

ELBS

Uveal Melanoma

- Perspectives/challenges in metastatic disease -

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Potential Conflicts of Interest

Scientific support:

BMS, Sanofi, Sunpharma

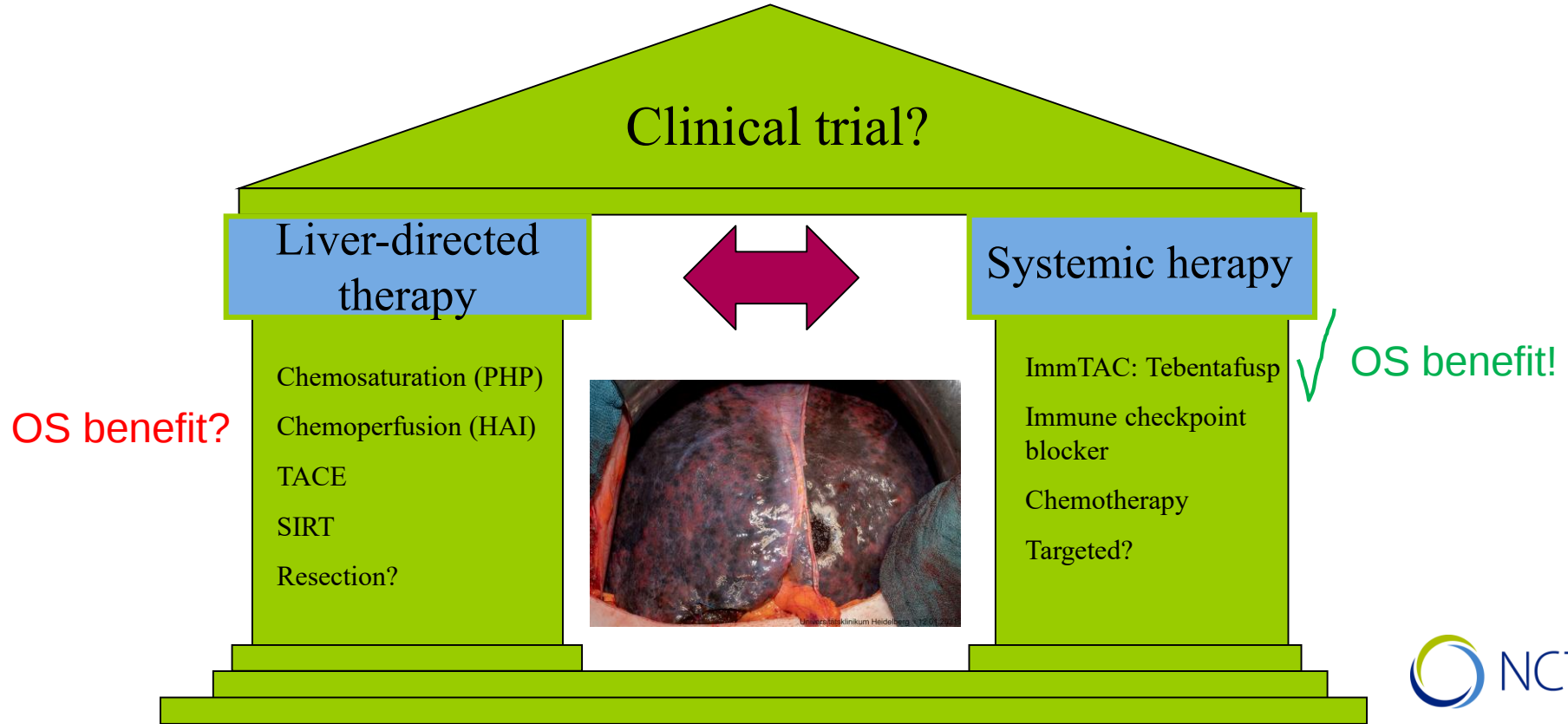
Talks:

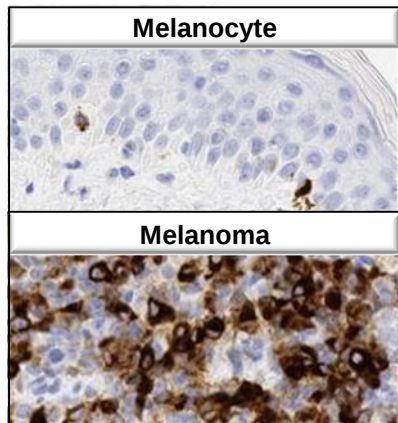
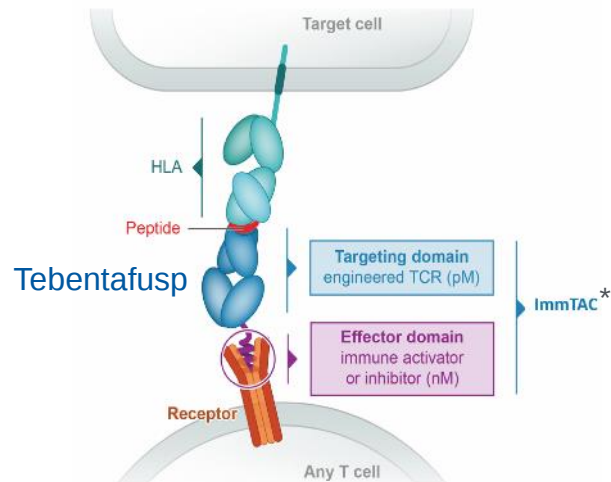
Almirall, Amgen, BMS, GSK, Immunocore, MSD, Novartis, Pfizer, Pierre Fabre, Roche, Sanofi, Sunpharma

Advisory Boards:

GSK, MSD, Pierre Fabre, Sunpharma

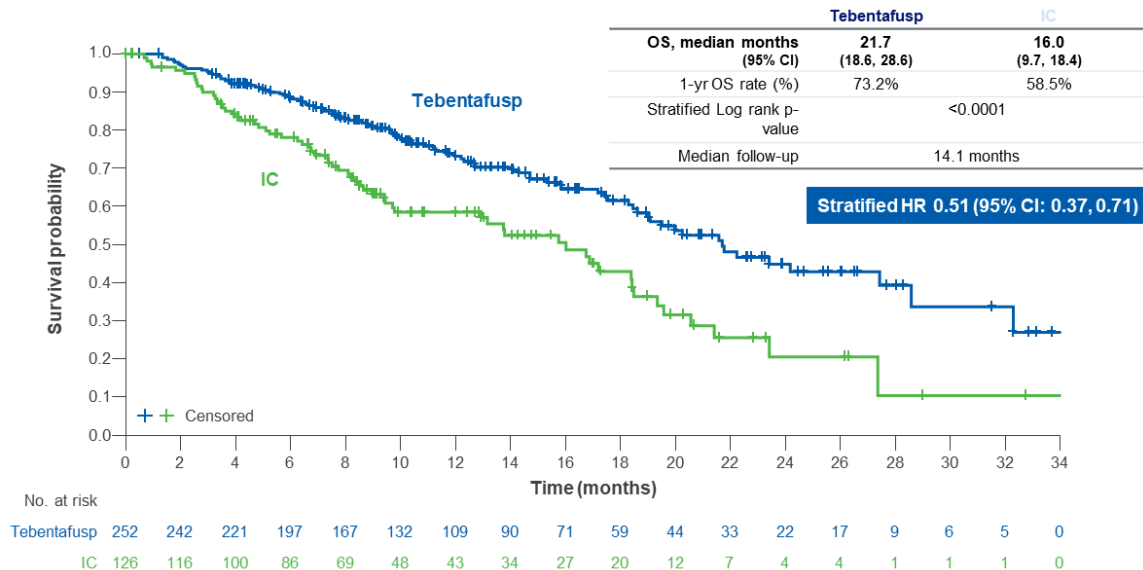
Treatment options in metastasized Uveal Melanoma





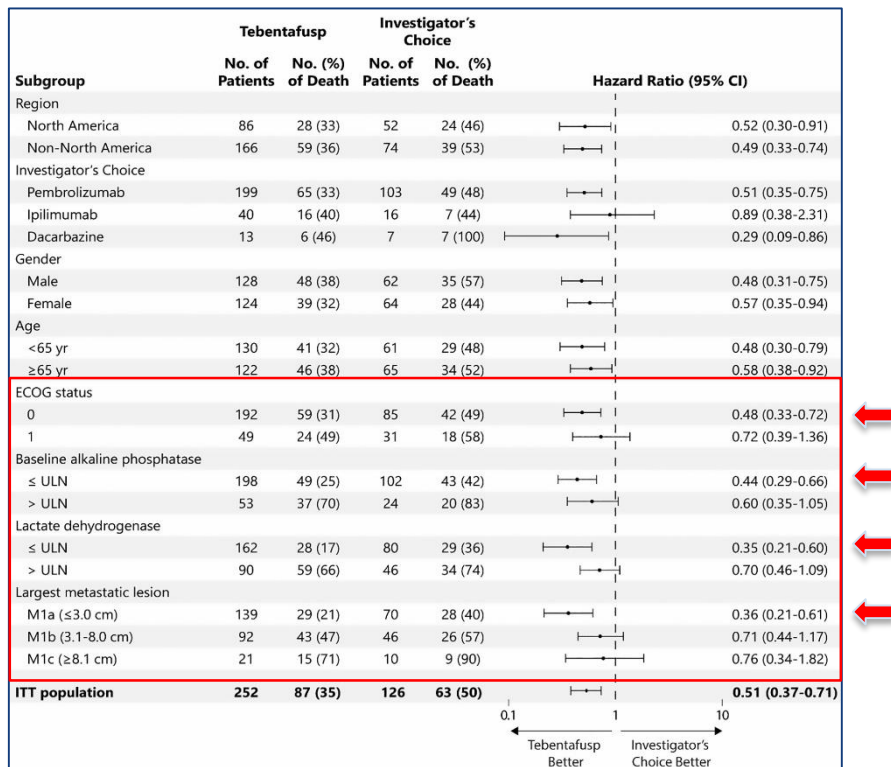
Tebentafusp

Phase 3: IMCgp100-202



Nathan, Hassel et al, NEJM 2021

Who benefits most?

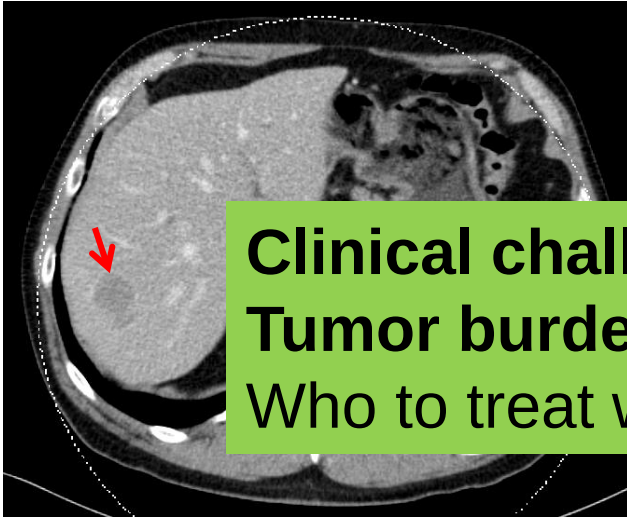


Uveal melanoma patients with:

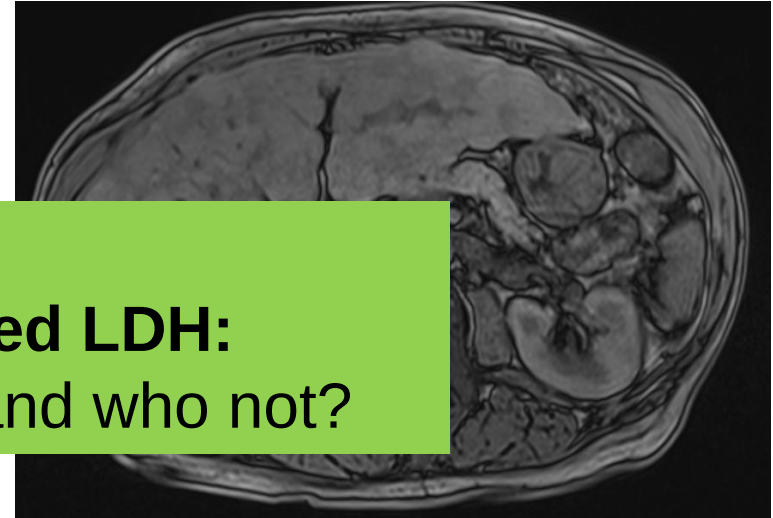
- ECOG 0
- Normal LDH
- M1a stage

➔ Low tumor burden!

Who to treat with tebentafusp



Clinical challenge:
Tumor burden - Elevated LDH:
Who to treat with Tebe and who not?



- Low tumor burden
- Normal LDH at inclusion
- Low dynamic of disease
- firstline

> 3 years with Tebe

- High tumor burden
- High LDH at inclusion
- High dynamic of disease
- $\geq$ 2ndline

died 4 months after Tebe initiation

Low response rate with Tebentafusp

ORR and DCR by investigator assessment

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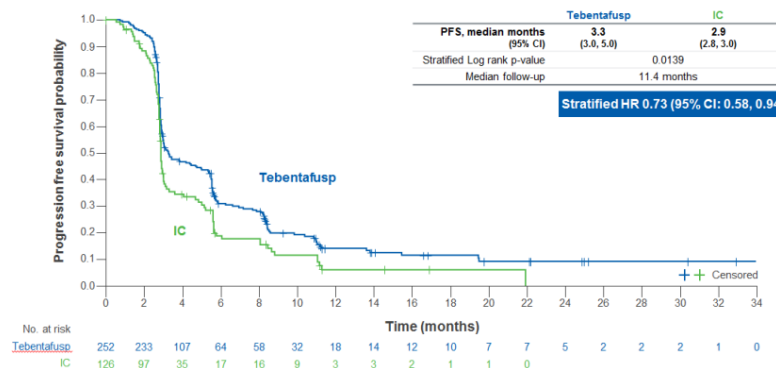
Best response to systemic therapy	<u>Tebentafusp</u> (n=252) n (%)	IC (n=126) n (%)
ORR*	23 (9) ←	6 (5)
CR	1 (0.4)	0
PR	22 (9)	6 (5)
SD	92 (37)	28 (22)
PD	131 (52) ←	78 (62)
Non-evaluable/not applicable	6 (2)	10 (8)
DCR ≥12 wks[†]	115 (46)	34 (27)

*Defined as CR, PR or SD for ≥12 weeks.
CR, complete response; DCR, disease control rate; PD, progressive disease; PR, partial response; SD, stable disease.

AACR ANNUAL MEETING 2021: APRIL 10-15, 2021 AND MAY 17-21, 2021

Progression free survival

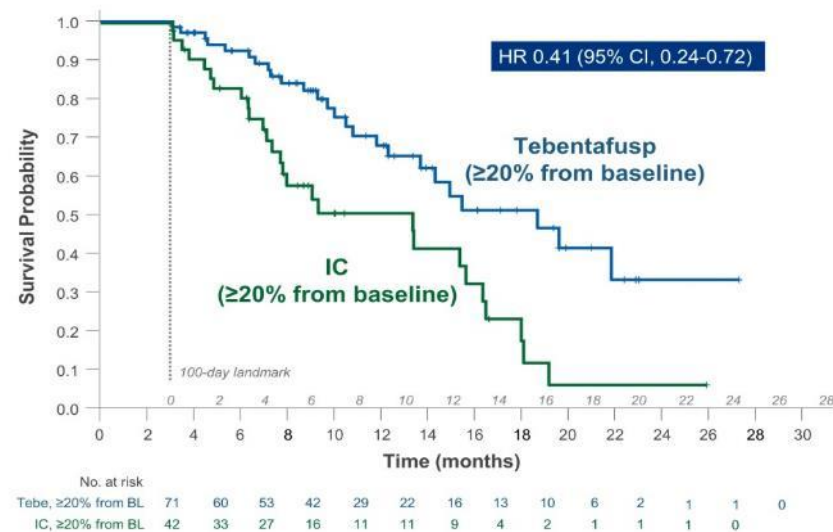
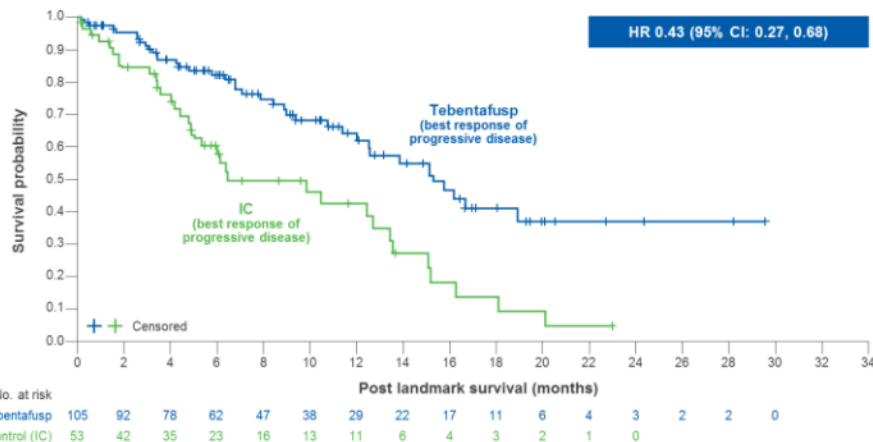
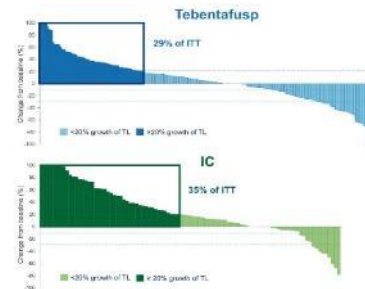
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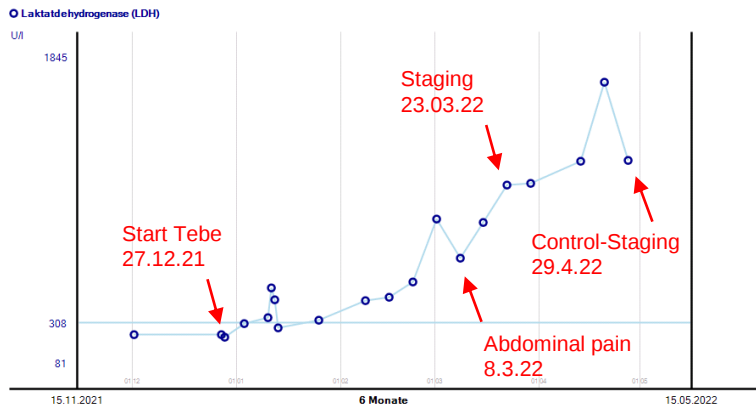
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Overall survival in patients with PD as best response

TBP in 53% Tebe and 17% Pembro patients

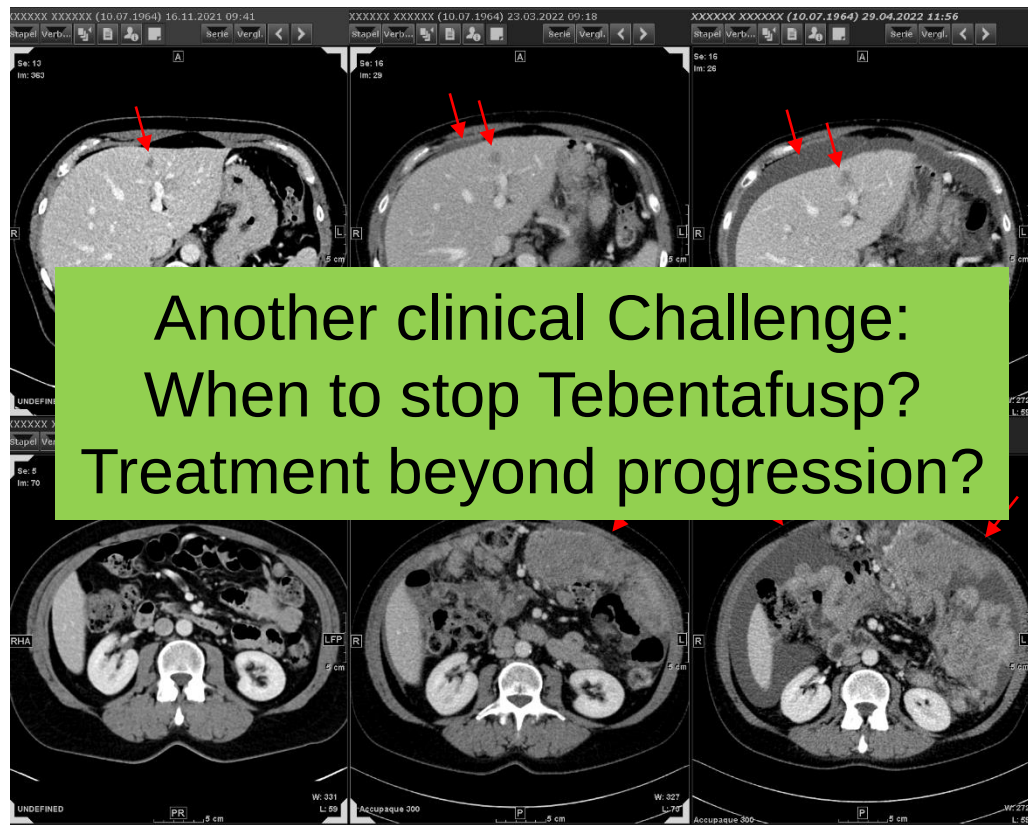


Clinical Case: 57yo female



- Start Tebe after 3x PHP
- Constantly rising LDH
- Good clinical condition
- Staging: stable liver mets, increasing peritoneal mets

16.11.21, LDH 243U/l 23.3.22, LDH 1100U/l 29.4.22, LDH 1200U/l



Liquid biopsy?

IMC-100-102

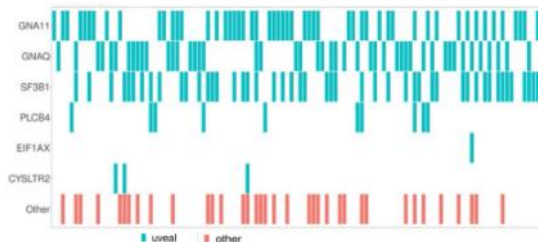
Eligibility

- Metastatic UM
- HLA-A*02:01 positive
- ≥ 1 prior therapy in metastatic setting
- Measurable disease

Treatment Regimen

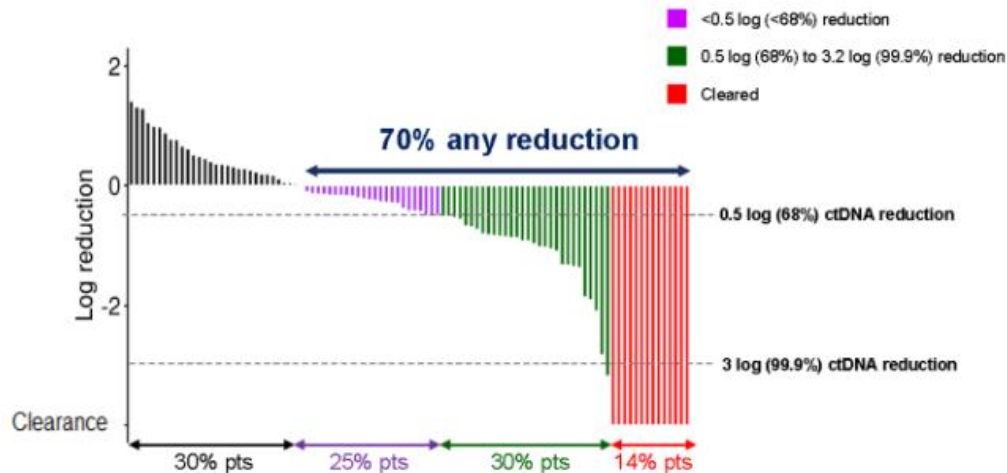


Almost all evaluable patients had mutations in known UM oncogenes



New custom panel to detect melanoma ctDNA using multiplex PCR followed by next gen sequencing, including UM specific: GNAQ, GNA11, SF3B1, PLCB4, CYSLTR2, EIF1AX

1. Yanikyan M, et al. *NEJM* 2017; 377(25): 2500-2501



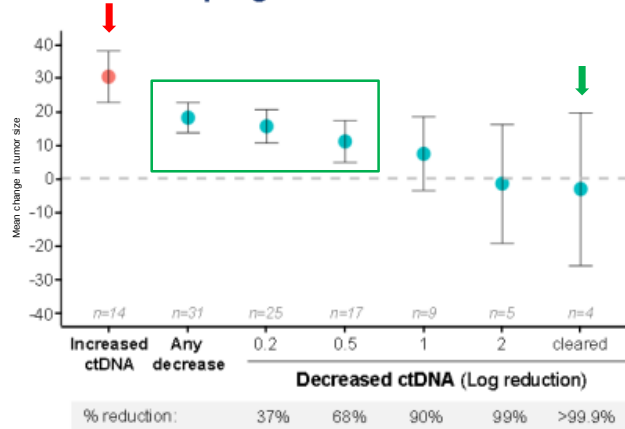
2021 ESMO congress

Shoushtari et al, #3600, ESMO 2021



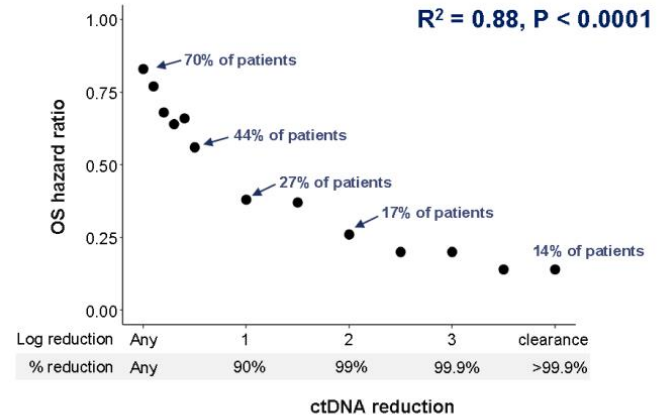
ctDNA better than RECIST?

Patients with best response of progressive disease

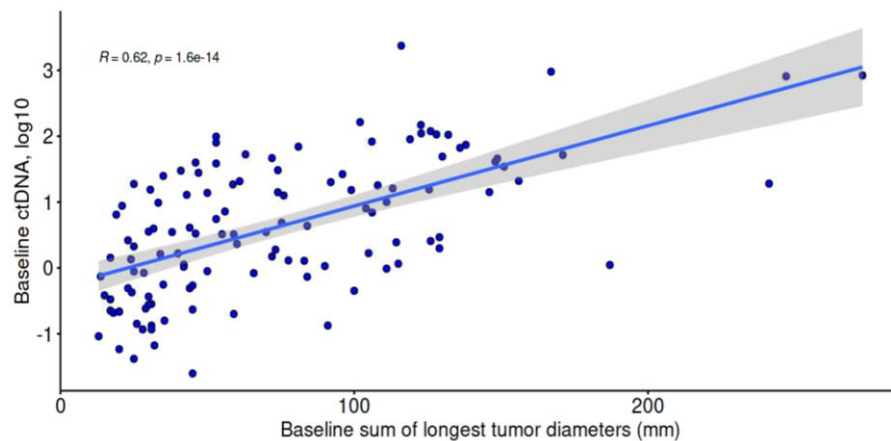


ctDNA reduction associated with less tumor growth

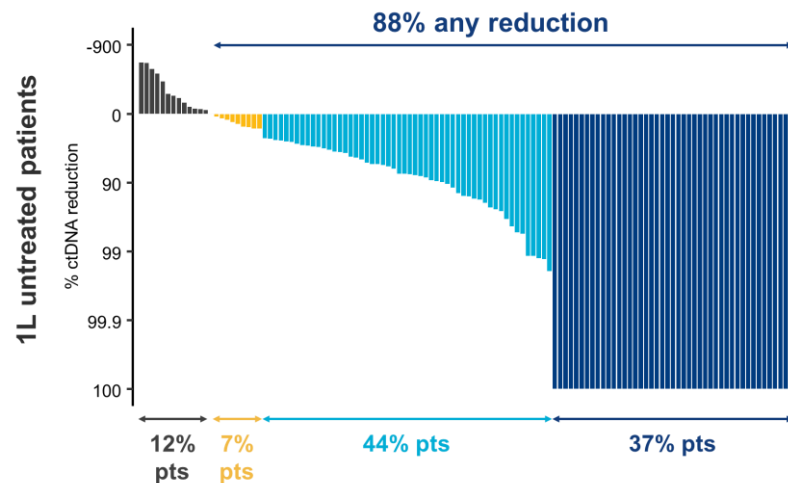
Linear correlation between ctDNA reduction and better OS



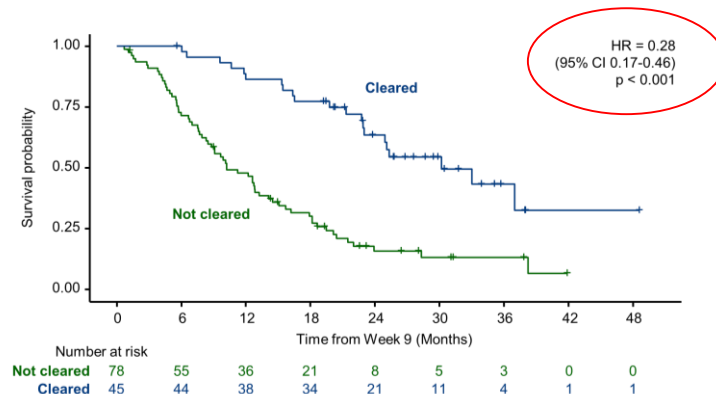
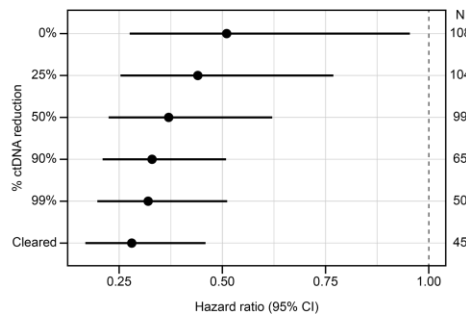
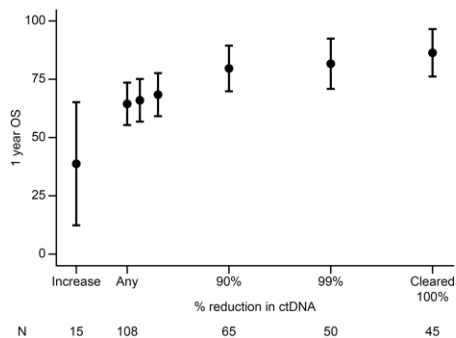
Similar results from phase 3 firstline trial



ctDNA levels at baseline correlated with tumor burden



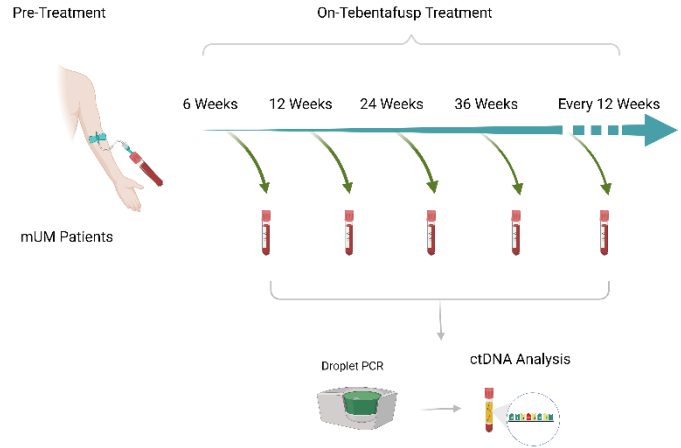
Clearance of ctDNA correlated with OS



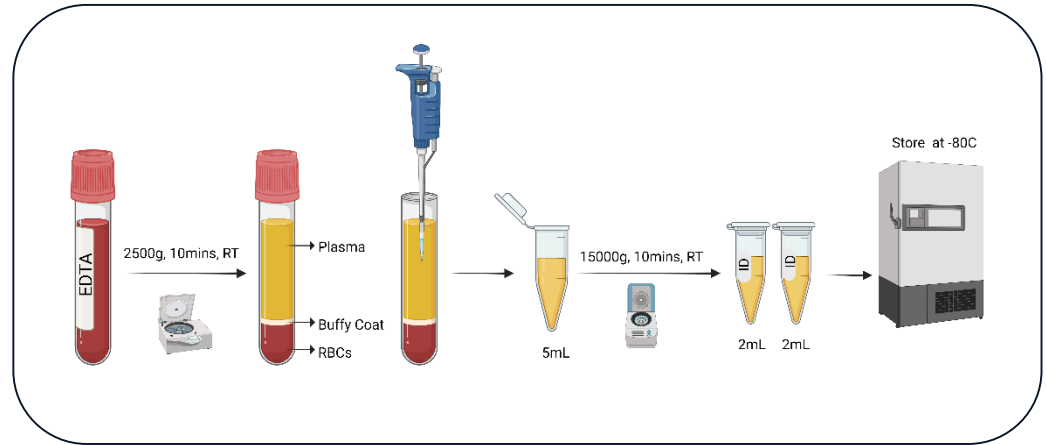


Monitoring ctDNA in mUM Patients Under Tebentafusp

Sample Collection Time Points



Blood Processing for Plasma ctDNA Analysis



Conclusions

- Liver-directed therapies without proven OS benefit
- Tebentafusp as first systemic agent with proven OS benefit
- Patients with low tumor burden (ECOG 0, normal LDH, M1a stage) benefit more
- However, also patients with high tumor burden benefit compared to IC
- Difficult clinical decision who not to treat with Tebe if HLA-0201+
- Chemosaturation to cut-down liver tumor burden?
- Difficult clinical decision who not to treat beyond progression (LDH? Liquid biopsy?)
- How to get liquid biopsy into clinical routine?

DECOG comittee
„Uveal melanoma“

Thank you!



Guy Ungerechts



Team Section DermatOncology

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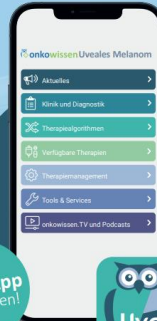
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ctDNA and minimal residual disease in UM

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UK

CONFLICT OF INTEREST STATEMENT

In the past two years I have received funding for advisory boards and/or speakers bureau from the following sources:

AZ

BMS

Esai

Ideaya

Immunocore

Ipsen

Medicenna

MSD

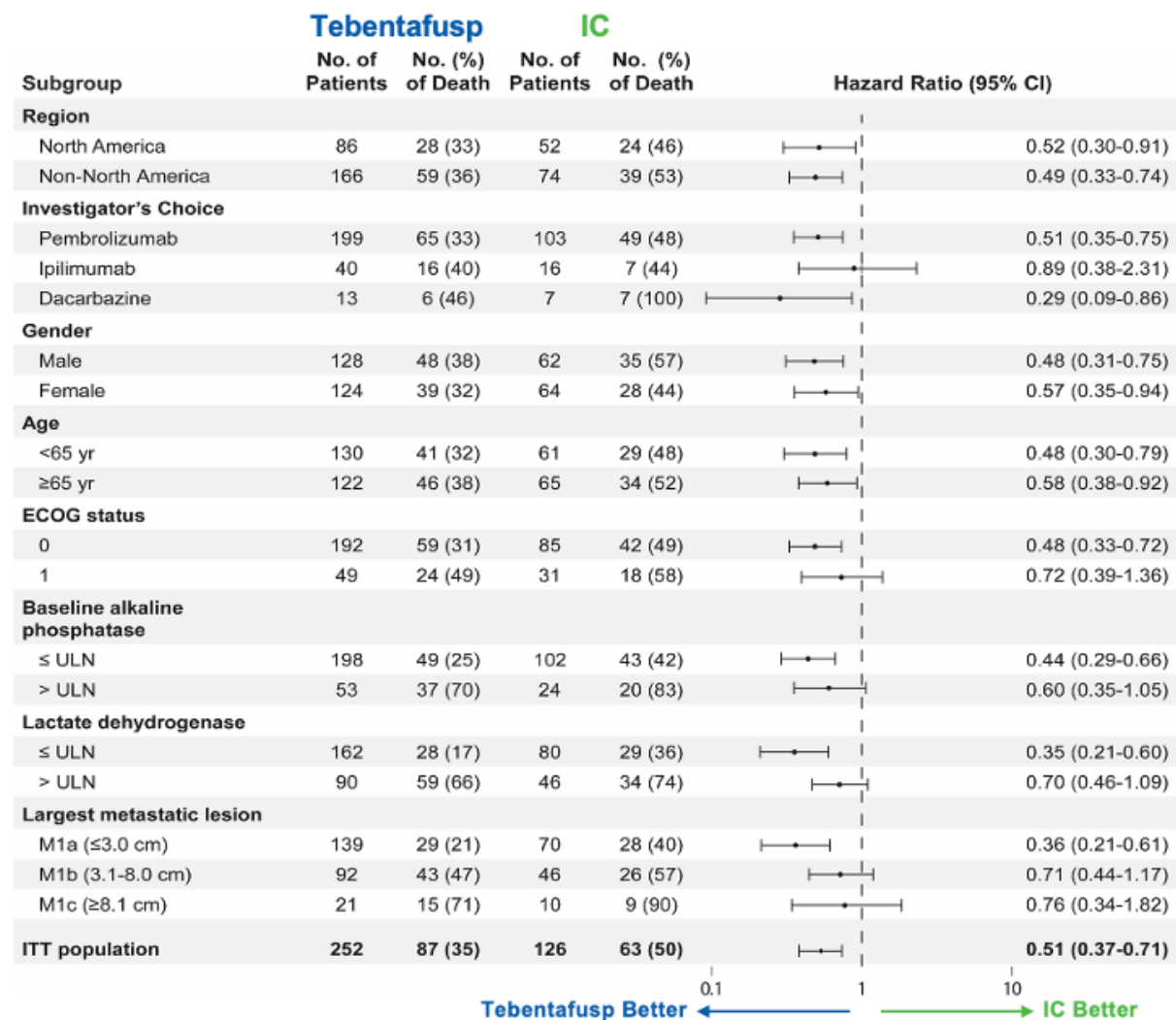
Merck

Novartis

Pfizer

4SC

IMCgp100-202 Pre-planned subset analysis

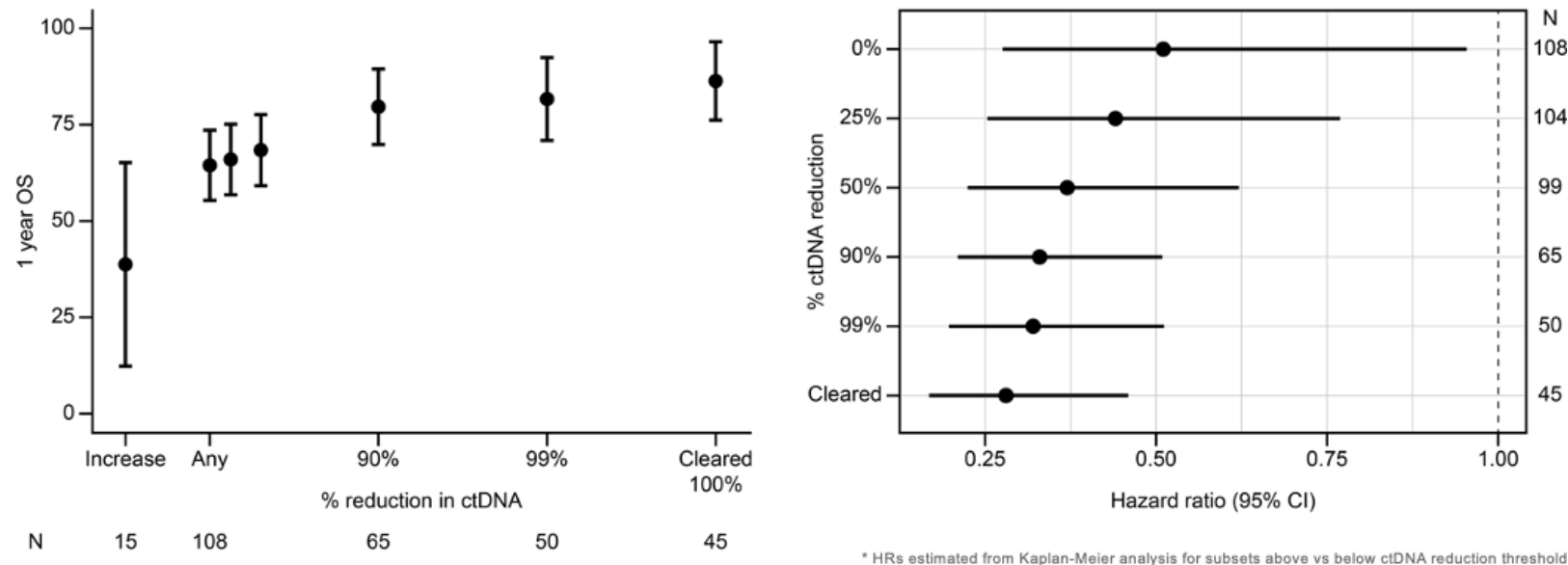


Early ctDNA reduction is associated with better overall survival in the Phase 3 trial of tebentafusp in previously untreated metastatic uveal melanoma

Ryan Sullivan¹, Laura Collins², Manuel Rodrigues³, Paul Nathan⁴, Jessica C. Hassel⁵, Reinhard Dummer⁶, John M. Kirkwood⁷, Piotr Rutkowski⁸, Jean-Francois Baurain⁹, Marcus O. Butler¹⁰, Max Schlaak¹¹, Sebastian Ochsenreither¹², Shaad E. Abdullah¹³, Chris Holland¹³, Koustubh Ranade¹³, Alexander N. Shoushtari¹⁴

¹Massachusetts General Hospital, Boston, MA, USA; ²Immunocore, Abingdon-on-Thames, UK; ³Institut Curie, Paris, France; ⁴Mount Vernon Cancer Centre – East and North Herts NHS Trust, Northwood, UK; ⁵University Hospital Heidelberg, Heidelberg, Germany; ⁶University of Zurich Hospital, Zurich, Switzerland; ⁷University of Pittsburgh Medical Center, Pittsburgh, PA, USA; ⁸Maria Skłodowska-Curie National Research Institute of Oncology, Warsaw, Poland; ⁹Institut Roi Albert II Cliniques Universitaires St-Luc, Université catholique de Louvain, Brussels, Belgium; ¹⁰Princess Margaret Cancer Centre, University of Toronto, Toronto, ON, Canada; ¹¹Charité – Universitätsmedizin Berlin, corporate member of Freie Universität Berlin and Humboldt-Universität zu Berlin, Berlin, Germany; ¹²Cancer Center, and Dept. of Hematology, Oncology and Tumor Immunology, Campus Benjamin Franklin, Charité – Universitätsmedizin Berlin; ¹³Immunocore, Rockville, MD, USA; ¹⁴Memorial Sloan-Kettering Cancer Center, New York, NY, USA

Figure 4. Deeper reductions in ctDNA were associated with progressively longer OS



- Lower baseline ctDNA levels were associated with longer overall survival (< median levels HR 0.43 (95% C.I. 0.28-0.67))
- 79/202 (39%) patients with undetectable ctDNA at baseline had 1-year OS of 90%
- 123/202 (61%) patients with detectable ctDNA at baseline had 1-year OS of 62%
- 15/123 (12%) patients without any reduction in ctDNA on tebentafusp had 1-year OS of 43%



TebeMRD

Full title: A Phase II Non-Randomized, Open-label, Multi-centre Study of the Safety and Efficacy of Tebentafusp in Melanoma with Molecular Relapsed Disease

Short title: Tebentafusp in MRD Melanoma

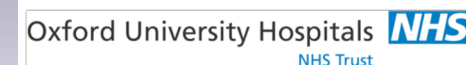
CI Prof Mark Middleton

Trial Management Group:

Prof. Paul Nathan, Dr. Joe Sacco



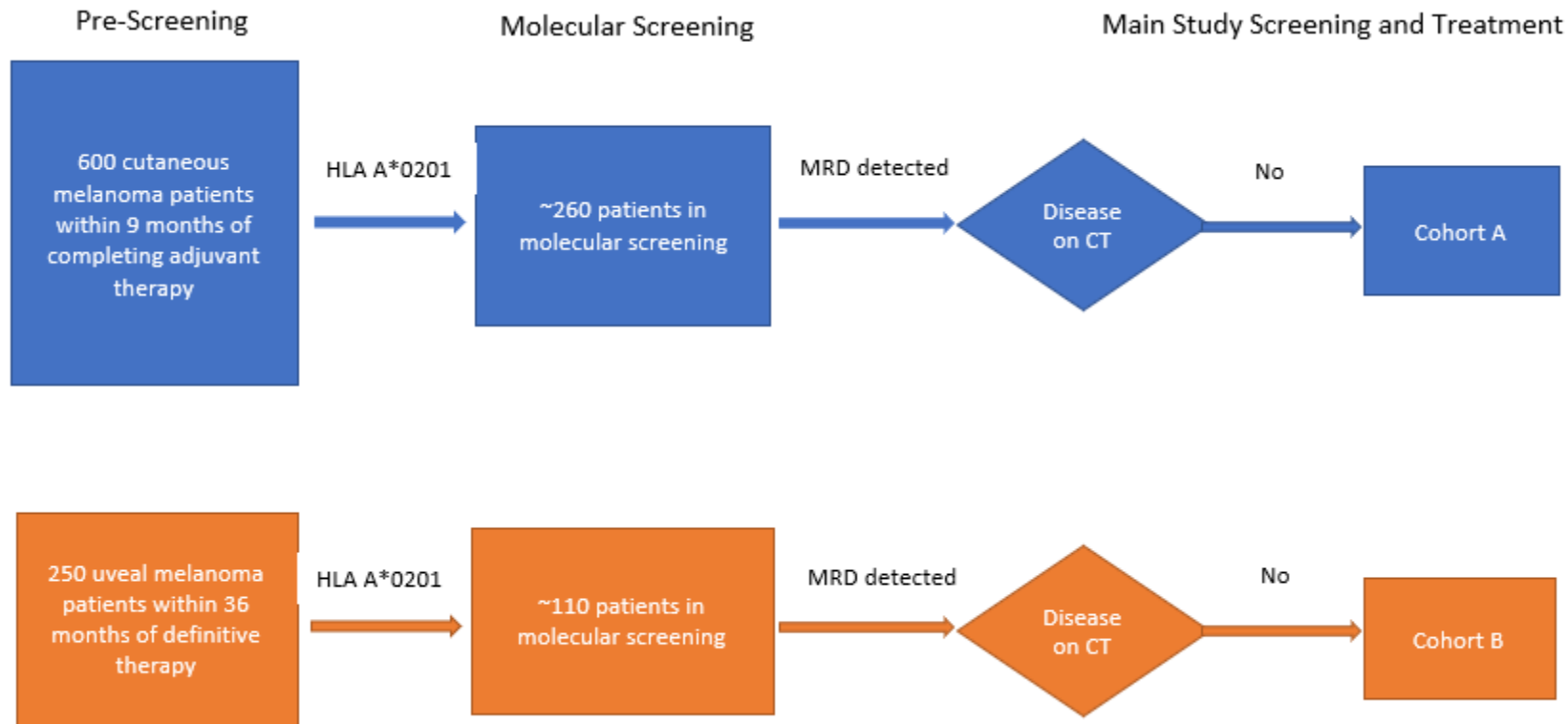
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TebeMRD

- Cutaneous and Uveal Cohorts
- HLA A2.01
- Following treatment of high risk disease with curative intent
- 850 patients screened – 3/12ly ctDNA for 1 year
- 50 patients with detected ctDNA Rx with tebentafusp for up to 6/12

Trial Design – TebeMRD



TebeMRD

- Primary endpoint
 - Molecular response rate – defined as pMR & cMR
- Secondary endpoints
 - Efficacy – RFS at 12 months, duration of MR, OS
 - Safety & tolerability
 - Molecular relapse rate – baseline, 6/12 & 12/12

Summary

- Strong evidence that fall in ctDNA correlates with OS benefit from tebentafusp. Likely superior to radiological endpoints.
- 40% of patients with radiological evidence of disease on 1st line phase III study were ctDNA –ve.
- TebeMRD investigates whether treatment with tebentafusp at the point of ctDNA molecularly defined relapse is associated with greater benefit



1st ELBS Clinical WG Uveal Melanoma Meeting 2023

Uveal Melanoma: Biology and genetics

Christoffer Gebhardt | 26th September 2023

Conflict of Interest/Disclosures

Advisory Roles for, and Honoraria as well as Travel Expenses by the following Companies:

Almirall

Amgen

Beiersdorf

BioNTech

Bristol-Myers Squibb

GlaxoSmithKline

Immunocore

Janssen

MSD Sharp & Dohme

Novartis

Pierre-Fabre

Roche

Sanofi Genzyme

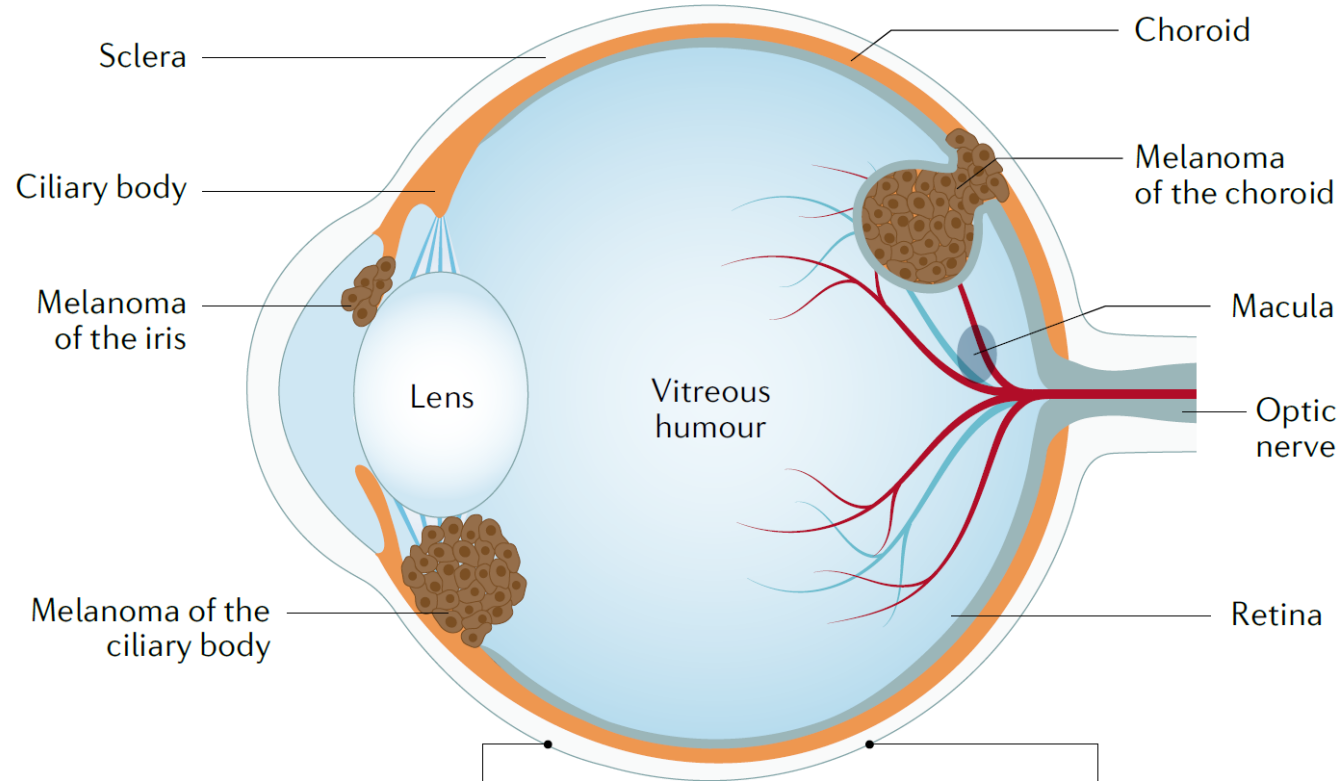
Sciomics

SUN Pharma

Sysmex/Inostix

Co-Founder of Dermagnostix and Dermagnostix R&D

Uveal Melanoma



Risk factors for developing UM

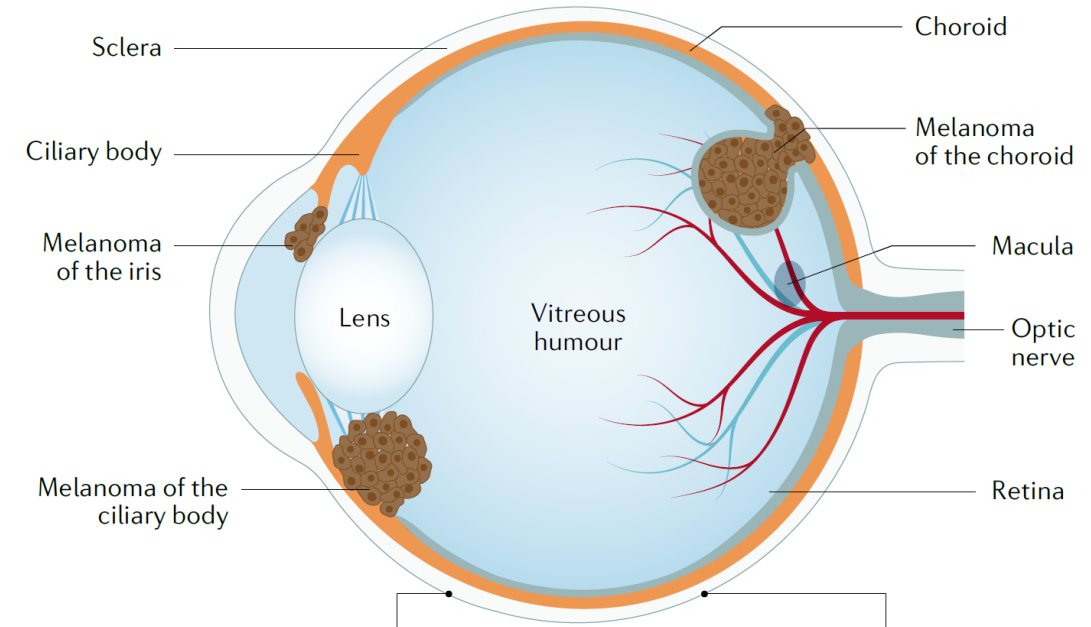
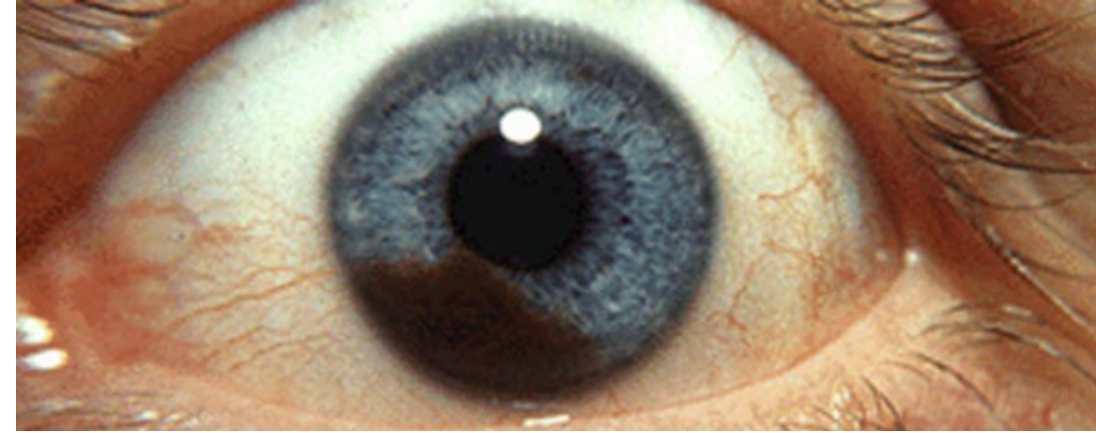
- Age 50–70 years
- Fair skin colour
- Many skin naevi
- Sensitivity to sunburn
- Northern European ancestry
- Light iris colour (blue or grey)
- Congenital ocular melanocytosis
- Melanocytoma
- Family member with cutaneous melanoma
- Family member with uveal melanoma
- Germline mutation in *BAP1*, *MLH1* or *PALB2*

Symptoms of UM

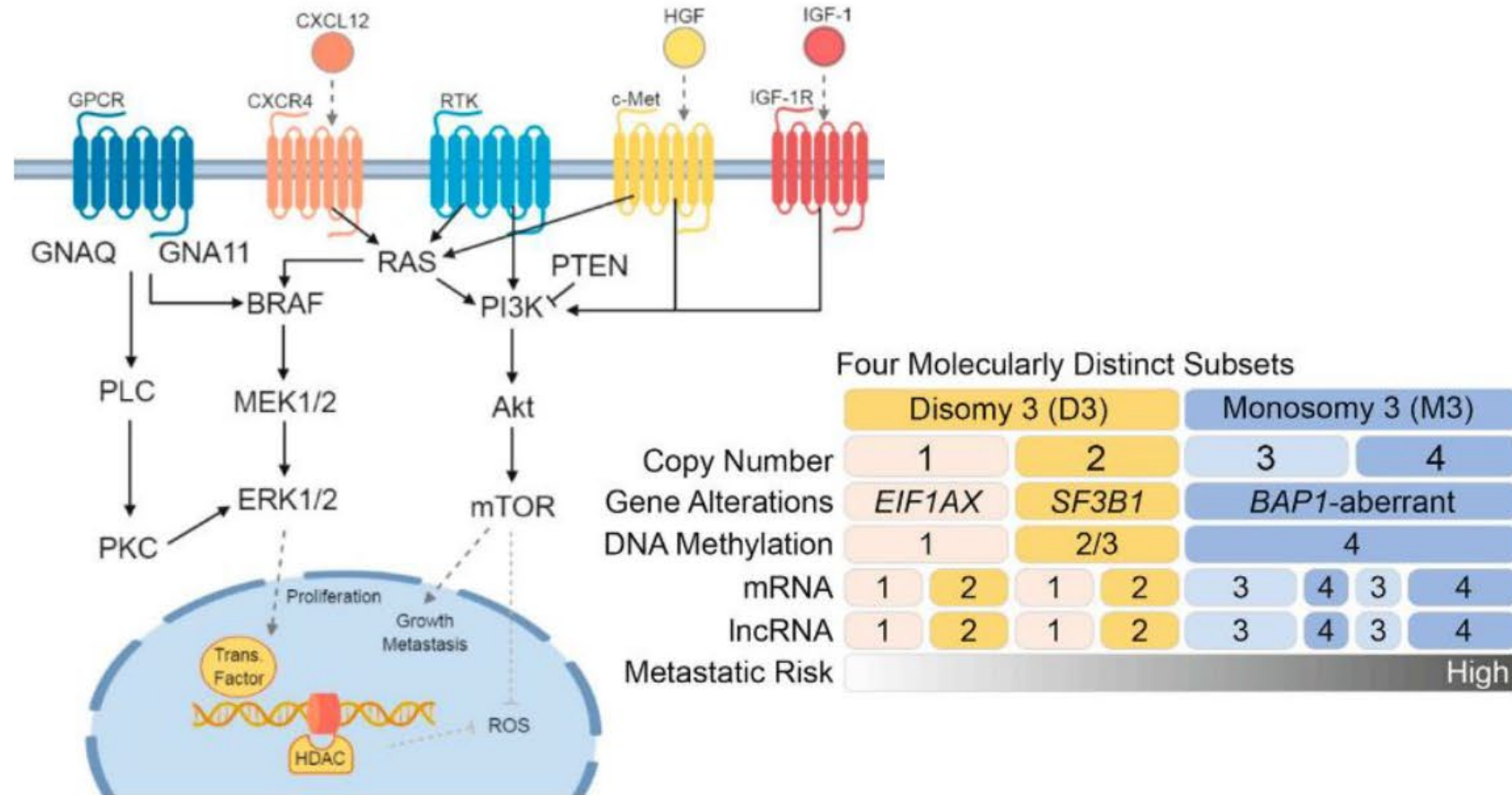
- Blurred or distorted vision
- Visual field loss
- Flashes of light
- Change in iris colour

Uveal Melanoma

- UM arises in the uveal tract:^{1,2}
 - Choroid (85–90%), ciliary body (5–8%) and iris (3–5%)
- The eye is an immune-privileged area, which affects how the immune system interacts with UM³
- **UM** can be considered an immunologically **‘cold’ tumor**, with a low response rate to treatment with ICIs⁴
 - Characterized by point mutations in the G-protein α -subunit coded for by the *GNAQ* and *GNA11* genes⁵
 - *GNAQ/GNA11* mutations have been found in 83–96% of patients⁵⁻⁸
- **CM**, in contrast, has many of the hallmarks of a **‘hot’ tumor** with a high mutational burden and response to ICIs⁴
 - CM has a different mutational profile versus UM, with mutations commonly seen in *BRAF*, *RAS* and *NF1*⁹



Uveal Melanoma has a different tumor biology than cutaneous melanoma



Uveal Melanoma has a different tumor biology than cutaneous melanoma

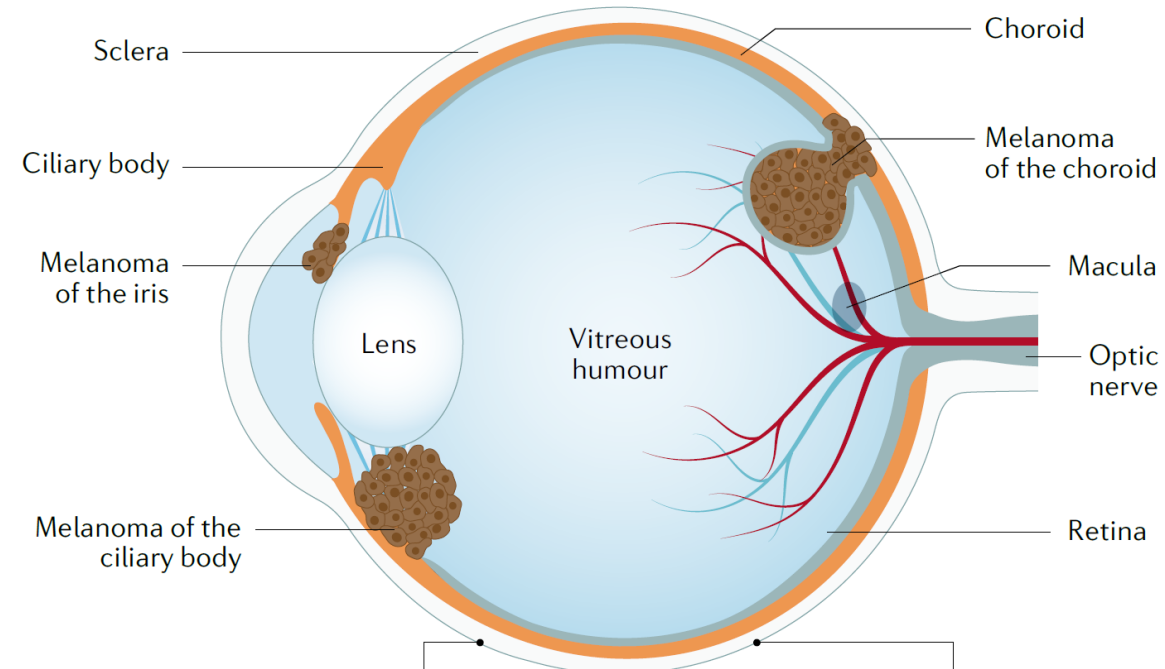
	UM	CM
Incidence	8,000 new cases worldwide/year ¹	230,000 new cases worldwide/year ¹
UV radiation-driven mutation	None ¹	Yes ¹
Familial inheritance	1–2% ¹	~10% ¹
Metastatic pattern	- Up to 50% develop metastases following successful treatment of the primary tumor^{1,2} - Predominantly liver^{1,4} - Hematogenous dissemination^{1,4}	15.5% develop metastases³ - Most common sites (in order): lungs, liver, bones, brain^{1,3,4} - Lymphatic and hematogenous spread^{1,4}
Immunogenicity	- Similar extent of immune cell infiltration in metastatic sites⁵ - Higher ratio in UM of: exhausted CD8+ T cells to cytotoxic T cells, to CD8+ T cells, and to Th1 cells⁵ - Lower infiltration of PD-1-positive lymphocytes in UM metastatic sites⁶ - Lower levels of PD-L1 in UM metastatic sites^{5,6}	
Genetic burden	Low genetic mutational burden ^{1,4}	High genetic mutational burden ^{1,4}
Associated genes	Distribution: ^{4,7–9} BRAF: 0% KIT: 0% GNAQ: ~63% GNA11: ~33% PLCB4:~2.5% CYSLTR2:~4% BAP1: ~60% SF3B1 ~25% EIF1AX: ~15%	Distribution: ^{4,7,10,11} BRAF: ~36% KIT: ~1.7% GNAQ: ~1.4% GNA11: ~1.3% NRAS: ~12% NF1: ~14% CDKN2A
Prognosis	Patients with metastases (mostly liver) ¹² have: - Median survival of 3–30 months^{12–15} 1-year survival rate of ~29–83%^{12,15} 5-year survival rate of <20%¹⁴	Patients with advanced/metastatic CM have: - Median survival of 4–>60 months^{16–20} 1-year survival rate of 36–81%^{16,21,22} 5-year survival rate of 10–70%^{17–20,22}
Responsiveness to immunotherapy	Low response rates to immunotherapy ^{1,4,13,23}	Higher response to ICIs (anti-CTLA4, anti-PD-1, anti-PD-L1) than UM, especially to ICI combination (up to 58% ORR) ^{1,24}
Targeted therapies	None ^{1,25}	Anti-BRAF, anti-MEK ¹

CTLA4: cytotoxic T-lymphocyte antigen 4; ICIs, immune checkpoint inhibitors; ORR, overall response rate; PD-1: programmed death-1; PD-L1: programmed death-ligand 1; UV: ultraviolet.

1. Pandiani C, et al. *Genes Dev* 2017;31:724–43; 2. Dogrusöz M, Jager MJ. *Asia-Pac J Ophthalmol* 2017;6:186–96; 3. Zbytek B, et al. *Expert Rev Dermatol* 2008;3:569–85; 4. Rodrigues M, et al. *Cancers (Basel)* 2019;11(7):1032; 5. Hoefsmit EP, et al. *J Immunother Cancer* 2020;8:e001501; 6. Qin Y, et al. *Oncoimmunol* 2017;6:e1321187; 7. Beadling C, et al. *Clin Cancer Res* 2008;14:6821–8; 8. Piperno-Neumann S, et al. *J Clin Oncol* 2014;32(Suppl.): Abstract 9043; 9. Robertson AG, et al. *Cancer Cell* 2018;33:151; 10. Akbani R, et al. *Cell* 2015;161:1681–96; 10. Shaughnessy M, et al. *J Transl Genet Genom* 2018;2:14; 12. Khoja L, et al. *Ann Oncol* 2019;30:1370–80; 13. Schank TE et al. *Cancers (Basel)* 2019;11:1048; 14. NCCN Guidelines, Version 3.2020 (Jan 15 2021), Uveal Melanoma. Available at: https://www.nccn.org/store/login/login.aspx?ReturnURL=https://www.nccn.org/professionals/physician_gls/pdf/uveal.pdf. Last accessed February 2021; 15. Rantala ES, et al. *Melanoma Res* 2019;29:561–8; 16. Larkin J, et al. *J Clin Oncol* 2018;36:383–90; 17. Larkin J, et al. *N Engl J Med* 2019;381:1535–46; 18. Robert C, et al. *Lancet Oncol* 2019;20:1239–51; 19. Robert C, et al. *J Clin Oncol* 2020;38:3937–46; 20. NCCN guidelines Version 2.2021 (Feb 19 2021), Melanoma: Cutaneous. Available at: https://www.nccn.org/professionals/physician_gls/pdf/cutaneous_melanoma.pdf. Last accessed April 2021 ; 21. Lebbé C, et al. *J Clin Oncol* 2019;37:867–76; 22. Wang Y, et al. *Medicine* 2021;100:4; 23. Pelster MS, et al. *J Clin Oncol* 2021;39:599–607; 24. Leonardi GC, et al. *Int J Oncol* 2020;57:609-18; 25. Croce M et al. *Cancers (Basel)* 2019;11:846.

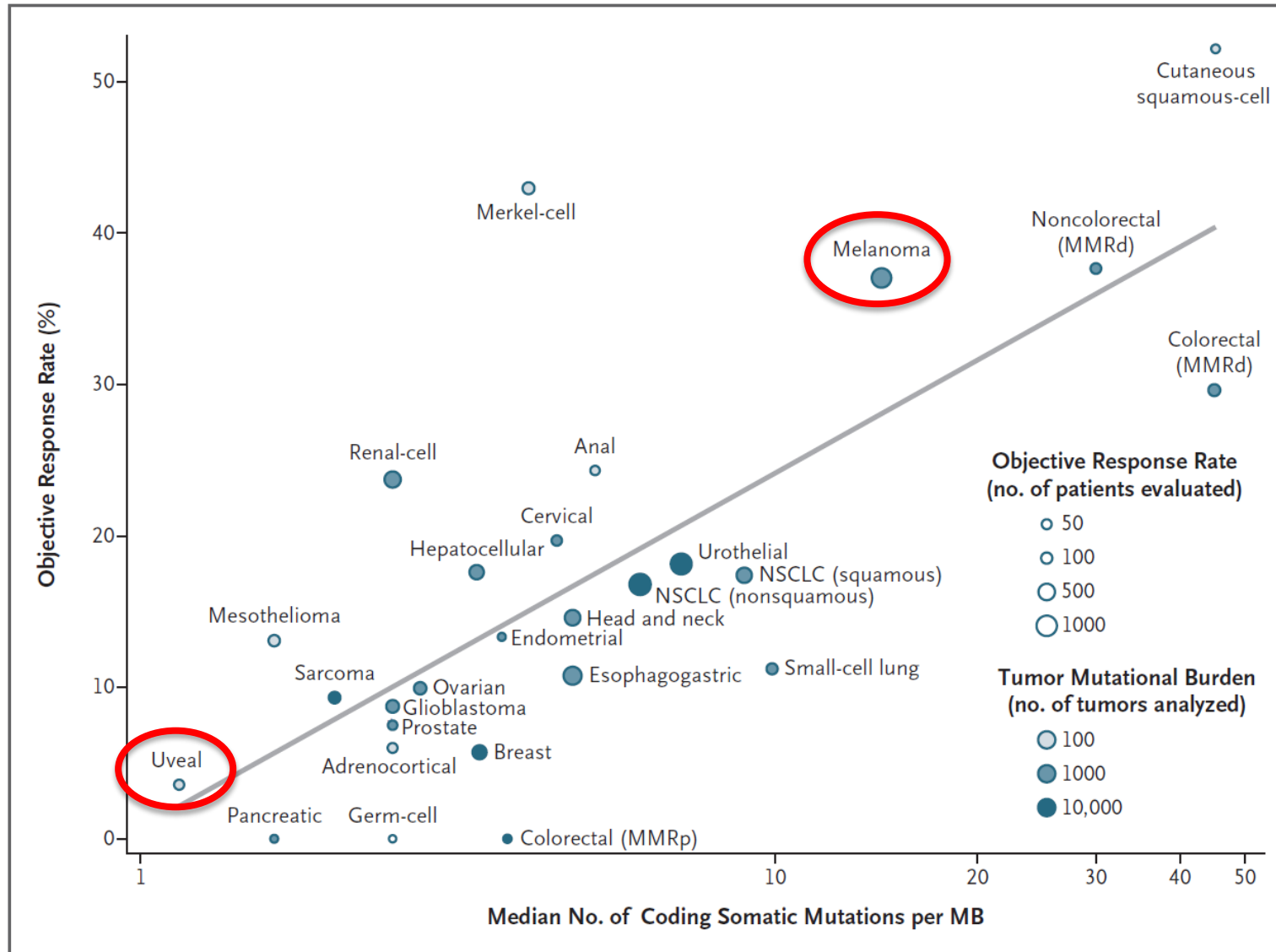
Uveal Melanoma: Treatment resistance in mUM

- **(Liver-)metastasis Resection**
- **Liver-directed therapy** (chemosaturation/ - perfusion, SIRT, TACE): limited RCT data, efficacy difficult to interpret
- **Targeted therapy:** no druggable targets, low efficacy (sorafenib, selumetinib)
- **Systemic chemotherapy:** low efficacy (mono-/polychemotherapy)
- **Systemic Immunotherapy:** ORR up to 17%, 12mo OS: 52% (NIVO+IPI), PD-1 shows similar efficacy



Jager et al., Nat Dis Primer 2020
Piulats, et al. *J Clin Oncol* 2021
Rantala, et al. *Melanoma Res* 2019
Khoja, et al. *Ann Oncol* 2019

Tumor mutational burden predicts response to ICB



UM has one of the lowest TMB of all tumor types, which likely explains the poor observed activity of CPIs in UM relative to their benefit in CM

Molecular prognostic testing to predict metastatic disease and stratify risk

- Identifying a high-risk mUM patient means they can be more aggressively monitored and treated¹⁻³
 - Significant heterogeneity exists in metastatic risk profiling for melanoma¹⁻³
 - Clinical, histopathologic and genetic variables can be assessed, often in combination.^{3,4}
- Most common approaches include:

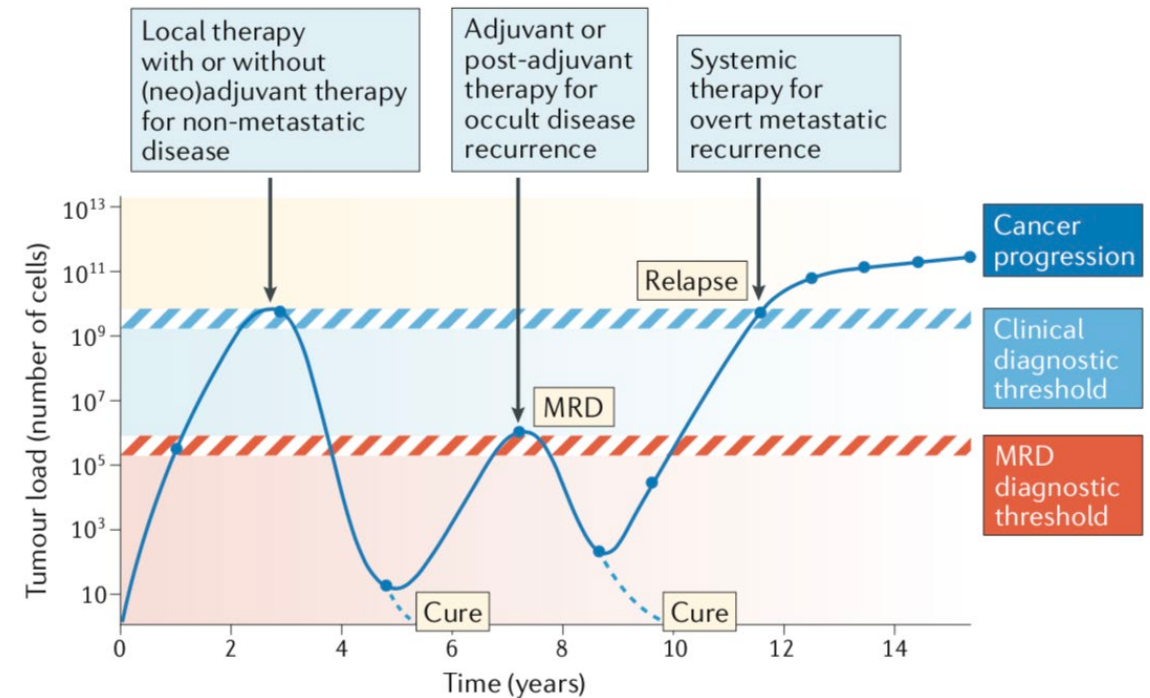
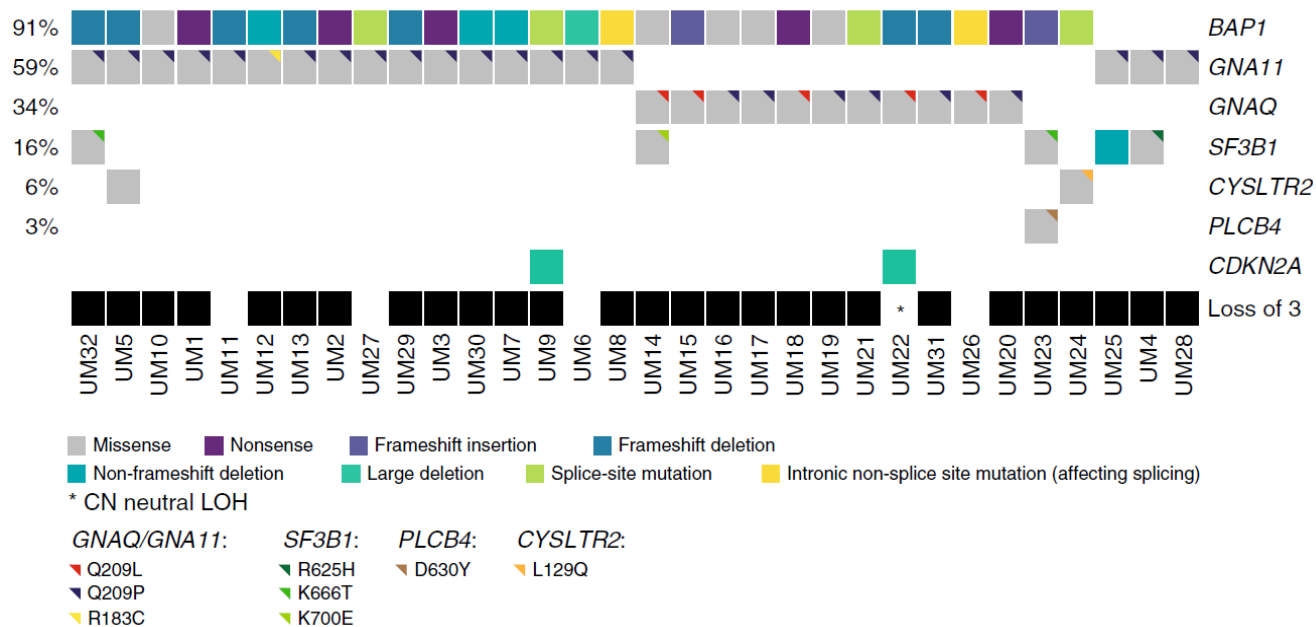
PRAME: An antigen recognized by cytolytic T lymphocytes and an independent prognostic UM biomarker. PRAME mRNA expression analysis identifies increased metastatic risk in patients with Stage I or disomy 3 tumors^{1,5}

DecisionDx™ – Melanoma (Castle Biosciences): A genetic expression profiling tool to analyze 31 genes within Stage I or II melanoma tumors to assess metastatic risk^{2,5}

LUMPO system (Liverpool Ocular Oncology Centre): Combines clinical, histologic and genetic data to enhance reliability of prediction of metastatic disease, while minimizing bias from missing data and competing risks. This system also estimates survival time³

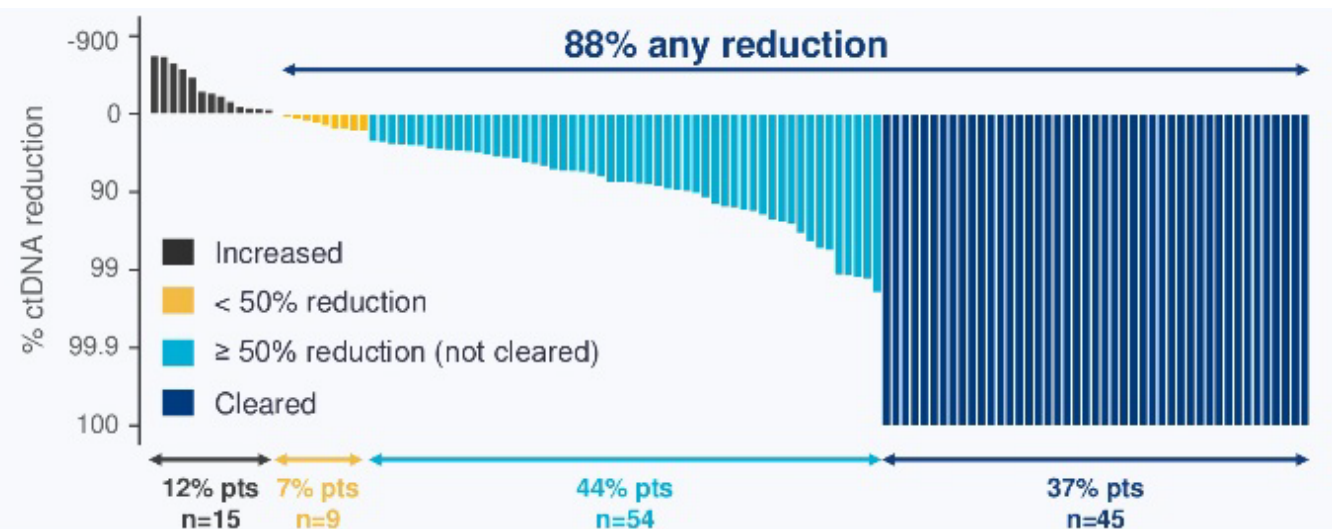
Therapy monitoring using Liquid Biopsy (ctDNA) in Uveal Melanoma

Driver mutations in UM

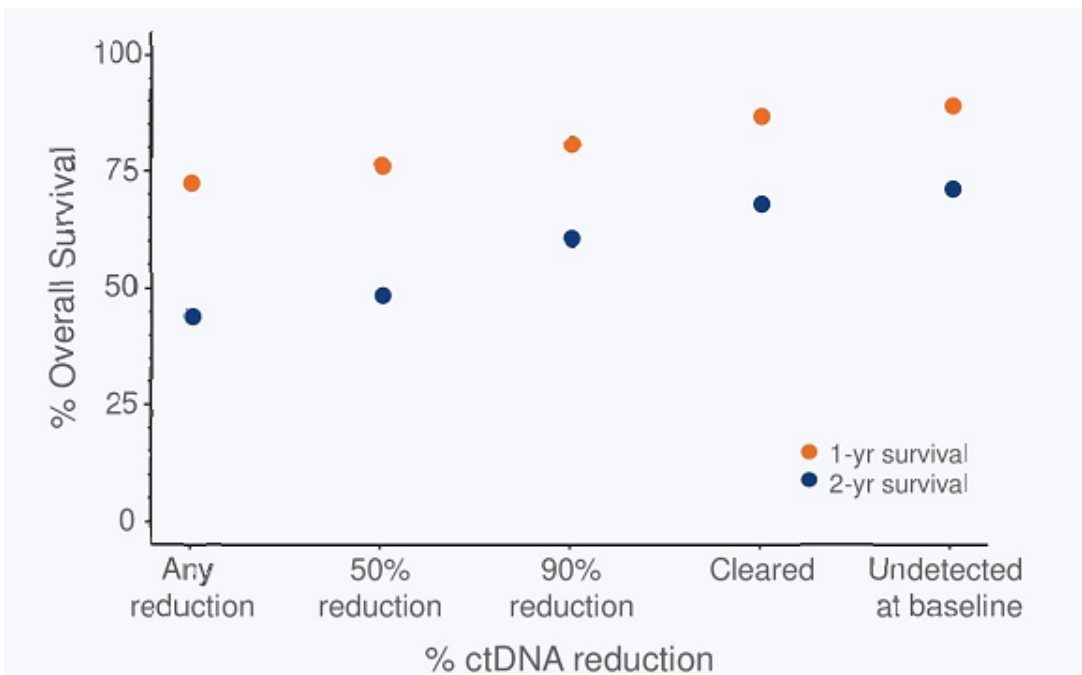


ctDNA dynamics as predictive marker for OS under tebentafusp

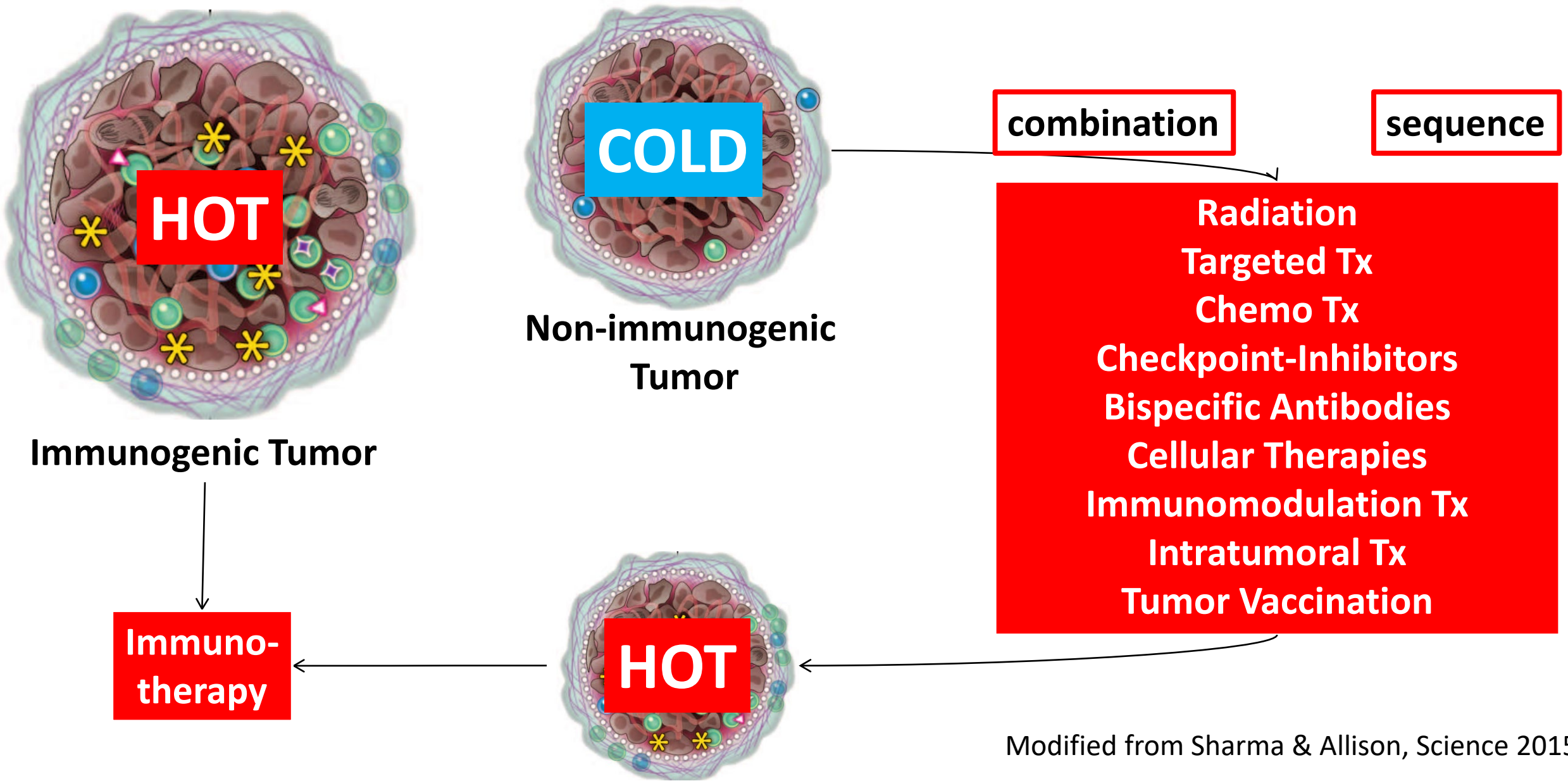
ctDNA reduction (baseline – week 9)
in 88% of all tebentafusp-treated patients



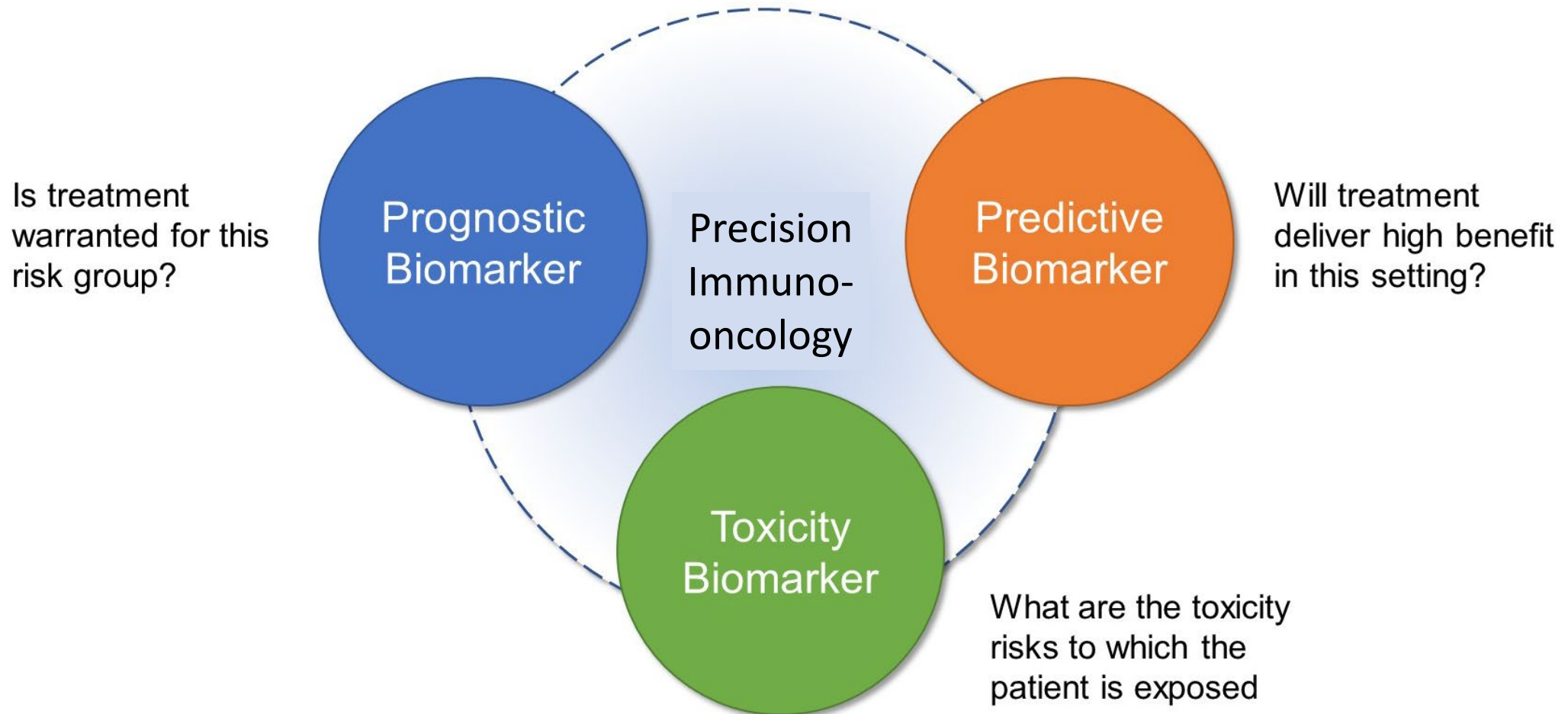
Deep ctDNA reduction (baseline – week 9)
correlates with prolonged OS



Overcoming resistance to immuno-therapy



Towards precision Immuno-oncology in mUM



Trial designs using ctDNA levels for therapeutic decisions

Prognosis

Identifying patients at risk of metastasis

Early Detection of Metastasis/MRD

Follow Up

Therapy selection

Prediction of resistance

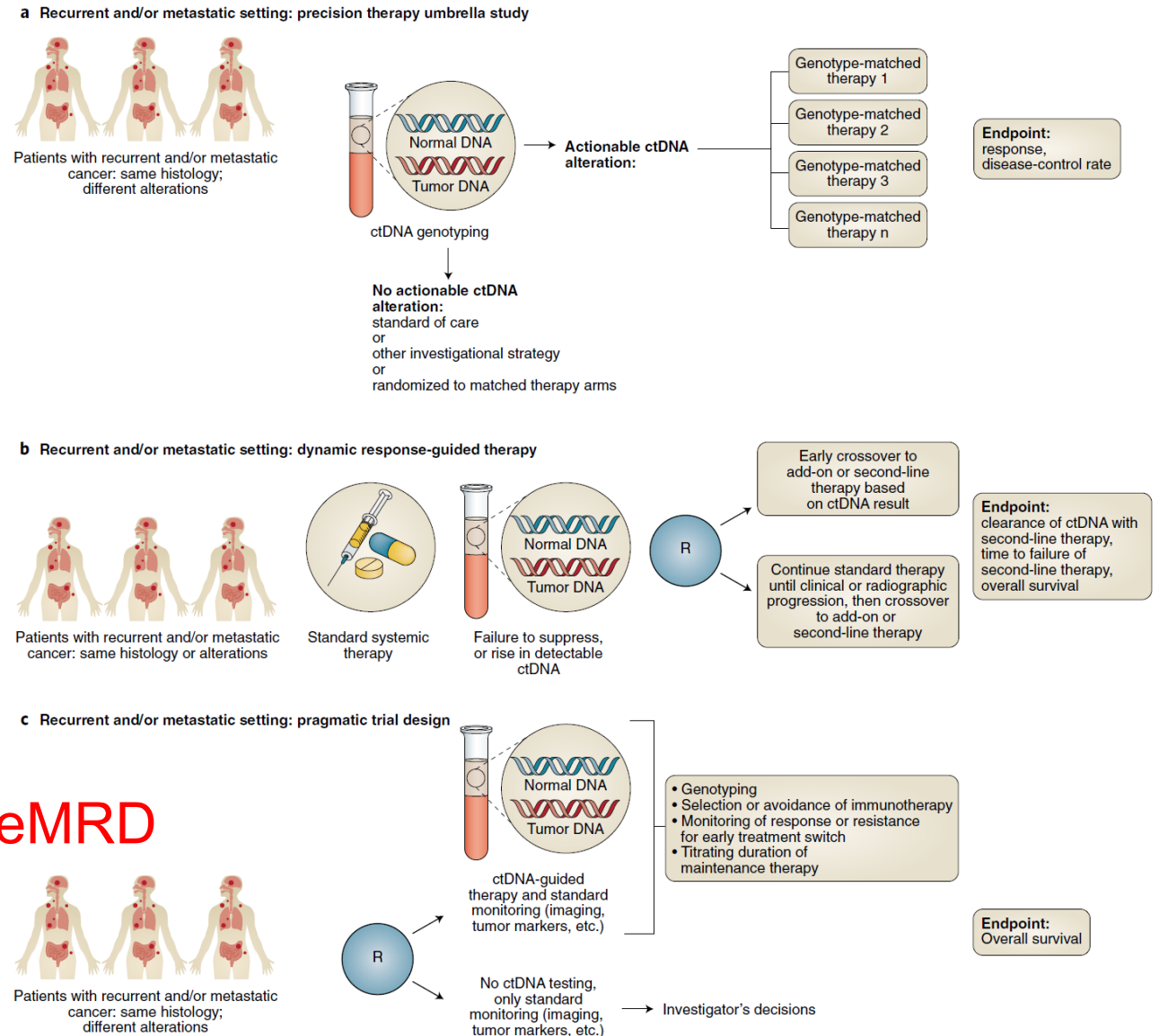
1L: Tebe OR IPI/NIVO

Therapy monitoring

Treatment beyond progression?

Identifying resistance

TebeMRD





ctDNA @Curie

Manuel Rodrigues, MD PhD
Shufang Renault, PhD
Marc-Henri Stern, MD PhD

Institut Curie
Paris



- ~320 new primary tumor patients
- ~70 new metastatic patients per year
- 108 patients treated with tebentafusp (early access)

ENSEMBLE, PRENONS LE
CANCER DE VITESSE

Proof of concept study: ctDNA & liver surgery (« ctDNA R0 » cohort)

53 patients
2-y trial
longitudinal ctDNA assessment for monitoring patients undergoing liver surgery.

ddPCR

GNAQ Q209P / Q209L / Q209R
GNA11 Q209L
GNAQ R183Q
GNA11 R183C

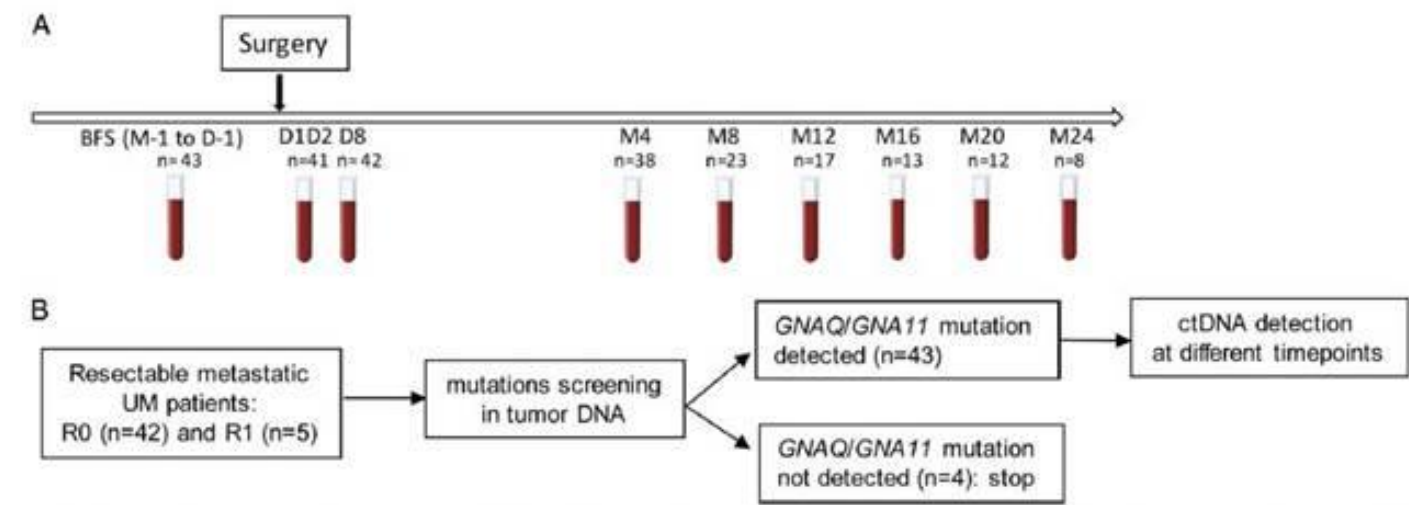
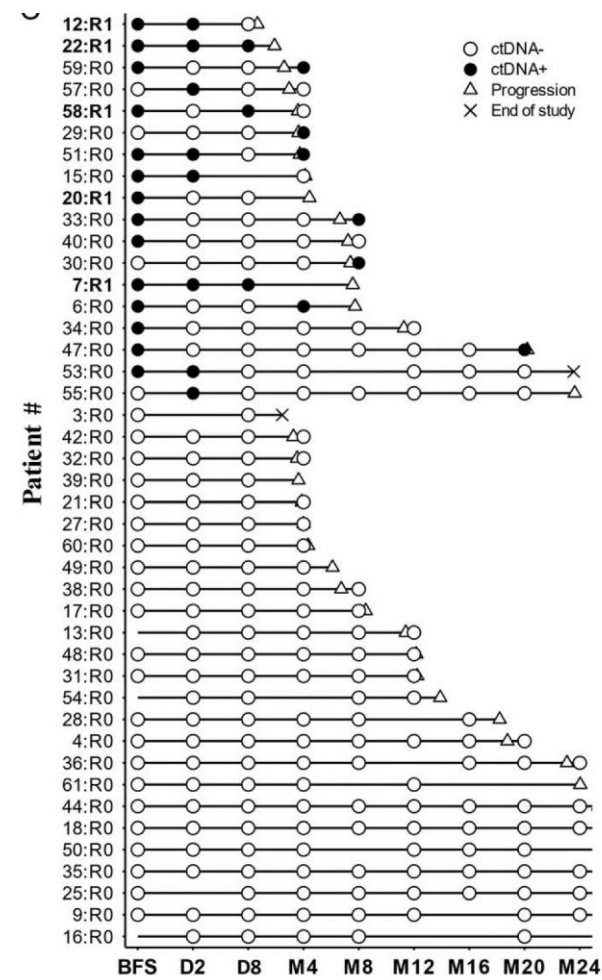
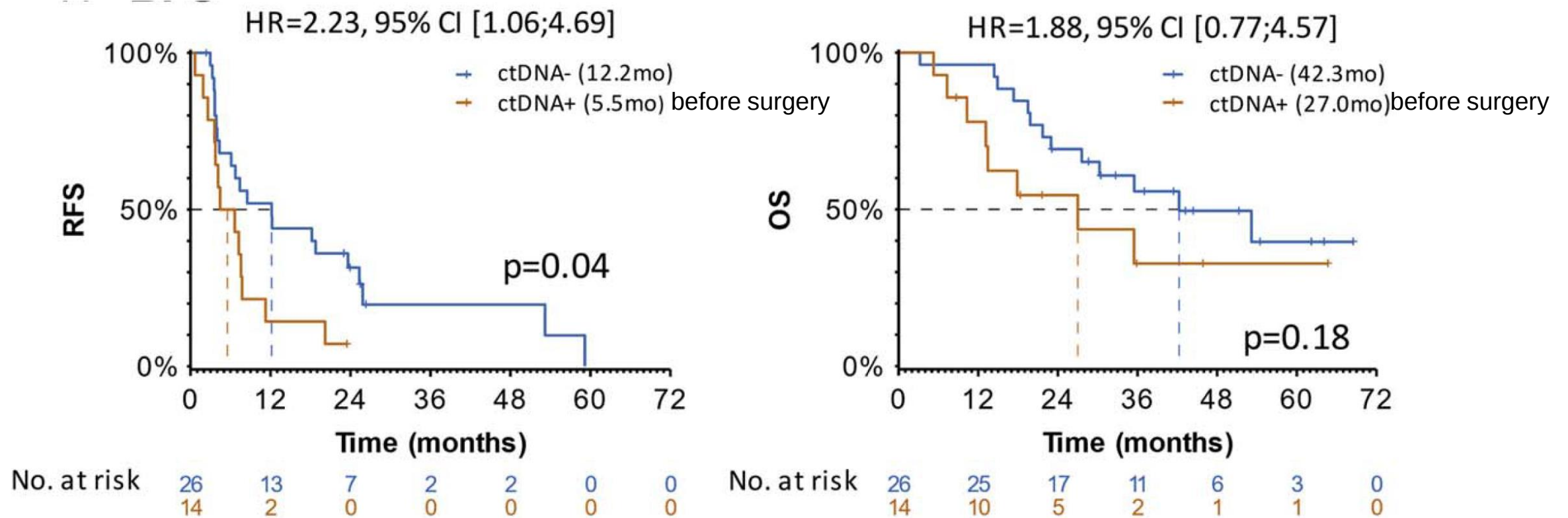


FIGURE 1. Blood sampling (A) and workflow of the study (B). BFS indicates before surgery; R0, microscopically complete resection; R1, microscopically incomplete resection.



Pascale Mariani
Shufang Renault
Aurore Rampanou
Marc-Henri Stern

Proof of concept study: ctDNA & liver surgery (« ctDNA R0 » cohort)



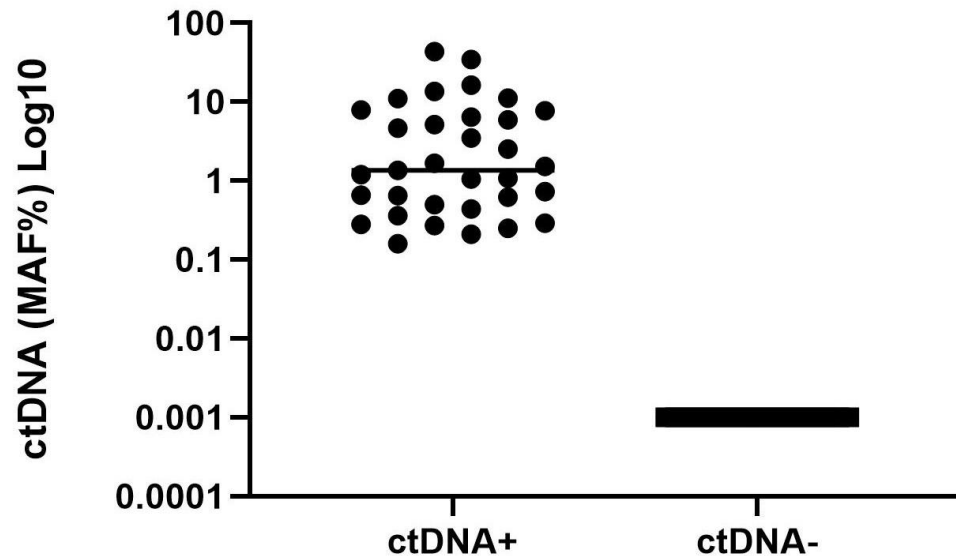
→ Now in routine to select patients for liver surgery

Proof of concept study: ctDNA & tebentafusp (« ALCINA » cohort)

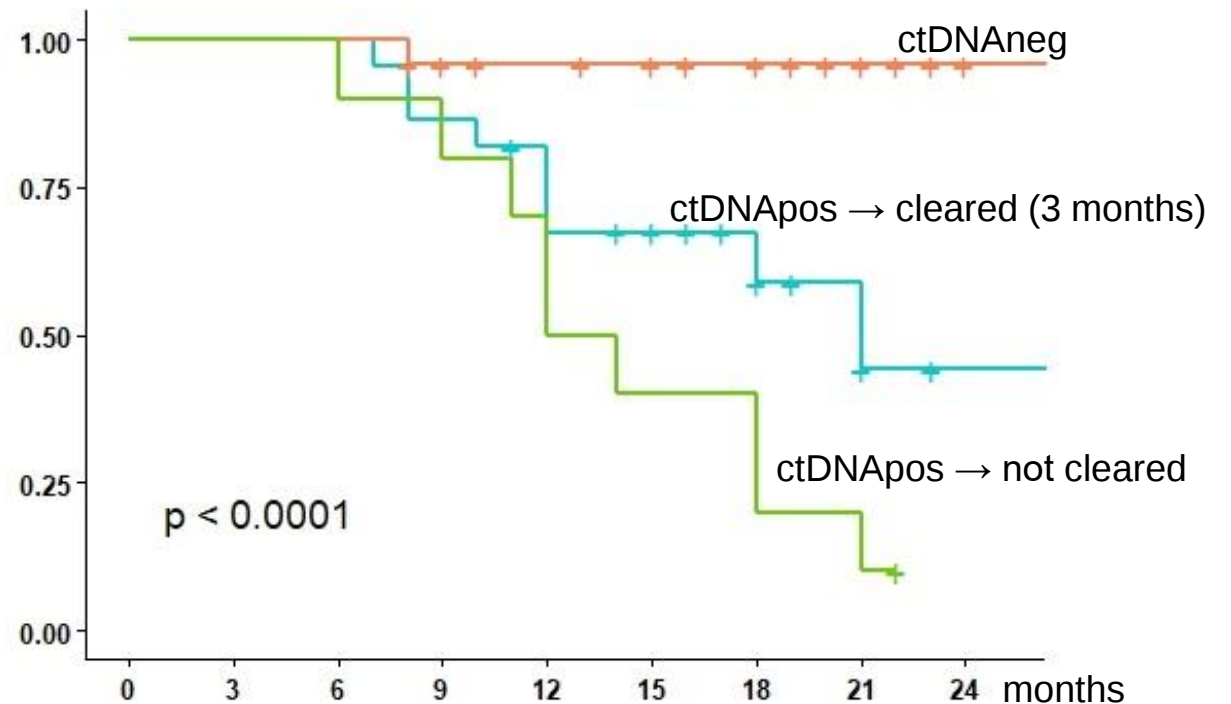
63 pts treated with tebentafusp in mUM with *GNAQ/GNA11* mutations

“real-life” study → smaller metastases (not RECIST)

Pts with detectable ctDNA at baseline = 62% pts

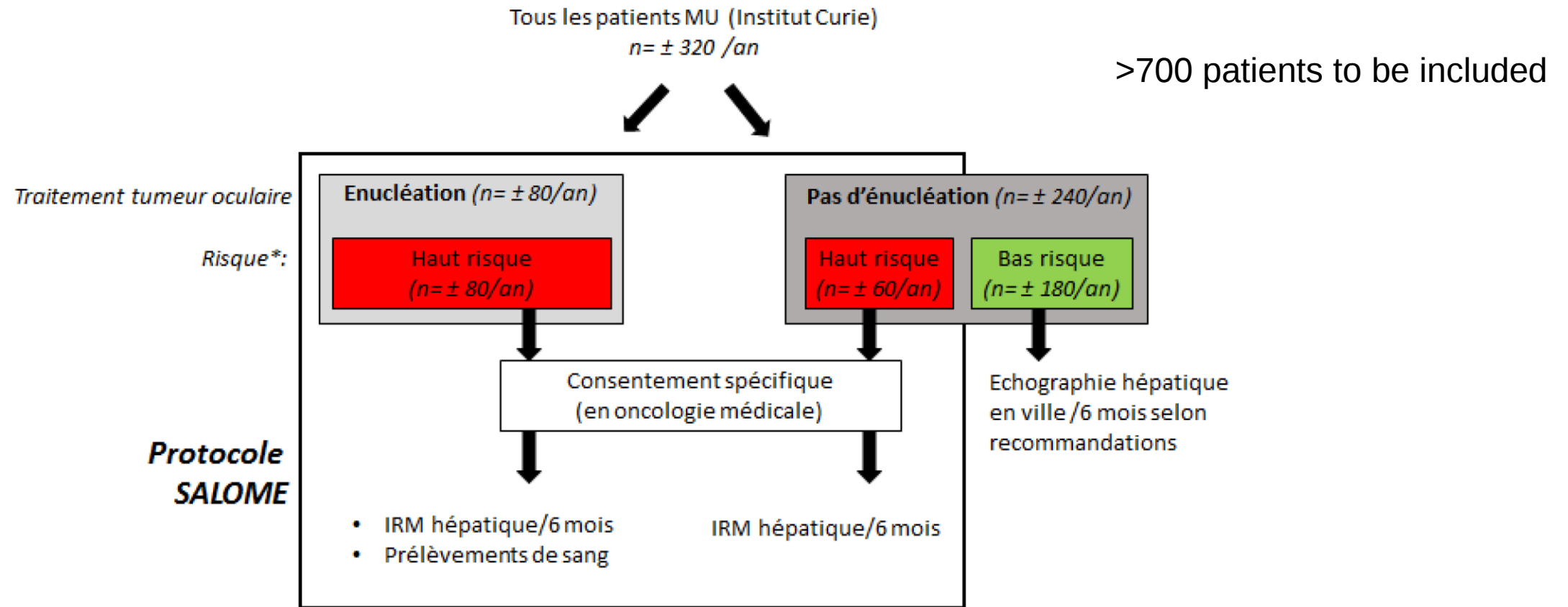


OS according to ctDNA at baseline



→ Now in routine to assess tebentafusp efficacy
→ Next: agnostic ctDNA

Ongoing studies – « SALOME » cohort (adjuvant setting)



* Définition du risque:

- **Haut risque** : T2b/c/d ou $\geq T3$ (AJCC Ophthalmic Oncology Task Force. International validation of the American Joint Committee on Cancer's 7th edition classification of uveal melanoma. JAMA Ophthalmol. 2015; 133: 376–383) OU anomalie du chromosome 3 et/ou du chromosome 8.
- **Bas risque** : tous les autres cas.
- Les patients dont le risque génomique est inconnu seront suivis selon leur risque clinique.
- Les patients énucléés sont à haut risque >95% et seront par définition suivis comme des patients à haut risque.

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 - Aurore Rampanou

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 - Emmanuelle Jeannot

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- Dr Manuel Rodrigues (medical oncology)
- Dr Sophie Piperno-Neumann (medical oncology)
- Dr Vincent Servois (radiology)
- Dr Toulosie Ramtohul (radiology)

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