



Identificación de perfiles moleculares en el CPRC hacia la medicina de la precisión

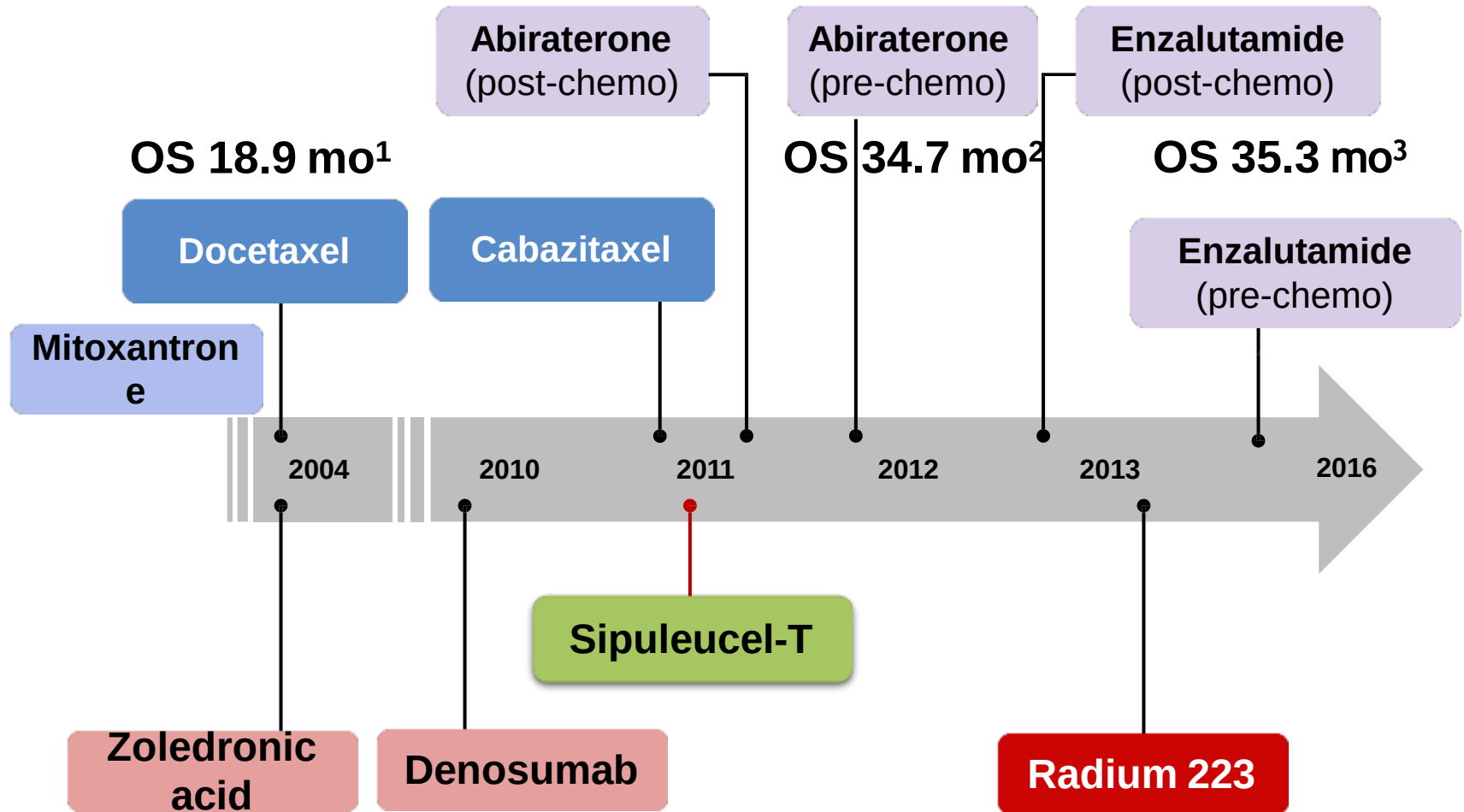
Dr. Antonio Ruiz Fernández

Adjunto Servicio Oncología Medica Área Hospitalaria Zafra-Llerena

INTRODUCCIÓN

- El receptor androgénico como mecanismo clave para la progresión del CPRC
- **El presente.** Las primeras terapias dirigidas al receptor androgénico.
- Principios de Biología Molecular.
- **Presente-Futuro.**
 - ✓ Paneles Genéticos. La búsqueda de Biomarcadores pronósticos y predictivos.
 - ✓ La Biopsia Líquida en la obtención de los Biomarcadores.
 - ✓ La inmunoterapia llega a mCPRC ?
 - ✓ Los primeros biomarcadores. BCRA2. **El tratamiento futuro** hacia la medicina de la precisión.

Cronología aprobación de fármacos en CPRCm



El receptor Androgénico como mecanismo clave para la progresión del CPRC

Andrógenos: papel que desempeñan en el cáncer de próstata

- El cáncer de próstata es un tumor sólido maligno de origen hormonal¹.
- Los andrógenos (testosterona y DHT) son los principales determinantes del crecimiento de las células prostáticas normales y cancerosas¹.
 - Entre el 90 y el 95 % de la testosterona circulante se produce en los testículos.
 - El 5-10 % restante lo sintetizan las glándulas suprarrenales.
- En el cáncer de próstata, los andrógenos y otras moléculas transmiten señales de crecimiento y proliferación a las células malignas².
 - Los andrógenos se unen a los RA nucleares activando genes diana que favorecen el crecimiento de las células prostáticas.
 - Por lo tanto, los andrógenos son los principales promotores de la progresión del cáncer de próstata.
- La reducción de los niveles de andrógenos induce la apoptosis de las células prostáticas malignas y normales².

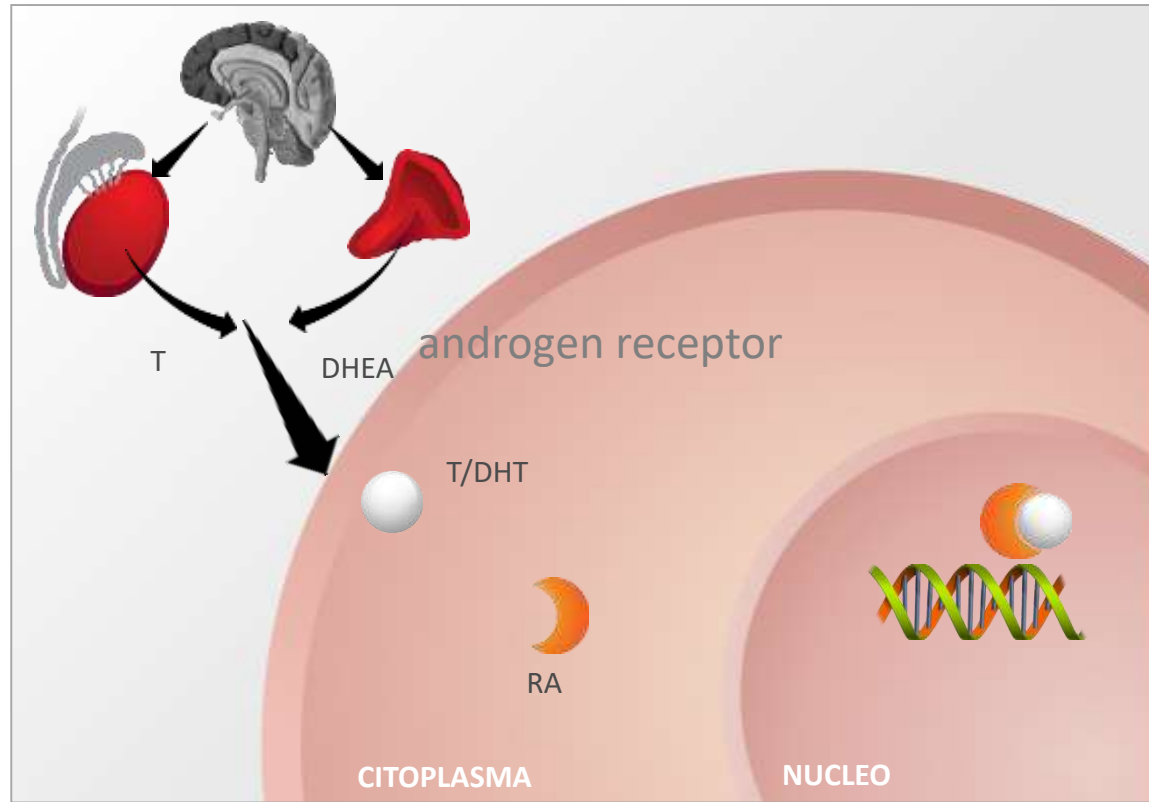
RA = receptor androgénico; DHT = dihidrotestosterona.

1. Hou X, Flaig TW. *Adv Urol* 2012 doi:10.1155/2012/9785312.

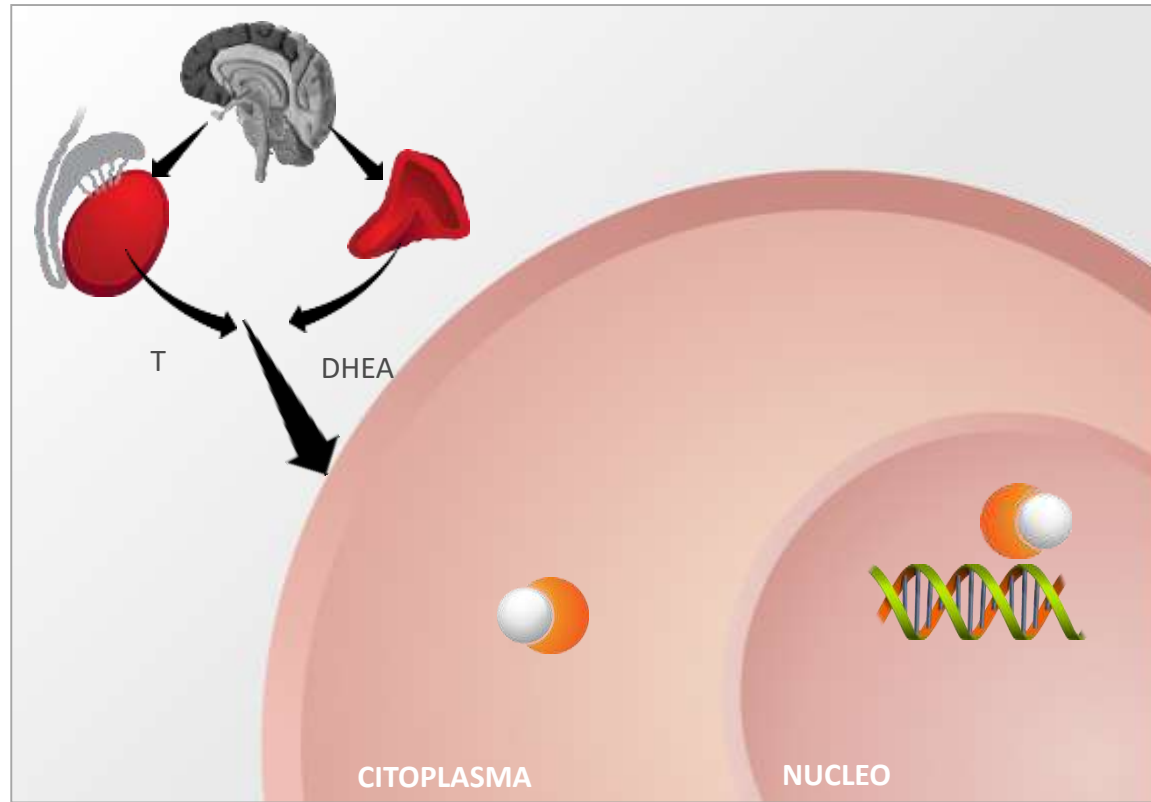
2. Hu R, et al. *Expert Rev Endocrinol Metab* 2010;5:753-64.

La vía de señalización intracelular del receptor androgénico (RA)

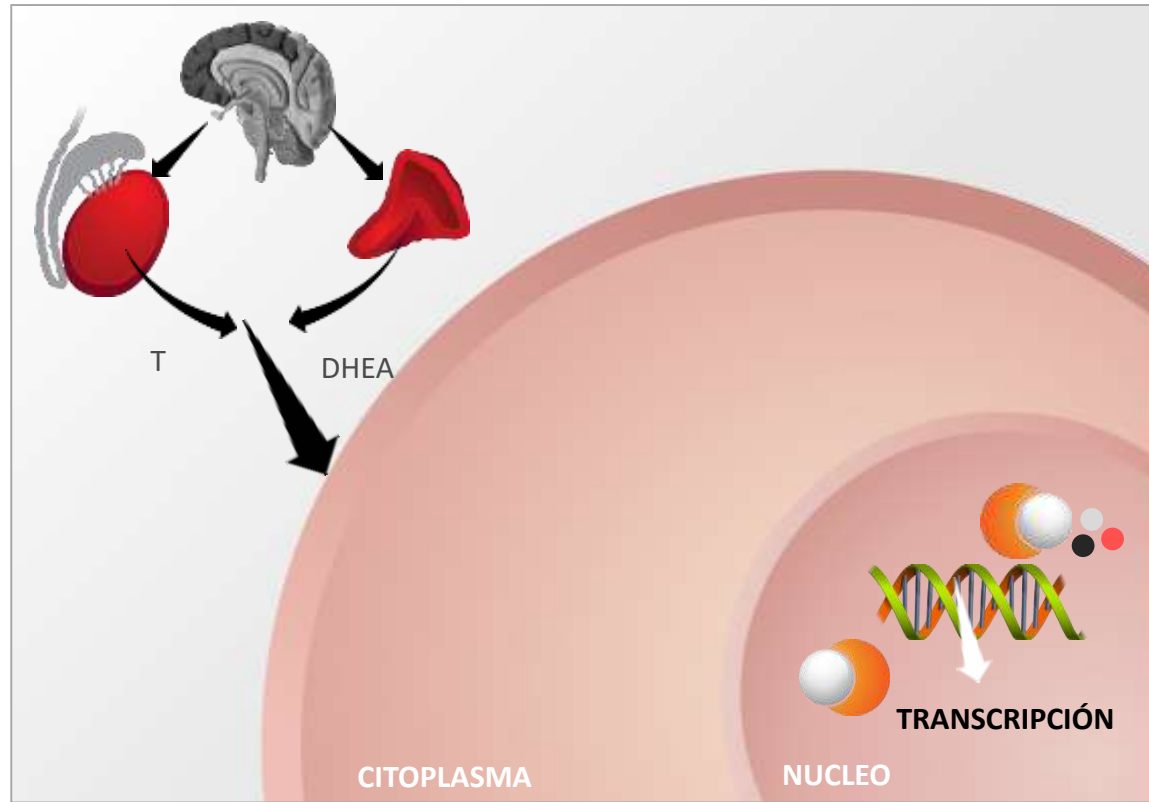
1. Unión Ligando-Receptor Androgénico



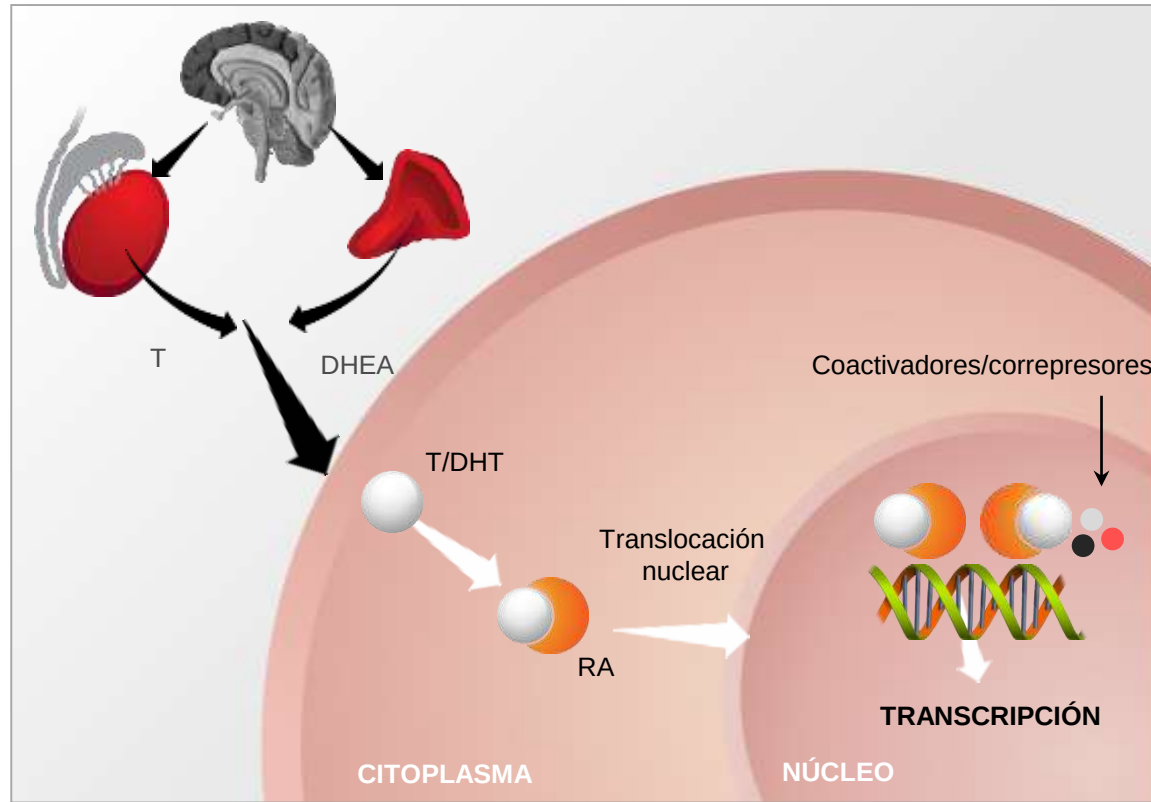
2. Translocación nuclear del Receptor Androgénico



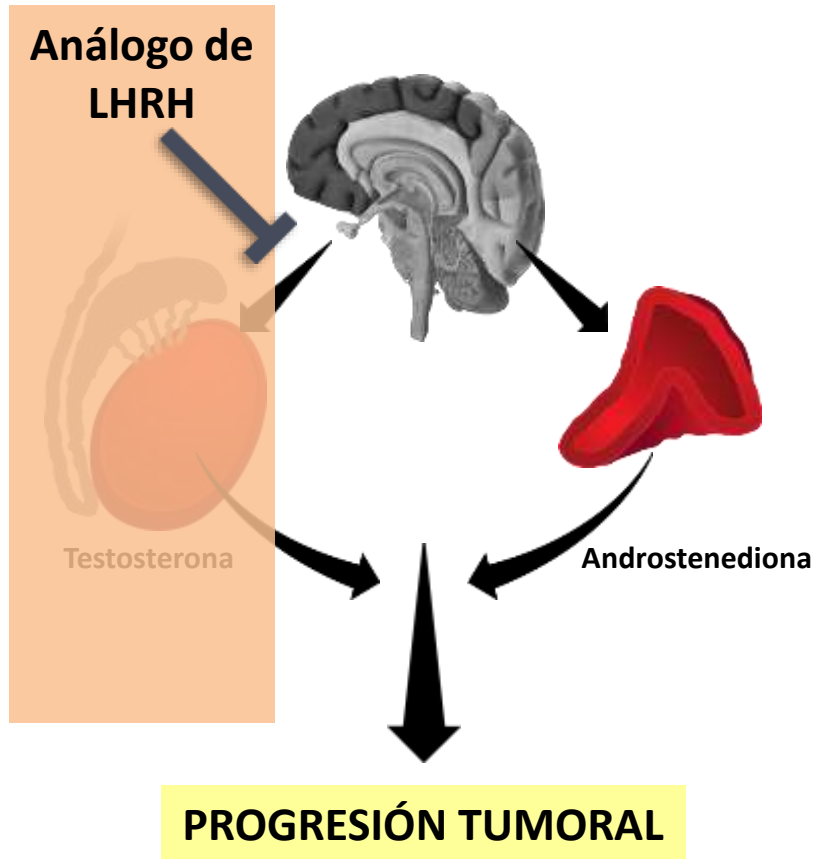
3. Unión Receptor Androgénico y transcripción génica



La vía de señalización intracelular del receptor androgénico (RA)



Cáncer de próstata resistente a la castración (CPRC)

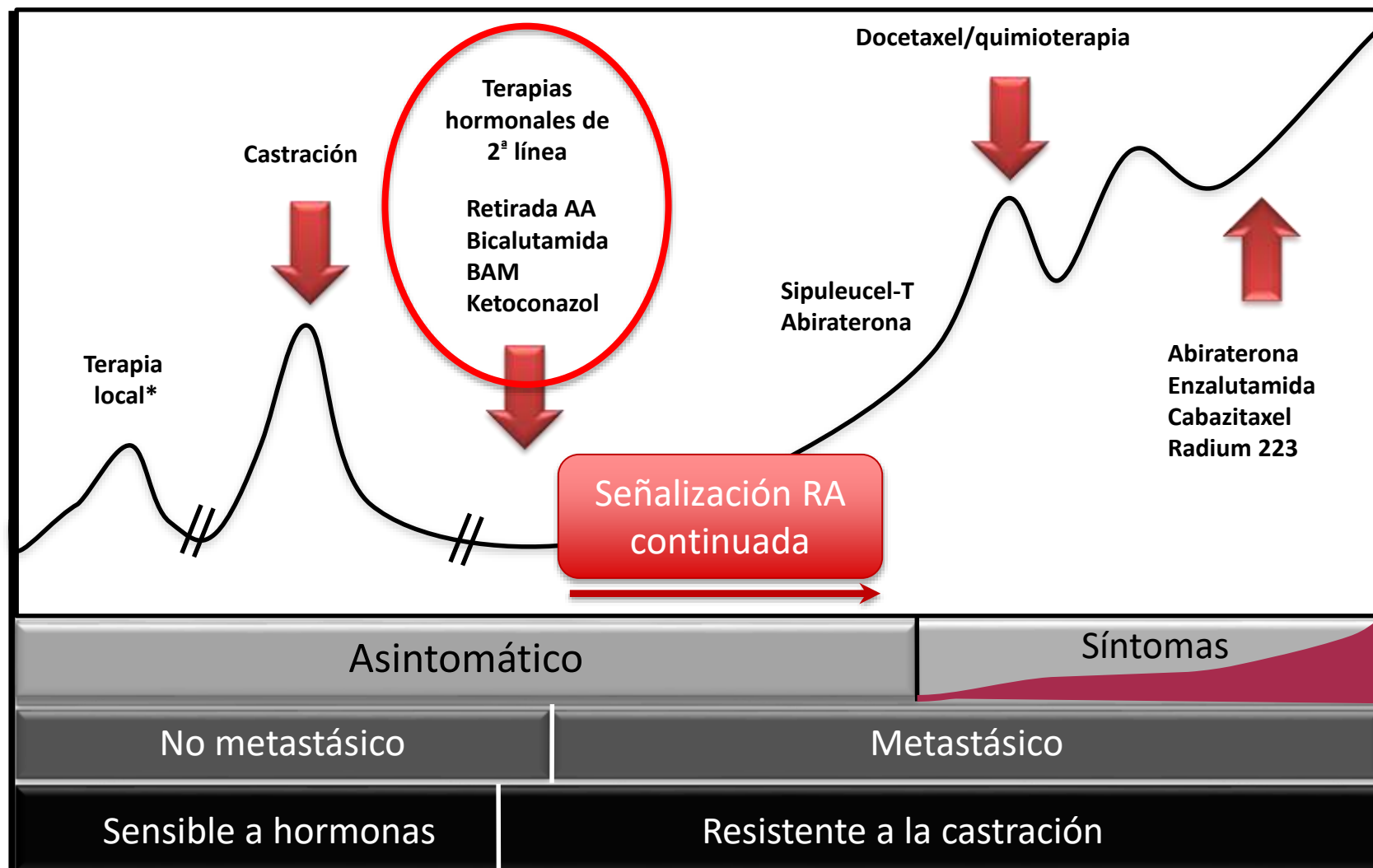


- **Progresión del tumor en un ambiente de privación de andrógenos.**
 - [Testosterona]p < 50 ng/dl).
 - Maniobra previa de retirada del antiandrógeno.
- **En la mayoría de los pacientes sigue funcionando el mecanismo del RA.**
 - Reactivación de la vía del RA.
- **Mecanismos de resistencia a castración:**
 - Dependientes del RA.
 - Independientes del RA

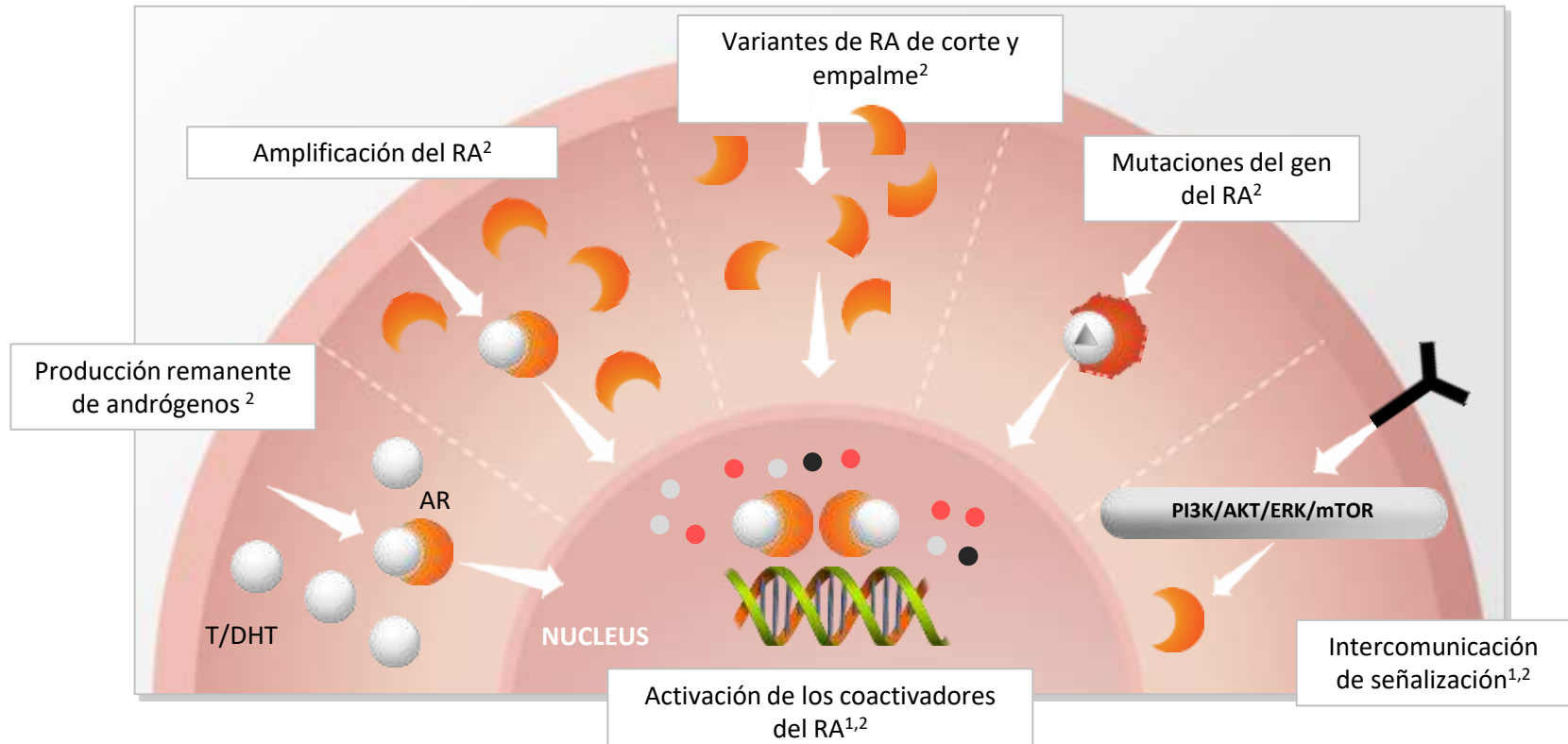
ACTH: corticotropina; LH: hormona luteinizante; LHRH: hormona liberadora de la hormona luteinizante

1. Hou X and Flaig TW. *Adv Urol* 2012; 978351. Epub ahead of print. 2. Heinlein CA and Chang C. *Endocr Rev* 2004; 25 (2): 276–308. 3. Ryan CJ and Tindall DJ. *J Clin Oncol* 2011; 29 (27): 3651–3658.

Evolución natural del Cáncer de Próstata



Múltiples procesos moleculares contribuyen a la señalización sostenida del RA :

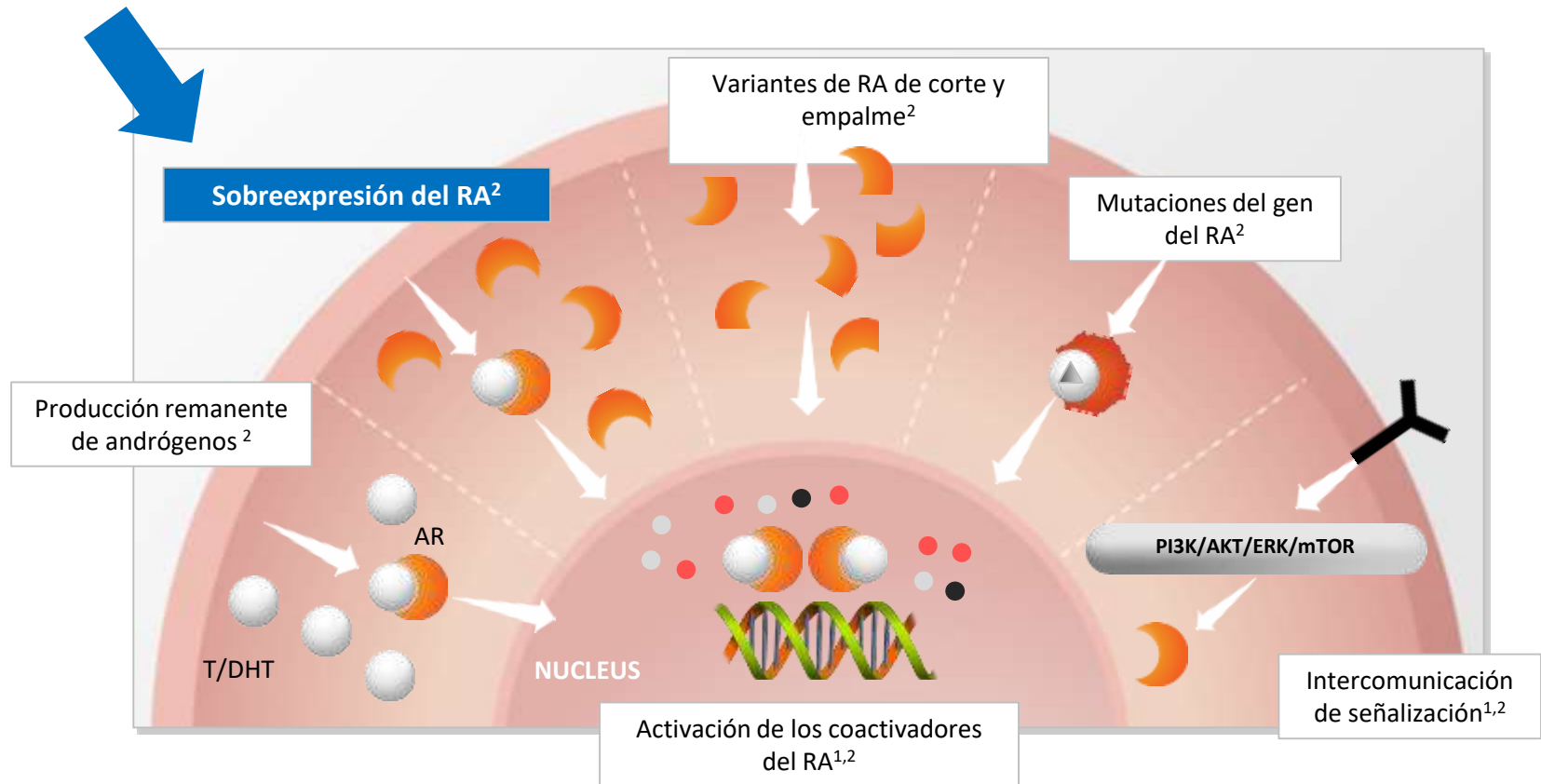


La señalización del RA es un mecanismo clave en la progresión del CPRC y es una diana lógica del tratamiento.

AR: androgen receptor; DHT: dihydrotestosterone; ERK: extracellular signal-regulated kinase; mTOR: mammalian target of rapamycin; PI3K=phosphatidylinositol-3 kinase; T: testosterone.

1. Heinlein CA, Chang C. Endocr Rev 2004;25: 276–308; 2. Hu R et al. Expert Rev Endocrinol Metab 2010;5:753–64.

Múltiples procesos moleculares contribuyen a la señalización sostenida del RA :

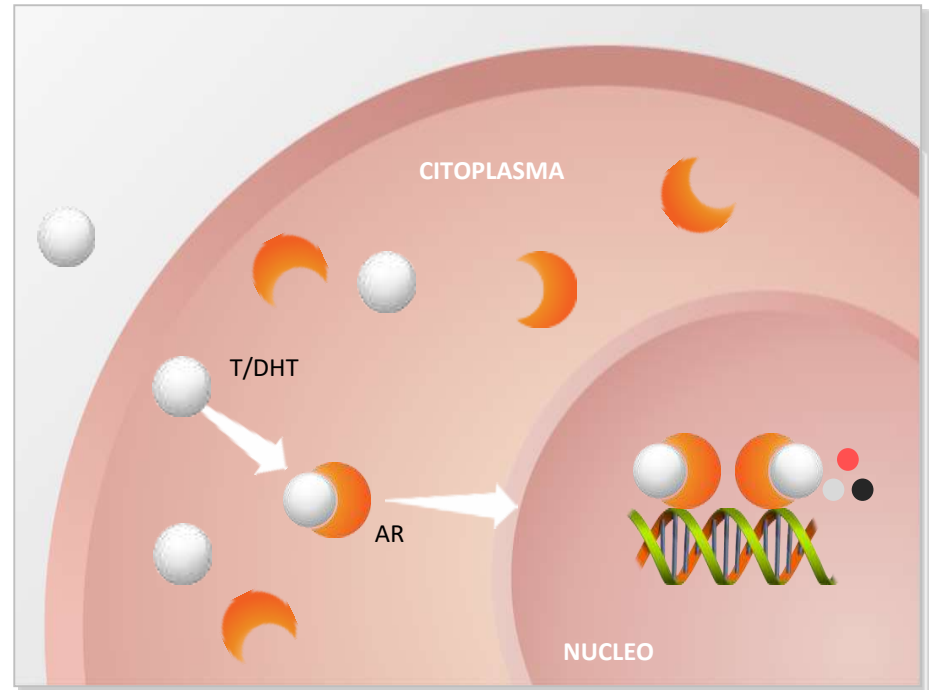


AR: androgen receptor; DHT: dihydrotestosterone; ERK: extracellular signal-regulated kinase; mTOR: mammalian target of rapamycin; PI3K=phosphatidylinositol-3 kinase; T: testosterone.

1. Heinlein CA, Chang C. Endocr Rev 2004;25: 276–308; 2. Hu R et al. Expert Rev Endocrinol Metab 2010;5:753–64.

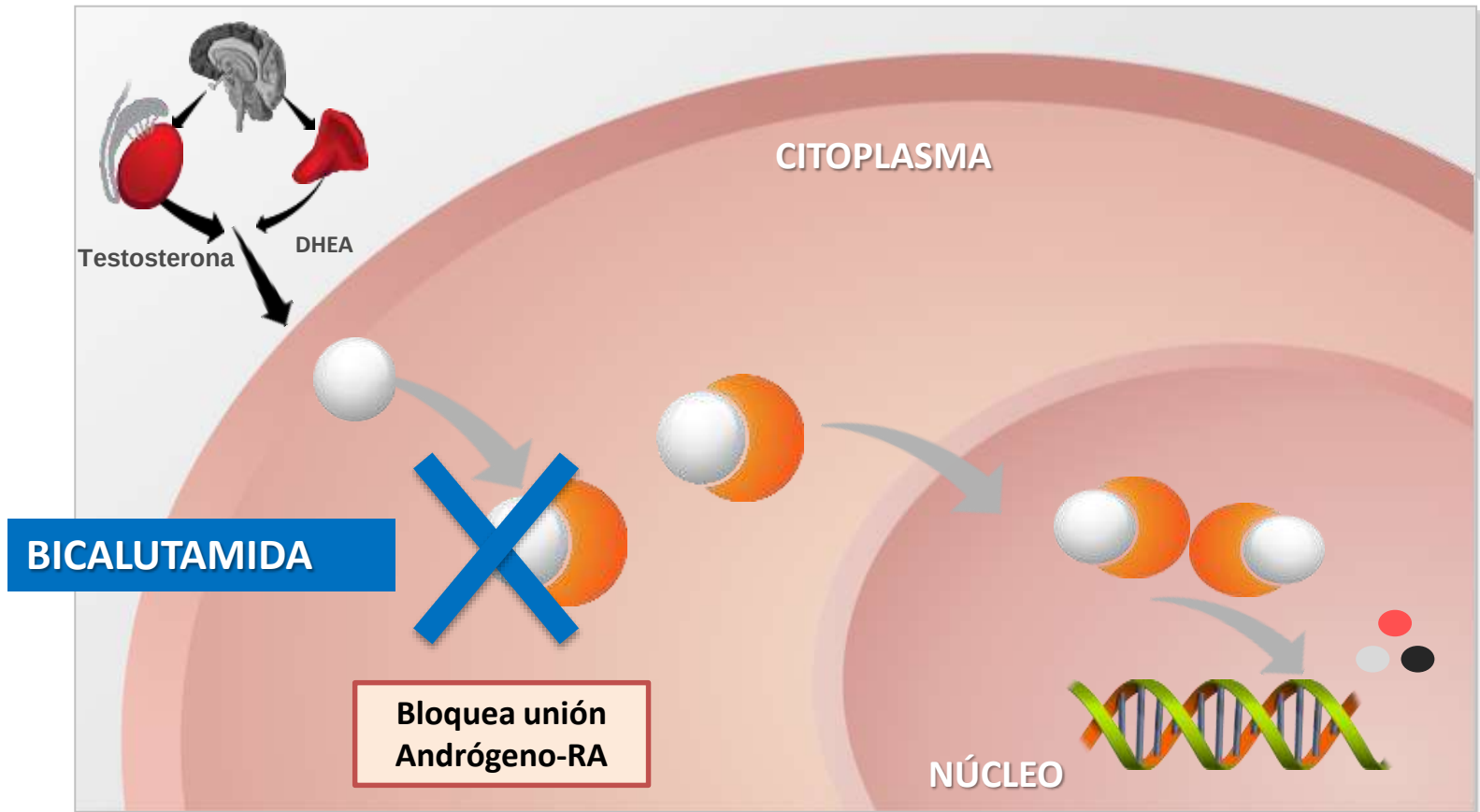
Sobreexpresión del RA

- La amplificación del gen del RA se ha detectado en el 50-85% de los casos de CPRC
- La sobreexpresión del RA lleva a:
 - Aumento de los niveles de proteína del RA
 - Elevada respuesta a bajos niveles de andrógenos.

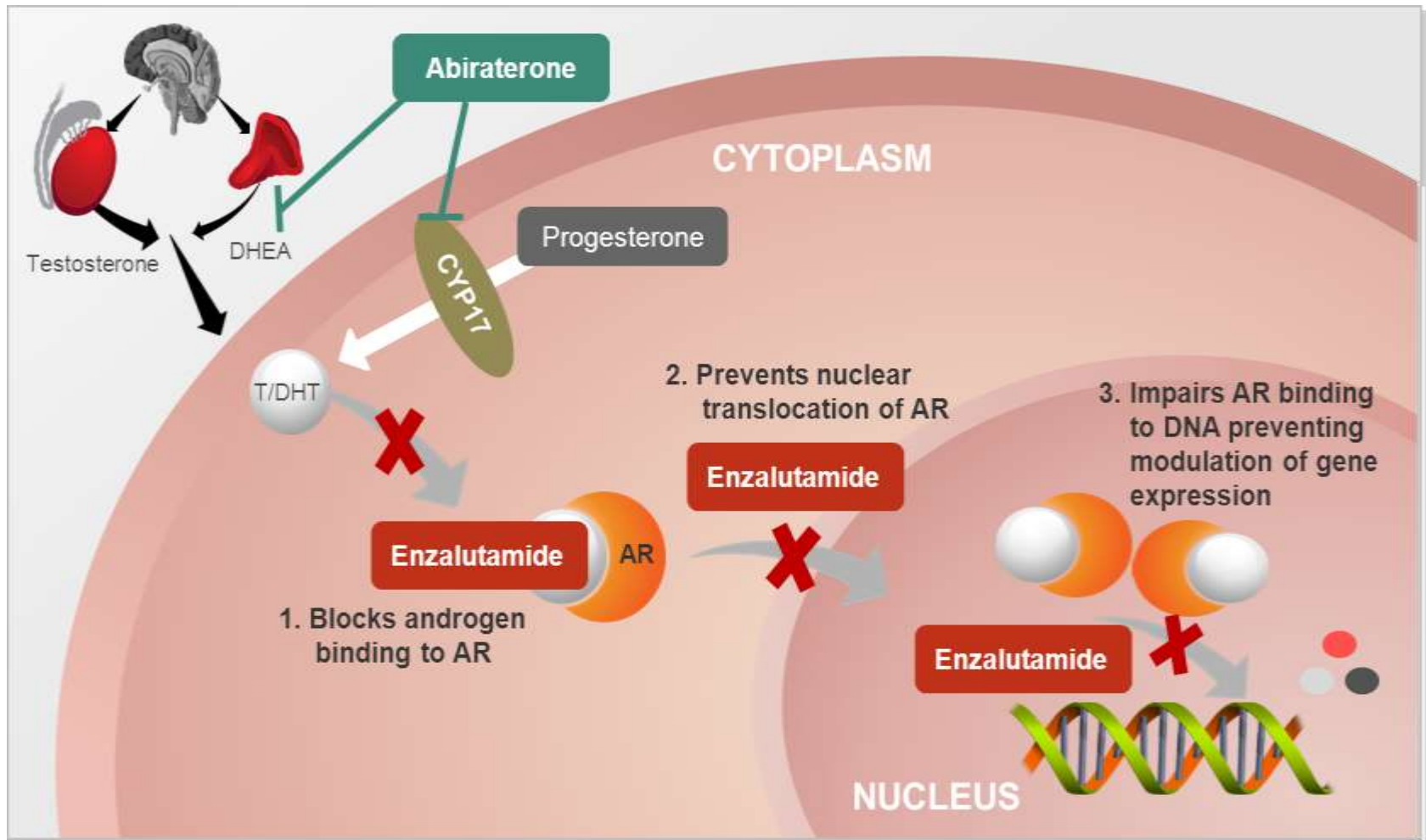


El presente. Las primeras terapias dirigidas al receptor androgénico

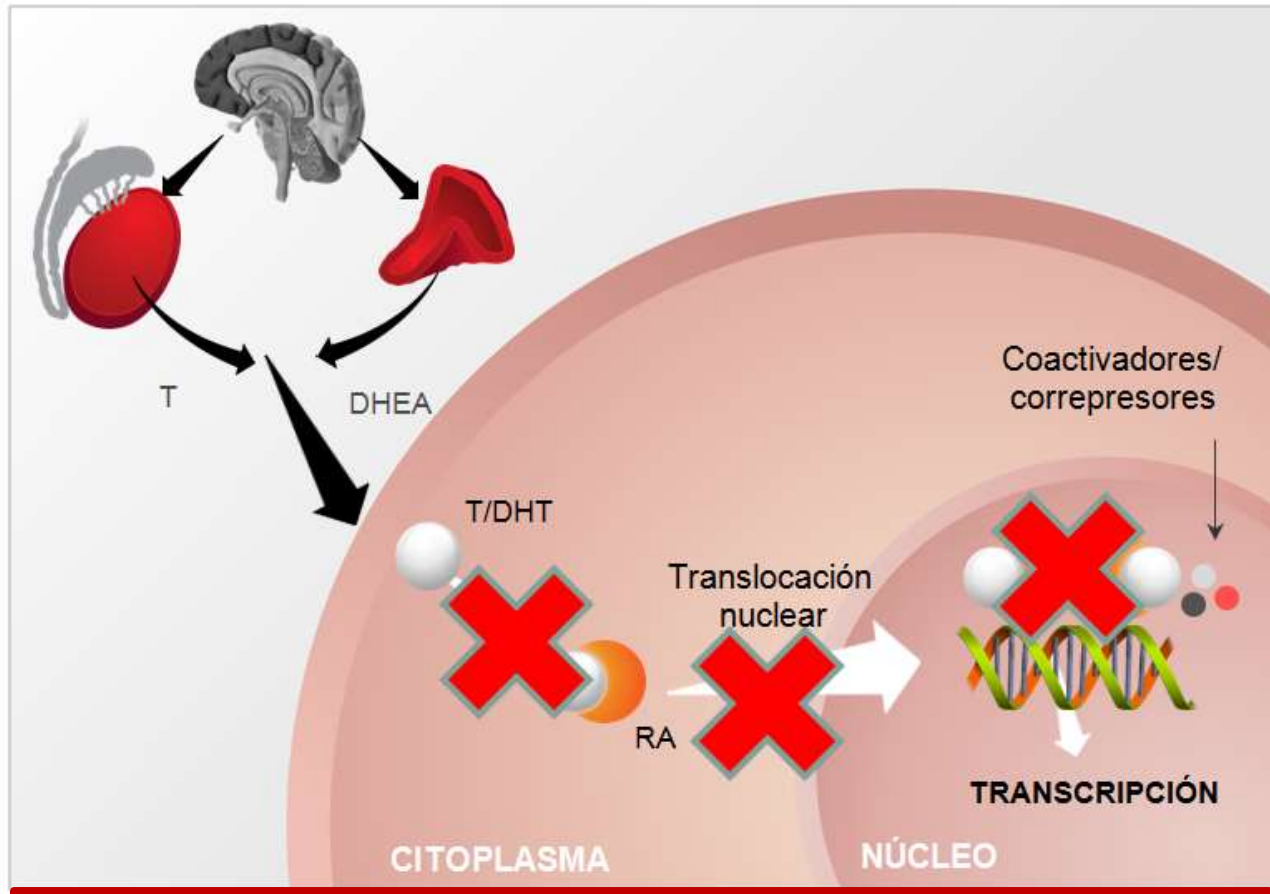
Mecanismo de Acción Antiandrógenos Clásicos



Enzalutamida y abiraterona inhiben la vía de señalización del Receptor Androgénico por mecanismos distintos



Mecanismo de Acción de enzalutamida



Enzalutamida

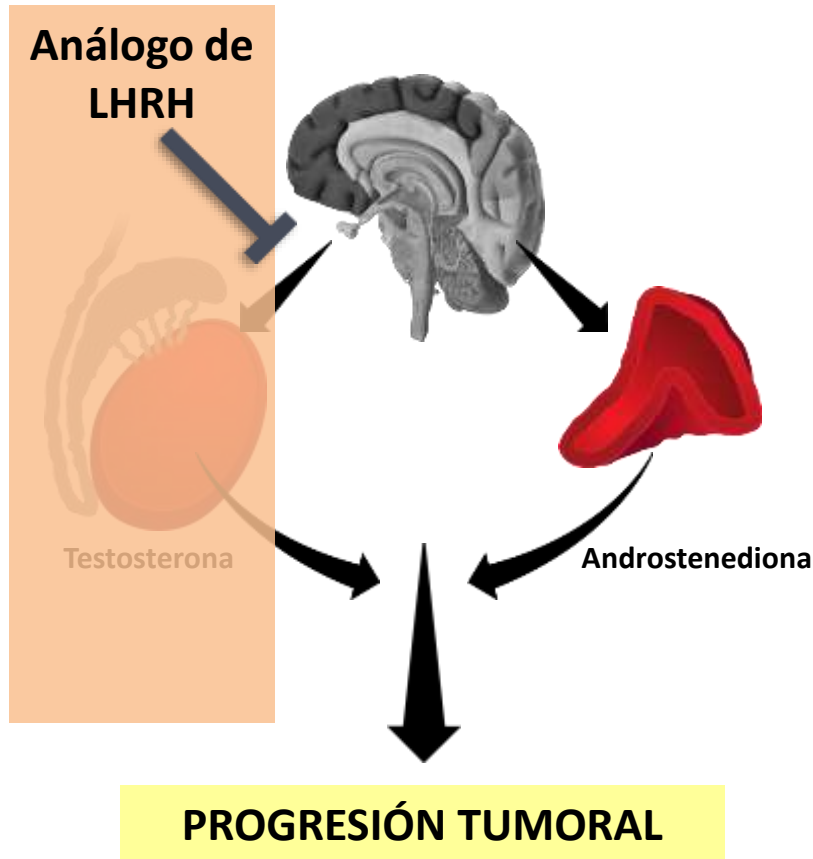


Enzalutamida inhibe la señalización del RA en 3 puntos:

- *Bloquea la unión del andrógeno al receptor*
- *Impide la translocación nuclear*
- *Bloquea la unión y activación al DNA*

¿RECORDAIS?

Cáncer de próstata resistente a la castración (CPRC)

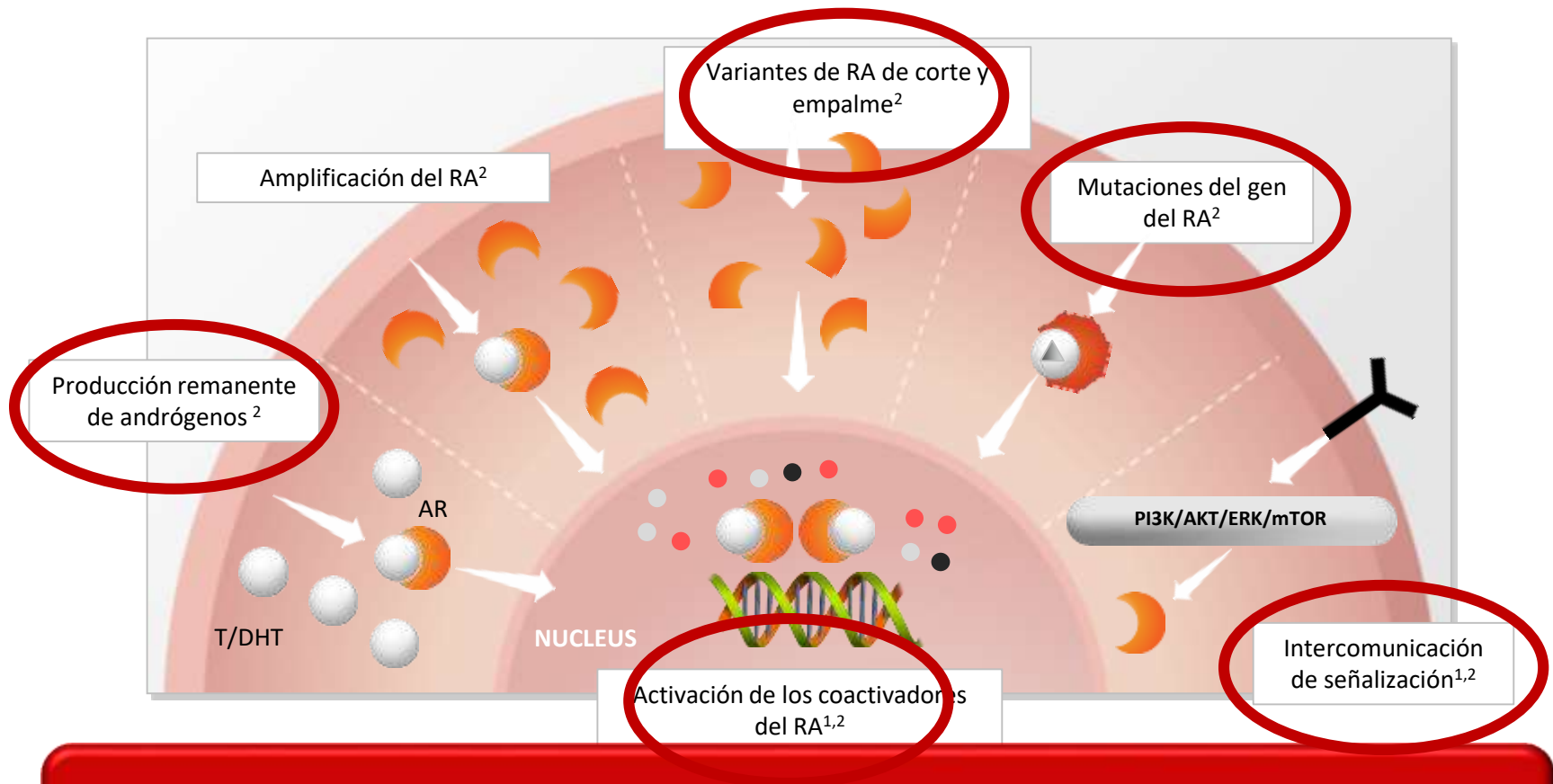


- **Progresión del tumor en un ambiente de privación de andrógenos.**
 - [Testosterona]p < 50 ng/dl).
 - Maniobra previa de retirada del antiandrógeno.
- **En la mayoría de los pacientes sigue funcionando el mecanismo del RA.**
 - Reactivación de la vía del RA.
- **Mecanismos de resistencia a castración:**
 - Dependientes del RA.
 - Independientes del RA

ACTH: corticotropina; LH: hormona luteinizante; LHRH: hormona liberadora de la hormona luteinizante

1. Hou X and Flaig TW. *Adv Urol* 2012; 978351. Epub ahead of print. 2. Heinlein CA and Chang C. *Endocr Rev* 2004; 25 (2): 276–308. 3. Ryan CJ and Tindall DJ. *J Clin Oncol* 2011; 29 (27): 3651–3658.

Múltiples procesos moleculares contribuyen a la señalización sostenida del RA :



La señalización del RA es un mecanismo clave en la progresión del CPRC y es una diana lógica del tratamiento.

Principios de Biología Molecular.

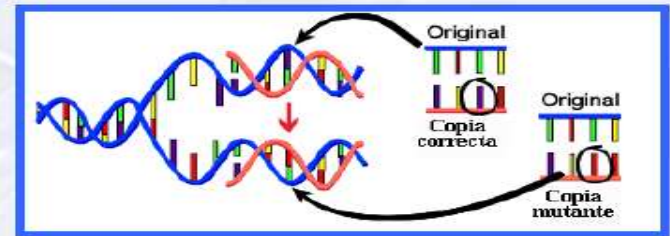
Genoma



- Codificado por ADN.
- El genoma comprende genes y non-coding sequences de ADN
- El ADN codificante se define como aquellas secuencias que se transcriben en mRNA y se traducen en proteínas; estas secuencias ocupan solo una fracción minoritaria del genoma (<2%).
- **ADN no codificante** esta compuesto por secuencias (98% del genoma) que no codifican para proteínas.
- Algunos noncoding DNA contienen genes para moléculas de RNA con funciones biológicas vitales (noncoding RNA, ribosomal RNA y transfer RNA).
- Humanos poseen **3 billones de bases** de información, dividido en 20,000 -22,000 genes, que se distribuyen en **24 cromosomas (22 autosomas mas los sexuales X e Y)**.

El ADN como material genético

- El material genético debe ser capaz de proporcionar una **fuente de variabilidad** a través del proceso de mutación.
- **Mutación**: cambio que afecta a la composición del ADN que tendrá su reflejo final en la proteína resultante.
- **Mutación línea germinal**: afecta a los gametos, pasará a la descendencia.
- **Mutación somática**: Alteración o cambio en la información genética (ADN) que ocurre después de la concepción.
- Este cambio podrá ser **beneficioso, neutral o perjudicial**.



Tipos de mutaciones

**Original
DNA
sequence**

GGG AGT GTA GAT CGT

**(a)
Base
substitution**

GGG AGT **GCA** GAT CGT

One codon changed

**(b)
Base
insertion**

GGG AGT **GT****T** AGA TCG T

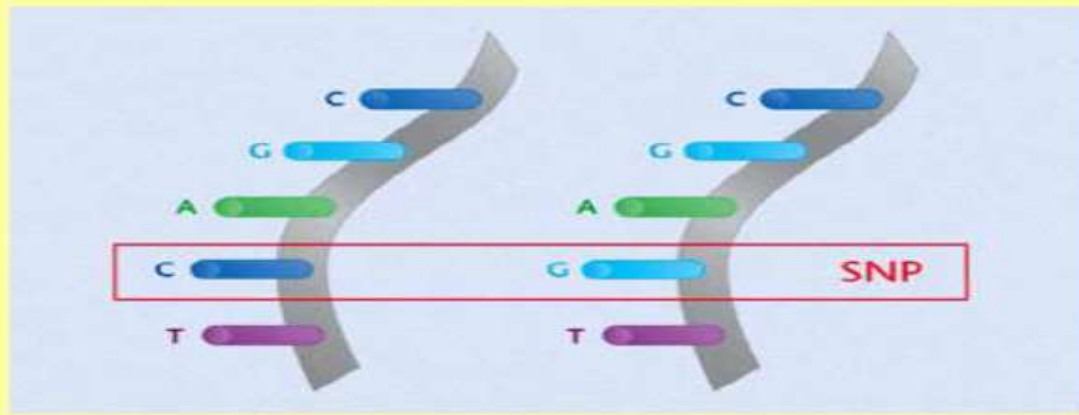
**(c)
Base
deletion**

GGG AGT **GAG** ATC GT

Fig. 17-03 Genetics, Second Edition © 2005 W.H. Freeman and Company

Mutaciones indel: Cambio de la secuencia de codones que debe ser leída durante la traducción, resultando en una proteína completamente diferente, que no suele ser funcional.

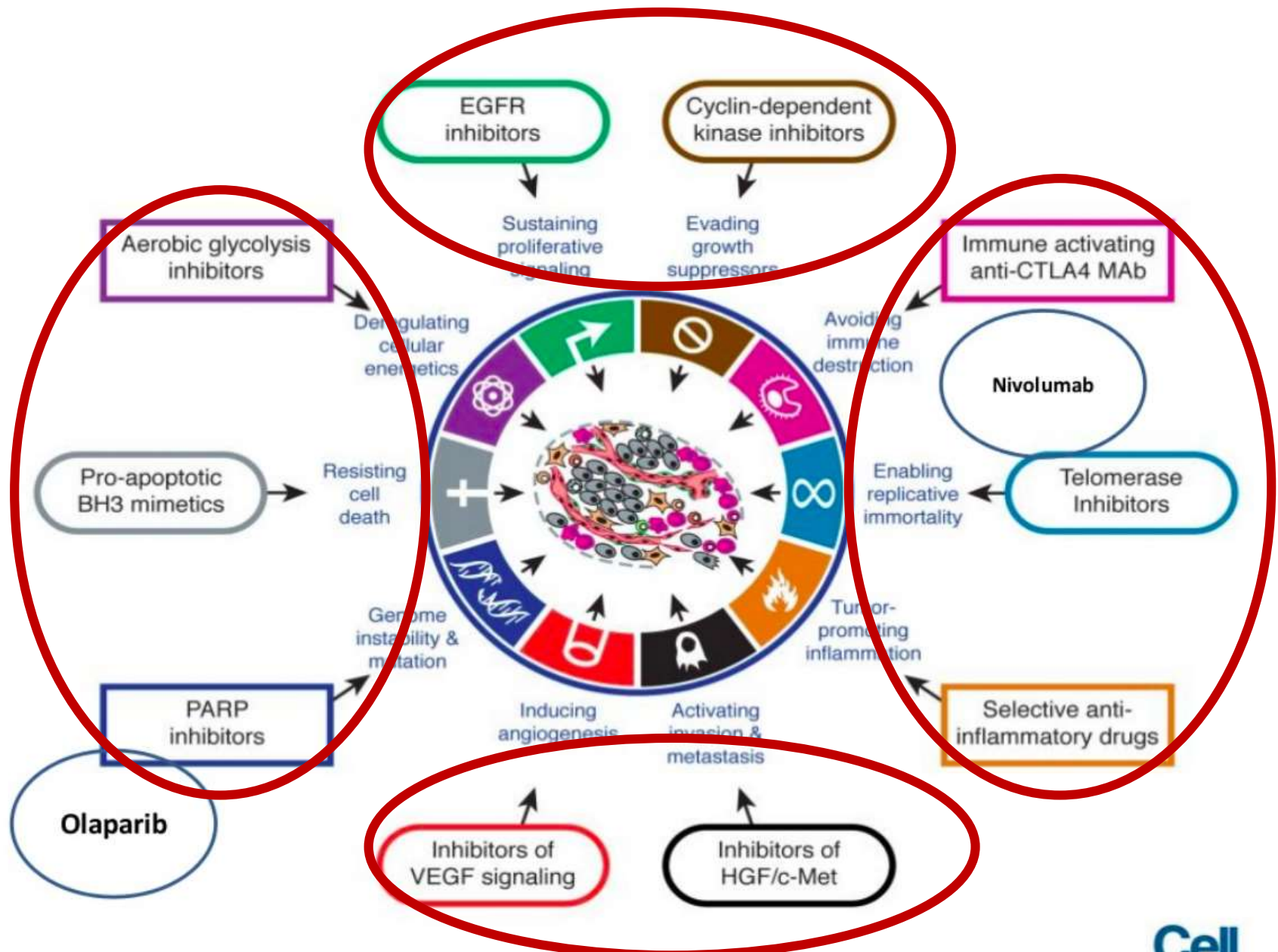
SNPs



SINGLE NUCLEOTIDE POLYMORPHISM



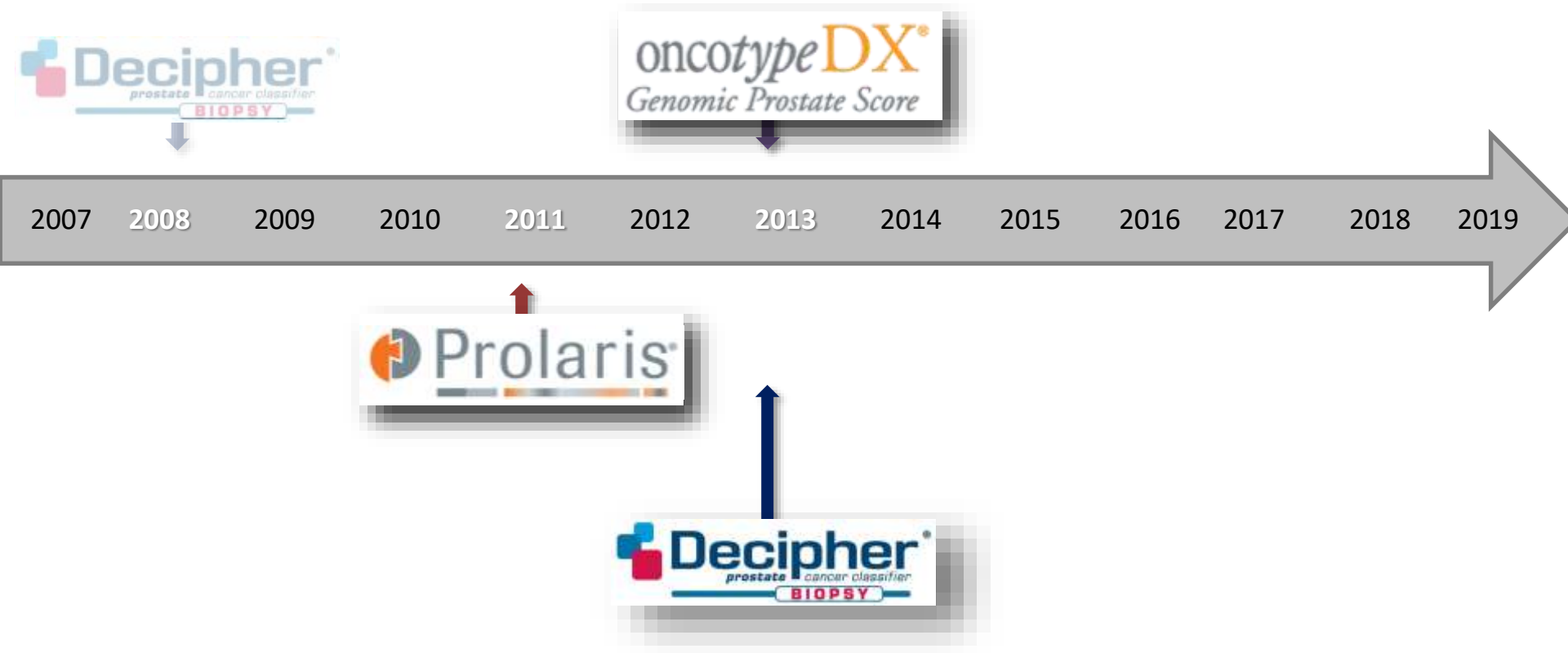
- Alrededor de 8 millones de SNPs en todo el genoma.
- Algunos pueden producir cambios en proteína:
 - susceptibilidad a padecer enfermedades
 - resistencia a tratamientos específicos



Hanahan and Weinberg, 2011

Paneles Genéticos.
**La búsqueda de Biomarcadores
pronósticos y predictivos**

Paneles genéticos en cáncer de próstata



Paneles genéticos en cáncer de próstata



Riesgo de tener CaP

Riesgo de CaP Desfavorable

Riesgo de Recidiva

Riesgo de Metástasis

Riesgo de Muerte



Resultados:

Table 4. Univariable and multivariable odds Ratios for CC, GC and GCC, and clinicopathologic variables.

	Univariable		Multivariable	
	Odds Ratio (95% CI)	P	Odds Ratio (95% CI)	P
GC	1.42 (1.28–1.60)	p<0.001	1.36 (1.16–1.60)	p<0.001
GCC	1.36 (1.21–1.53)	p<0.001	n.a	n.a
CC	1.35 (1.15–1.59)	p<0.001	n.a	n.a
Pre-operative PSA	0.99 (0.77–1.26)	0.92	0.75 (0.52–1.07)	0.11
Pathologic Gleason Score ≥8	3.02 (1.61–5.68)	p<0.001	1.91 (0.85–4.33)	0.12
Seminal Vesicle Invasion	2.44 (1.30–4.58)	0.01	1.93 (0.79–4.73)	0.15
Tumor Volume	1.02 (0.97–1.06)	0.44	0.97 (0.92–1.04)	0.42
Lymph Node Involvement	1.69 (0.74–3.88)	0.21	1.42 (0.41–4.96)	0.58
Positive Surgical Margins	1.05 (0.57–1.93)	0.87	0.93 (0.40–2.17)	0.87
Extra-capsular Extension	2.01 (1.18–3.73)	0.03	1.00 (0.45–2.20)	0.99

Odd ratios for multivariable classifiers are adjusted as indicated in the Materials and Methods. CC: clinical-only classifier. GC: genomic classifier. GCC: integrated genomic-clinical classifier.

doi:10.1371/journal.pone.0066855.t004



Genomic Dx (Bioinformatics Laboratory)
10335 Science Center Drive, Suite 240
San Diego, CA 92121
Tel: 1.855.732.1601 | Fax: 1.855.324.2788
customersupport@genomedi.com | www.genomedi.com

DECIPHER BIOPSY REPORT

PATIENT DETAILS

Patient Name:
MRN/ID:
Date of Birth: 1
Date of Biopsy:
Pathology Laboratory:
Address:

ORDER INFORMATION

Order Date:
Specimen Received Date:
Genomic Dx Accession ID:
Specimen ID:
Ordering Physician:
Circulating Hospital Name:
Circulating Hospital Address:

CLINICAL DETAILS

PSA, Most Recent (ng/mL): 7.1
Specimen Type: Needle Biopsy

NCCH Risk Category: Intermediate Risk
of Positive Cores: 11 (11 of 42 Cores)

Gleason Score: 3+4
Clinical Stage: T1c

YOUR DECIPHER RESULT: GENOMIC LOW RISK

DECIPHER SCORE: 0.23

Risk at RP - Percent Likelihood

High Grade Disease (primary Gleason grade 4 or 5)	12.5%
5-Year Metastases	1.0%
10-Year Prostate Cancer Specific Mortality	1.9%

Clinical studies have shown that men with a Decipher low risk score have a favorable prognosis. Men considering active surveillance with a Decipher low risk score may be suitable candidates for active surveillance. Men considering a definitive therapy may have excellent outcomes when treated with local therapy alone.^{1,4}

¹ Average clinical risk score for the average cohort of men with a 5-year post radical prostatectomy (RP) outcome. The average cumulative incidence of metastases was 6.0% at 5 years post RP, as reported by Kaplan et al., 2012 when analyzed as a cohort of 1,473 men with intermediate and high risk clinical features who received RP as first line treatment of the Mayo Clinic between 2003 and 2009.

⁴ **Prognosis of high grade disease (primary Gleason grade 4 or 5) at RP.** Decipher uses the genomic risk score to predict the probability of primary Gleason grade 4 or 5 disease upon pathologic examination of the RP. Probabilities were generated using a logistic regression model in a prospective cohort of 2,352 prostate cancer patients.¹ The model is adjusted using a prevalence of 25% for a finding of primary Gleason grade 4 or 5 at RP among NCCH low-, intermediate- and high-risk patients. ¹ Han et al., 2016 study found Decipher Biopsy predicted high-grade disease at RP with an AUC of 0.71. The percent likelihood for this endpoint ranges from 8.4-61%.

Five-year probability of metastases endpoint. Decipher uses the genomic risk score to predict the 5-year probability of metastases from the time of RP. Probabilities were generated from a Cox proportional hazards model based upon a cohort of 1,210 men with intermediate and high risk clinical features with a median of 5.8 years of follow-up.¹ In a secondary cohort, Han et al., 2016 reported that Decipher Biopsy predicted 5-year metastases with an AUC of 0.67.² Subsequently, in another independent cohort, Decipher Biopsy predicted 5-year metastases with an AUC of 0.78 and similarly found that Decipher Biopsy was the only significant variable that predicted metastases over 5-year endpoint analysis with clinical risk factors.³ The percent likelihood for this endpoint ranges from 0.5-6.6%. The average correlation between Decipher Biopsy and RP specimen is 50%. Needle biopsy sampling accuracy impacts outcomes.

10-year probability of prostate cancer specific mortality (PCSM) endpoint. Decipher uses the genomic risk score to predict the 10-year probability of PCSM from the time of RP. Probabilities are generated from a logistic regression model based upon a cohort of 827 patients with T1c prostate cancer treated with 19 years post RP. These probabilities are adjusted for a PCSM incidence rate of 20% at 10 years post RP.¹ All men in the study had at least 10 years of follow-up. In a retrospective study with a biopsy cohort of 200 patients, Decipher was a significant predictor of PCSM at diagnosis with a hazard ratio (HR) of 1.08 (95% CI 1.03-1.13) per 1% increase for Decipher score (p=0.001). The percent likelihood for this endpoint ranges from 0.1-0.3%.



Manejo sugerido

Decipher Biopsy
Alto Riesgo

Pronóstico desfavorable- puede no ser un candidato adecuado para vigilancia activa y puede beneficiarse de **intensificación con terapia multimodal**.

Decipher Biopsy
Riesgo Intermedio

Pronóstico intermedio- existe un riesgo clínico y pronóstico intermedio. Dependiendo de esperanza de vida y estado general, pueden beneficiarse de **intensificación con terapia multimodal**.

Decipher Biopsy
Bajo Riesgo

Pronóstico favorable- puede ser un candidato adecuado para **vigilancia activa** y puede tener excelentes resultados cuando se trata sólo con **terapia local**, como cirugía o radioterapia.

PROLARIS® BIOPSY TEST RESULT

ORDERING PHYSICIAN
Bob Doctor MD

123 Grand Ave
Las Vegas, NV 89109

Joe Pathologist MD

CLINICOPATHOLOGIC FEATURES USED FOR ANALYSIS

Patient Age at Diagnosis:	65	% Positive Cores:	25% (3/12)
PSA Prior to This Biopsy:	9.0	Gleason Score:	3+3=6
Clinical T Stage:	T1c	AUA Risk:	Low

SPECIMEN

Specimen Type: Tissue Block
Tissue: Prostate
Biopsy Date: Mar 13, 2015
TRF Received: Jun 10, 2016

Sample Received: Jun 10, 2016
Accession Date: Jun 10, 2016
Report Date: Aug 15, 2016

PATIENT

Name: Patient, Dave
Date of Birth: Oct 26, 1946
Patient ID: 12654
Gender: Male

Accession #: 07000106-BLD
Requisition #: 07000106

Block(s) Analyzed: 1234546821-8

Less Aggressive
Than Average AUA¹ Low Risk

PROLARIS SCORE 2.5

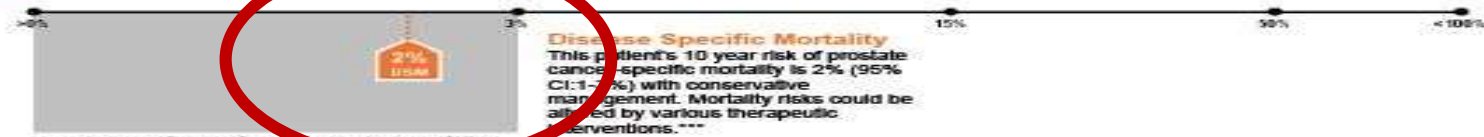
US Distribution Percentile: 9%
(For AUA Low Risk)

Interpretation: 9% of patients in the AUA Low Risk* category have a lower Prolaris Score



Mortality Risk

Mortality Risk: 2% 10-Year Prostate Cancer-Specific



In a clinical study estimating 10-year prostate cancer-specific mortality risks for patients using conservative management, there were no observed prostate cancer deaths in patients with a predefined clinical risk score (CCP combined with CAPRA) corresponding to a 3.2% (95% CI 2.0, 5.2%) prostate cancer-specific mortality risk. **

CONFIDENTIAL



MM58234



Myriad Genetic Laboratories, Inc.
320 Wakara Way, Salt Lake City, Utah 84108 | 800.469.7423 | Fax 801.584.3615
prolaris.com

Genomic Prostate Score (GPS) Report

oncotype^{DX}
Genomic Prostate Score

PATIENT-LAST-NAME, FIRST-NAME I.

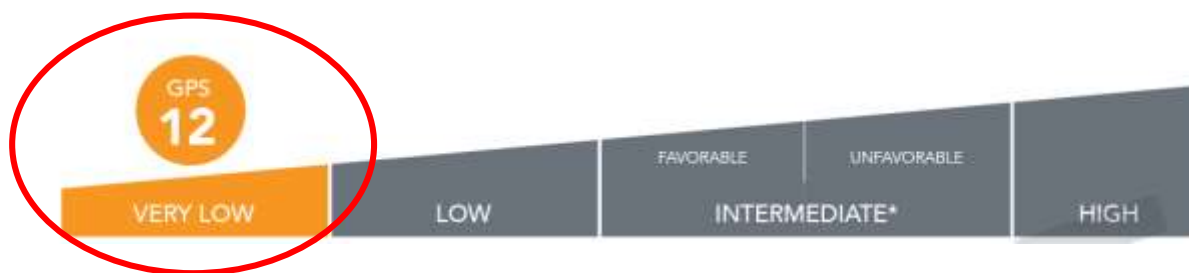
Date of Birth: 01-Jan-1950

Gender: Male

Report Number: OR000123456-01

Report Date: 22-May-2017

Ordering Physician: Dr. First-Name I. Ordering-Physician-Last-Name

GPS + NCCN^{®1}: Very Low Risk

Clinical Interpretation

Clinical Endpoints[†]

Individualized Risk (95% Confidence Interval [CI])

The combination of GPS and clinical features predicts that this patient's risk is consistent with **NCCN Very Low Risk** disease.[‡]

In a clinical validation study including patients with NCCN Very Low, Low, and Intermediate Risk, **no patient with a GPS result <20 had metastasis or died** from prostate cancer within 10 years.[†]

Prostate Cancer Death
Within 10 Years

<1%

(95% CI: <1% - <1%)

0%

100%

Metastasis Within
10 Years[§]

1%

(95% CI: <1% - 4%)

0%

100%

Adverse Pathology

(Gleason $\geq$ 4+3 and/or pT3+)

16%

(95% CI: 12% - 21%)

0%

100%

Paneles genómicos en Cáncer de próstata

	Estudio “pivotal”		Validaciones posteriores		¿Costes? ¹
Test	Tejido	Objetivo (predicción)	Tejido	Objetivo (predicción)	
Genomic Classifier Decipher®	Prostatectomía	• Metástasis a 5 años	Biopsia	Vigilancia activa ² , SLM, SCE, SG	US\$ 4250
			Prostatectomía	Patología adversa ² , SLM, SCE, SG, Respuesta a Radioterapia (adyuvante/rescate), Asociación a MRI, N(+) en PSMA	
Panel de 22 genes implicados en Proliferación del ciclo celular, Adhesión y movilidad, Inmuno-modulación, y Señalización del RA.					
Cell-cycle proliferation Prolaris®	Prostatectomía RTU (WW)	• Recidiva bioquímica • Muerte	Biopsia	SLRB ^a , Patología adversa, MRI	US\$ 3400
			Prostatectomía	SLRB ^a .	
Firma de 46 genes implicados en exclusiva en la Proliferación del ciclo celular.					
Genomic prostate score Oncotype®	Prostatectomía Biopsia	• AP adversa (pT3 / GG3)	Biopsia	Patología adversa	US\$ 3825
			Prostatectomía	SLM, SCE, MRI,	
Panel de 17 genes implicados en Respuesta estromal, Organización celular, Proliferación, Epitelio basal, Respuesta al estrés, Y Señalización del RA.					

¹ Davis JW. Novel commercially available genomic tests for prostate cancer: a roadmap to understanding their clinical impact. BJU Int. 114(3), 320–322 (2014).

² Identifican no candidatos a VA por asociar GC a patología adversa, pero el estudio lo realizan sobre prostatectomías.

**Prostate Cancer**

Version 4.2019 — August 19, 2019

NCCN.org

| Guidelines for Patients® available at www.nccn.org/patients[Continue](#)[NCCN Guidelines Index](#)
[Table of Contents](#)
[Discussion](#)**NCCN Guidelines Version 4.2019
Prostate Cancer**

Printed by AngelBacquet on 11/10/2019 2:22:48 AM. For personal use only. Not approved for distribution. Copyright © 2019 National Comprehensive Cancer Network, Inc., All Rights Reserved.

Table 1. Available Tissue-Based Tests for Prostate Cancer Risk Stratification/Prognosis

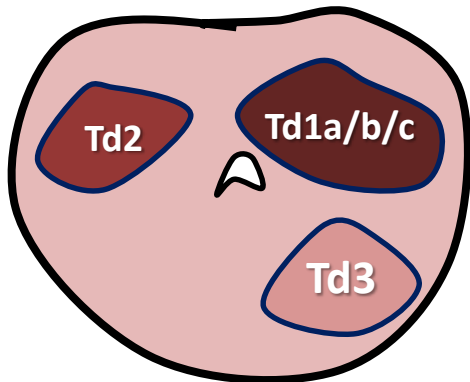
Test	Platform	Populations Studied	Outcome(s) Reported (Test independently predicts)	Selected References	Molecular Diagnostic Services Program (MoDX) Recommendations
Decipher	Whole-transcriptome 1.4M RNA expression (44,000 genes) oligonucleotide microarray optimized for FFPE tissue	- Post radical prostatectomy (RP), adverse pathology/high-risk features - Post RP, biochemical recurrence - Post RP, adjuvant, or salvage radiation - Biopsy, localized prostate cancer post RP or EBRT	- Metastasis - Prostate cancer-specific mortality - Postoperative radiation sensitivity (PORTOS) - Metastasis - Prostate cancer-specific mortality - PORTOS - Metastasis - Prostate cancer-specific mortality - PORTOS - Metastasis - Prostate cancer-specific mortality - Gleason grade at disease at RP - Adverse pathologic feature at RP	140,143,144,243,671,731-743	- Cover post-biopsy for NCCN very-low- and low-risk prostate cancer in patients with at least 10 years life expectancy who have not received treatment for prostate cancer and are candidates for active surveillance or definitive therapy - Cover post-RP for 1) pT2 with positive margins; 2) any pT3 disease; 3) rising PSA (above nadir)
KI-67	IHC	- Biopsy, intermediate- to high-risk treated with EBRT - Biopsy, conservatively managed (active surveillance)	- Metastasis - Prostate cancer-specific mortality	744-747	Not recommended
OncoType DX Prostate	Quantitative RT-PCR for 12 prostate cancer-related genes and 5 housekeeping controls	Biopsy, low- to intermediate-risk treated with RP	Non-organ-confined pT3 or Gleason grade 4 disease on RP	138,748-751	Cover post-biopsy for NCCN very-low-, low-risk, and favorable intermediate-risk prostate cancer in patients with at least 10 years life expectancy who have not received treatment for prostate cancer and are candidates for active surveillance or definitive therapy
Prostate	Quantitative RT-PCR for 31 cell cycle-related genes and 15 housekeeping controls	- Transurethral resection of the prostate (TURP), conservatively managed (active surveillance) - Biopsy, conservatively managed (active surveillance) - Biopsy, localized prostate cancer - Biopsy, intermediate-risk treated with EBRT - RP, node-negative localized prostate cancer	- Prostate cancer-specific mortality - Prostate cancer-specific mortality - Biochemical recurrence - Metastasis - Biochemical recurrence - Biochemical recurrence	135-138,752-754	Cover post-biopsy for NCCN very-low-, low-risk, and favorable intermediate-risk prostate cancer in patients with at least 10 years life expectancy who have not received treatment for prostate cancer and are candidates for active surveillance or definitive therapy
ProMark	Multiplex immunofluorescent staining of 8 proteins	Biopsy, Gleason grade 3+3 or 3+4	Non-organ-confined pT3 or Gleason pattern 4 disease on RP	755	Cover post-biopsy for NCCN very-low- and low-risk prostate cancer in patients with at least 10 years life expectancy who have not received treatment for prostate cancer and are candidates for active surveillance or definitive therapy
PTEN	Fluorescence in situ hybridization or IHC	- TURP, conservatively managed (active surveillance) - Biopsy, Gleason grade 3+3 - RP, high-risk localized disease	- Prostate cancer-specific mortality - Upgrading to Gleason pattern 4 on RP - Biochemical recurrence	756-760	Not recommended

Intratumoral and Intertumoral Genomic Heterogeneity of Multifocal Localized Prostate Cancer Impacts Molecular Classifications and Genomic Prognosticators

Lei Wei^{a,1}, Jianmin Wang^{a,1}, Erika Lampert^b, Simon Schlanger^b, Adam D. DePriest^c, Qiang Hu^b, Eduardo Cortes Gomez^b, Mitsuko Murakami^b, Sean T. Glenn^c, Jeffrey Conroy^d, Carl Morrison^e, Gissou Azabdaftari^e, James L. Mohler^f, Song Liu^{a,1}, Hannelore V. Heemers^{a,b,h,i,*}

N = 4 p., 26 cilindros

Pieza de prostatectomía,
fijada en parafina



Td1a/b/c: Heterogeneidad intratumoral
Td2/3/4/5: Heterogeneidad intertumoral

DEBILIDADES DE LOS PANELES GENETICOS

Oncotype DX (12 genes)	Prolaris (31 genes)	Decipher (22 genes)
AZGP1	FOXM1	LASP1
KLK2	CDC20	IQGAP3
SRD5A2	CDKN3	NFIB
FAM13C	CDC2	S1PR4
FLNC	KIF11	THBS2
GSN	KIAA0101	ANO7
TPM2	NUSAP1	PCDH7
GSTM2	CENPF	MYBPC1
TPX2	ASPM	EPPK1
BGN	BUB1B	TSBP
COL1A1	RRM2	PBX1
SFRP4	DLGAP5	NUSAP1
	BIRC5	ZWILCH
	KIF20A	UBE2C
	PLK1	CAMK2N1
	TOP2A	RABGAP1
	TK1	PCAT-32
	PBK	GLYATL1P4/PCAT-80
	ASF1B	TNFRSF19
	C18orf24	
	RAD54L	
	PTTG1	
	CDCA3	
	MCM10	
	PRC1	
	DTL	
	CEP55	
	RAD51	
	CENPM	
	CDCA8	
	ORC6L	

Prostate cancer genomics: comparing results from three molecular assays

Syed Alam, MD,¹ Joseph Tortora, MS,² Ilene Staff, PhD,² Tara McLaughlin, PhD,¹ Joseph Wagner, MD¹

¹Urology Division, Hartford Healthcare Medical Group, Hartford Hospital, Hartford, Connecticut, USA

²Hartford Hospital Research Program, Hartford Hospital, Hartford, Connecticut, USA

Valoración del Índice Kappa	
Valor de k	Fuerza de la concordancia
< 0.20	Pobre
0.21 - 0.40	Débil
0.41 - 0.60	Moderada
0.61 - 0.80	Buena
0.81 - 1.00	Muy buena

TABLE 3. Agreement between tests

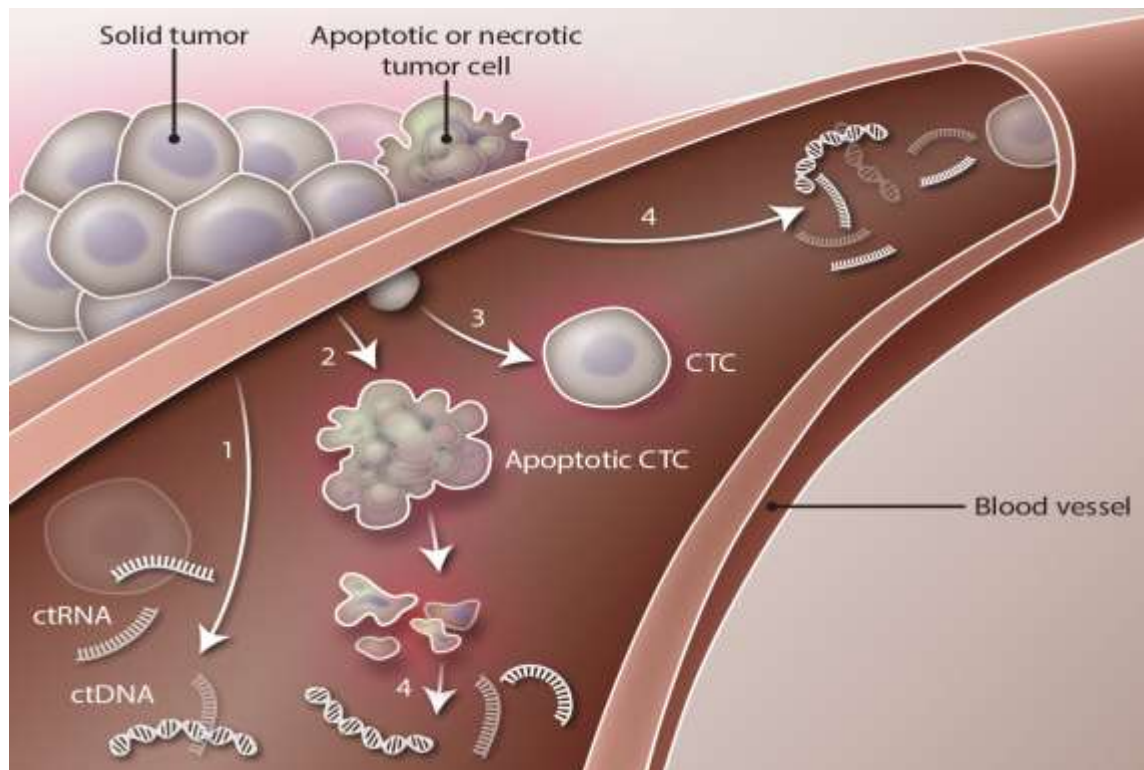
Tests	n	% agree	kappa	p value	Δ Definition favorable test
Decipher vs. Prolaris	12	67%	0.31	0.276	62%
Prolaris vs. Oncotype DX	8	75%	0.39	0.168	89%
Decipher vs. Oncotype DX	2	50%	-	-	50%

TABLE 4. Agreement between each test and results of surgical pathology

Patient	Gleason group	Stage	Margin	Oncotype	Prolaris	Decipher
1	2	T3aN0	Negative	Unfavorable	-	Unfavorable
2	2	T2cN0	Negative	Unfavorable	-	Favorable
3	2	T3bN1	Positive	-	Unfavorable	Favorable
4	2	T2cN0	Negative	-	Favorable	Favorable
5	2	T2cN0	Positive	-	Favorable	Unfavorable
6	2	T3bN0	Negative	-	Favorable	Favorable
7	3	T3bN0	Negative	-	Unfavorable	Favorable
8	4	T3bN0	Positive	-	Unfavorable	Unfavorable

La Biopsia Líquida en la obtención de los Biomarcadores

“test minimamente invasivo que se realiza utilizando un fluido corporal (sangre, orina, semen, líquido cefalorraquídeo) y que permite la detección de biomarcadores celulares o moleculares con el fin de obtener información diagnóstica, pronóstica o predictiva de la enfermedad”



cfDNA

CTCs

mRNA

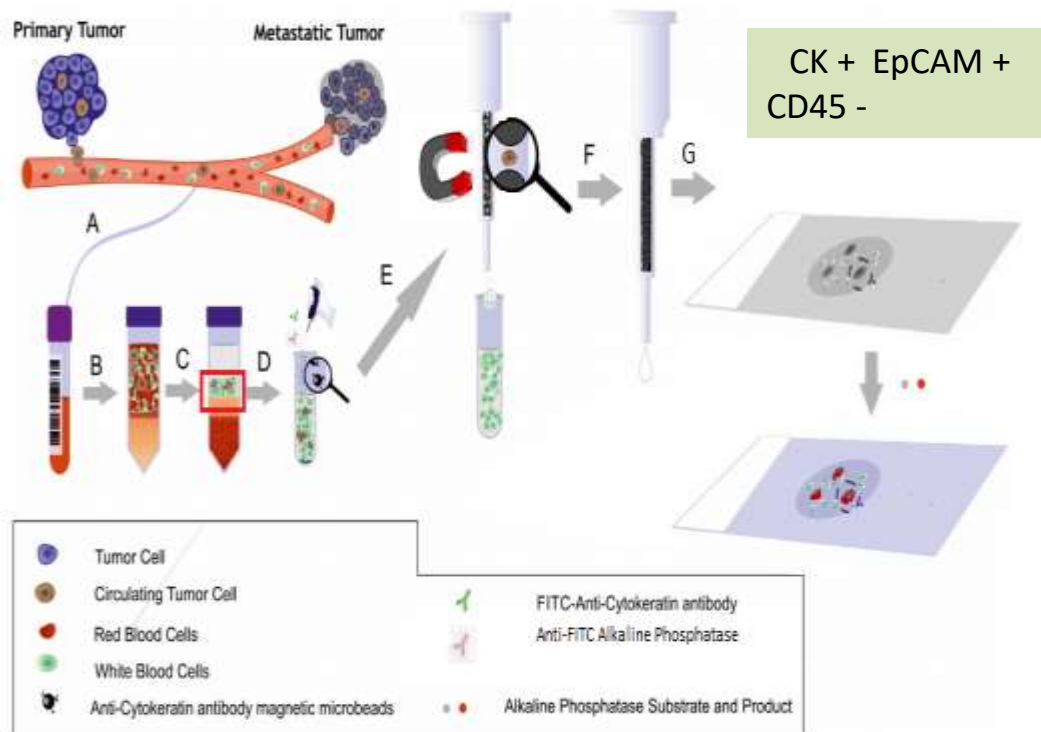
Exoxomas

Proteínas

(...)

¿Cómo se detectan las CTCs?

- 1.- **Fase enriquecimiento** (centrifugado, lavado, conservación)
- 2.- **Fase aislamiento** (inmunomagnético) CKs y marcadores epiteliales
- 3.- **Fase selección** (manual, semiautomática, automática)



breast cancer: >5 CTCs/7.5 mL
colorectal cancer: >3 CTCs/7.5 mL
prostate cancer: >5 CTCs/7.5 mL

**SLP y SG
en pacientes CPRCm**

¿Cómo se regula la aprobación de los test de biopsia líquida?

1- Regulación CIENTÍFICO-ADMINISTRATIVA



2- Regulación CIENTÍFICO-TÉCNICA

Estándares metodológicos

3- Regulación ADMINISTRATIVO-SANITARIA

Países, Comunidades Autónomas, administraciones locales, hospitales....

*SelectMDx*TM *for Prostate Cancer*

Es un algoritmo basado en:

- Niveles ARNm de **HOXC6 y DLX1**
- Edad
- PSA
- Vol próstata
- Tacto rectal
- Antecedentes familiares

Detection of High-grade Prostate Cancer Using a Urinary Molecular Biomarker-Based Risk Score.

Van Neste L¹, Hendriks RJ², Dijkstra S², Trooskens G³, Cornel EB⁴, Jannink SA³, de Jong H³, Hessels D³, Smit FP³, Melchers WJ⁵, Leyten GH⁶, de Reijke TM⁷, Vergunst H⁸, Kil P⁹, Knipscheer BC¹⁰, Hulsbergen-van de Kaa CA¹¹, Mulders PF², van Oort IM², Van Criekinge W¹², Schalken JA¹³.

Características	Cohorte A (desarrollo)	Cohorte B (validación)
n (905) (Pcts a (re)biop por PSA≥3, DRE an)	519	386
1ª biopsia n (%)	410 (79)	322 (89)
PSAD, ng/ml . ml (log ₁₀ ; Q1-Q3; 0,0-0,25)	115 (22,3)	85 (22,0)
DRE anormal (%)	199 (38,3)	119 (31,3)
Diagnóstico (Gleason (%))	111 (21,4)	81 (21,0)
GS≤6, n (%)	103 (48,6)	91 (50,0)
GS 7, n (%)	58 (27,4)	51 (28,3)
GS 8-10, n (%)	51 (24,1)	39 (21,6)

HOXC6 – DLX1

AUC: 0.76

S: 91%
E: 36%

SelectMdx

AUC: 0.86

EVITA en un **53%**
BIOPSIAS INNECESARIAS

↑ VPN

El score de SelectMDx puede estimar:

90% de probabilidad que un paciente NO tiene CaP.

98% de probabilidad que un paciente NO tiene CaP de alto riesgo (Gleason 7 o más)

Caracterización



Factor Predictivo???

ARV-7

ARV7 parece un **claro marcador predictivo** de respuesta a Enza o Abi

Controversia: METODOLOGÍA DE DETECCIÓN

THE NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

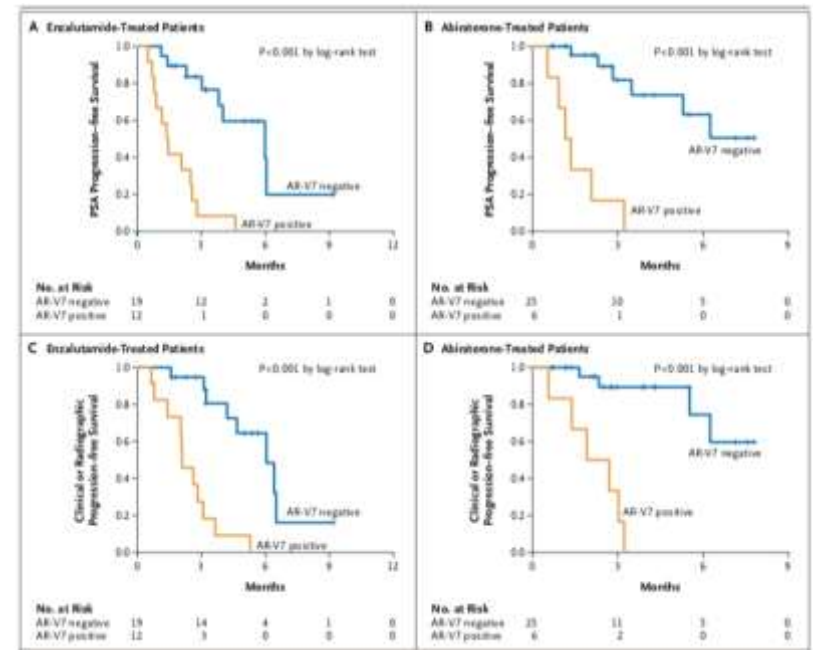
AR-V7 and Resistance to Enzalutamide and Abiraterone in Prostate Cancer

Emmanuel S. Antonarakis, M.D., Changxue Lu, Ph.D., Hao Wang, Ph.D.,
Brandon Luber, Sc.M., Mary Nakazawa, M.H.S., Jeffrey C. Roeser, B.S.,
Yan Chen, Ph.D., Tabrez A. Mohammad, Ph.D., Yidong Chen, Ph.D.,

62 pacientes
SLP PSA, SLP Rx, SG

ARV7

¿Primer test biopsia líquida
que cambia el manejo terapéutico
del cáncer de próstata avanzado?



qPCR CTCs detectadas por CellSearch

Standardization of CTC AR-V7 PCR assay and evaluation of its role in castration resistant prostate cancer progression.

Tommasi S¹, Pilato B¹, Carella C², Lasorella A^{1,3}, Danza K¹, Vallini I³, De Summa S¹, Naglieri E².

 **Author information**



ClinicalTrials.gov identifier: NCT02269982

J Clin Oncol. 2019 May 1;37(13):1120-1129. doi: 10.1200/JCO.18.01731. Epub 2019 Mar 13.

Prospective Multicenter Validation of Androgen Receptor Splice Variant 7 and Hormone Therapy Resistance in High-Risk Castration-Resistant Prostate Cancer: The PROPHECY Study.

Armstrong AJ¹, Halabi S¹, Luo J², Nanus DM³, Giannakakou P³, Szmulewitz RZ⁴, Danila DC^{3,5}, Healy P¹, Anand M¹, Rothwell CJ¹, Rasmussen J¹, Thornburg B¹, Berry WR¹, Wilder RS¹, Lu C², Chen Y², Silberstein JL², Kemeny G¹, Galletti G³, Somarelli JA¹, Gupta S¹, Gregory SG¹, Scher HI^{3,5}, Dittamore R⁶, Tagawa ST³, Antonarakis ES², George DJ¹.

118 pacientes

Johns Hopkins University
modified-AdnaTest **CTC AR-V7**
mRNA assay

Epic Sciences CTC
nuclear-specific AR-V7
protein assay.

Acuerdo
82%

Detection of AR-V7 in CTCs by two blood-based assays is independently associated with shorter PFS and OS with abiraterone or enzalutamide, and such men with mCRPC should be offered alternative treatments.

...Vía rápida...

THERANOS

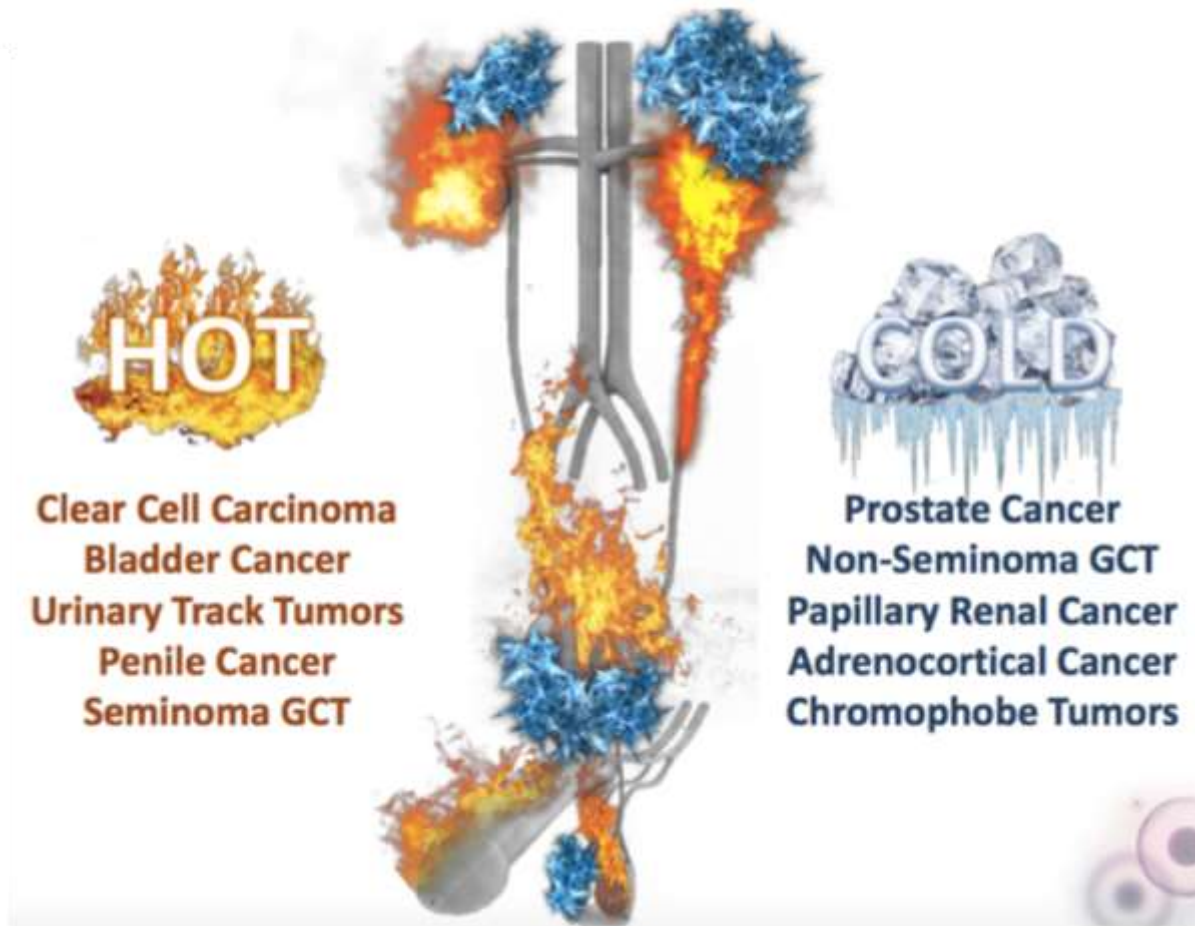
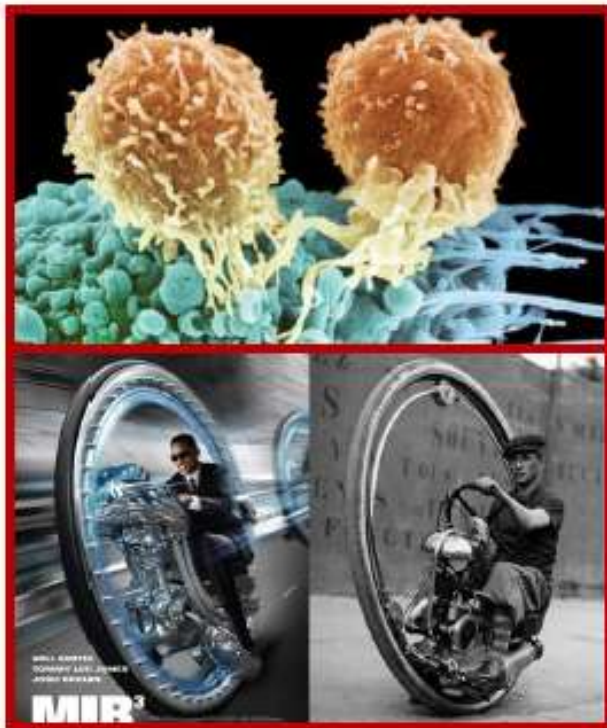


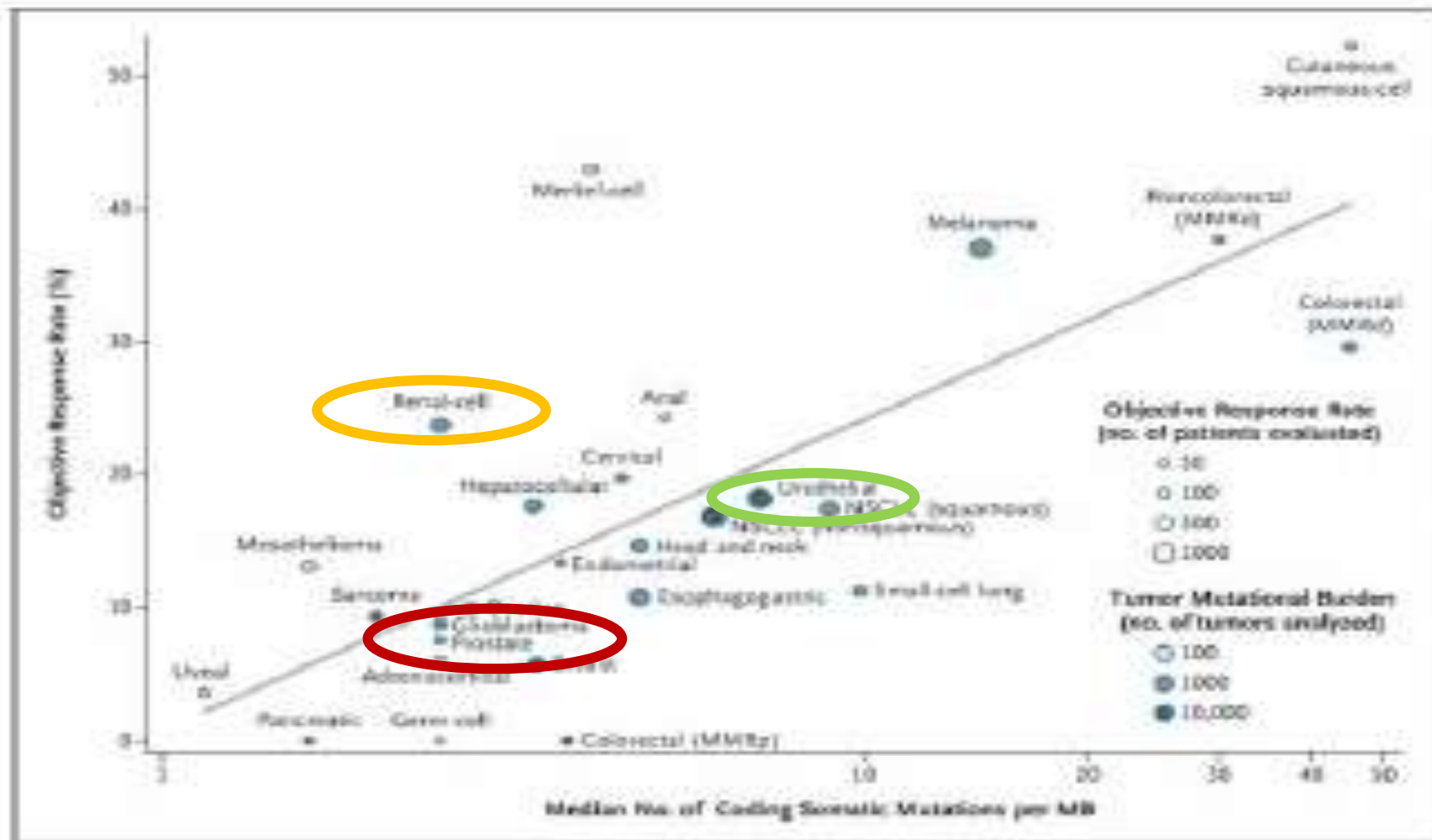
CANCER INTERCEPT DETECT TEST



La inmunoterapia llega al CPRC

Inmunoterapia





Correlation between Tumor Mutational Burden and Objective Response Rate with Anti-PD-1 or Anti-PD-L1



NCCN Guidelines Version 4.2019

Prostate Cancer

SYSTEMIC THERAPY FOR M1 CRPC: ADENOCARCINOMA WITH VISCERAL METASTASES ^{xx,bbb}

FIRST-LINE TREATMENT

- Docetaxel^{yy} (category 1)
- Enzalutamide^u (category 1)
- Abiraterone^u with prednisone
- Abiraterone^u with methylprednisolone
- Clinical trial
- Mitoxantrone with prednisone^{yy,ggg}
- Other secondary hormone therapy^u

Prior therapy
enzalutamide/
abiraterone^{ccc,ddd,hhh}

Prior therapy
docetaxel^{ddd,hhh}

SECOND-LINE TREATMENT

- Docetaxel^{yy} (category 1)
 - If not previously received:
 - ▶ Abiraterone^u with prednisone
 - ▶ Abiraterone^u with methylprednisolone
 - ▶ Enzalutamide^u
 - ▶ Cabazitaxel^{yy}
 - Pembrolizumab for MSI-H or dMMR^{yy} (category 2B)
 - Clinical trial
 - Other secondary hormone therapy^u
 - Best supportive care
-
- Abiraterone^u with prednisone (category 1)
 - Enzalutamide^u (category 1)
 - Cabazitaxel^{yy} (category 1)
 - Abiraterone^u with methylprednisolone
 - Pembrolizumab for MSI-H or dMMR^{yy} (category 2B)
 - Clinical trial
 - Docetaxel rechallenge^{yy,fff}
 - Mitoxantrone with prednisone^{yy,ggg}
 - Other secondary hormone therapy^u
 - Best supportive care

SUBSEQUENT TREATMENT (category 2B)

- At progression^{ddd,hhh}:
- If not previously received:
 - ▶ Enzalutamide^u
 - ▶ Cabazitaxel^{yy}
 - ▶ Abiraterone^u with prednisone
 - ▶ Abiraterone^u with methylprednisolone
 - ▶ Mitoxantrone with prednisone^{ggg}
 - ▶ Pembrolizumab for MSI-H or dMMR^{yy}
 - Clinical trial
 - Docetaxel rechallenge^{yy,fff}
 - Other secondary hormone therapy^u
 - Best supportive care

^u See [Principles of Androgen Deprivation Therapy \(PROS-F\)](#)

^{yy} See [Principles of Immunotherapy and Chemotherapy \(PROS-G\)](#)

^{xx} Visceral metastases refers to liver, lung, adrenal, peritoneal, and brain metastases. Soft tissue/lymph node sites are not considered visceral metastases.

^{bbb} Patients can continue through all treatment options listed. Best supportive care is always an appropriate option.

^{ccc} Consider AR-V7 testing to help guide selection of therapy ([See Discussion](#))

^{ddd} Workup for progression should include chest CT, bone imaging, and abdominal/pelvic CT or abdominal/pelvic MRI with and without contrast. Consider metastatic lesion biopsy. [See Principles of Imaging \(PROS-B\)](#) and [Discussion](#)

^{fff} Patients who received docetaxel with ADT in the metastatic castration-naïve setting can be considered for docetaxel rechallenge in the CRPC setting.

^{ggg} Mitoxantrone with prednisone is for palliation in symptomatic patients who cannot tolerate other therapies.

^{hhh} Patients treated with systemic therapy for non-visceral metastases ([see PROS-17](#)) should proceed to a different subsequent therapy for patients with visceral metastases based on their previous treatment. If small cell neuroendocrine is found, [see PROS-18](#)

Note: All recommendations are category 2A unless otherwise indicated.

Clinical Trials: NCCN believes that the best management of any patient with cancer is in a clinical trial. Participation in clinical trials is especially encouraged.



National
Comprehensive
Cancer
Network®

NCCN Guidelines Version 2.2020 Kidney Cancer

[NCCN Guidelines Index](#)
[Table of Contents](#)
[Discussion](#)

PRINCIPLES OF SYSTEMIC THERAPY FOR RELAPSE OR STAGE IV DISEASE

FIRST-LINE THERAPY FOR CLEAR CELL HISTOLOGY			
Risk	Preferred regimens	Other recommended regimens	Useful under certain circumstances
Favorable ^a	- Axitinib + pembrolizumab - Pazopanib - Sunitinib	- Ipilimumab + nivolumab - Cabozantinib (category 2B) - Axitinib + avelumab	- Active surveillance^b - Axitinib (category 2B) - High-dose IL-2^c
Poor/ intermediate ^a	- Ipilimumab + nivolumab (category 1) - Axitinib + pembrolizumab (category 1) - Cabozantinib	- Pazopanib - Sunitinib - Axitinib + avelumab	- Axitinib (category 2B) - High-dose IL-2^c - Temsirolimus^d

SUBSEQUENT THERAPY FOR CLEAR CELL HISTOLOGY		
Preferred regimens	Other recommended regimens	Useful under certain circumstances
- Cabozantinib (category 1) - Nivolumab (category 1) - Ipilimumab + nivolumab	- Axitinib (category 1) - Ipilimumab + everolimus (category 1) - Axitinib + pembrolizumab - Everolimus - Pazopanib - Sunitinib - Axitinib + avelumab (category 3)	- Bevacizumab or biosimilar^e (category 2B) - Sorafenib (category 2B) - High-dose IL-2 for selected patients^c (category 2B) - Temsirolimus^d (category 2B)

^a See Risk Models to Direct Treatment (IMDC criteria) (KID-D).

^b Rini BI, Dorr TB, Elson P, et al. Active surveillance in metastatic renal-cell carcinoma: a prospective, phase 2 trial. *Lancet Oncol* 2016;17:1317-1324.

^c Patients with excellent performance status and normal organ function.

^d The poor risk model used in the global ARCC trial to direct treatment with temsirolimus included at least 3 of the following 6 predictors of short survival: <1 year from the time of diagnosis to start of systemic therapy, Karnofsky performance status score 60–70, hemoglobin <LLN, corrected calcium greater than 10 mg/dL, LDH >1.5 times the ULN, and metastasis in multiple organs. Hudes G, Carducci M, Tomczak P, et al. Temsirolimus, interferon alfa, or both for advanced renal-cell carcinoma. *N Engl J Med* 2007;356:2271-2281.

^e Biosimilar options include: bevacizumab-awwb, bevacizumab-bvzr.

Note: All recommendations are category 2A unless otherwise indicated.

Clinical Trials: NCCN believes that the best management of any patient with cancer is in a clinical trial. Participation in clinical trials is especially encouraged.



PRINCIPLES OF SYSTEMIC THERAPY

First-line systemic therapy for locally advanced or metastatic disease (Stage IV)	
Cisplatin eligible	Preferred regimens - Gemcitabine and cisplatin⁴ (category 1) - DDMVAC with growth factor support (category 1)^{2,8}
Cisplatin ineligible	Preferred regimens - Gemcitabine and carboplatin¹¹ - Atezolizumab¹² (only for patients whose tumors express PD-L1^a or who are not eligible for any platinum-containing chemotherapy regardless of PD-L1 expression) - Pembrolizumab¹³ (only for patients whose tumors express PD-L1^b or who are not eligible for any platinum-containing chemotherapy regardless of PD-L1 expression) Other recommended regimens - Gemcitabine¹⁴ - Gemcitabine and paclitaxel¹⁵ Useful under certain circumstances - Ifosfamide, doxorubicin, and gemcitabine¹⁶ (for patients with good kidney function and good PS)

- The presence of both non-nodal metastases and ECOG performance score ≥ 2 strongly predict poor outcome with chemotherapy. Patients without these adverse prognostic factors have the greatest benefit from chemotherapy. The impact of these factors in relation to immune checkpoint inhibition is not fully defined, but they remain poor prognostic indicators in general.
- For most patients, the risks of adding paclitaxel to gemcitabine and cisplatin outweigh the limited benefit seen in the randomized trial.¹⁷
- A substantial proportion of patients cannot receive cisplatin-based chemotherapy due to renal impairment or other comorbidities.
 - Participation in clinical trials of new or more tolerable therapy is recommended.

^a Atezolizumab: SP142 assay. PD-L1–stained tumor-infiltrating immune cells covering $\geq 5\%$ of the tumor area.

^b Pembrolizumab: 22C3 antibody assay. Combined Positive Score (CPS) ≥ 10 .

Note: All recommendations are category 2A unless otherwise indicated.

Clinical Trials: NCCN believes that the best management of any patient with cancer is in a clinical trial. Participation in clinical trials is especially encouraged.



NCCN Guidelines Version 3.2020

Bladder Cancer

PRINCIPLES OF SYSTEMIC THERAPY

Second-line systemic therapy for locally advanced or metastatic disease (Stage IV) (post-platinum) ^c Participation in clinical trials of new agents is recommended.	
Preferred regimen • Pembrolizumab (category 1) ¹⁶	Other recommended regimens • Paclitaxel ²⁶ or docetaxel ²⁷ • Gemcitabine ¹⁴
Alternative preferred regimens • Immune checkpoint inhibitor ▶ Atezolizumab ^{19,20} ▶ Nivolumab ²¹ ▶ Durvalumab ²² ▶ Avelumab ^{23,24} ▶ Enfortumab vedotin ²⁵	Useful in certain circumstances based on prior medical therapy • Ifosfamide, doxorubicin, and gemcitabine ¹⁶ • Gemcitabine and paclitaxel ¹⁵ • Gemcitabine and cisplatin ⁴ • DDMVAC with growth factor support ²

Second-line systemic therapy for locally advanced or metastatic disease (Stage IV) (post-checkpoint inhibitor) Participation in clinical trials of new agents is recommended.	
Preferred regimen for cisplatin ineligible, chemotherapy naïve • Gemcitabine/carboplatin	Other recommended regimens • Erdafitinib ^{d,25} • Paclitaxel or docetaxel ²⁷ • Gemcitabine ¹⁴
Preferred regimens for cisplatin eligible, chemotherapy naïve • Gemcitabine and cisplatin ⁴ • DDMVAC with growth factor support ²	Useful in certain circumstances based on prior medical therapy • Ifosfamide, doxorubicin, and gemcitabine ¹⁶ • Gemcitabine and paclitaxel ¹⁵

^c If PFS >12 months after platinum (eg, cisplatin or carboplatin), consider re-treatment with platinum if the patient is still platinum eligible.

^d Only for patients with susceptible *FGFR3* or *FGFR2* genetic alterations.

Note: All recommendations are category 2A unless otherwise indicated.

Clinical Trials: NCCN believes that the best management of any patient with cancer is in a clinical trial. Participation in clinical trials is especially encouraged.

[Continued](#)
[References](#)

BL-G
3 OF 7

ESOU20

17th Meeting of the
EAU Section of
Oncological Urology

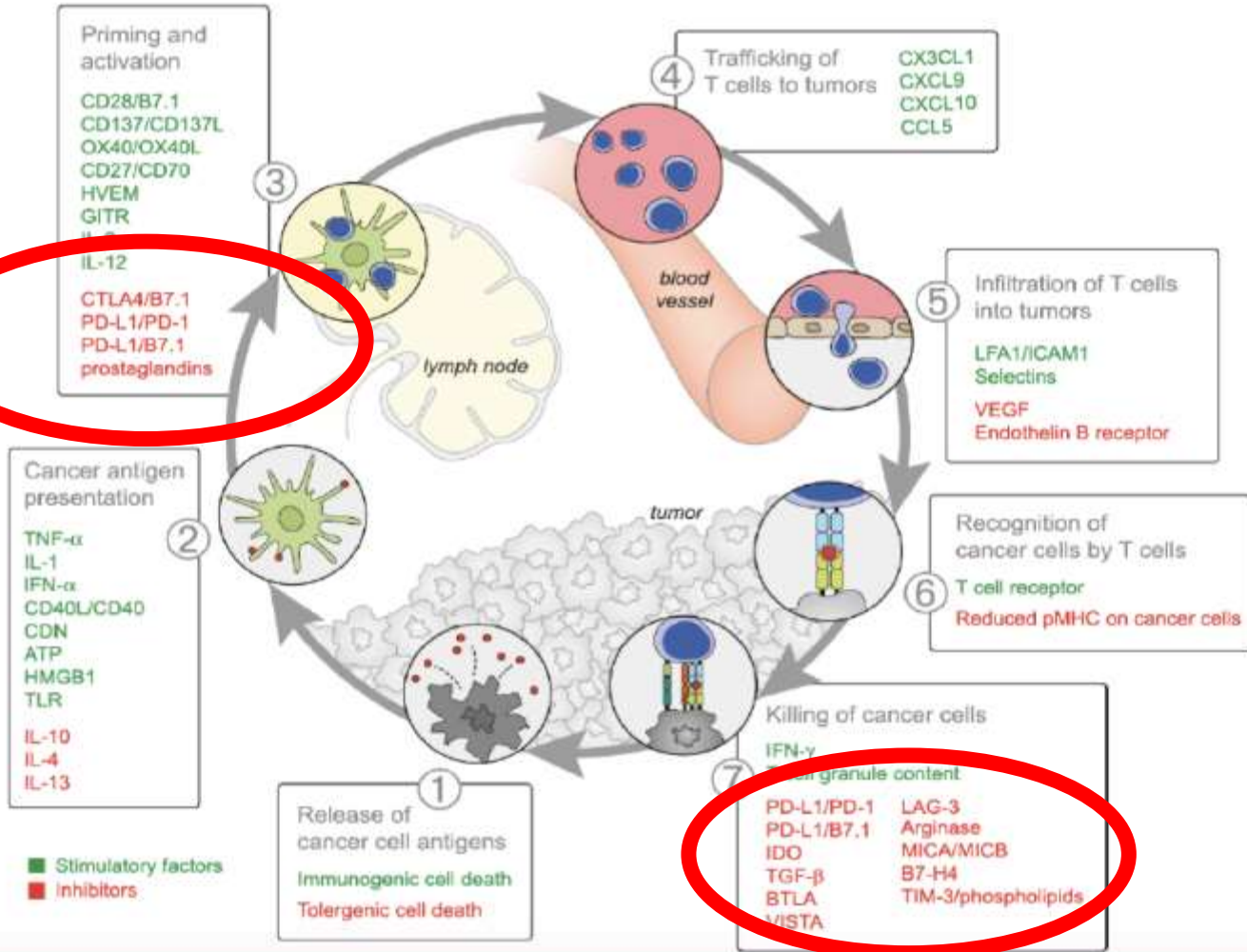
17-19 January 2020
Dublin, Ireland

Immunotherapy in PCa

- I do agree with Nicholas: IO is quite disappointing in prostate cancer
- Though, do we have some reasons to expect better outcomes ?



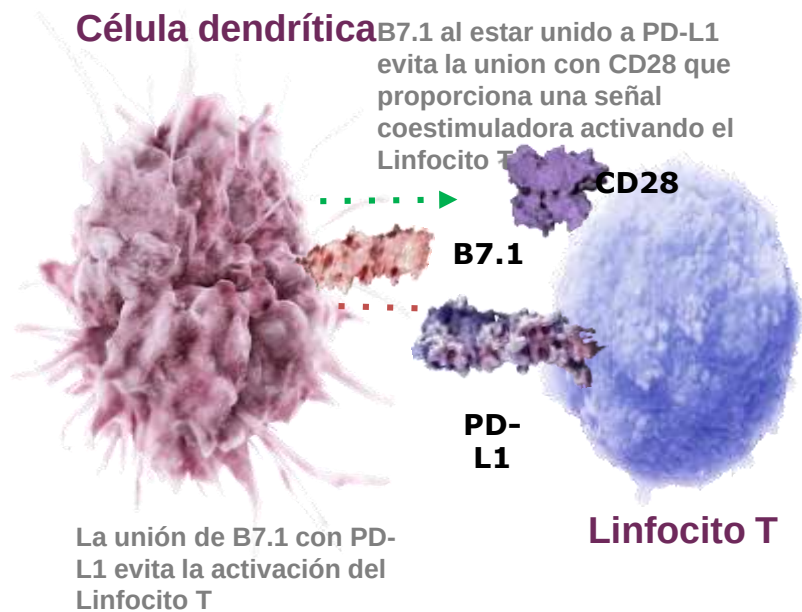
Activatory and Inhibitory Signals



3

Ganglio linfático:

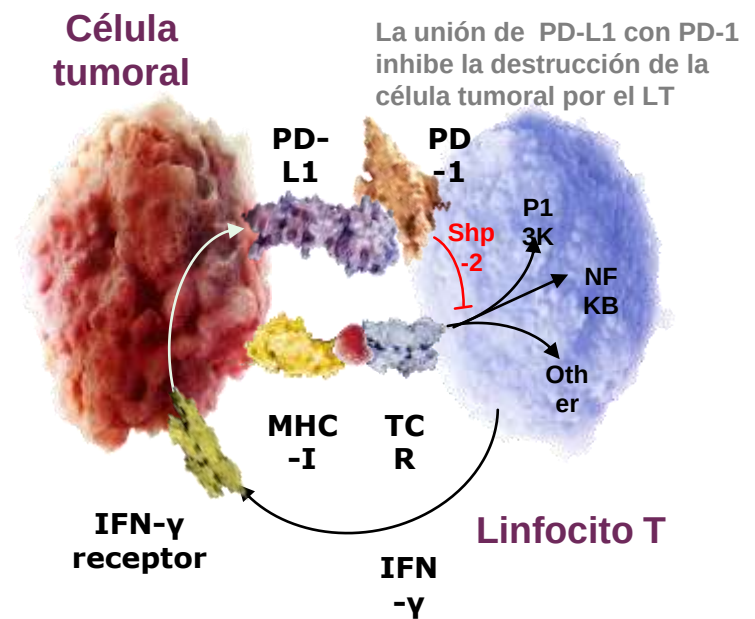
PD-L1 inhibe la activación del LT

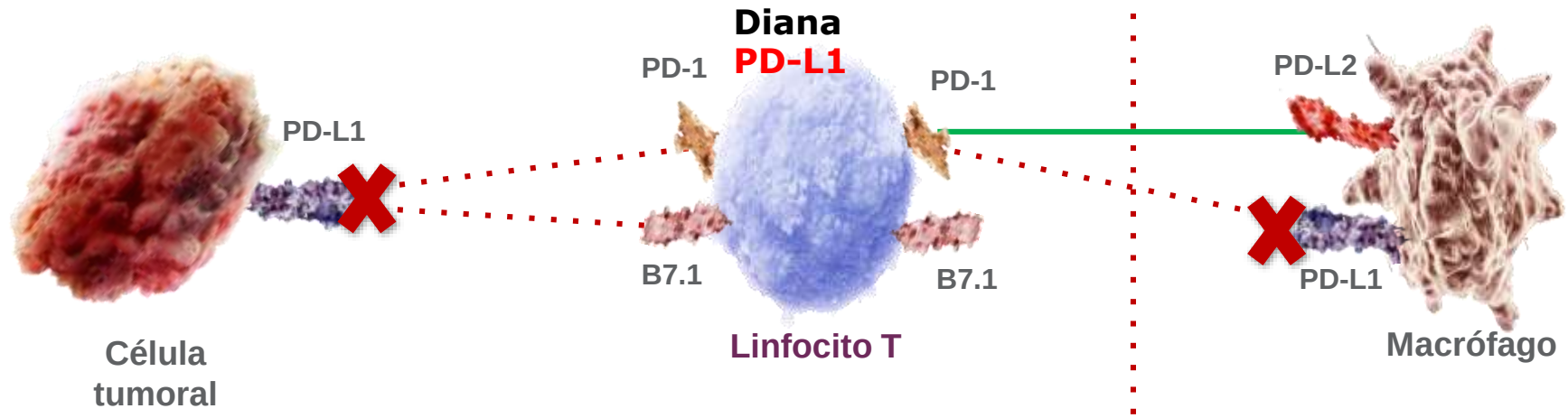


7

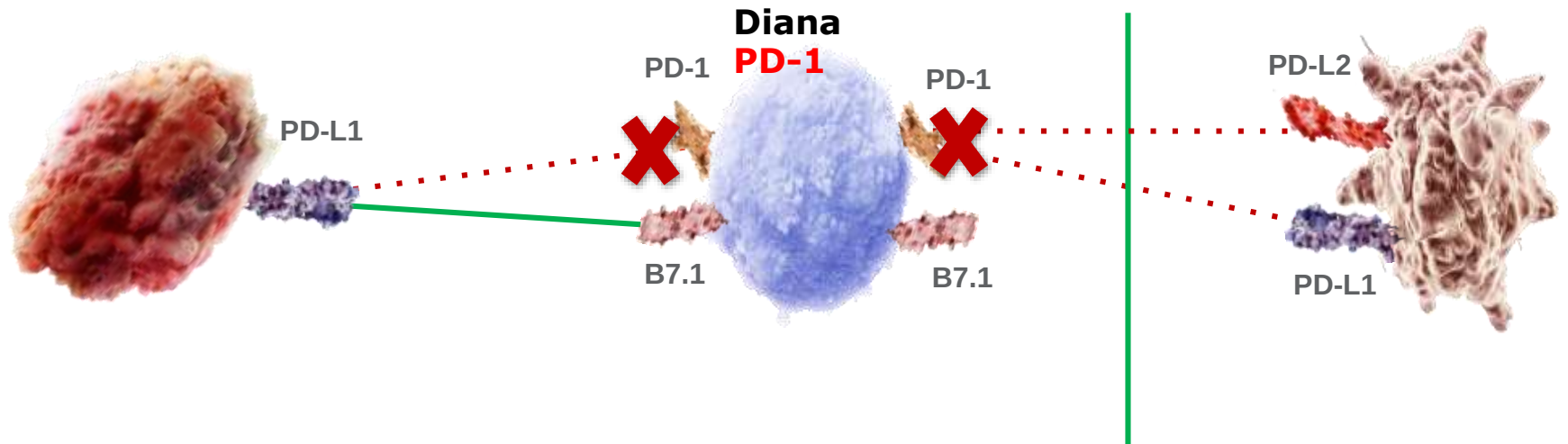
Microambiente tumoral:

PD-L1 inhibe la destrucción de la célula tumoral por el LT





Los inhibidores PD-L1 and PD-1 tienen diferentes lugares de activación, que generan diferencias en la activación del linfocito T

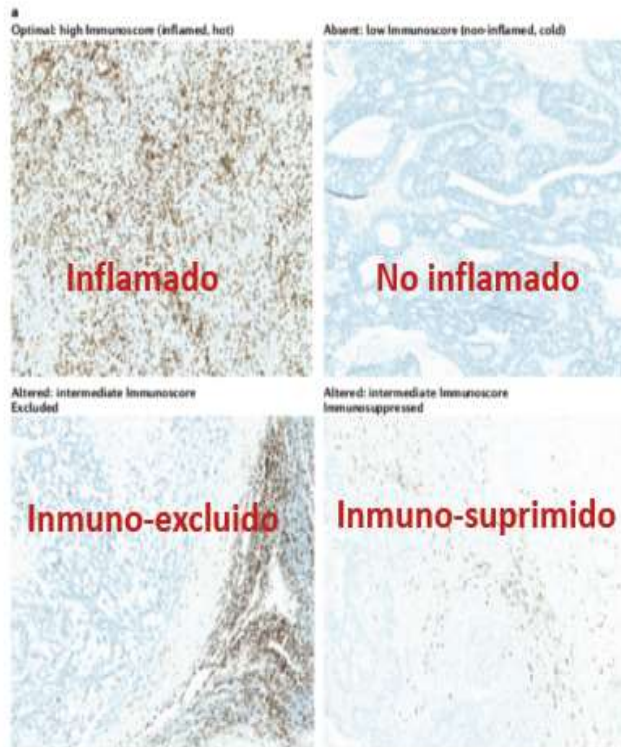


Approaches to treat immune hot, altered and cold tumours with combination immunotherapies

Jérôme Galon* and Daniela Bruni



Galon et al. Nat Rev Drug Discov 2019

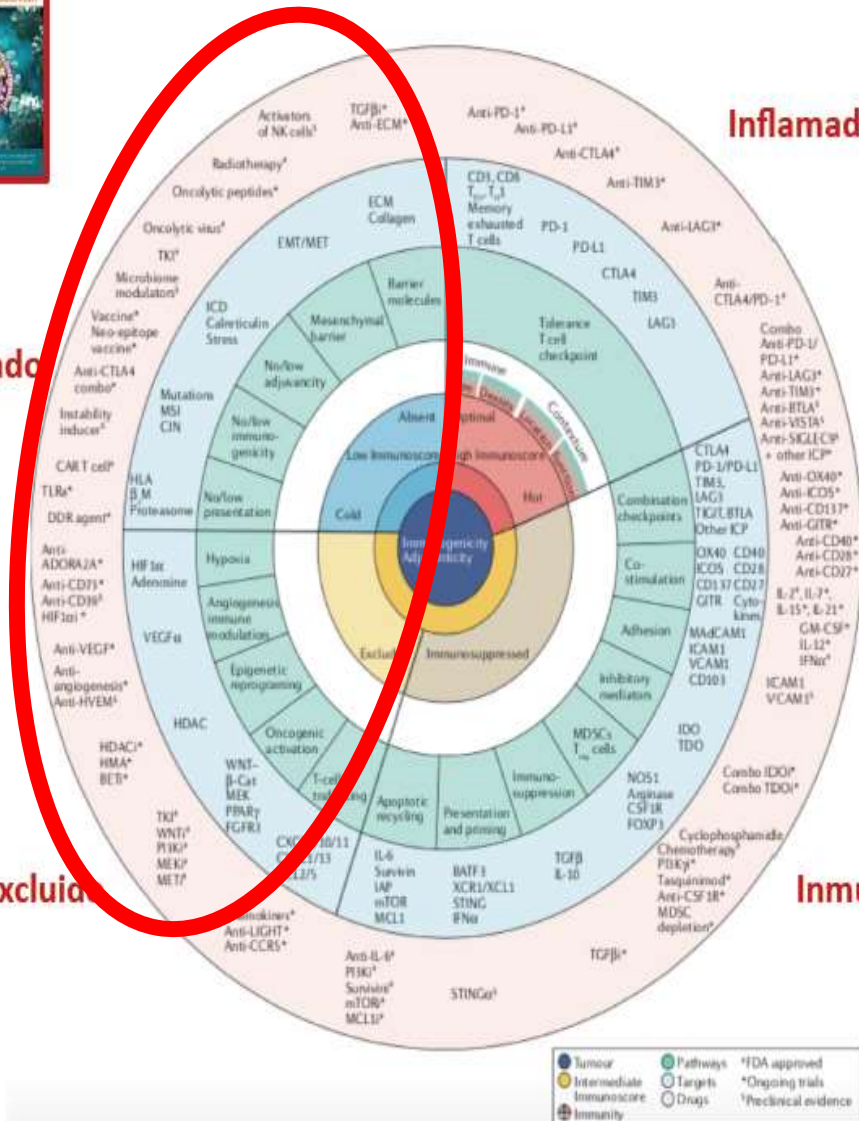


No inflamado

Immuno-excludido

Inflamado

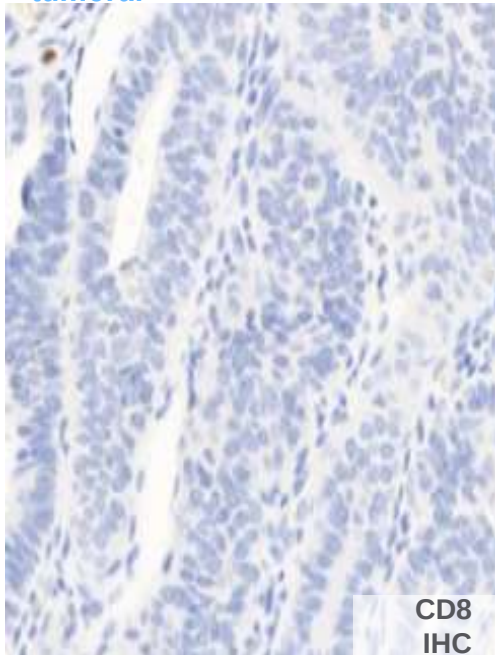
Immuno-suprimido



Absent Low Immunoscore	Altered Intermediate Immunoscore	Optimal High Immunoscore
Cold Non-inflamed	Excluded CT-LA, HI-IM	Immunosuppressed Hot Inflamed

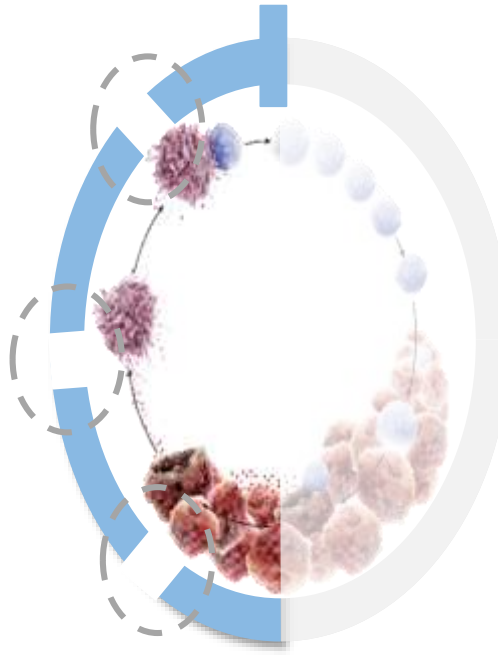
Response to T cell checkpoint inhibition

Se caracteriza por una falta de linfocitos T CD8+ en el estroma o parénquima tumoral



Fenotipo No inflamado

Estos tumores suelen tener un defecto en los estadios primarios del ciclo inmune del cáncer (paso 1-3)



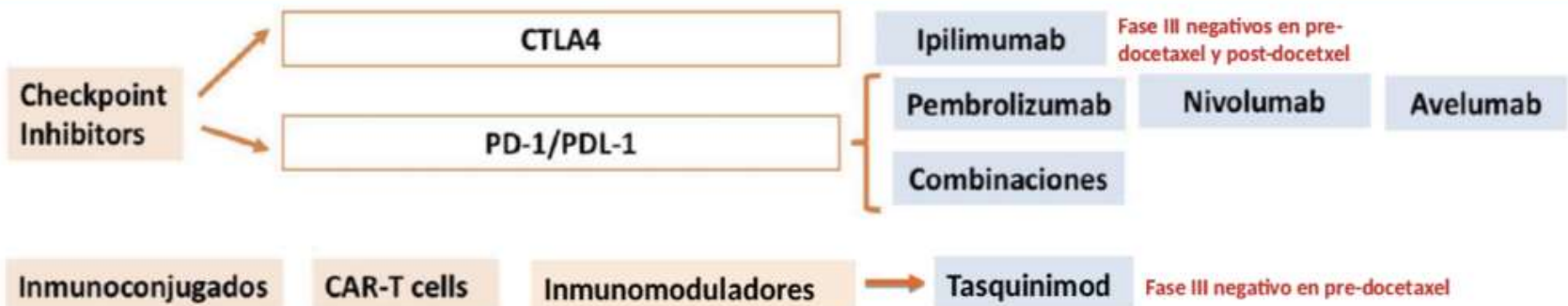
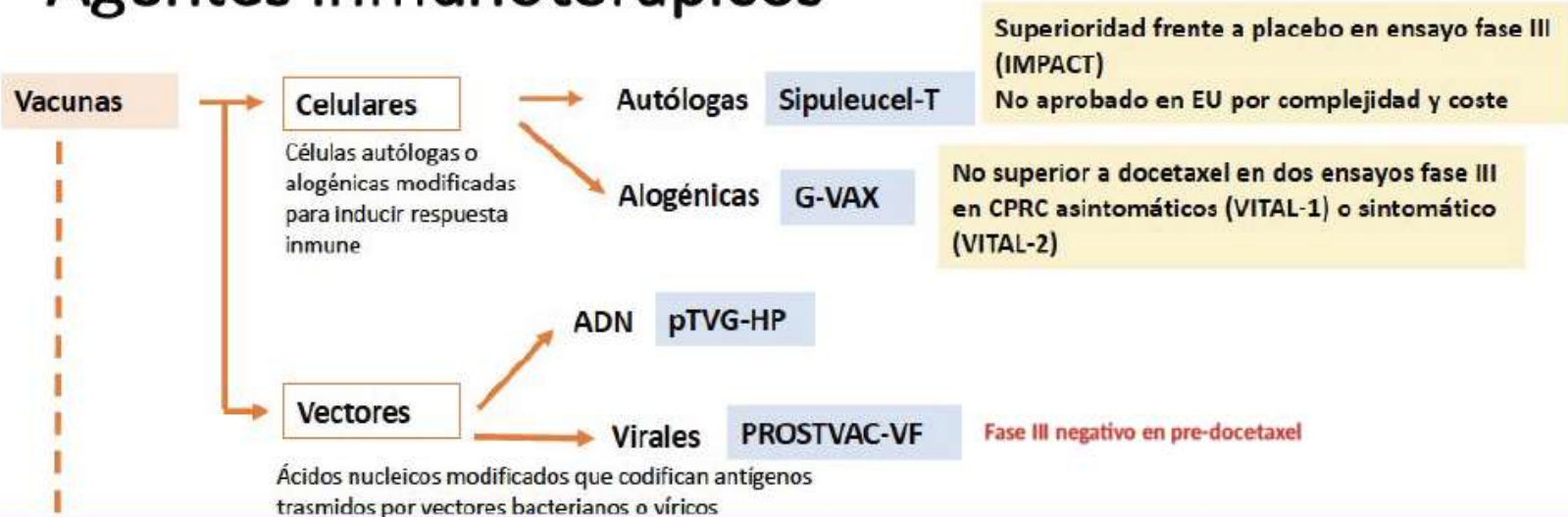
Probablemente debido a fallos en los pasos 1-3 del ciclo inmune del cáncer⁴

Mecanismos asociados con el escape al Sistema inmune que perjudican a la generación de linfocitos T

- 1 Escape debido a la baja inmunogenicidad del tumor**
(ej: carga mutacional del tumor reducida, MHC-I bajo)
- 2 Escape debido a la reducida maduración de las células dendríticas**
(ej: numerosos factores solubles derivados del cáncer o células inmunes inhibitorias)
- 3 Escape debido a la baja activación de linfocitos T**
(ej: falta de interacciones coestimuladoras entre las células dendríticas y los linfocitos T, como OX40 o 4-1BB, limitada producción de IL-2 o sobreexpresión de CTLA-4)

I N M U N O T E R A P I A

Agentes inmunoterápicos



Initial Results From a Phase 2 Study of Nivolumab Plus Ipilimumab for the Treatment of Metastatic Castration-Resistant Prostate Cancer (CheckMate 650)

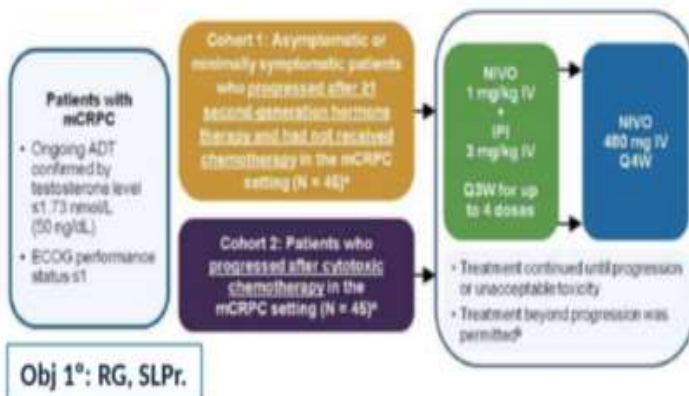
CPRCm

➤ Inhibidores de *checkpoint*.

❖ Combinaciones.

Nivolumab + Ipilimumab

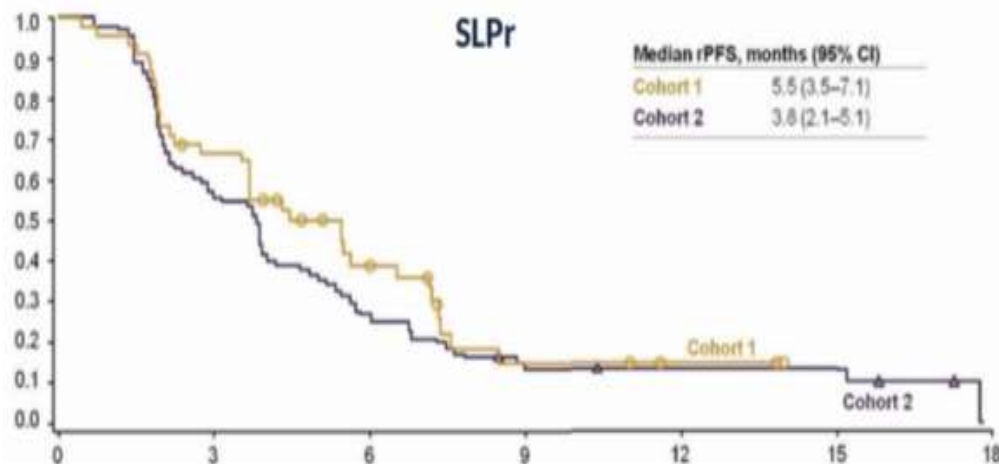
CA209-650.



RG

Objective response (measurable disease only) ^a	Cohort 1 (N = 32)	Cohort 2 (N = 30)
Confirmed ORR, n (%)	8 (25.0)	3 (10.0)
95% CI	11.5–43.4	2.1–26.5
Best overall response, n (%)		
Complete response	2 (6.3) ^b	2 (6.7)
Partial response	6 (18.8) ^c	1 (3.3)
Stable disease	13 (40.6)	11 (36.7)
Progressive disease	9 (28.1)	13 (43.3)
Unable to determine	2 (6.3)	3 (10.0)
Median time to response, months (Q1–Q3)	1.9 (1.9–2.8)	2.1 (1.9–7.4)

SLPr



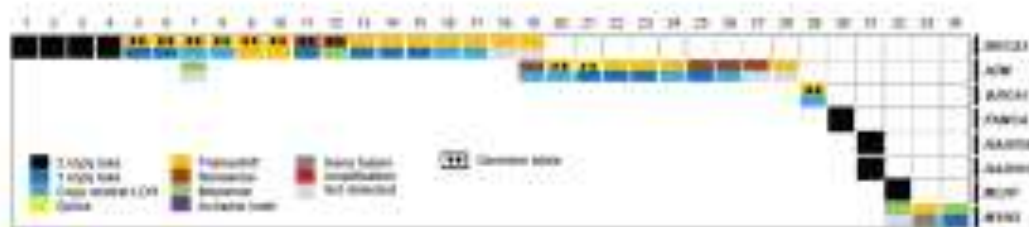
Immunotherapy of prostate cancer combination therapy



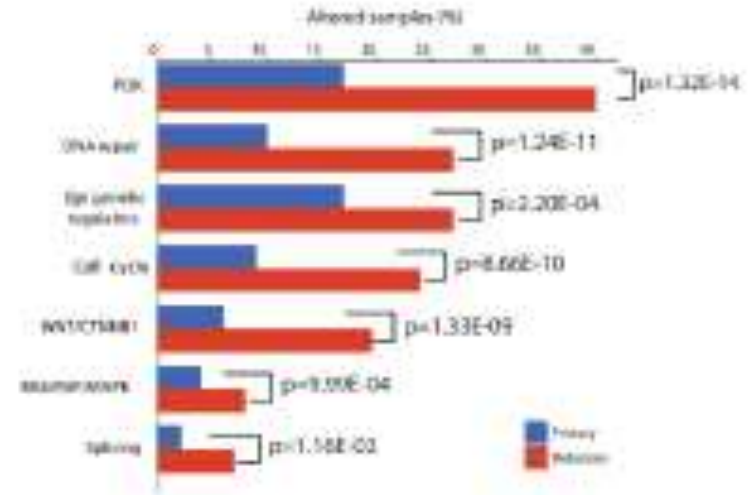
Anti-PD-1/ anti-PDL-1	combined with	remarks
nivolumab	Vaccines (eg Prostavac)	
nivolumab	ipilimumab	Neoantigen DNA vaccine, Prostavac, Biomarker driven, Immunogenic signature, enzalutamide
pembrolizumab	Olaparib, docetaxel, enzalutamide	Phase III planned
pembrolizumab	DNA vaccines, Radium-223, ADX531-142, CPI-444	
atezolizumab	Radium-223, Sipuleucel-T, Enzalutamide	Phase III study enzalutamide
avelumab	Sipuleucel-T	
Durvalumab	Tremulimumab	
Regn2810	Ipilimumab (intraprostatic)	+ stereotactic RT

**Los primeros biomarcadores. BCRA2.
El tratamiento futuro hacia la medicina de la
precisión.**

Prevalence of DDR defects in prostate cancer

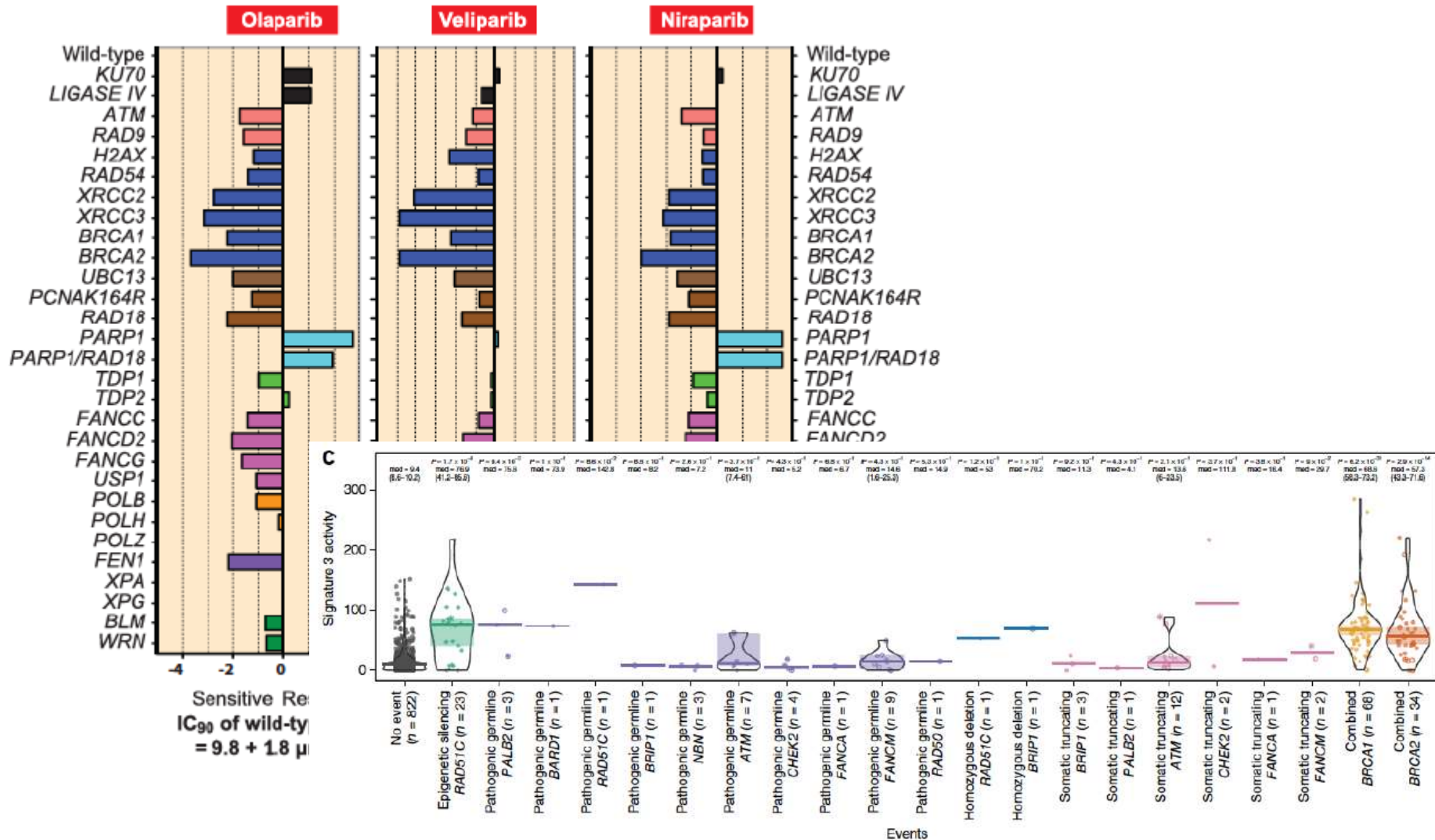


- DNA repair defects identified in 23% of mCRPC
- 8% germline mutations
- 13% BRCA2

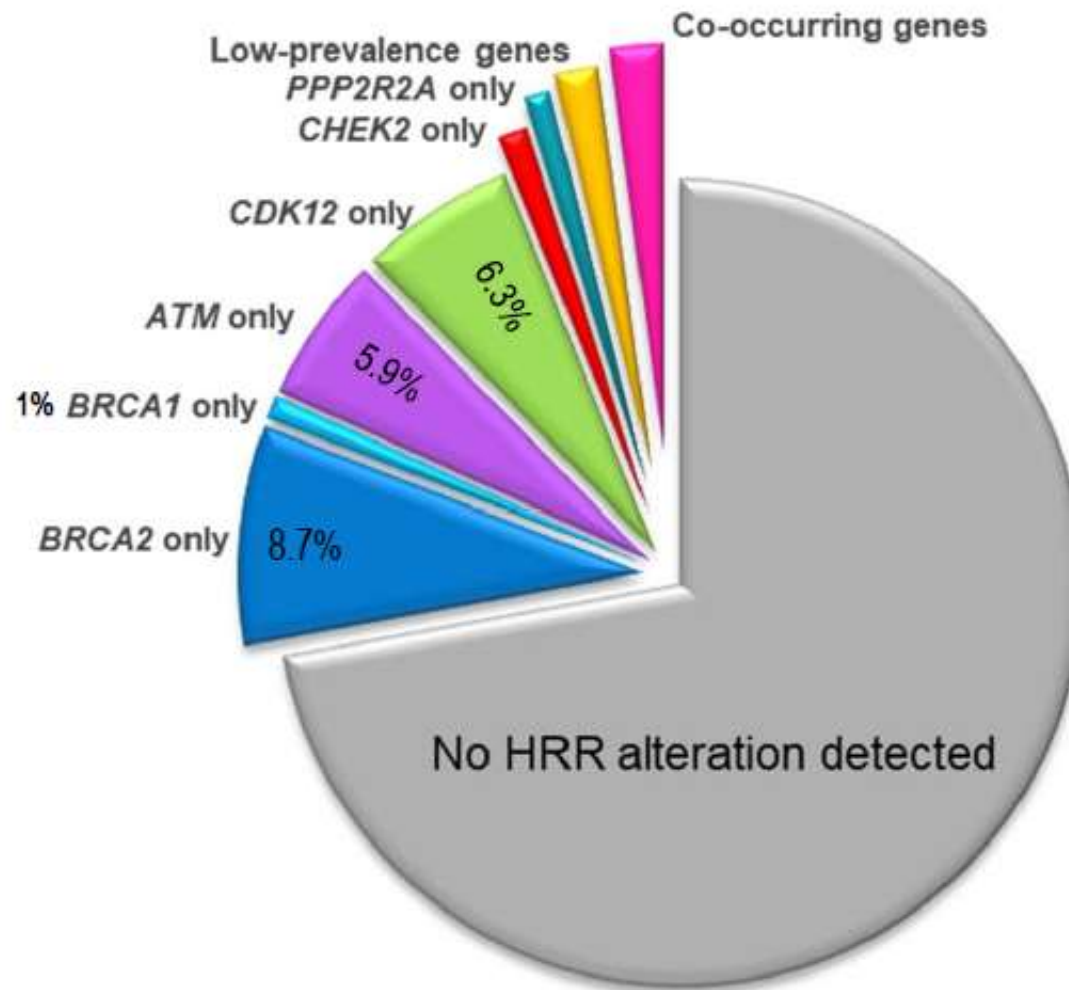


PARP inhibitors:

Different magnitude of sensitization



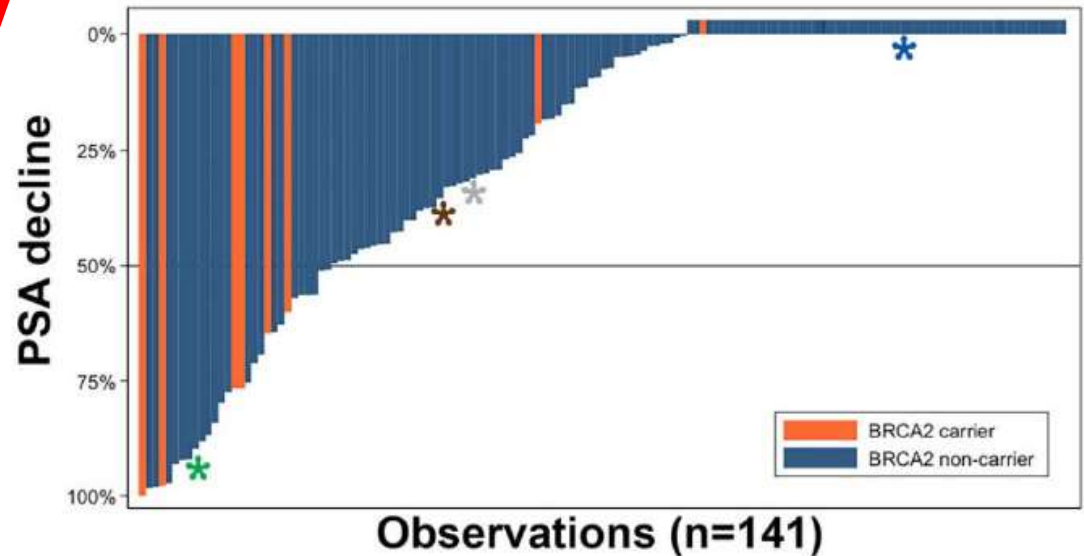
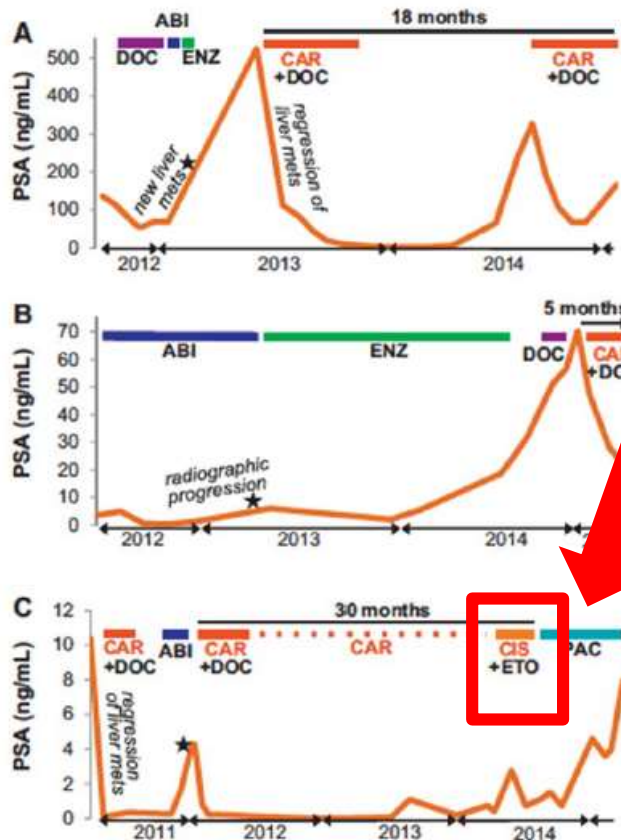
PREVALENCE OF ALTERATIONS IDENTIFIED IN PROFOUND STUDY



Phase II studies with PARPi presented at ESMO2019

	TRITON 2	GALAHAD
Drug	Rucaparib 600 mg bid	Niraparib 300 mg qd
Study Design	Phase II	Phase II
Population	mCRPC Progression to ARSi and taxane	mCRPC Progression to ARSi and taxane
Primary objective	ORR and PSA response (≥50% decline) in pts with DDR alterations	ORR in patients with Bi- allelic BRCA1/2 alterations
Specimen tested	Plasma or tumor tissue Central/local	Plasma Central
Test used	FoundationOne® FoundationACT® Local	Resolution-HRD®
Genes screened	<i>ATM, BARD1, BRCA1, BRCA2, BRIP1, CDK12, CHEK2, FANCA, NBN, PALB2, RAD51, RAD51B, RAD51C, RAD51D, RAD54L</i>	<i>ATM, BRCA1, BRCA2, BRIP1, CHEK2, FANCA, HDAC2, PALB2</i>
Genomic alteration required	Mono- Bi- allelic DDR alterations	Bi-allelic DDR alterations

gBRCA2 mutations and platine-based chemotherapy



CONCLUSIONES

- La señalización del RA es un mecanismo clave en la progresión del CPRC y es una diana lógica de tratamiento. Así surgieron las primeras terapias dirigidas contra receptores androgénicos.
- Múltiples procesos moleculares contribuyen a la señalización sostenida del RA algunos dependientes del RA y otros independientes.
- Es importante el conocimiento de las mutaciones genómicas para identificar nuevas rutas de tratamiento y así bloquear las diversas vías de escape que presentan las células tumorales.
- Los paneles genéticos se basan en un conjunto de biomarcadores que tienen como objetivos ser pronósticos y predictivos de la enfermedad y dirigir a tratamientos selectivos individualizados.
- La biopsia líquida es un buen procedimiento para la obtención de estos biomarcadores útiles en tumores como el CPRC poco accesibles y que precisan de controles periódicos. Precisa todavía de una regulación científico-técnica-administrativa.
- El CPRC es un tumor frío, aunque puede existir un futuro prometedor con las combinaciones de inmunoterapia y sobre todo a las combinaciones de esta con otras terapias dirigidas.
- Terapias dirigidas a Biomarcadores como BCRA2, ATM o CDK12 han provocado la aparición de nuevas terapias altamente precisas, dirigidas y que superan a los tratamientos gold standar. Han abierto nuevas expectativas al tratamiento futuro del CPRC

MUCHAS GRACIAS