasingly, fresh frozen (FF) tissues and cell block preparations not processed by FFPE methods. Although testing on cell block specimens is not recommended for most companion diagnostic assays, testing on these specimen types is highly recommended when FFPE is not available.
  - ▶ A major limitation in obtaining molecular testing results for NSCLC occurs when minimal tissue is available for testing. A minimum of 100 mg of tumor tissue is generally considered sufficient to enable all appropriate testing.
  - ▶ When tissue is minimal, laboratories should deploy techniques to maximize tissue for molecular testing, including the use of microdissection and other protocols for small biopsies, including “up-front” slide sectioning for diagnostic and predictive testing.
- Testing Methodologies
  - ▶ Appropriate possible testing methodologies are indicated below for each analyte separately; however, several methodologies are generally considered for use:
    - ◊ Next-generation sequencing (NGS) is used in clinical laboratories. Not all types of alterations are detected by individual NGS assays and it is important to be familiar with the types of alterations identifiable in individual assays or combination(s) of assays.
    - ◊ It is recommended at this time that when feasible, testing be performed via a broad, panel-based approach, most typically performed by NGS. For patients who, in broad panel testing don't have identifiable driver oncogenes (especially EGFR, KRAS, and ALK), testing for these genes should be performed, to maximize detection of fusion events.
    - ◊ Real-time polymerase chain reaction (PCR) can be used in a highly targeted fashion (specific alterations), but when a broad panel of alterations is deployed, only those specific alterations that are targeted by the assay are assessed.
    - ◊ Sanger sequencing requires the greatest degree of tumor enrichment. Unmodified Sanger sequencing is not recommended for testing in tumor samples with less than 25% to 30% tumor after enrichment and is not appropriate for testing for gene rearrangements (eg, resistance mutations) is important. If Sanger sequencing is utilized, tumor enrichment is essential.
    - ◊ Other methodologies may be utilized, including multiplex approaches not listed above.
    - ◊ Fluorescence in situ hybridization (FISH) analysis is utilized for many assays examining gene rearrangements.
    - ◊ IHC is specifically utilized for some specific analytes, and can be a useful surrogate or screening assay for others.

**Plusieurs marqueurs :**

- Mutation EGFR (2009)
- Mutation exon 20 (2019)
- Translocation ALK (2011)
- Translocation ROS-1 (2012)
- Mutation BRAF V600E
- Fusion NTRK
- Mutation KRAS G12C
- NTRK
- MET perte exon 14
- RET

**Plusieurs méthodes :**

- NGS = Next Generation Sequencing
- PCR = Real-Time Polymerase Chain Reaction
- FISH = Hybridation In Situ par Fluorescence
- IHC = Immuno Histo Chimie

**PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS**

**EGFR**

**Targets for Analysis**

In general, the mutations/alterations described below are seen in a non-overlapping fashion, although between 1%–3% of NSCLC may harbor concurrent alterations.

▶ **EGFR (Epidermal Growth Factor Receptor) Gene Mutations:** EGFR is a receptor tyrosine kinase normally found on the surface of epithelial cells and is often overexpressed in a variety of human malignancies.

- ◊ The most commonly to oral EGFR tyrosine kinase inhibitors should not be treated.
- ◊ Molecular testing for EGFR mutations are available for adjuvant and metastatic disease.
- ◊ Many of the less common mutations, including exon 19 insertions, have been studied in clinical trials. The most studied patients is in the adjuvant setting.
- ◊ **EGFR exon 20 (EGFR Exon 20 Insertions)**—Knowledge of the specific mutation is important.
  - EGFR p.T790M is not a resistance mutation in first-generation EGFR TKI therapy.
  - Third-generation TKI therapy is indicated for possible germline EGFR mutations.
  - Most other EGFR Exon 20 mutations are associated with resistance to EGFR TKI therapy.

- **Plusieurs mutations ont été individualisées ayant une fonction activatrice répondant au TKI anti-EGFR**
  - Délétion au niveau de l'**exon 19** (environ 45% des mutations) au niveau des amino-acides 747-749,
  - Substitution sur un seul nucléotide de l'**exon 21**, substituant une arginine à la place d'une leucine au codon 858 (**L858R**), (ceci représente environ 40% des mutations),
  - Mutations **exon 18**, plus rares (10 %)
  - Insertion **exon 19 P.L861Q, P.G719X, P.S768I**

▪ These are generally associated with lack of response to EGFR TKI therapy, with select exceptions:

– **p.A763\_Y764insFOEA** is associated with sensitivity to TKI therapy

– **p. A763\_Y764insL858R**

▪ For this reason, the use of EGFR TKI therapy is indicated for EGFR Exon 20 insertions.

**Méthodes de recherche = Real time PCR, Sanger sequencing, NGS**

- ◊ As use of NGS testing increases, additional EGFR variants are increasingly identified; however, the clinical implications of individual alterations are unlikely to be well established.
- ◊ Some clinicopathologic features—such as smoking status, ethnicity, and histology—are associated with the presence of an EGFR mutation; however, these features should not be utilized in selecting patients for testing.
- ◊ **Testing Methodologies:** Real-time PCR, Sanger sequencing (ideally paired with tumor enrichment), and NGS are the most commonly deployed methodologies for examining EGFR mutation status.

**EGFR**

## PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS

... targets for Analysis  
In general, the mutations/alterations described are often accompanied by concurrent alterations.

- ▶ **EGFR** (Epidermal Growth Factor Receptor) is often overexpressed in a variety of NSCLC subtypes.
- ◊ The most commonly described mutation associated with sensitivity to oral EGFR tyrosine kinase inhibitor (TKI) therapy is *p.L858R*.
- ◊ Molecular testing for *EGFR* mutation status is available for adjuvant treatment decisions.
- ◊ Many of the less commonly observed alterations in *EGFR*, which cumulatively account for ~10% of *EGFR*-mutation positive NSCLC (ie, *exon 19* insertions, *p.L861Q*, *p.G719X*, *p.S768I*) are also associated with responsiveness to *EGFR* TKI therapy, although the number of studied patients is lower.

Mutation résistance aux TKI = Anomalie exon 20 :

- Insertion exon 20, *p.A763\_Y764insFQEA*, *p.A763\_Y764insLQEA*
- Mutation T790M (résistance acquise, secondaire après prise TKI)

◊ *EGFR exon 20 (EGFR<sub>ex20</sub>)* mutations are a heterogeneous group, some of which are responsive to targeted therapy and that require detailed knowledge of the specific alteration.

▶ *EGFR p.T790M* is most commonly observed as a mutation that arises in response to and as a mechanism of resistance to first- and second-generation *EGFR* TKI. In patients with progression on first- or second-generation TKI with *p.T790M* as the primary mechanism of resistance, third-generation TKIs are typically efficacious. If *p.T790M* is observed in the absence of prior *EGFR* TKI therapy, genetic counseling and possible germline genetic testing is warranted.

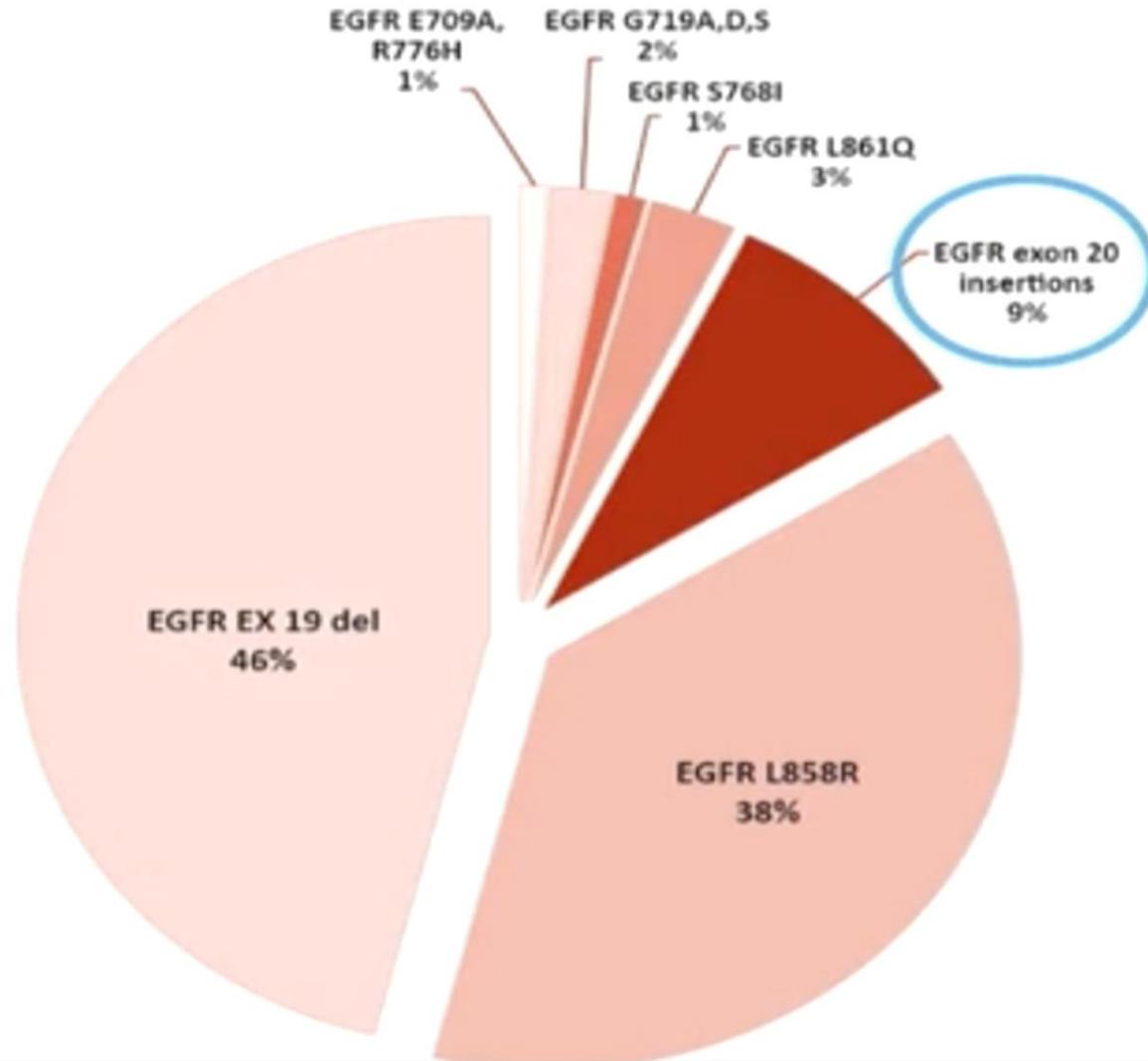
- Most other *EGFR<sub>ex20</sub>* alterations are a diverse group of in-frame duplication or insertion mutations.
  - These are generally associated with lack of response to *EGFR* TKI therapy, with select exceptions:
    - ◊ *p.A763\_Y764insFQEA* is associated with sensitivity to TKI therapy
    - ◊ *p.A763\_Y764insLQEA* may be associated with sensitivity to TKI therapy
  - For this reason, the specific sequence of *EGFR<sub>ex20</sub>* insertion mutations is important, and some assays will identify the presence of an *EGFR<sub>ex20</sub>* insertion without specifying the sequence. In this scenario, additional testing to further clarify the *EGFR<sub>ex20</sub>* insertion is indicated.

◊ As use of NGS testing increases, additional *EGFR* variants are increasingly identified; however, the clinical implications of individual alterations are unlikely to be well established.

◊ Some clinicopathologic features—such as smoking status, ethnicity, and histology—are associated with the presence of an *EGFR* mutation; however, these features should not be utilized in selecting patients for testing.

◊ Testing Methodologies: Real-time PCR, Sanger sequencing (ideally paired with tumor enrichment), and NGS are the most commonly deployed methodologies for examining *EGFR* mutation status.

# EGFR Mutated Cases



- Amivantamab
- Mobocetinib
- Osimertinib
- Poziotinib

## PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS

• **Molecular Targets for Analysis (continued)**

- ▶ **ALK (Anaplastic Lymphoma Kinase) Gene Rearrangements:** ALK rearrangements lead to constitutive activation and inappropriate signaling through the ALK kinase domain. The most common fusion partner seen with ALK is echinoderm microtubule-associated protein 3 (EML4). Numerous other fusion partners have been identified. The presence of an ALK rearrangement is associated with improved outcomes. Some clinicopathologic features—such as smoking status and histology—however, these features should not be utilized in selecting patients for testing. Testing Methodologies: FISH break-apart probe methodology can be used for screening strategy. FDA-approved IHC (ALK [D5F3] CDx Assay) is available. Numerous NGS methodologies can detect ALK fusions. Targeted NGS methodologies with novel partners.

ALK

**Translocation ALK-EML4:**  
**FISH :** méthode de detection standard = Hybridation in situ par fluorescence (FISH break-apart probe)  
**IHC :** validé avec Ac D5F3  
 2019, utiliser comme pre screening et si + Varifier avec FISH  
 2020, pas de confirmation par FISH.

- ▶ **ROS1 (ROS proto-oncogene 1) Gene Rearrangements:** ROS1 is a receptor tyrosine kinase that can be rearranged in NSCLC, resulting in dysregulation and inappropriate signaling through the ROS1 kinase domain. Numerous fusion partners are seen with ROS1, and common fusion partners include ESRB1, CD74, and HNRNBA. The presence of a ROS1 rearrangement is associated with responsiveness to tyrosine kinase inhibitors. Some clinicopathologic features—such as smoking status and histology—however, these features should not be utilized in selecting patients for testing. Testing Methodologies: FISH break-apart probe methodology can be used for screening strategy. IHC for ROS1 fusions has low sensitivity. Numerous NGS methodologies can detect ROS1 fusions, although DNA-based NGS may under-detect ROS1 fusions. Targeted real-time PCR assays are utilized in some settings, although they are unlikely to detect fusions with novel partners.

ROS1

**Tanslocation ROS-1**  
**Méthode de detection standard = IHC a confirmer**  
**FISH (Hybridation in situ par fluorescence).**

- ▶ **BRAF (B-Raf proto-oncogene) point mutations:** BRAF is a serine/threonine kinase that is part of the canonical MAP/ERK signaling pathway. Activating mutations in BRAF result in unregulated signaling through the MAP/ERK pathway. Mutations in BRAF can be seen in NSCLC. The presence of a specific mutation resulting in a change in BRAF activity is associated with responsiveness to combined therapy with oral inhibitors of BRAF and MEK. Numerous other mutations in BRAF are observed in NSCLC, and the impact of those mutations on their effect on signaling is unclear.

BRAF

**Mutation BRAF V600E**  
**PCR, Sanger sequencing, NGS**

Testing Methodologies: Real-time PCR, Sanger sequencing (ideally paired with tumor enrichment), and NGS are the most commonly deployed methodologies for examining BRAF mutation status. While an anti-BRAF p.V600E-specific monoclonal antibody is commercially available, and some studies have examined utilizing this approach, it should only be deployed after extensive validation.

- ▶ **KRAS (KRAS proto-oncogene) point mutations:** KRAS is a G-protein with intrinsic GTPase activity, and activating mutations result in unregulated signaling through the MAP/ERK pathway. Mutations in KRAS are most commonly seen at codon 12, and the presence of a KRAS mutation is prognostic of poor survival. Numerous other mutations in KRAS have been associated with reduced response to EGFR tyrosine kinase inhibitors due to the low probability of overlapping targetable alterations. Patients with KRAS mutations are unlikely to benefit from further molecular testing.

KRAS

**Mutation KRAS = Plus fréquente des mutations, F mauvais pronostic**  
**F de non réponse aux TKI classique**

### PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS

• Molecular Targets for Analysis (continued)

▶ **MET** (mesenchymal-epithelial transition) **exon 14 (METex14)** skipping variants: MET is a receptor tyrosine kinase. A mutation in NSCLC. Loss of **METex14** leads to dysregulation and inappropriate signaling. Presence of **METex14** skipping mutation is associated with responsiveness to oral MET TKIs.

**MET exon 14 =**

- NGS
- RNA NGS

◊ Testing Methodologies: NGS-based testing is the primary method for detection of **METex14** skipping events, with RNA-based NGS demonstrating improvement in detection. IHC is not a method for detection of **METex14** skipping.

▶ **RET** (rearranged during transfection) Gene Rearrangements: RET is a receptor tyrosine kinase that can be rearranged in NSCLC, resulting in dysregulation and inappropriate signaling through the RET kinase domain.

**Rearrangement RET =**

- FISH
- RNA NGS

◊ The presence of a **RET** rearrangement is associated with responsiveness to oral RET TKIs regardless of fusion partner. Common fusion partners are *KIF5B*, *NCOA4*, and *CCDC6*; however, numerous other fusion partners have been identified.

◊ Testing Methodologies: FISH break-apart probe methodology can be deployed; however, it may under-detect some fusions. Reverse-transcriptase PCR assays are utilized in some settings, although they are unlikely to detect fusions with novel partners. NGS-based methodology has a high specificity, and RNA-based NGS is preferable to DNA-based NGS for fusion detection.

▶ **NTRK1/2/3** (neurotrophic tyrosine receptor kinase) gene fusions

◊ **NTRK** are tyrosine receptor kinases that are rarely rearranged in NSCLC as well as in other tumor types, resulting in dysregulation and inappropriate signaling.

**Fusion NTRK = FISH, IHC, PCR, NGS**

◊ Numerous fusion partners have been identified. To date, no specific clinicopathologic features, other than absence of other driver alterations, have been identified in association with **NTRK** fusions.

◊ Point mutations in **NTRK1/2/3** are generally non-activating and have not been studied in association with targeted therapy.

◊ Testing Methodologies: Various methodologies can be used to detect **NTRK1/2/3** gene fusions, including: FISH, IHC, PCR, and NGS; false negatives may occur. IHC methods are complicated by baseline expression in some tissues. FISH testing may require at least 3 probe sets for full analysis. NGS testing can detect a broad range of alterations. DNA-based NGS may under-detect **NTRK1** and **NTRK3** fusions.

**En cas de progression  
s/TKI**

## PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS

## • Testing in the Setting of Progression on Targeted Therapy:

▶ For many of the above listed analytes, there is growing recognition of the molecular mechanisms of resistance to therapy. Re-testing of a sample from a tumor that is actively progressing while exposed to targeted therapy can shed light on appropriate next therapeutic steps:

→ ◊ For patients with an underlying *EGFR* sensitizing mutation who have been treated with *EGFR* TKI, minimum appropriate testing includes high-sensitivity evaluation for *p.T790M*; when there is no evidence of *p.T790M*, testing for alternate mechanisms of resistance (*MET* amplification, *ERBB2* amplification) may be used to direct patients for additional therapies. The presence of *p.T790M* can direct patients to third-generation *EGFR* TKI therapy.

– Assays for the detection of *EGFR p.T790M* should be designed to have an analytic sensitivity of a minimum of 5% allelic fraction. The original sensitizing mutation can be utilized as an internal control in many assays to determine whether a *p.T790M* is within the range of detection if present as a sub-clonal event.

→ ◊ For patients with underlying *ALK* rearrangement who have been treated with *ALK* TKI, it is unclear whether identification of specific tyrosine kinase domain mutation can identify appropriate next steps in therapy, although some preliminary data suggest that specific kinase domain mutations can impact next line of therapy.

**PD-L1****PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS**

- **PD-L1 (Programmed Death Ligand 1):** PD-L1 is a co-regulatory molecule that can be expressed on tumor cells and inhibit T-cell-mediated cell death. T-cells express PD-1, a negative regulator, which binds to ligands including PD-L1 (CD274) or PD-L2 (CD273). In the presence of PD-L1, T-cell activity is suppressed.
  - ▶ Checkpoint inhibitor antibodies block the PD-1 and PD-L1 interaction, thereby improving the antitumor effects of endogenous T cells.
  - ▶ IHC for PD-L1 can be utilized to identify disease most likely to respond to first-line anti PD-1/PD-L1.
    - ◇ Various antibody clones have been developed for IHC analysis of PD-L1 expression, and while several show relative equivalence, some do not.
    - ◇ Interpretation of PD-L1 IHC in NSCLC is typically focused on the proportion of tumor cells expressing membranous staining at any level and therefore is a linear variable; scoring systems may be different in other tumor types.
    - ◇ The FDA-approved companion diagnostic for PD-L1 guides utilization of pembrolizumab in patients with NSCLC and is based on the tumor proportion score (TPS). TPS is the percentage of viable tumor cells showing partial or complete membrane staining at any intensity.
    - ◇ The definition of positive and negative testing is dependent on the individual antibody and platform deployed, which may be unique to each checkpoint inhibitor therapy. The potential for multiple different assays for PD-L1 has raised concern among both pathologists and oncologists.
    - ◇ Although PD-L1 expression can be elevated in patients with an oncogenic driver, targeted therapy for the oncogenic driver should take precedence over treatment with an immune checkpoint inhibitor.

**Testing PDL-1:****==> IHC****==> cut of > 50 % L1  
> 1 - 49 %****TPS :****% de cellules cancéreuses viables qui exprime  
une fixation partielle ou complète au niveau de  
la membrane cellulaire quelle que soit**

**PRINCIPLES OF MOLECULAR AND BIOMARKER ANALYSIS****• Plasma Cell-Free/Circulating Tumor DNA Testing:**

- ▶ Cell-free/circulating tumor DNA testing should not be used in lieu of a histologic tissue diagnosis.
- ▶ Some laboratories offer testing for molecular alterations examining nucleic acids in peripheral circulation, most commonly in processed plasma (sometimes referred to as "liquid biopsy").
- ▶ Studies have demonstrated cell-free tumor DNA testing to generally have very high specificity, but significantly compromised sensitivity, with up to 30% false-negative rate.
- ▶ Standards for analytical performance characteristics of cell-free tumor DNA have not been established, and in contrast to tissue-based testing, no guidelines exist regarding the recommended performance characteristics of this type of testing.
- ▶ Cell-free tumor DNA testing can identify alterations that are unrelated to a lesion of interest, for example, clonal hematopoiesis of indeterminate potential (CHIP).
- ▶ The use of cell-free/circulating tumor DNA testing can be considered in specific clinical circumstances, most notably:
  - ◇ If a patient is medically unfit for invasive tissue sampling
  - ◇ In the initial diagnostic setting, if following pathologic confirmation of a NSCLC diagnosis there is insufficient material for molecular analysis, cell-free/circulating tumor DNA should be used only if follow-up tissue-based analysis is planned for all patients in which an oncogenic driver is not identified ([see NSCL-18](#) for oncogenic drivers with available targeted therapy options).

**ADN tumoral circulant:**

**A utiliser si :**

- Patient ne pouvant pas supporter un geste invasif

# NCCN Guidelines Version 7.2021

## Non-Small Cell Lung Cancer

### CLINICAL PRESENTATION

### HISTOLOGIC SUBTYPE<sup>a</sup>

### BIOMARKER TESTING<sup>mm</sup>

Advanced or metastatic disease

- Establish histologic subtype<sup>a</sup> with adequate tissue for molecular testing (consider rebiopsy<sup>ll</sup> if appropriate)
- Smoking cessation counseling
- Integrate palliative care<sup>c</sup> ([See NCCN Guidelines for Palliative Care](#))

- Adenocarcinoma
- Large cell
- NSCLC not otherwise specified (NOS)

Squamous cell carcinoma

- Molecular testing, including:
  - ▶ *EGFR* mutation (category 1), *ALK* (category 1), *KRAS*, *ROS1*, *BRAF*, *NTRK1/2/3*, *MET* exon 14 skipping, *RET*
  - ▶ Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)

- Consider molecular testing, including:<sup>oo</sup>
  - ▶ *EGFR* mutation, *ALK*, *KRAS*, *ROS1*, *BRAF*, *NTRK1/2/3*, *MET* exon 14 skipping, *RET*
  - ▶ Testing should be conducted as part of broad molecular profiling<sup>nn</sup>
- PD-L1 testing (category 1)

### Quels tests moléculaires :

- Mutation *EGFR*
- Translocation *ALK*
- Translocation *ROS1*
- Mutation *BRAF* V600E
- perte *MET* exon 14
- Réarrangement *RET*
- Fusion gènes *NTRK* 1, 2, 3
- *KRAS* G12C
- Surexpression *PDL1*

### Pour qui :

- *ADK*
- Carcinome à grandes cellules
- *CBNPC NOS*
- Carcinome épidermoïde : si non ou peu fumeur < 15 p/a, si petite Bx. si histologie mixte.

# NCCN Guidelines Version 2.2021 Non-Small Cell Lung Cancer

1

2

3

4

## CLINICAL PRESENTATION

## HISTOLOGIC SUBTYPE<sup>a</sup>

## BIOMARKER TESTING<sup>ll</sup>

## TESTING RESULTS<sup>ij</sup>

- Establish histologic subtype<sup>a</sup> with adequate tissue for molecular testing (consider rebiopsy<sup>kk</sup> if appropriate)
- Smoking cessation counseling
- Integrate palliative care<sup>c</sup> (See [NCCN Guidelines for Palliative Care](#))

- Adenocarcinoma
- Large cell
- NSCLC not otherwise specified (NOS)

Squamous cell carcinoma

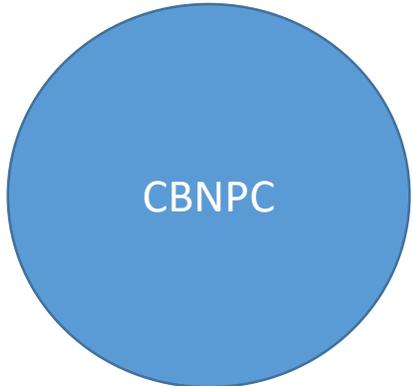
- Molecular testing, including:
  - *EGFR* mutation (category 1), *ALK* (category 1), *ROS1*, *BRAF*, *NTRK1/2/3*, *MET* exon 14 skipping, *RET*
  - Testing should be conducted as part of broad molecular profiling<sup>mmm</sup>
- PD-L1 testing (category 1)

- Consider molecular testing, including:<sup>nn</sup>
  - *EGFR* mutation, *ALK*, *ROS1*, *BRAF*, *NTRK1/2/3*, *MET* exon 14 skipping, *RET*
  - Testing should be conducted as part of broad molecular profiling<sup>mmm</sup>
- PD-L1 testing (category 1)

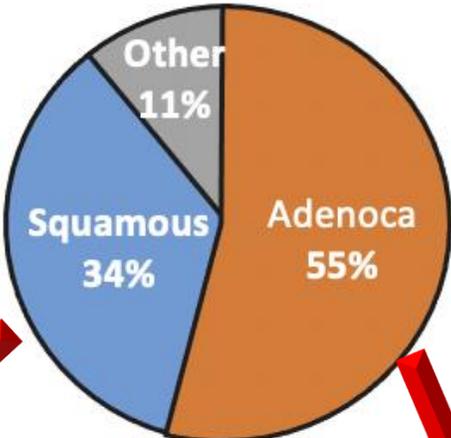
- Sensitizing *EGFR* mutation positive (see [NSCL-19](#))
- *ALK* positive (see [NSCL-22](#))
- *ROS1* positive (see [NSCL-25](#))
- *BRAF* V600E positive (see [NSCL-26](#))
- *MET* exon 14 skipping mutation positive (see [NSCL-28](#))
- *RET* positive (see [NSCL-29](#))
- PD-L1 ≥1% and *EGFR*, *ALK*, *ROS1*, *BRAF*, *MET* exon 14 skipping mutation, and *RET* negative<sup>ii</sup> (see [NSCL-30](#))
- PD-L1 <1% and *EGFR*, *ALK*, *ROS1*, *BRAF*, *MET* exon 14 skipping mutation, and *RET* negative<sup>ii</sup> (see [NSCL-32](#))
- Sensitizing *EGFR* mutation positive (see [NSCL-19](#))
- *ALK* positive (see [NSCL-22](#))
- *ROS1* positive (see [NSCL-25](#))
- *BRAF* V600E positive (see [NSCL-26](#))
- *MET* exon 14 skipping mutation positive (see [NSCL-28](#))
- *RET* positive (see [NSCL-29](#))
- PD-L1 ≥1% and *EGFR*, *ALK*, *ROS1*, *BRAF*, *MET* exon 14 skipping mutation, and *RET* negative<sup>ii</sup> (see [NSCL-30](#))
- PD-L1 <1% and *EGFR*, *ALK*, *ROS1*, *BRAF*, *MET* exon 14 skipping mutation, and *RET* negative<sup>ii</sup> (see [NSCL-33](#))

Advanced or metastatic disease

# Drivers moléculaires



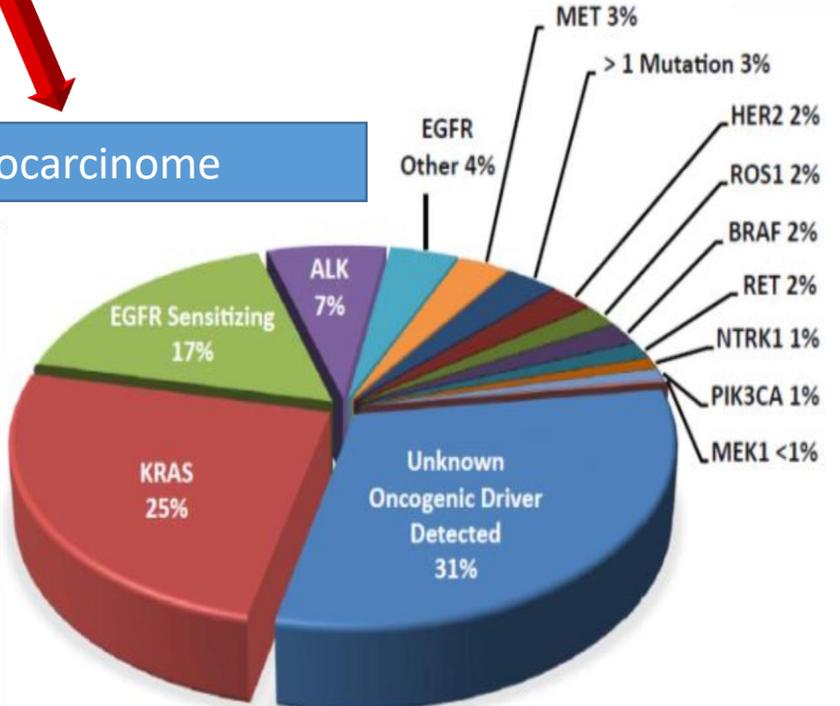
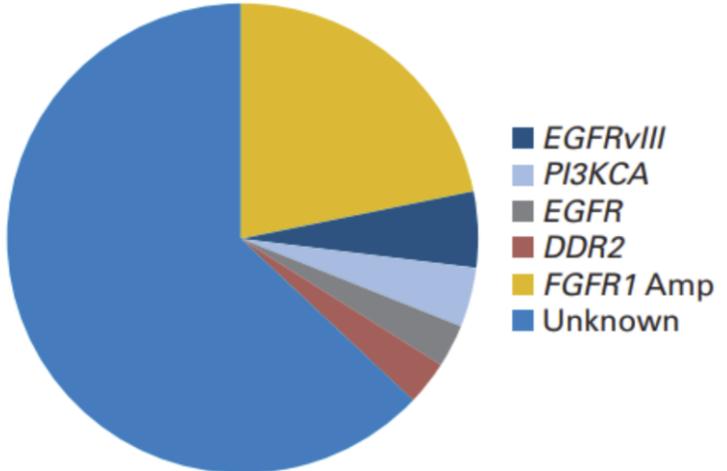
1990 - 2004



2004 - 2009

Carcinome épidermoïde

Adénocarcinome



2020

**PDL-  
1**

- Adénocarcinome
- Carcinome à grandes cellules
- CBNPC NOS
- Carcinome épidermoïde :  
si non ou peu fumeur < 15 p/a  
si petite biopsie  
si histologie mixte

Carcinome épidermoïde

**Drivers  
moléculaires**

Patients

**Avec** drivers moléculaires

PDL-1 < 1 %, ≥ 1-49 %, ≥ 50 %



**Thérapies ciblées**

+/- anti-angiogénique

Patients

**sans** drivers moléculaires

PDL-1 < 1 %



**Chimiothérapie +/-  
anti-angiogéniques**

Patients

**sans** drivers moléculaires

PDL-1 ≥ 1-49 %



**Immunothérapie +**

**Chimiothérapie**

+/- Anti-angiogénique

Patients

**sans** drivers moléculaires

PDL-1 ≥ 50 %



**Immunothérapie**

IO + CT +/- anti-angiogénique

Patients  
Avec drivers moléculaires  
PDL-1 < 1 %, ≥ 1-49 %, ≥ 50 %

# Arbre décisionnel 2021, 1<sup>ère</sup> ligne de traitement

C non épidermoïde

C épidermoïde

EGFR  
Exon 19

ALK

ROS-1

BRAF

NTRK

MET

RET

Osimertinib  
Erlotinib  
Afatinib  
Gefitinib  
Dacomitinib  
Erlo + Ramu  
Erlo + Beva

Alectinib  
Brigatinib  
Lorlatinib  
Ceritinib  
Crizotinib

Entrectinib  
Crizotinib  
Ceritinib

Dabrafenib +  
Trametinib  
Vemurafenib

Larotrectinib  
Entrectinib

Capmatinib  
Tepotinib  
crizotinib

Selpercatinib  
Pralsetinib  
Cabozantinib  
Vandetanib

EGFR  
Exon 20

L1 = CT

L2 = Amivantamab  
ou Mobocertinib

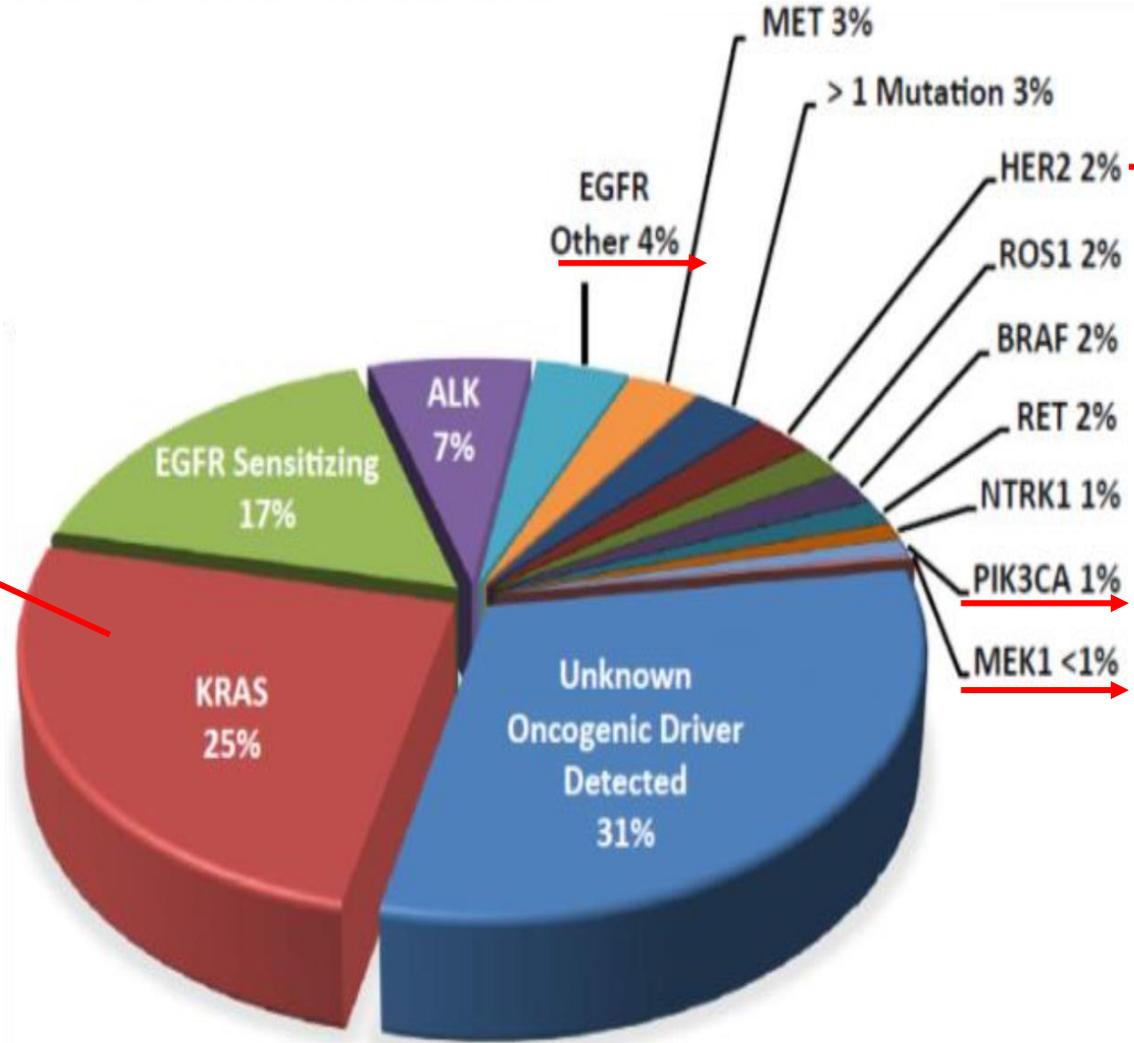
KRAS  
G12C

L1 = CT

L2 = Sotorasib

# Adenocarcinoma

Inhibiteur de la mutation KRAS G12C :  
- Sotorasib (AMG 510)  
- Adagrasib



Anti-Her2:  
- Pozotinib  
- Trastuzumab Deruxtecan

Patients  
sans drivers moléculaires  
PDL-1 < 1 %

# Arbre décisionnel 2021, 1<sup>ère</sup> ligne de traitement

## C non épidermoïde

## C épidermoïde

### Contraindications to PD-1 or PD-L1 inhibitors<sup>c</sup>

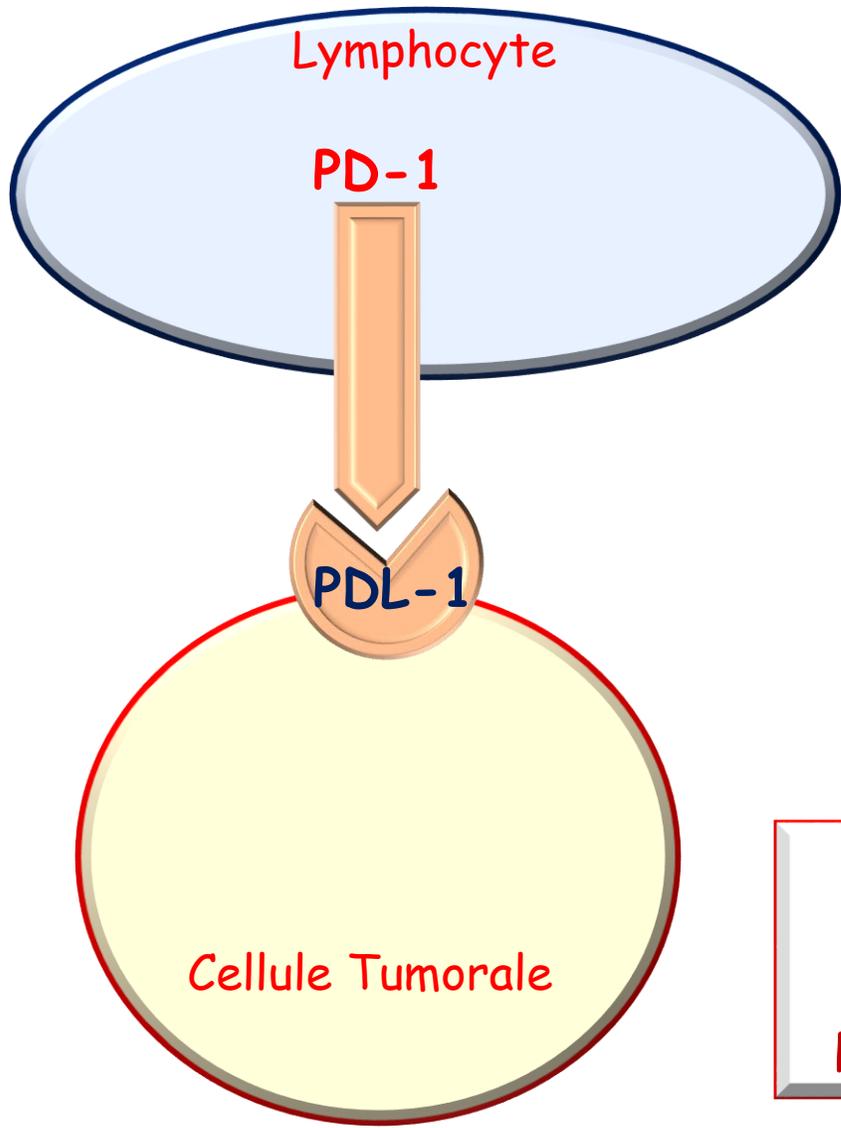
#### Useful in Certain Circumstances

- Bevacizumab<sup>e</sup>/carboplatin/paclitaxel (category 1)<sup>7,f,g,h</sup>
- Bevacizumab<sup>e</sup>/carboplatin/pemetrexed<sup>7,8,f,g,h</sup>
- Bevacizumab<sup>e</sup>/cisplatin/pemetrexed<sup>9,f,g,h</sup>
- Carboplatin/albumin-bound paclitaxel (category 1)<sup>10</sup>
- Carboplatin/docetaxel (category 1)<sup>11</sup>
- Carboplatin/etoposide (category 1)<sup>12,13</sup>
- Carboplatin/gemcitabine (category 1)<sup>14</sup>
- Carboplatin/paclitaxel (category 1)<sup>15</sup>
- Carboplatin/pemetrexed (category 1)<sup>16</sup>
- Cisplatin/docetaxel (category 1)<sup>11</sup>
- Cisplatin/etoposide (category 1)<sup>17</sup>
- Cisplatin/gemcitabine (category 1)<sup>15,18</sup>
- Cisplatin/paclitaxel (category 1)<sup>19</sup>
- Cisplatin/pemetrexed (category 1)<sup>18</sup>
- Gemcitabine/docetaxel (category 1)<sup>20</sup>
- Gemcitabine/vinorelbine (category 1)<sup>21</sup>

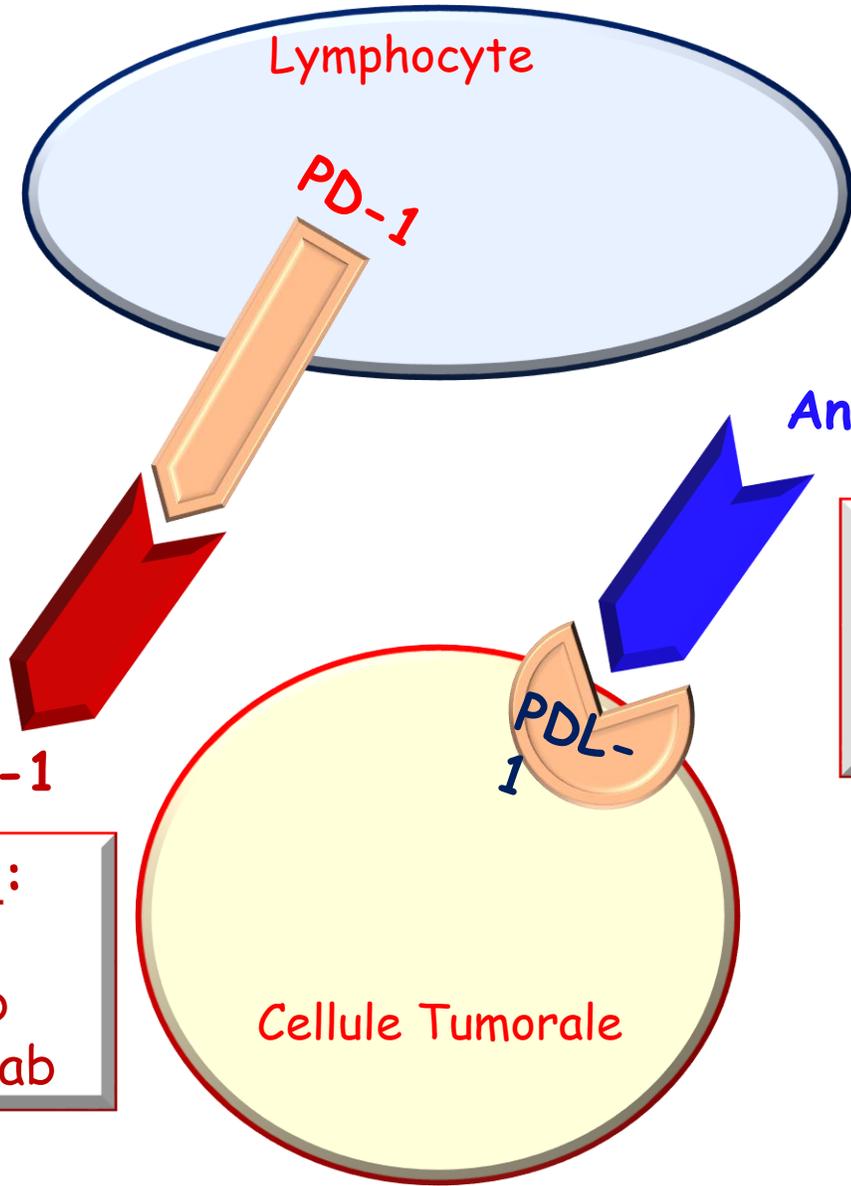
### Contraindications to PD-1 or PD-L1 inhibitors<sup>c</sup>

#### Useful in Certain Circumstances

- Carboplatin/albumin-bound paclitaxel (category 1)<sup>9</sup>
- Carboplatin/docetaxel (category 1)<sup>11</sup>
- Carboplatin/gemcitabine (category 1)<sup>14</sup>
- Carboplatin/paclitaxel (category 1)<sup>15</sup>
- Cisplatin/docetaxel (category 1)<sup>11</sup>
- Cisplatin/etoposide (category 1)<sup>17</sup>
- Cisplatin/gemcitabine (category 1)<sup>15,18</sup>
- Cisplatin/paclitaxel (category 1)<sup>19</sup>
- Gemcitabine/docetaxel (category 1)<sup>20</sup>
- Gemcitabine/vinorelbine (category 1)<sup>21</sup>



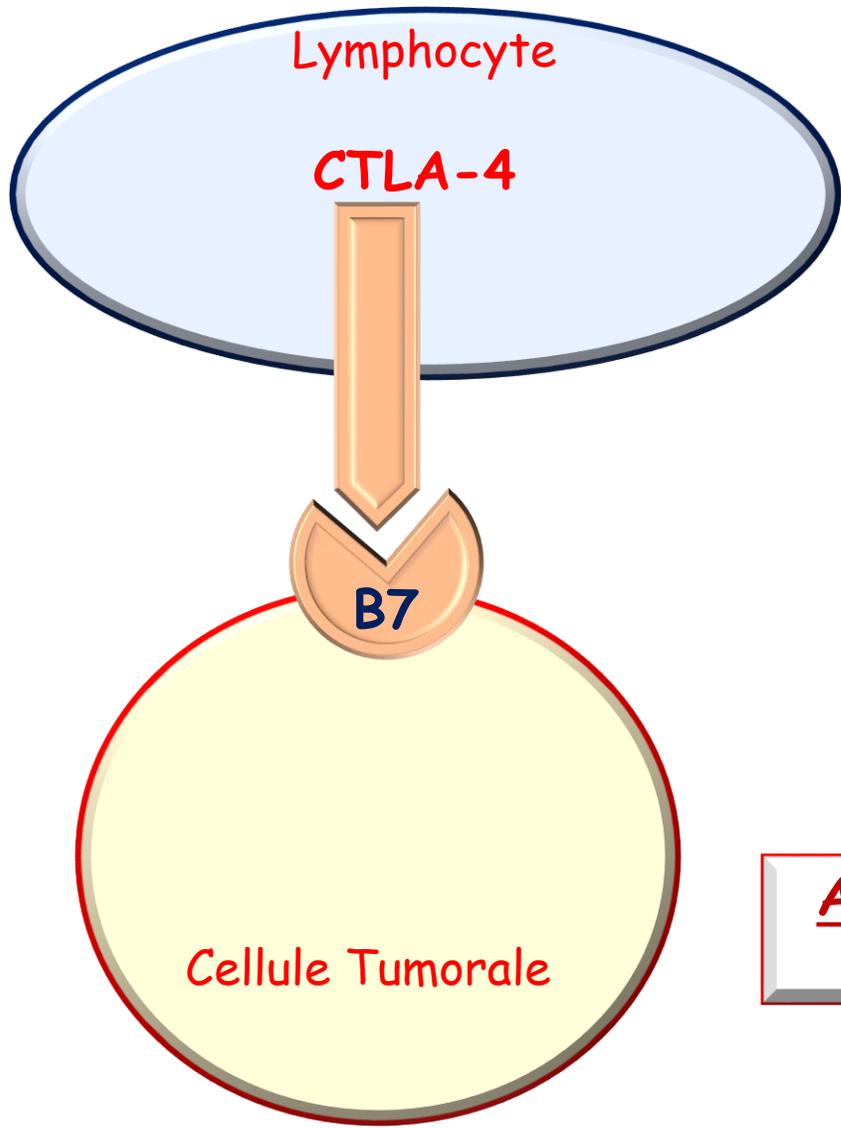
La liaison PD-1 et PDL-1 empêche l'activation lymphocytaire.



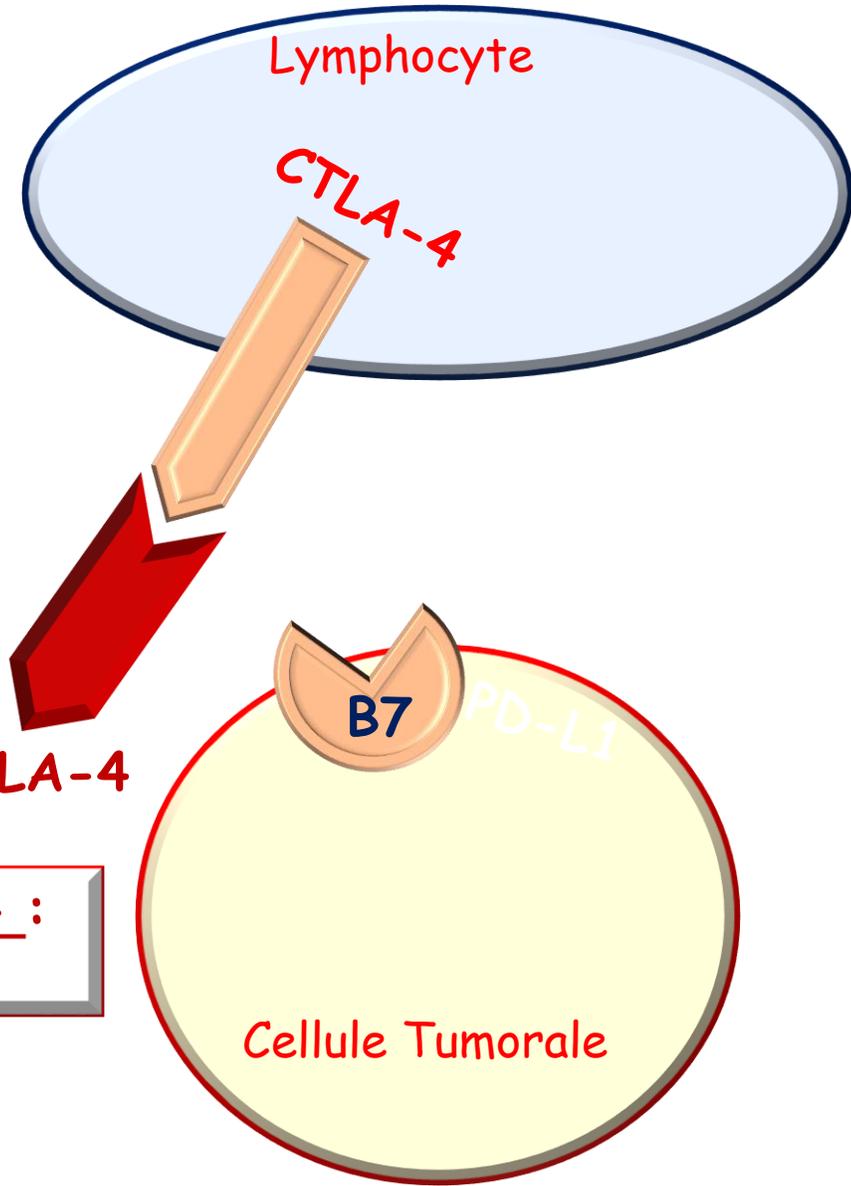
Anti PD-1 :  
Nivolumab  
Cemiplimab  
Pembrolizumab

Anti PDL-1 :  
Atezolizumab  
Avelumab  
Durvalumab

En bloquant la liaison PDL-1 et PD-1  
Le lymphocyte est réactivé.



La liaison CTLA-4 au B7 empêche l'activation lymphocytaire.



Anti CTLA-4 :  
Ipilimumab

En bloquant la liaison CTLA-4 au B7 Le lymphocyte est réactivé.

Patients  
sans drivers moléculaires  
PDL-1 ≥ 1- 49 %

# NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

PD-L1 EXPRESSION POSITIVE (≥1%–49%)<sup>mm</sup>

FIRST-LINE THERAPY<sup>PP</sup>

PD-L1 expression  
positive (≥1%–49%)  
and negative  
for actionable  
molecular  
markers and no  
contraindications  
to PD-1 or PD-L1  
inhibitors<sup>III</sup>

PS 0–2

Adenocarcinoma,  
large cell, NSCLC  
NOS

Squamous cell  
carcinoma

- **Preferred**  
(Carboplatin or cisplatin) + pemetrexed + pembrolizumab (category 1)
- **Other Recommended**  
Carboplatin + paclitaxel + bevacizumab<sup>tt</sup> + atezolizumab (category 1)  
or  
Carboplatin + albumin-bound paclitaxel + atezolizumab  
or  
Nivolumab + ipilimumab + pemetrexed + (carboplatin or cisplatin) (category 1)
- **Useful in Certain Circumstances**  
Nivolumab + ipilimumab (category 1)  
or  
Pembrolizumab (category 2B)<sup>ttt</sup>

- **Preferred**  
Carboplatin + (paclitaxel or albumin-bound paclitaxel) + pembrolizumab (category 1)
- **Other Recommended**  
Nivolumab + ipilimumab + paclitaxel + carboplatin (category 1)
- **Useful in Certain Circumstances**  
Nivolumab + ipilimumab (category 1)  
or  
Pembrolizumab (category 2B)<sup>ttt</sup>

[See PD-L1 expression positive \(≥50%\) NSCLC-33](#)

Patients sans drivers moléculaires  
 PDL-1 ≥ 1- 49 %

Guidelines Version 7.2021  
 Non-Small Cell Lung Cancer

PD-L1 EXPRESSION POSITIVE (≥1%–49%)<sup>mmm</sup>

FIRST-LINE THERAPY<sup>pp</sup>

PD-L1 expression positive (≥1%–49%) and negative for actionable molecular markers and no contraindications to PD-1 or PD-L1 inhibitors<sup>lll</sup>

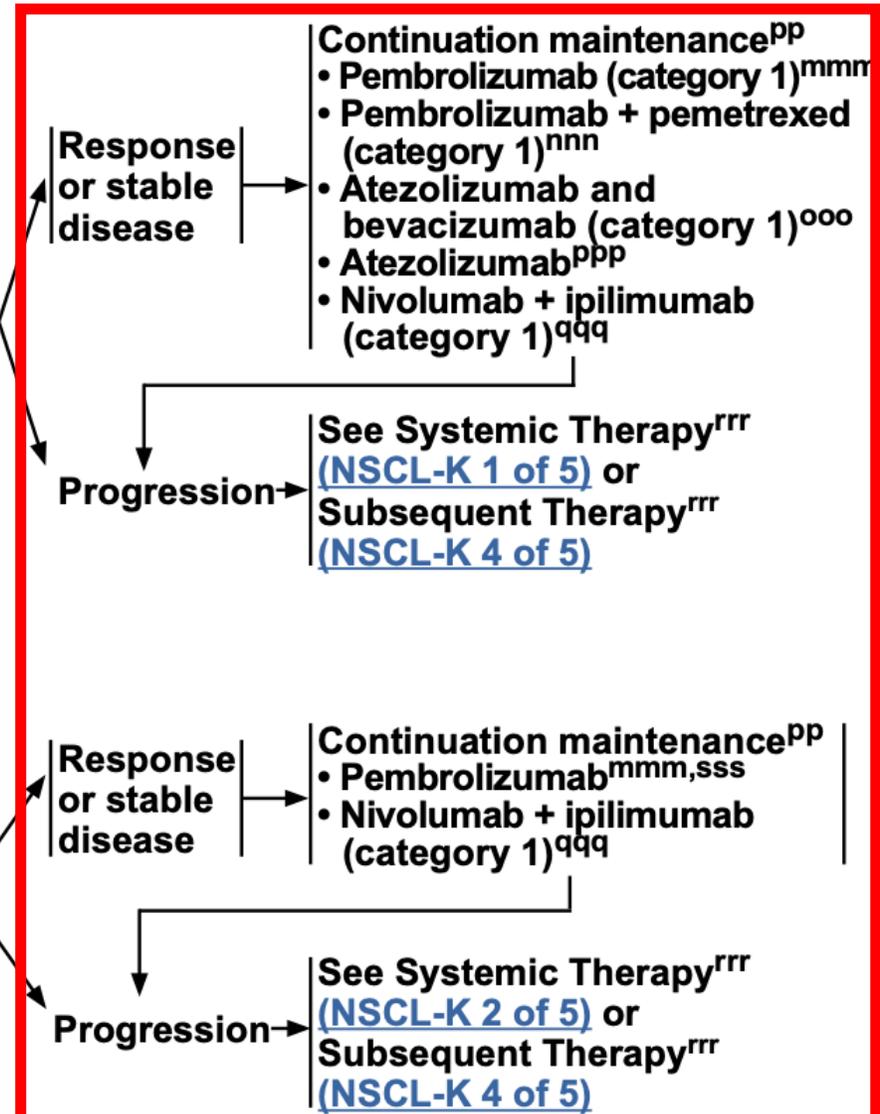
PS 0–2

Adenocarcinoma, large cell, NSCLC NOS

Squamous cell carcinoma

- **Preferred**  
 (Carboplatin or cisplatin) + pemetrexed + pembrolizumab (category 1)
- **Other Recommended**  
 Carboplatin + paclitaxel + bevacizumab<sup>tt</sup> + atezolizumab (category 1)  
 or  
 Carboplatin + albumin-bound paclitaxel + atezolizumab  
 or  
 Nivolumab + ipilimumab + pemetrexed + (carboplatin or cisplatin) (category 1)
- **Useful in Certain Circumstances**  
 Nivolumab + ipilimumab (category 1)  
 or  
 Pembrolizumab (category 2B)<sup>ttt</sup>

- **Preferred**  
 Carboplatin + (paclitaxel or albumin-bound paclitaxel) + pembrolizumab (category 1)
- **Other Recommended**  
 Nivolumab + ipilimumab + paclitaxel + carboplatin (category 1)
- **Useful in Certain Circumstances**  
 Nivolumab + ipilimumab (category 1)  
 or  
 Pembrolizumab (category 2B)<sup>ttt</sup>



See PD-L1 expression positive (≥50%) NSCL-33

Patients  
sans drivers moléculaires  
PDL-1 ≥ 50 %

# NCCN Guidelines Version 7.2021 Non-Small Cell Lung Cancer

PD-L1 EXPRESSION POSITIVE (≥50%)<sup>mm</sup>

FIRST-LINE THERAPY<sup>pp</sup>

PD-L1 expression positive (≥50%) and negative for actionable molecular markers and no contraindications to PD-1 or PD-L1 inhibitors<sup>lll</sup>

PS 0-2

Adenocarcinoma, large cell, NSCLC NOS

Squamous cell carcinoma

- **Preferred**  
Pembrolizumab (category 1)  
or  
(Carboplatin or cisplatin) + pemetrexed + pembrolizumab (category 1)  
or  
Atezolizumab (category 1)  
or  
Cemiplimab-rwlc (category 1)
- **Other Recommended**  
Carboplatin + paclitaxel + bevacizumab<sup>tt</sup> + atezolizumab (category 1)  
or  
Carboplatin + albumin-bound paclitaxel + atezolizumab  
or  
Nivolumab + ipilimumab + pemetrexed + (carboplatin or cisplatin) (category 1)
- **Useful in Certain Circumstances**  
Nivolumab + ipilimumab (category 1)
- **Preferred**  
Pembrolizumab (category 1)  
or  
Carboplatin + (paclitaxel or albumin-bound paclitaxel) + pembrolizumab (category 1)  
or  
Atezolizumab (category 1)  
or  
Cemiplimab-rwlc (category 1)
- **Other Recommended**  
Nivolumab + ipilimumab + paclitaxel + carboplatin (category 1)
- **Useful in Certain Circumstances**  
Nivolumab + ipilimumab (category 1)

[See PD-L1 expression positive \(≥1%–49%\) NSCLC-34](#)

## PD-L1 EXPRESSION POSITIVE (≥50%)<sup>mm</sup>

## FIRST-LINE THERAPY<sup>pp</sup>

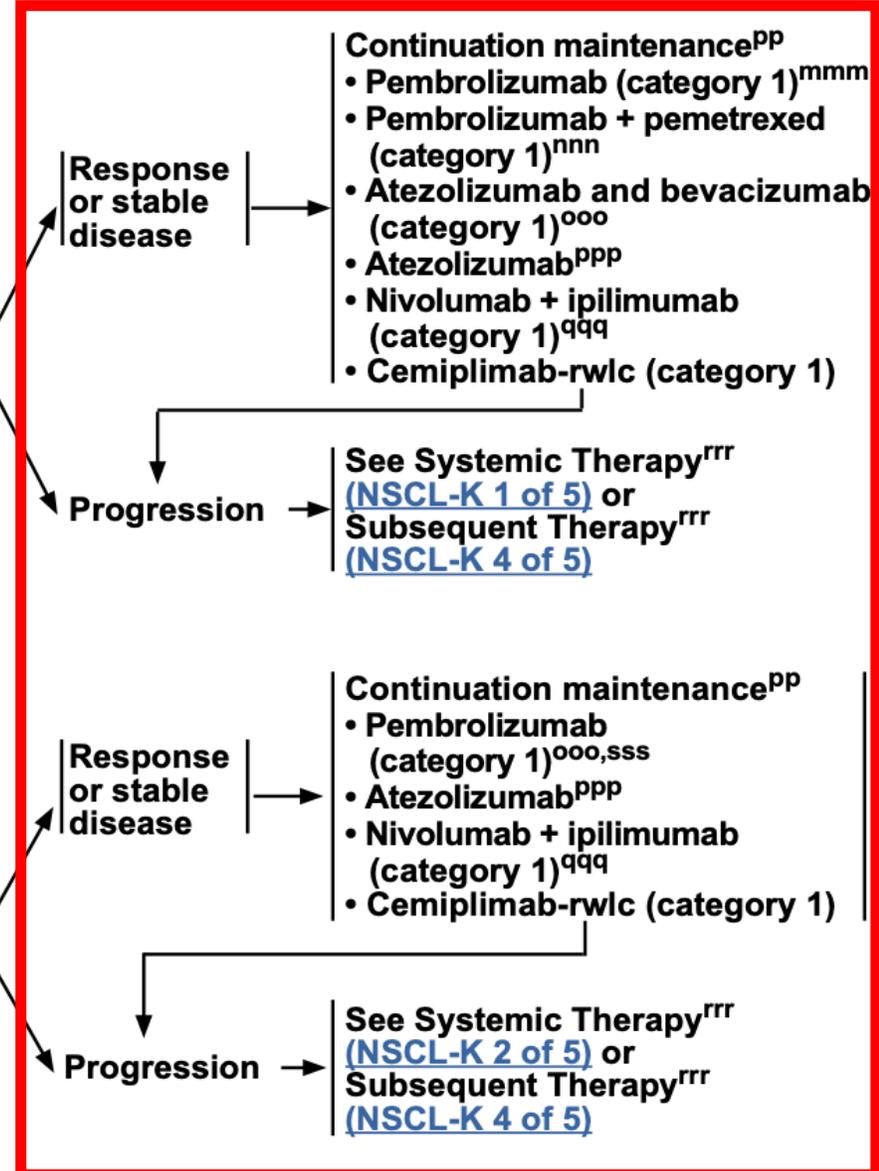
PD-L1 expression positive (≥50%) and negative for actionable molecular markers and no contraindications to PD-1 or PD-L1 inhibitors<sup>iii</sup>

Adenocarcinoma, large cell, NSCLC NOS

PS 0-2

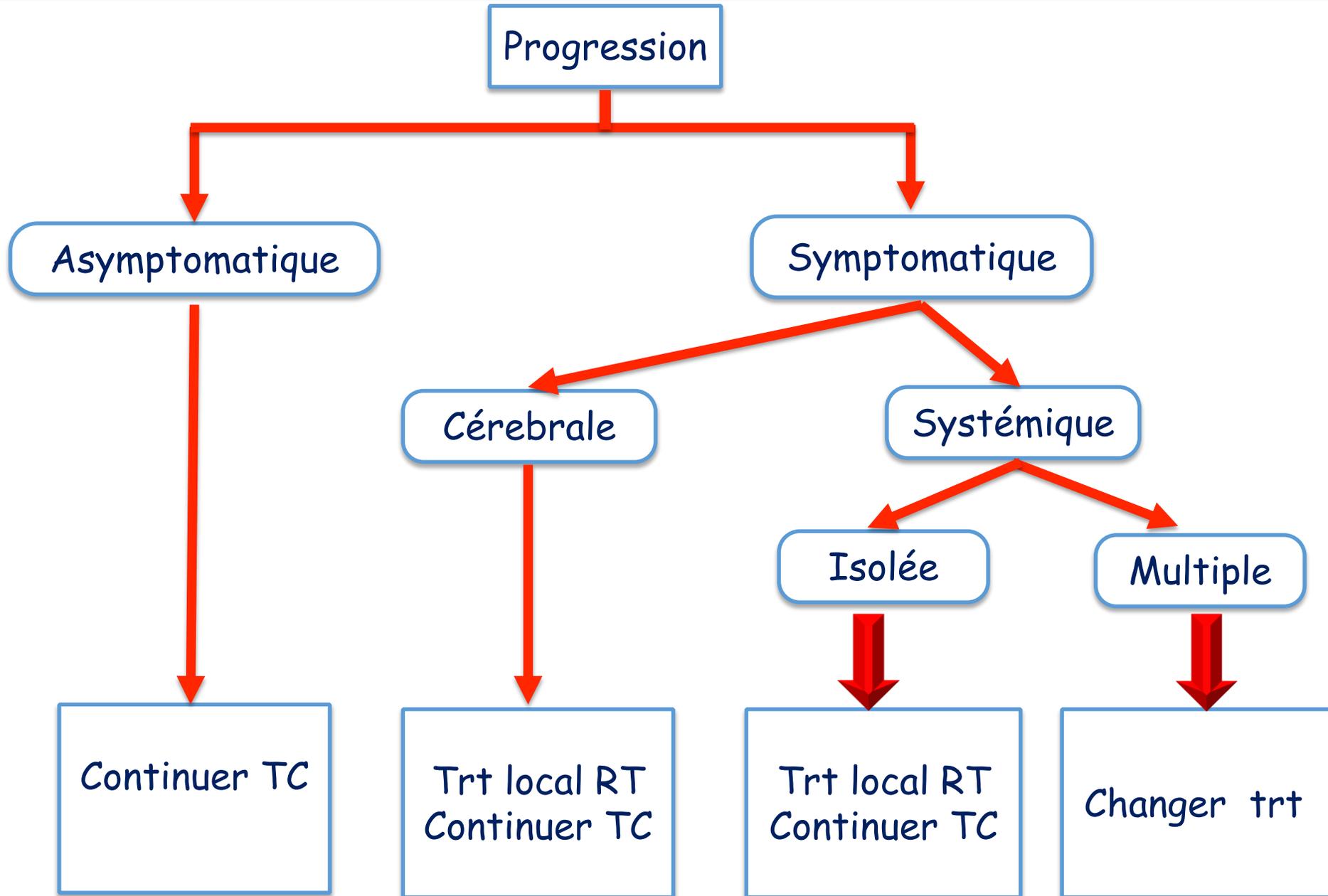
Squamous cell carcinoma

- **Preferred**  
Pembrolizumab (category 1)  
or  
(Carboplatin or cisplatin) + pemetrexed + pembrolizumab (category 1)  
or  
Atezolizumab (category 1)  
or  
Cemiplimab-rwlc (category 1)
  - **Other Recommended**  
Carboplatin + paclitaxel + bevacizumab<sup>tt</sup> + atezolizumab (category 1)  
or  
Carboplatin + albumin-bound paclitaxel + atezolizumab  
or  
Nivolumab + ipilimumab + pemetrexed + (carboplatin or cisplatin) (category 1)
  - **Useful in Certain Circumstances**  
Nivolumab + ipilimumab (category 1)
- 
- **Preferred**  
Pembrolizumab (category 1)  
or  
Carboplatin + (paclitaxel or albumin-bound paclitaxel) + pembrolizumab (category 1)  
or  
Atezolizumab (category 1)  
or  
Cemiplimab-rwlc (category 1)
  - **Other Recommended**  
Nivolumab + ipilimumab + paclitaxel + carboplatin (category 1)
  - **Useful in Certain Circumstances**  
Nivolumab + ipilimumab (category 1)



[See PD-L1 expression positive \(≥1%–49%\) NSCL-34](#)

# Stratégie thérapeutique / patient sous T Ciblée



**Conclusion**