



11th International Conference of Contemporary Oncology
Poznań 15.03.2019

Molecular classification of colorectal cancer (CRC)

Andrzej Deptała

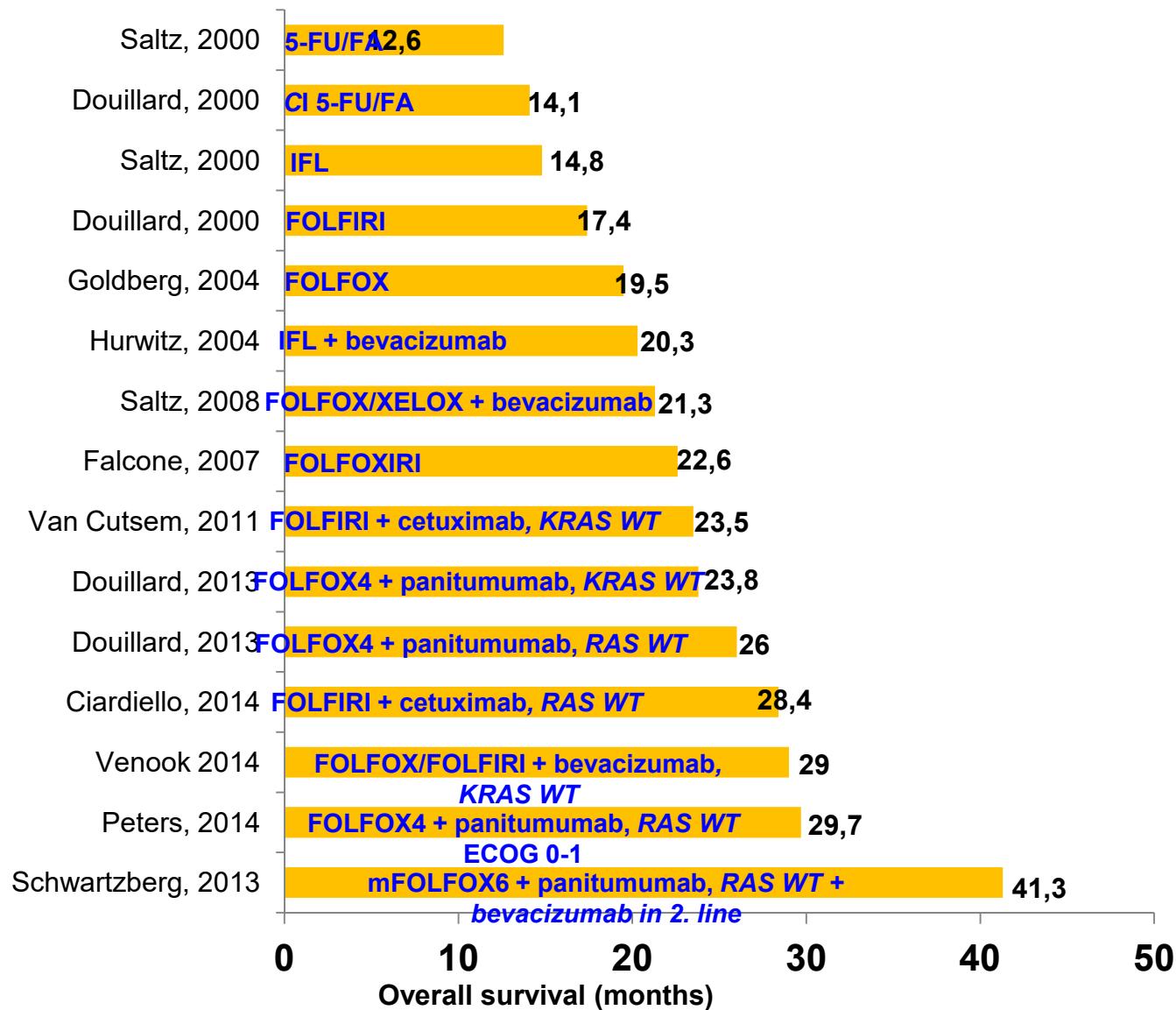


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Central Clinical Hospital of the MSWiA in Warsaw**

**Department of Cancer Prevention
Faculty of Health Sciences
Medical University of Warsaw**



Progress in the treatment of patients with metastatic colorectal cancer (mCRC)



Overview of molecular subtyping

Single marker molecular subtyping

- KRAS/NRAS
- BRAF
- MSI-H
- HER2 amplification
- Fusions

RNA-based molecular subtyping

- Consensus molecular subtypes
- Intrinsic subtyping

Immune subtyping

- Immune quantification
- Tumor mutation burden

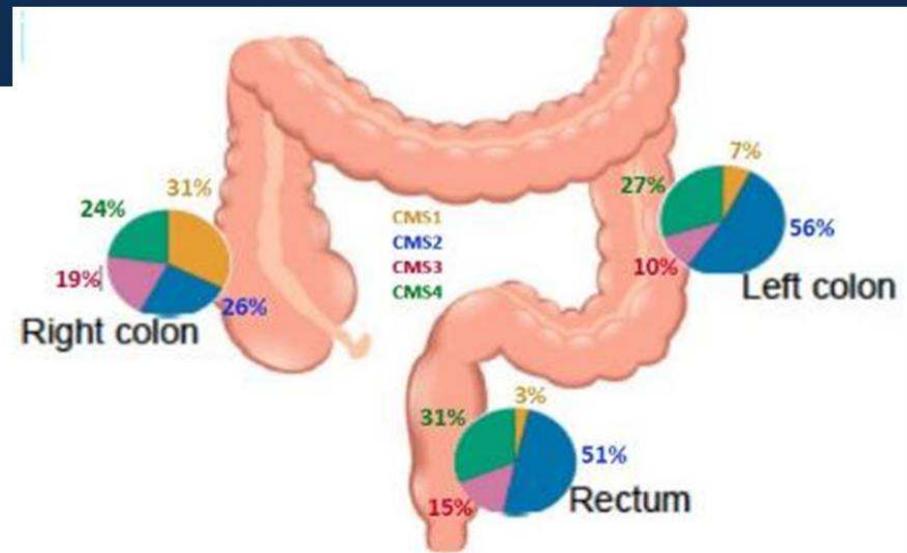
Proposed taxonomy of CRC reflecting biological difference in the gene expression-based molecular subtypes

The consensus molecular subtypes of colorectal cancer

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NATURE MEDICINE VOLUME 21 | NUMBER 11 | NOVEMBER 2015

CMS1 MSI immune	CMS2 Canonical	CMS3 Metabolic	CMS4 Mesenchymal
14%	37%	13%	23%
MSI, CIMP high, hypermutation	SCNA high	Mixed MSI status, SCNA low, CIMP low	SCNA high
BRAF mutations		KRAS mutations	
Immune infiltration and activation	WNT and MYC activation	Metabolic deregulation	Stromal infiltration, TGF-β activation, angiogenesis
Worse survival after relapse			Worse relapse-free and overall survival



PRESENTED AT: 2019 Gastrointestinal Cancers Symposium | #GI19

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Proposed taxonomy of CRC reflecting biological difference in the gene expression-based molecular subtypes

Tabela 25.1. Molekularne i kliniczne cechy raka jelita grubego wg klasyfikacji CMS i sugerowane na tej podstawie możliwości prowadzenia badań nad terapią personalizowaną

Parametr	CMS1 (ang. MSI immune – MSI immunogenny)	CMS2 (ang. canonical – podstawowy nabłonkowy)	CMS3 (ang. metabolic – związany z deregulacją szlaków metabolicznych)	CMS4 (ang. mesenchymal – zależny od mikrośrodowiska)
częstość występowania	~15	~40	~15	~30
lokalizacja guza w jelicie grubym	z przewagą w prawej połowie okrężnicy	z przewagą w lewej połowie okrężnicy i w odbytnicy	z przewagą w prawej połowie okrężnicy	z przewagą w lewej połowie okrężnicy i w odbytnicy
właściwości komórek raka	hiperzmutowane i hipermetylowane, MSI, przewaga zmutowanego BRAF	CIN, MSS, EGFR+, nadmierna regulacja ERBB2	mieszany MSI/MSS, CIN, deregulacja szlaków metabolicznych, przewaga zmutowanego KRAS	MSS, CIN, zaburzone interakcje pomiędzy komórkami nabłonkowymi i podścieliskowymi
właściwości mikrośrodowiska	liczne limfocyty T cytotoksyczne i pomocnicze oraz komórki NK	niewielkie nacieki komórek immunologicznych i podścieliskowych	niewielkie nacieki komórek immunologicznych i podścieliskowych	liczne komórki podścieliska, liczne komórki supresorowe: limfocyty T-reg, limfocyty B, komórki pochodzenia mieloidalnego
rokowanie	dłuższy RFS, po nawrocie krótszy OS	dłuższe RFS i OS	dłuższe RFS i OS	krótsze RFS i OS
leki do wykorzystania w badaniach klinicznych	inhibitatory PD-1 i PD-L1; inhibitatory szlaku sygnałowego BRAF	inhibitatory szlaku sygnałowego poprzez EGFR, hamowanie nadekspresji białek z rodziny HER	inhibitatory szlaków sygnałowych pan-RAF i MEK w skojarzeniu z inhibitorami torów enzymatycznych	kojarzenie leków immunostymulujących z lekami immunosupresyjnymi

What are the CMS strengths?

CMS1: Immunogenic Tumors

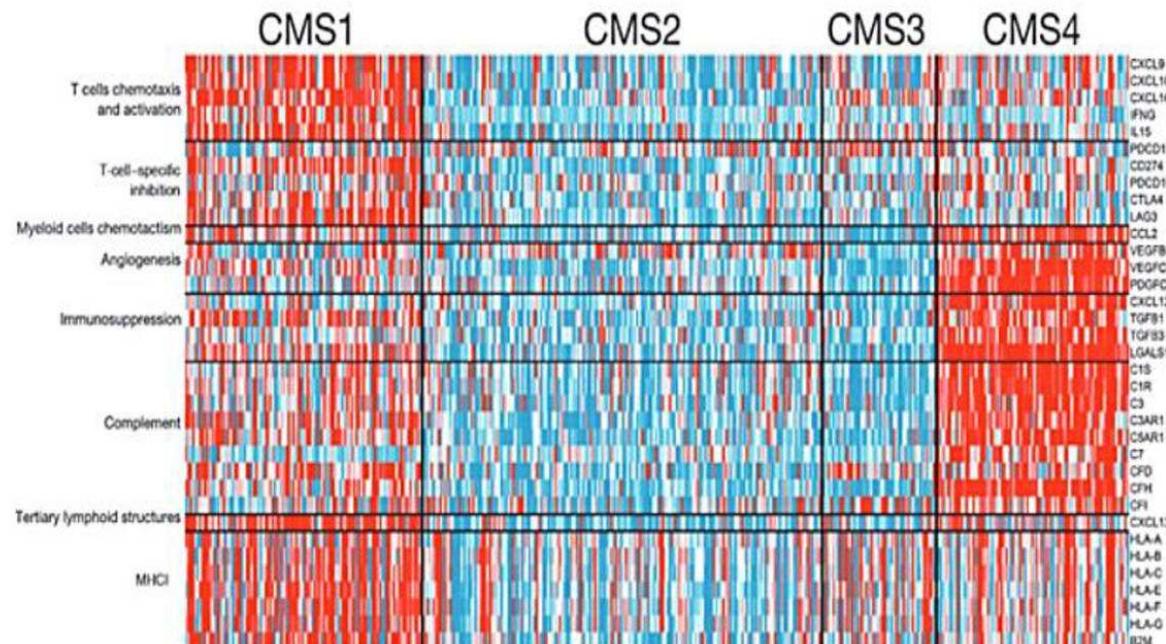
Infiltrating activated lymphocytes

CMS2/3: Immune Desert

No evidence of immune activation

CMS4: Immune Excluded

Immune system is engaged, but microenvironment prevents activity

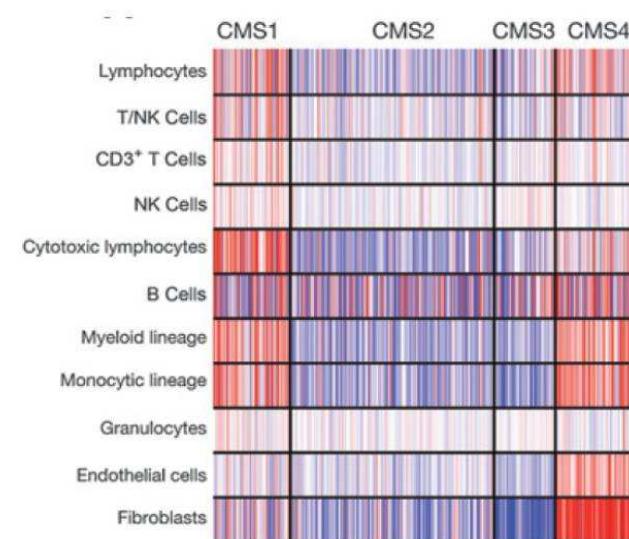


Becht et al CCR '16

What are the CMS strengths?

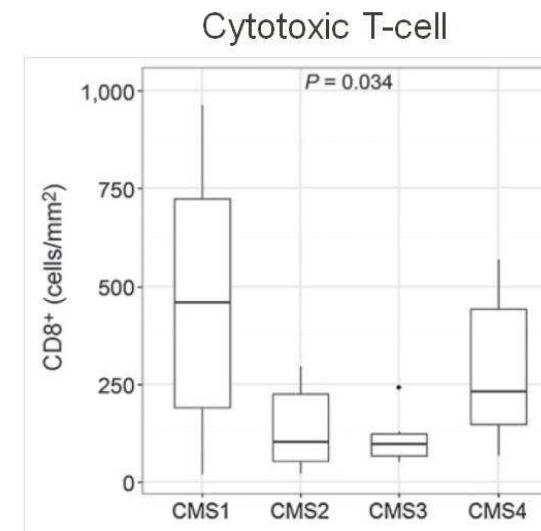
CMS1: Immunogenic Tumors

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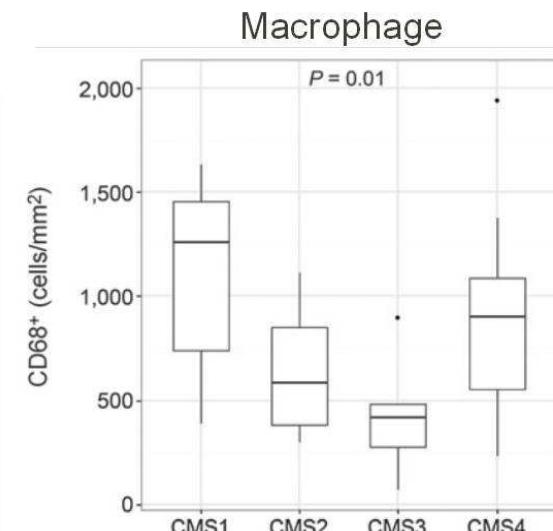
CMS2/3: Immune Desert

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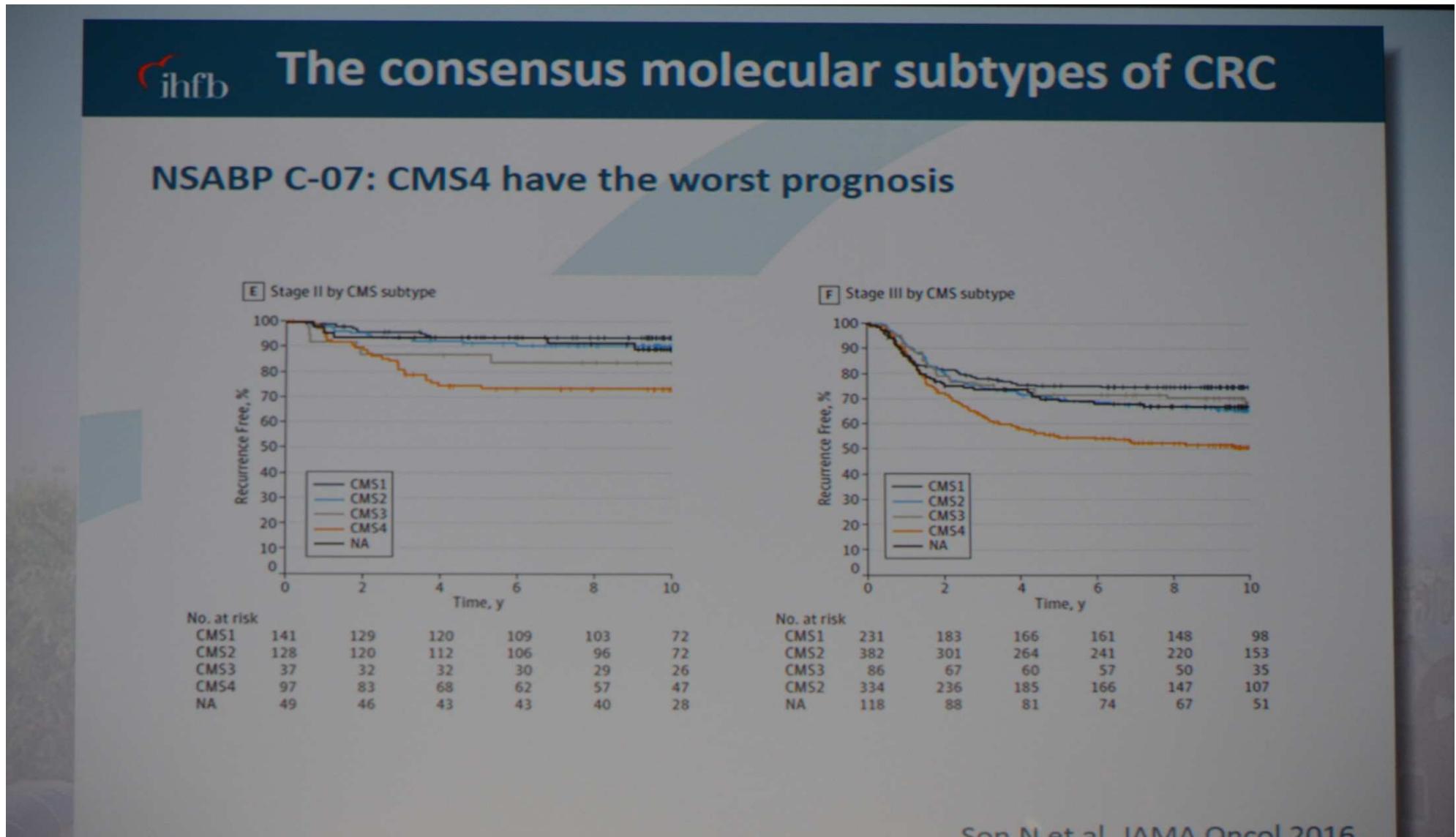


CMS4 has a moderate cytotoxic T-cell infiltrate, but high myeloid, TGF-β signaling

Becht et al CCR '16, Lai et al CCR '18

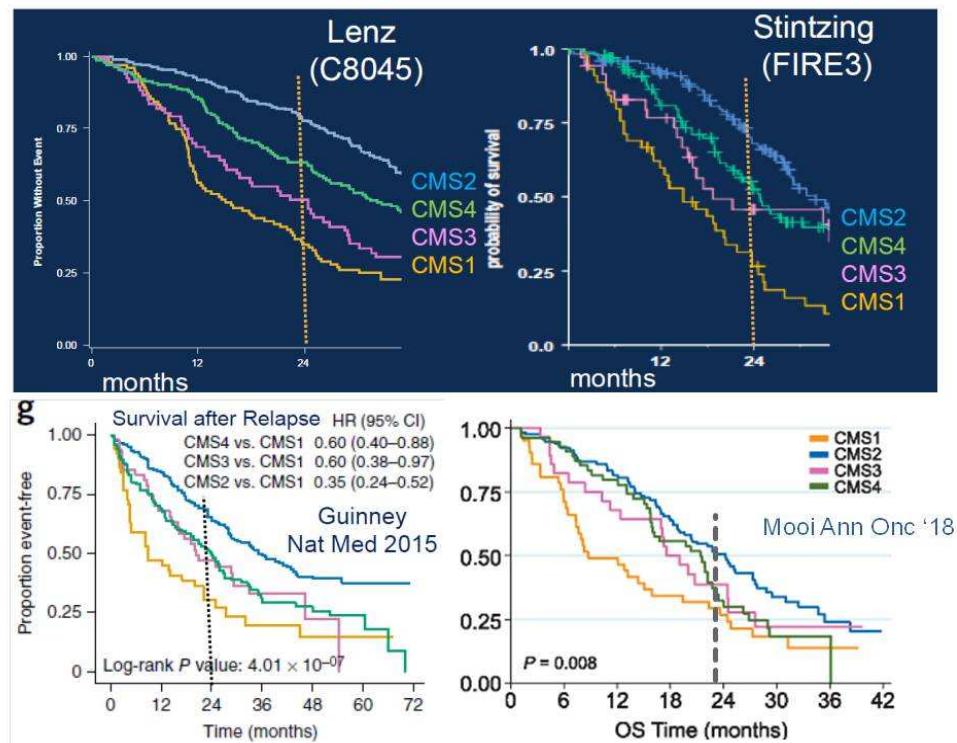
Presented By Scott Kopetz at 2019 Gastrointestinal Cancer Symposium

CMS has consistent prognostic information in CRC



CMS has consistent prognostic information in mCRC

Despite being designed agnostic to outcomes, strong prognostic information.



Median overall survival: Differs from 15 months (CMS1) to 40 months (CMS2)

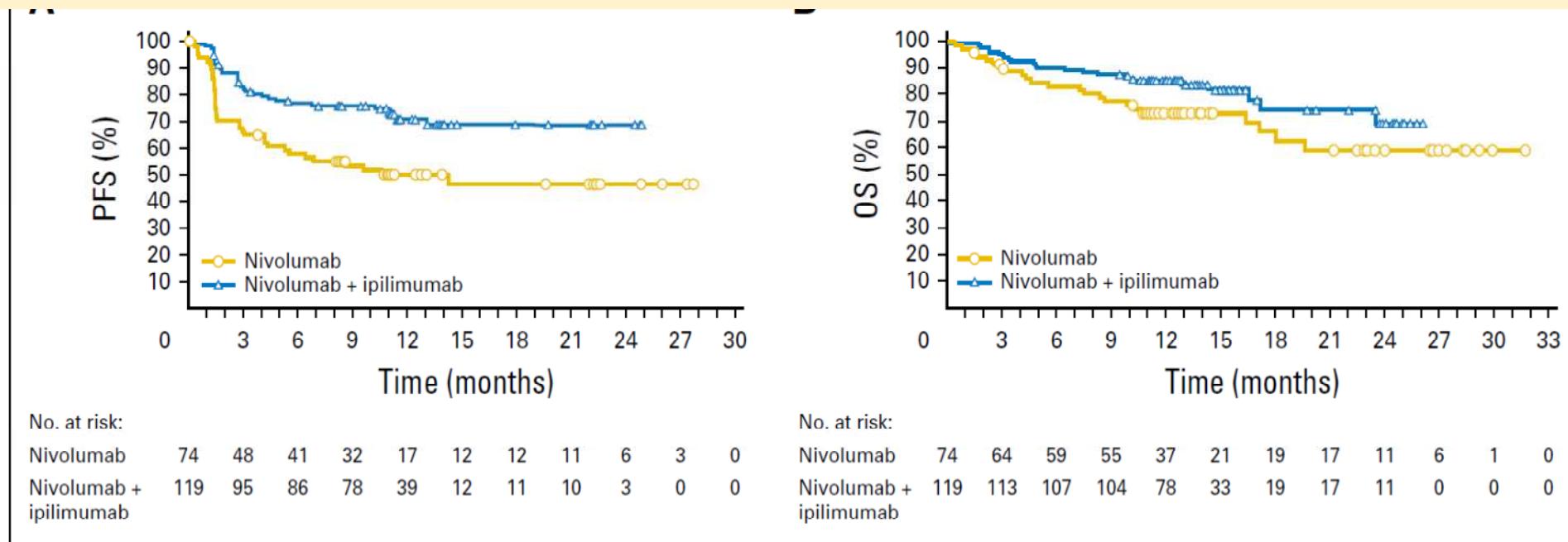
Median 1st Line Progression-free survival: Differs from 5.7 months (CMS1) to 14.1 months (CMS2)

Slide courtesy of Wells Messersmith, ASCO '17

Presented By Scott Kopetz at 2019 Gastrointestinal Cancer Symposium

CMS is a predictive factor in mCRC: CMS1 may benefit from anti-PD-1 ± anti-CTLA-4 agents

Durable Clinical Benefit With Nivolumab Plus Ipilimumab in DNA Mismatch Repair–Deficient/Microsatellite Instability–High Metastatic Colorectal Cancer

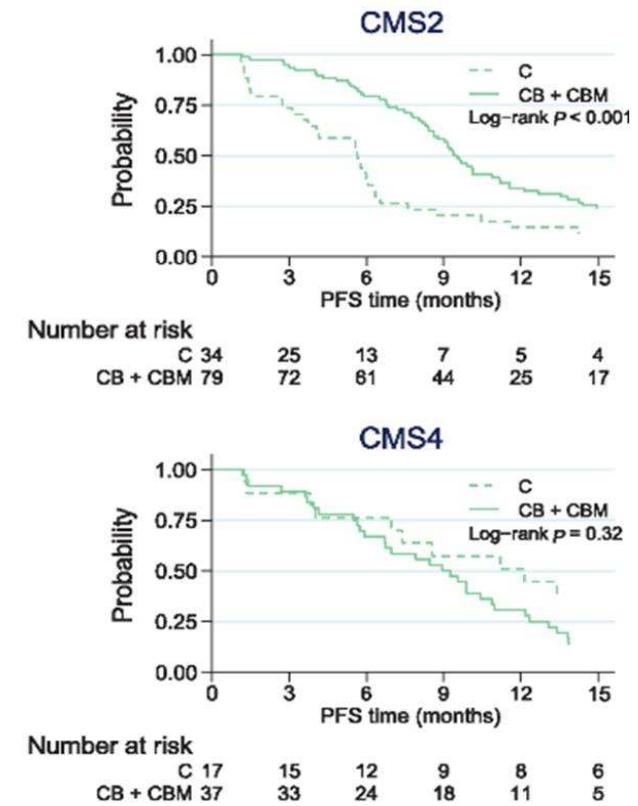
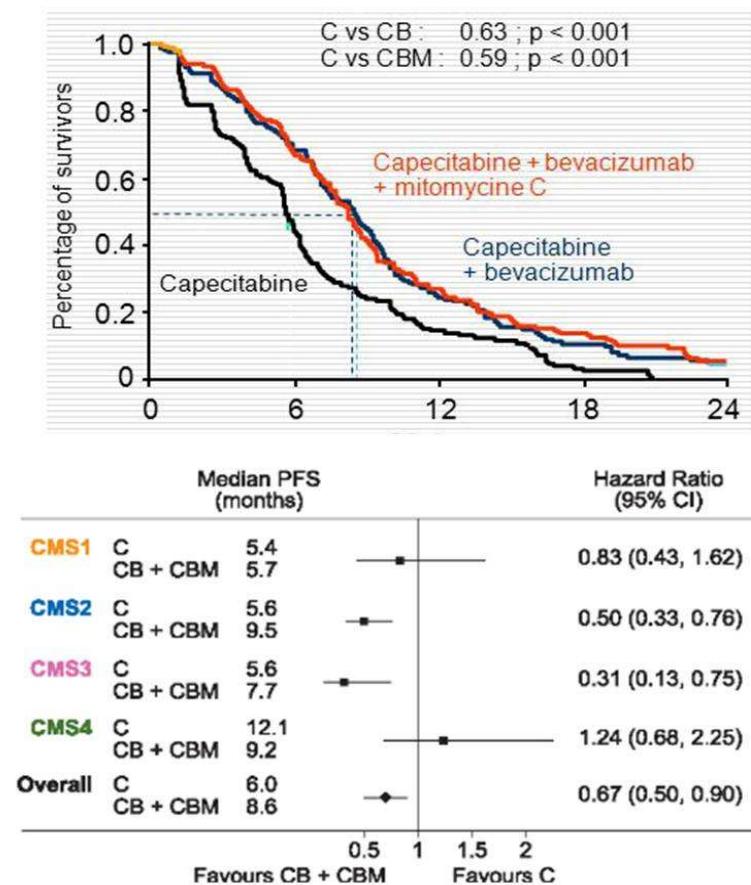


On study from first dose to data cutoff (July 2017).

CMS is predictive factor in mCRC: CMS2 and CMS3 may benefit from addition of bevacizumab

**CMS2/3 may
Benefit from
Addition of
Bevacizumab**

**AGITG MAX
Trial**



Mooi et al Annals Onc '18

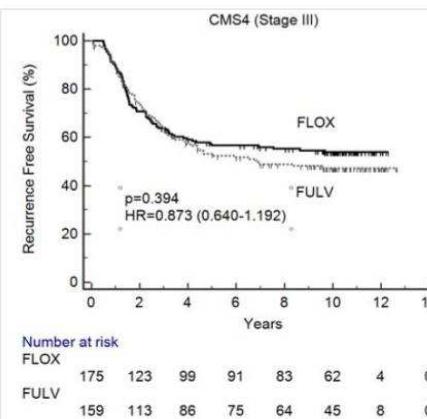
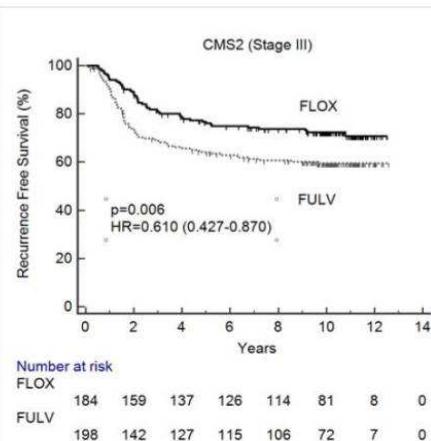
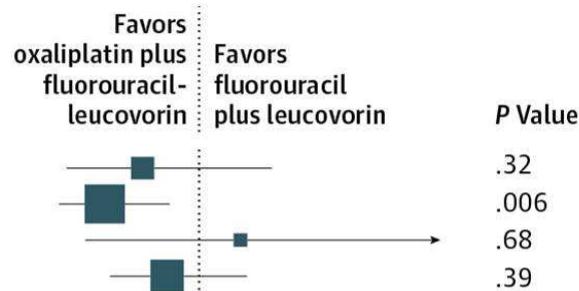
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CMS has predictive information in mCRC: Limited benefit with oxaliplatin?

Mesenchymal CMS4 : Limited Benefit with Oxaliplatin?

C-07 study of FLOX vs FULV

Subtypes	No. of Patients	HR (95% CI)
CMS1	231	0.77 (0.46-1.29)
CMS2	382	0.61 (0.43-0.87)
CMS3	86	1.17 (0.54-2.53)
CMS4	334	0.87 (0.64-1.19)

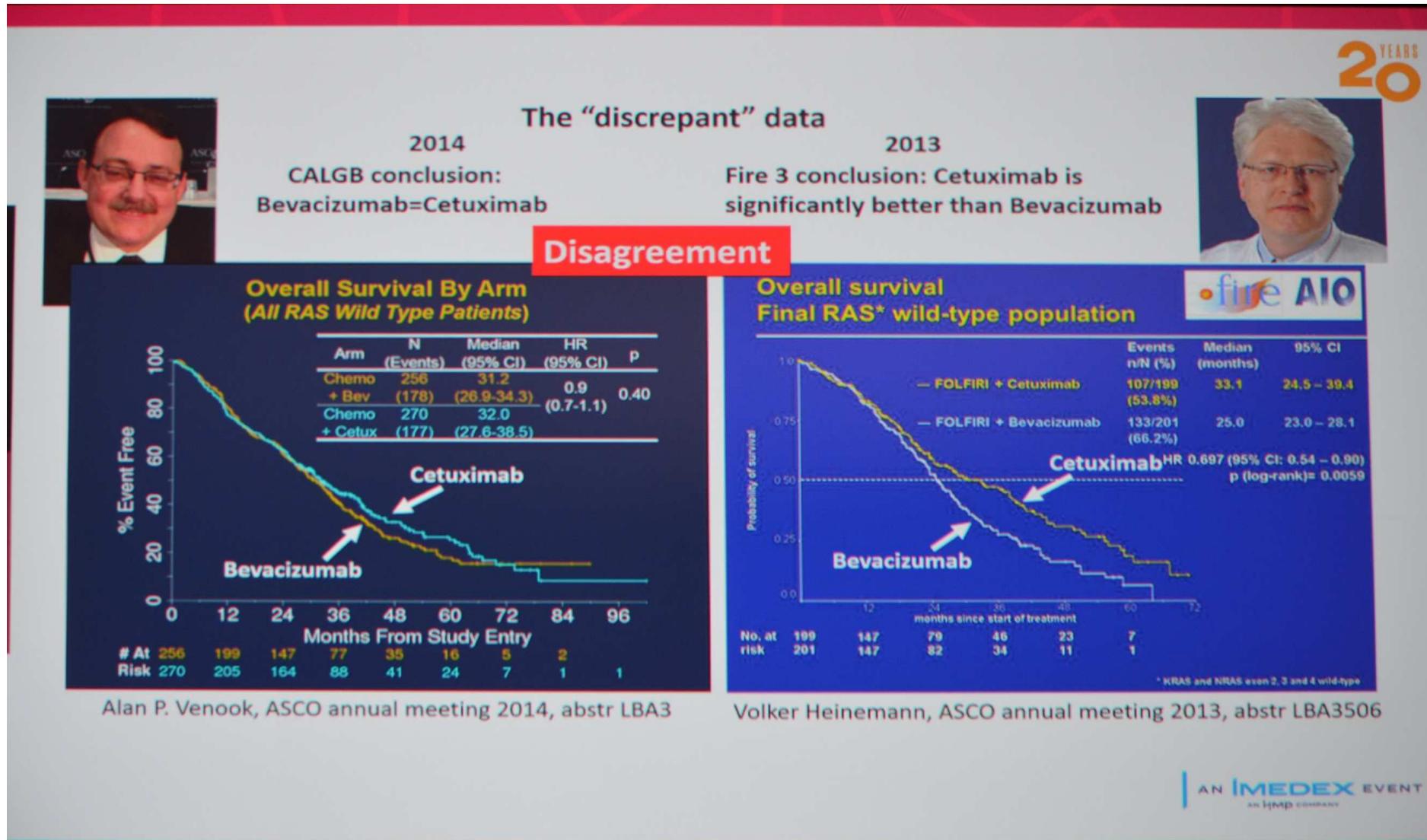


Are there other subgroups or oxali-specific signatures that would perform better?

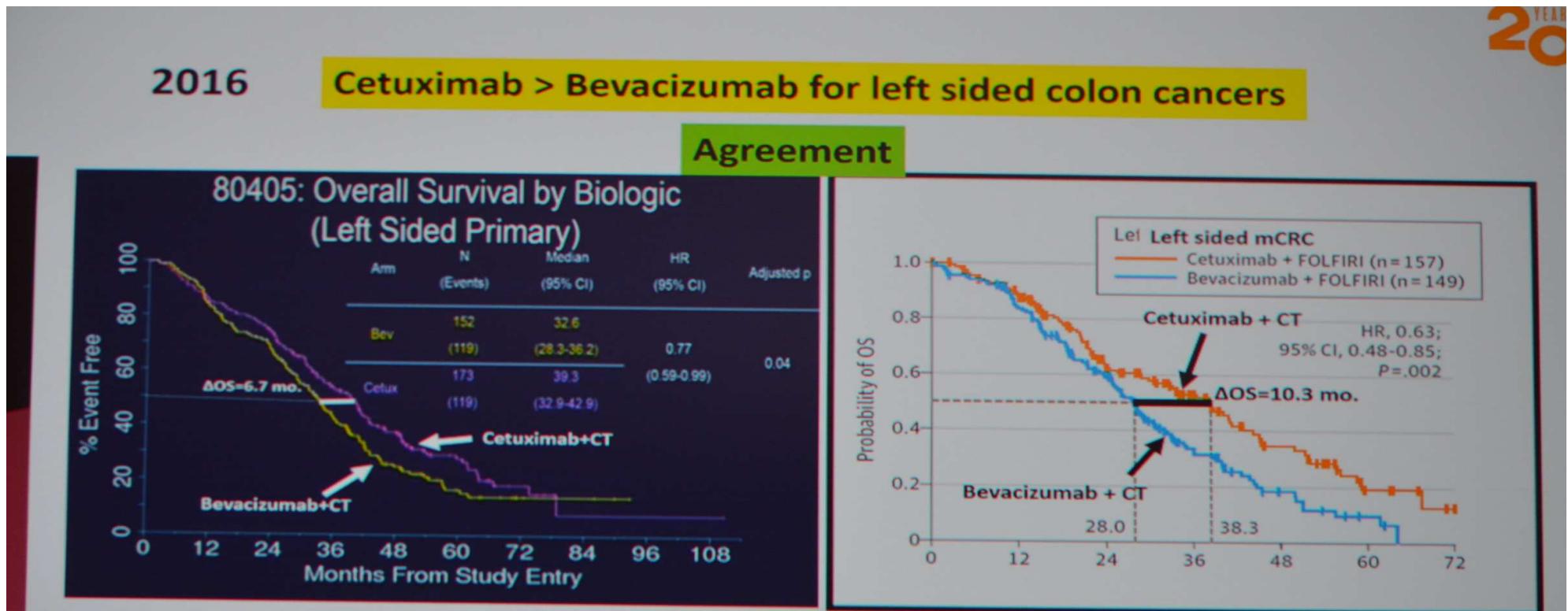
Song et al JAMA Onc '17

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CMS-dependent of explaining the "unexplainable" – analysis



CMS-dependent of explaining the "unexplainable" – analysis



Venook AP, et al. ASCO 2016 (Abstract No. 3504)

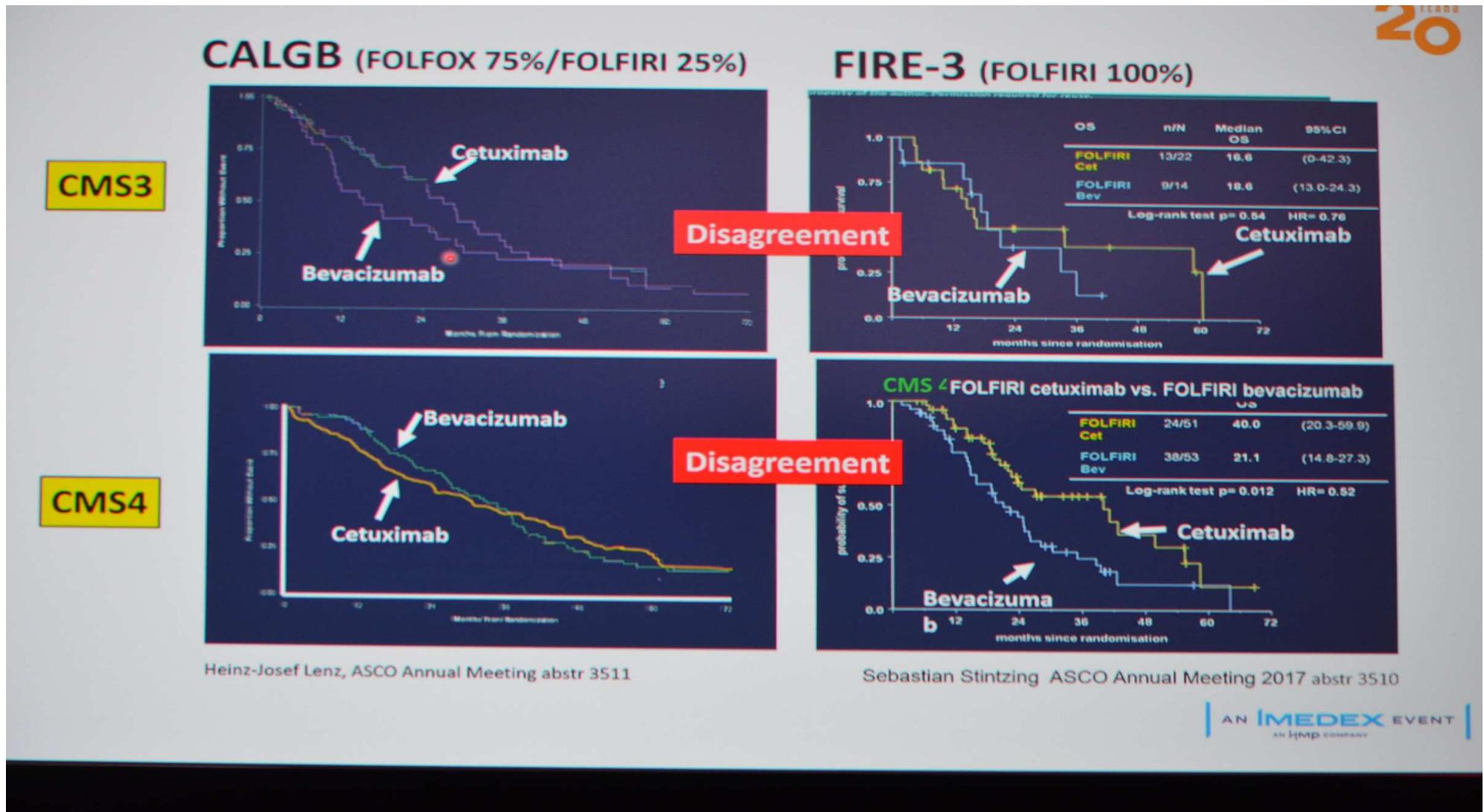
1. Tejpar S, et al. JAMA Oncol 2017;3(2):194–201;
2. Heinemann V et al. Lancet Oncol 2014;15:1065–1075

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CMS-dependent of explaining the "unexplainable" – analysis



CMS-dependent of explaining the "unexplainable" – analysis



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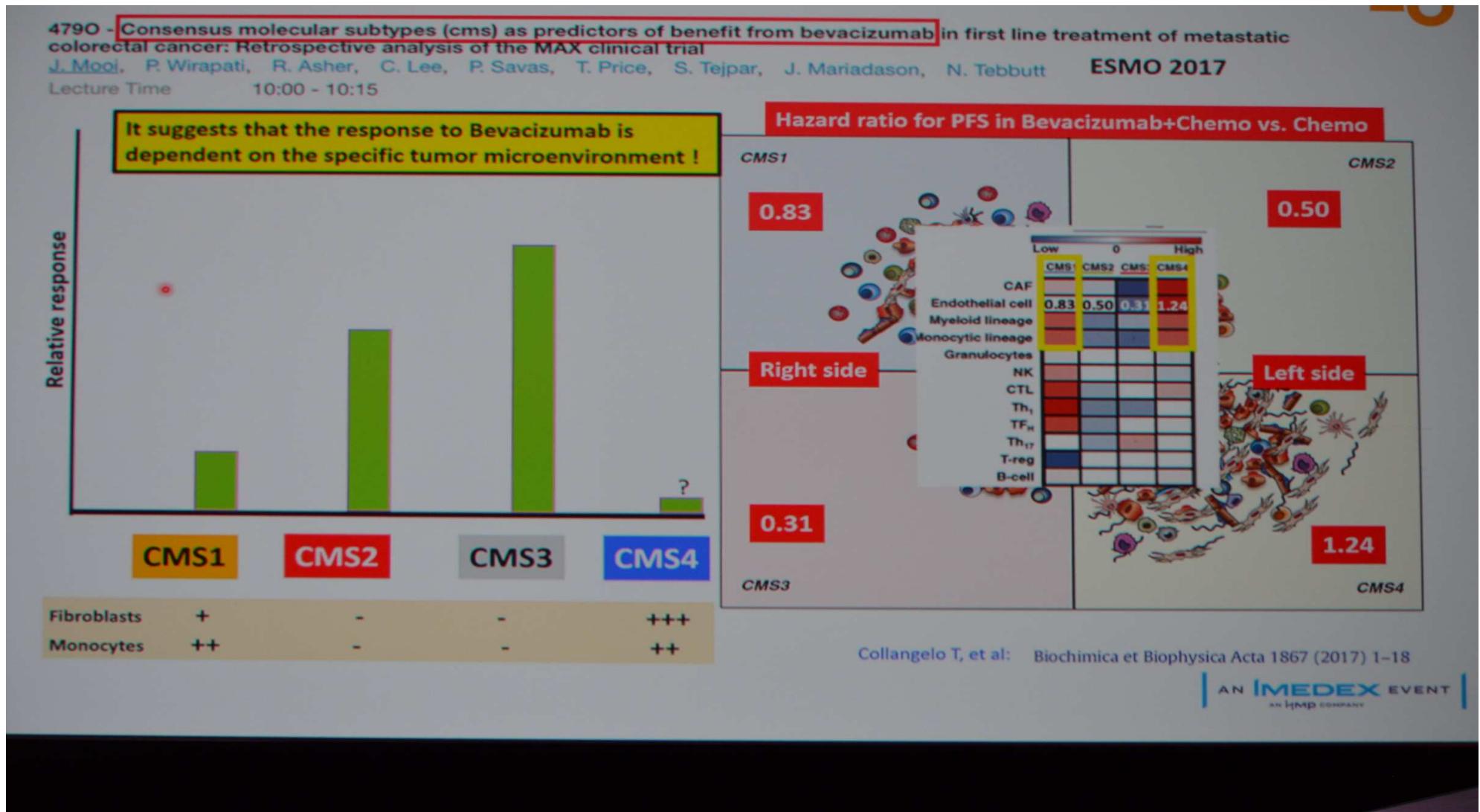
The discrepant data in overall survival (OS) may be interpreted as a interplay between:

- *Biological agents*
- *Chemotherapy regimen*
- *Microenvironment influence*

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Is a difference in response to biologics between the CMS subtypes?

CMS-dependent of explaining the "unexplainable" – analysis



CMS-dependent of explaining the "unexplainable" – analysis

Cancer associated fibroblasts and tumor associated macrophages (TAM) mediate resistance to Bevacizumab by release of alternative pro-angiogenic factors !

The diagram illustrates the mechanisms of resistance to Bevacizumab. Bevacizumab is shown targeting PDGF-C produced by Cancer Associated Fibroblasts (CAF). Additionally, TAM (Tumor Associated Macrophages) produce several other pro-angiogenic factors: Sema4D, IL-12, b-FGF, TNF α , ADM, uPA, and YKL-40, which contribute to angiogenesis.

PDGF-C Mediates the Angiogenic and Tumorigenic Properties of Fibroblasts Associated with Tumors Refractory to Anti-VEGF Treatment

Yongping Crawford,¹ Ian Kasman,¹ Lanlan Yu,¹ Cuiling Zhong,¹ Xiumin Wu,¹ Zora Modrusan,¹ Josh Kaminker,¹ and Napoleone Ferrara^{1*}
*Genentech, Inc., 1 DNA Way, South San Francisco, CA 94080, USA
Correspondence: nf@gene.com
DOI 10.1016/j.ccr.2008.12.004
Cancer Cell 15, 21–34, January 6, 2009

Tumor-Associated Fibroblasts as “Trojan Horse” Mediators of Resistance to Anti-VEGF Therapy

Giulio Francia,^{1,*} Urban Emmenegger,^{1,2,3} and Robert S. Kerbel^{1,4}
Cancer Cell 15, January 6, 2009 ©2009 Elsevier Inc. 3

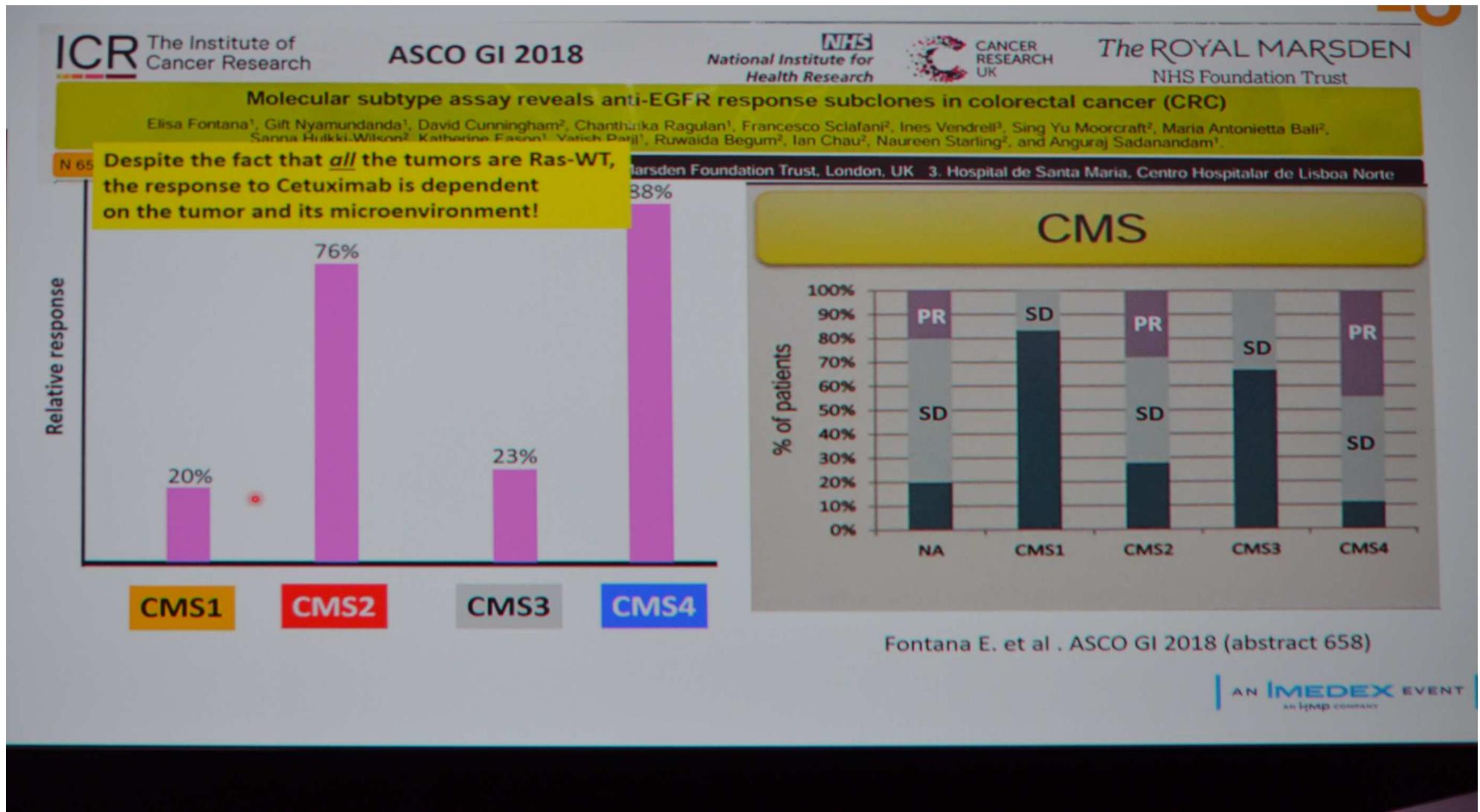
Role of tumor associated macrophages in tumor angiogenesis and lymphangiogenesis

Vladimir Riabov^{1,2*}, Alexandru Gudima^{1,2*}, Nan Wang¹, Amanda Mickley^{1,3}, Alexander Orekhov² and Julie Khyshkowska^{1,2,3*}
Front Physiol 5:75, 2014

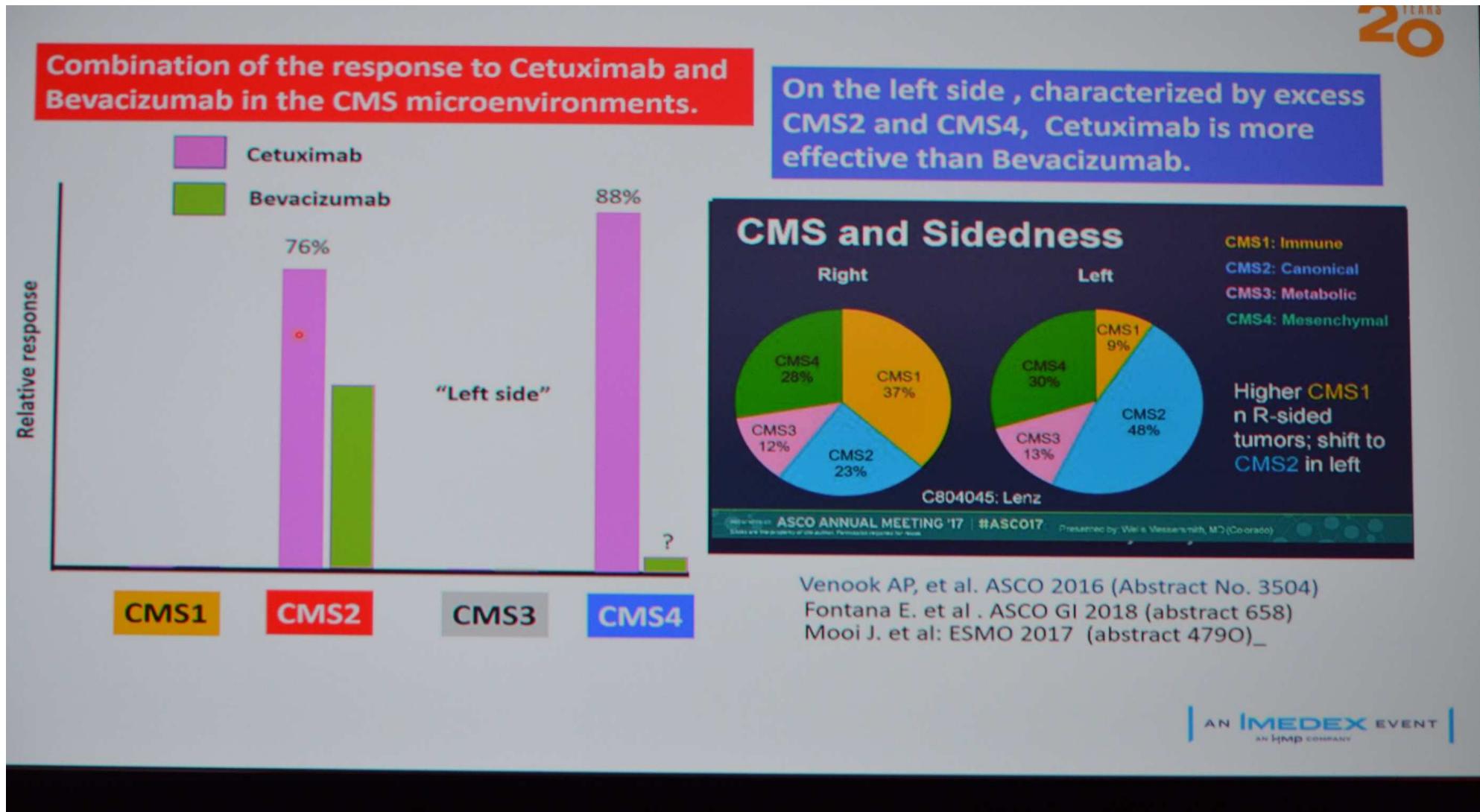
Thus, Bevacizumab is not enough to prevent angiogenesis in a fibroblast and monocyte rich microenvironment.

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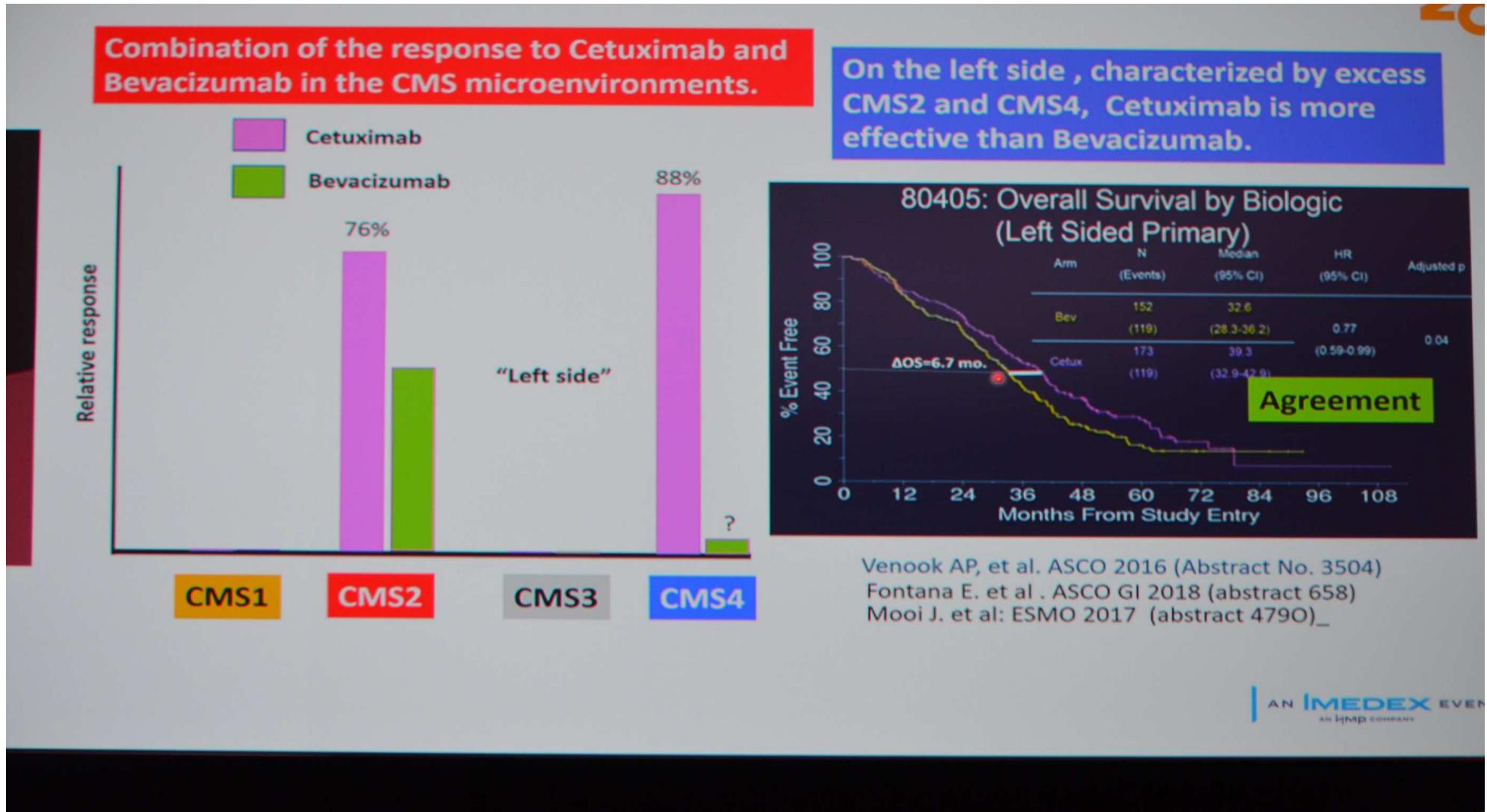
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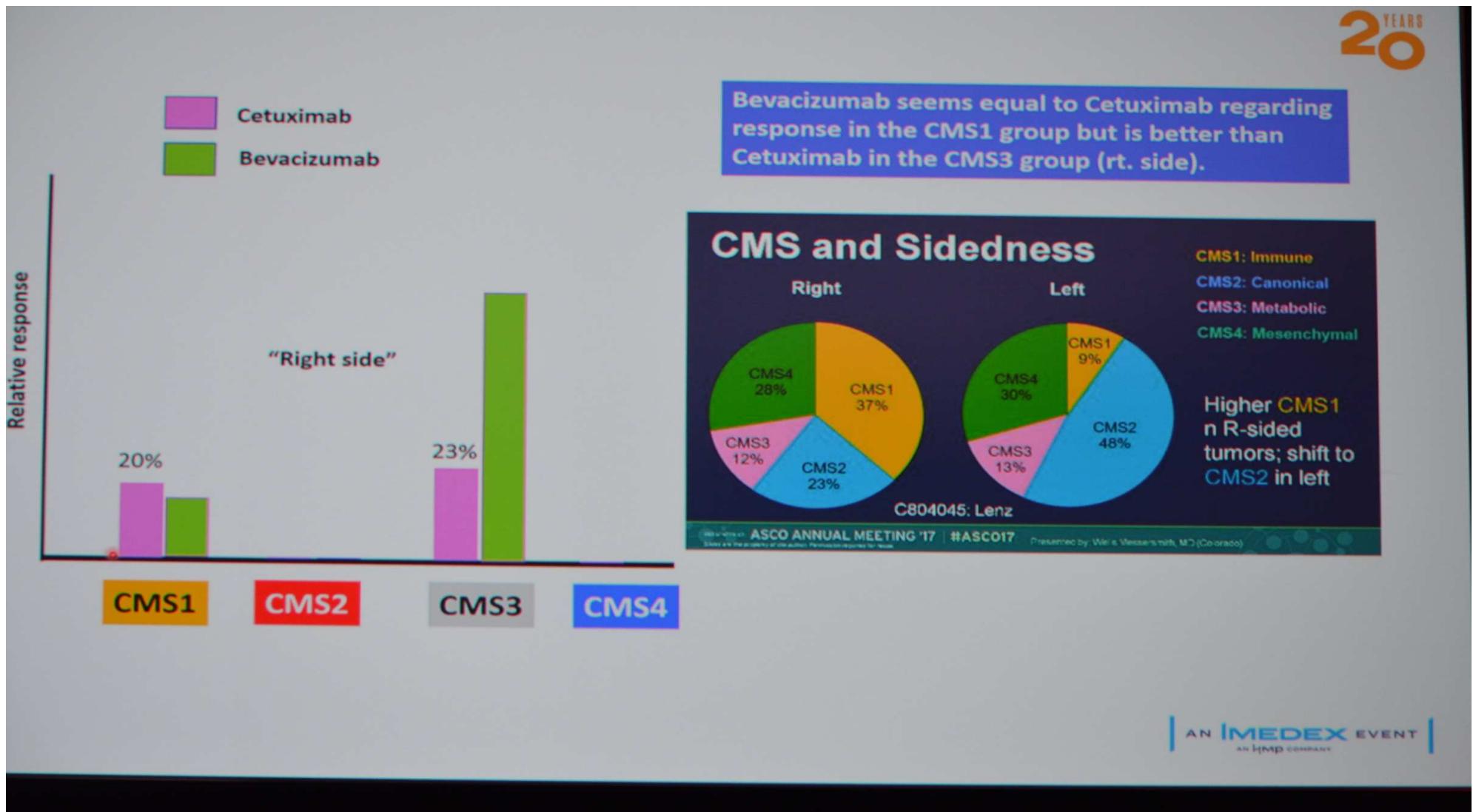
CMS-dependent of explaining the "unexplainable" – analysis



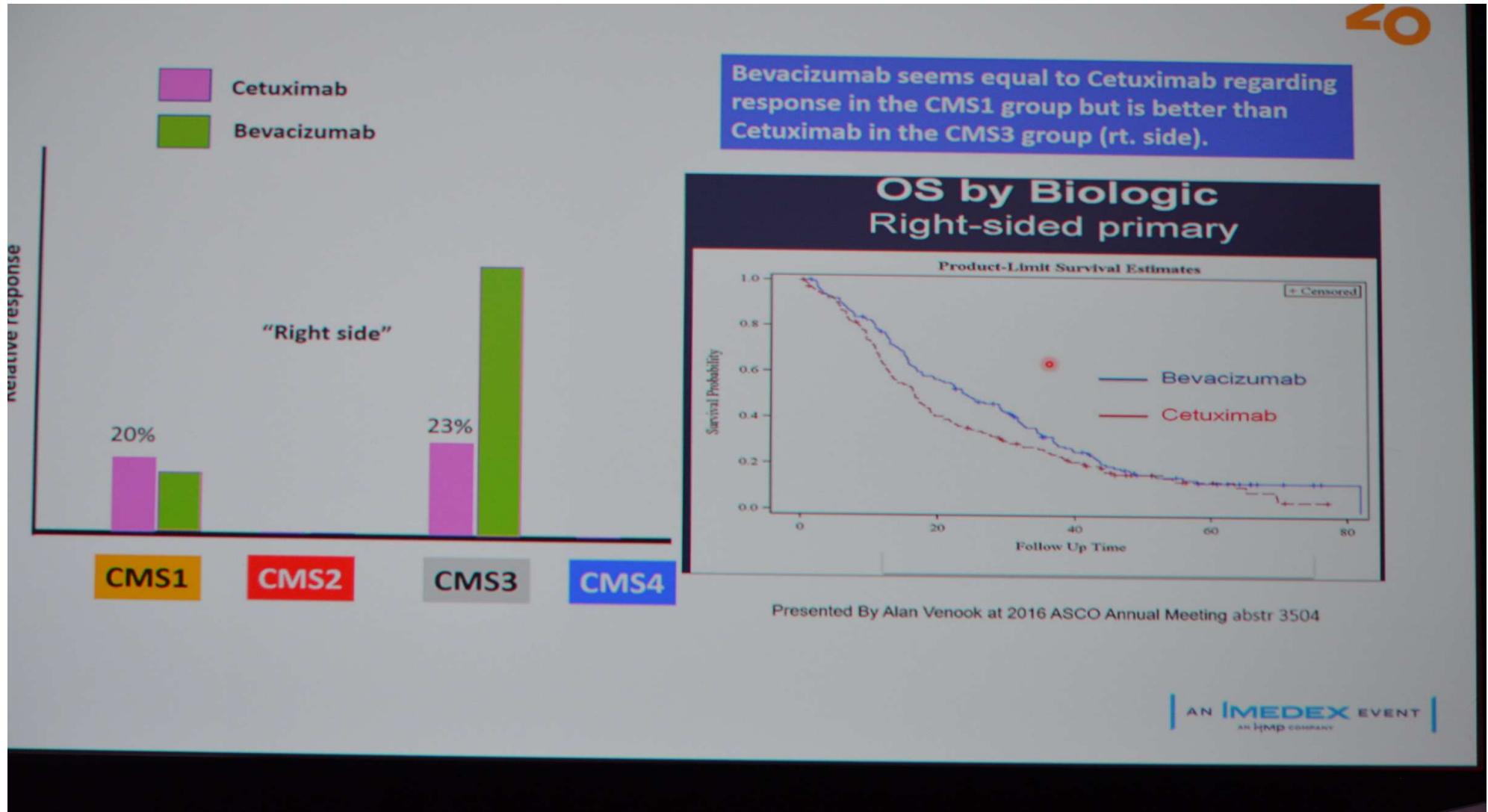
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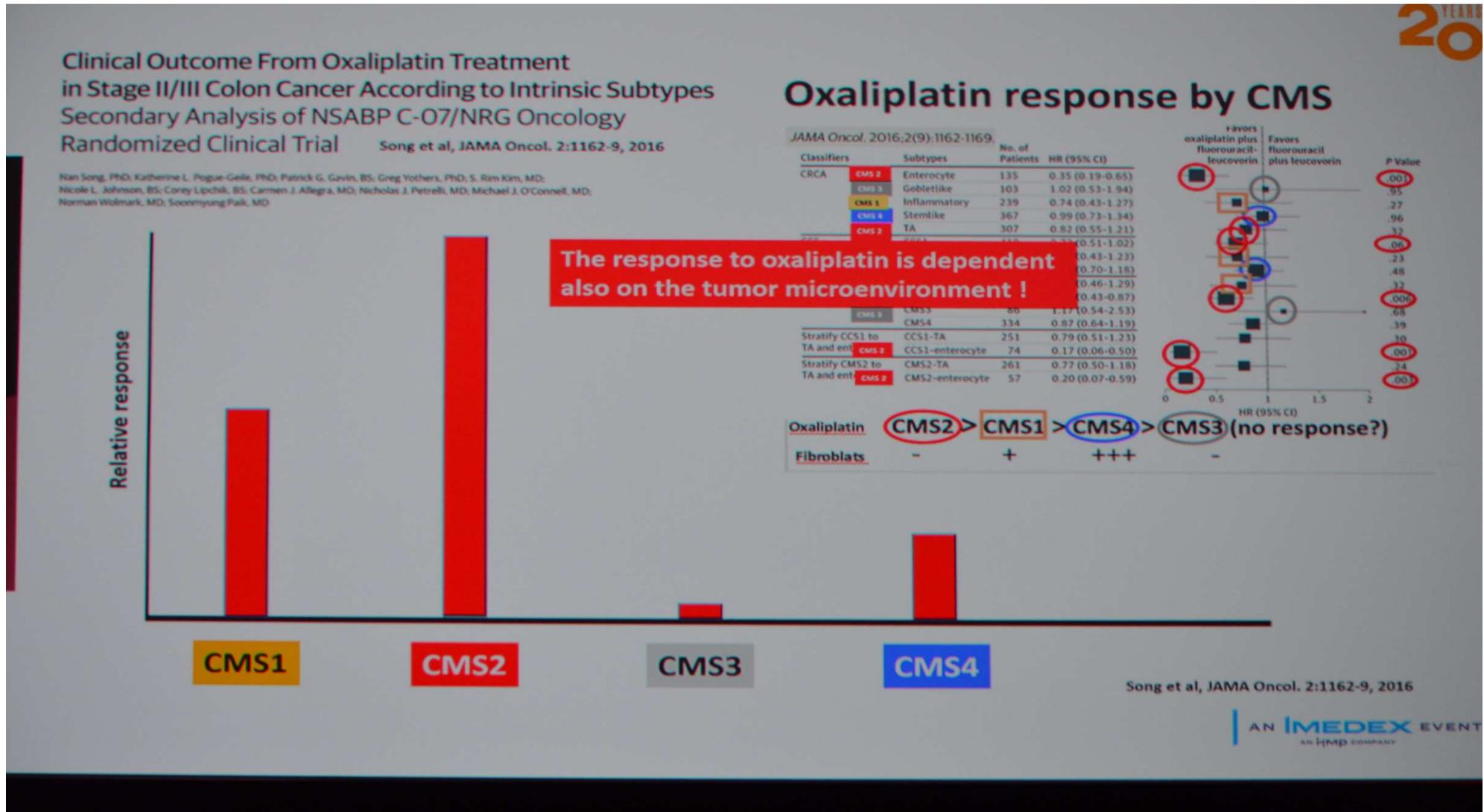
Is a difference in response to chemotherapy between the CMS subtypes?

GERCOR study

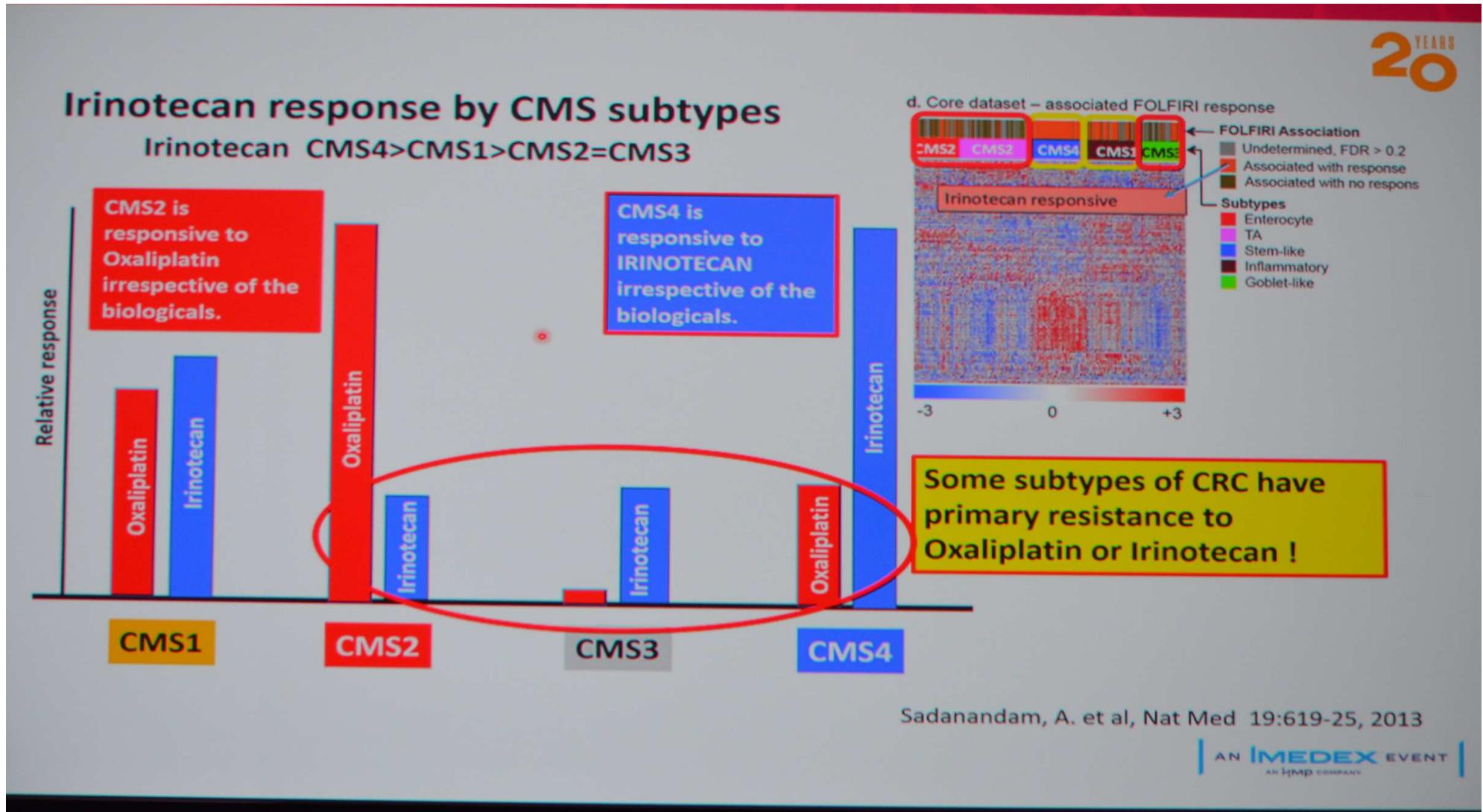
For ~15 years we consider that FOLFOX and FOLFIRI have identical clinical effects

	1-szy rzut → 2-gi rzut FOLFIRI → FOLFOX 6 n=109	1-szy rzut → 2-gi rzut FOLFOX 6 → FOLFIRI n=111	p
OR (CR+PR) (%)			
I linia leczenia	56	54	NS
II linia leczenia	15	4	0,05
Mediana PFS (miesiące)			
I linia leczenia	8,5	8,0	0,26
II linia leczenia	4,2	2,5	0,003
Mediana OS (miesiące)	21,5	20,6	0,99

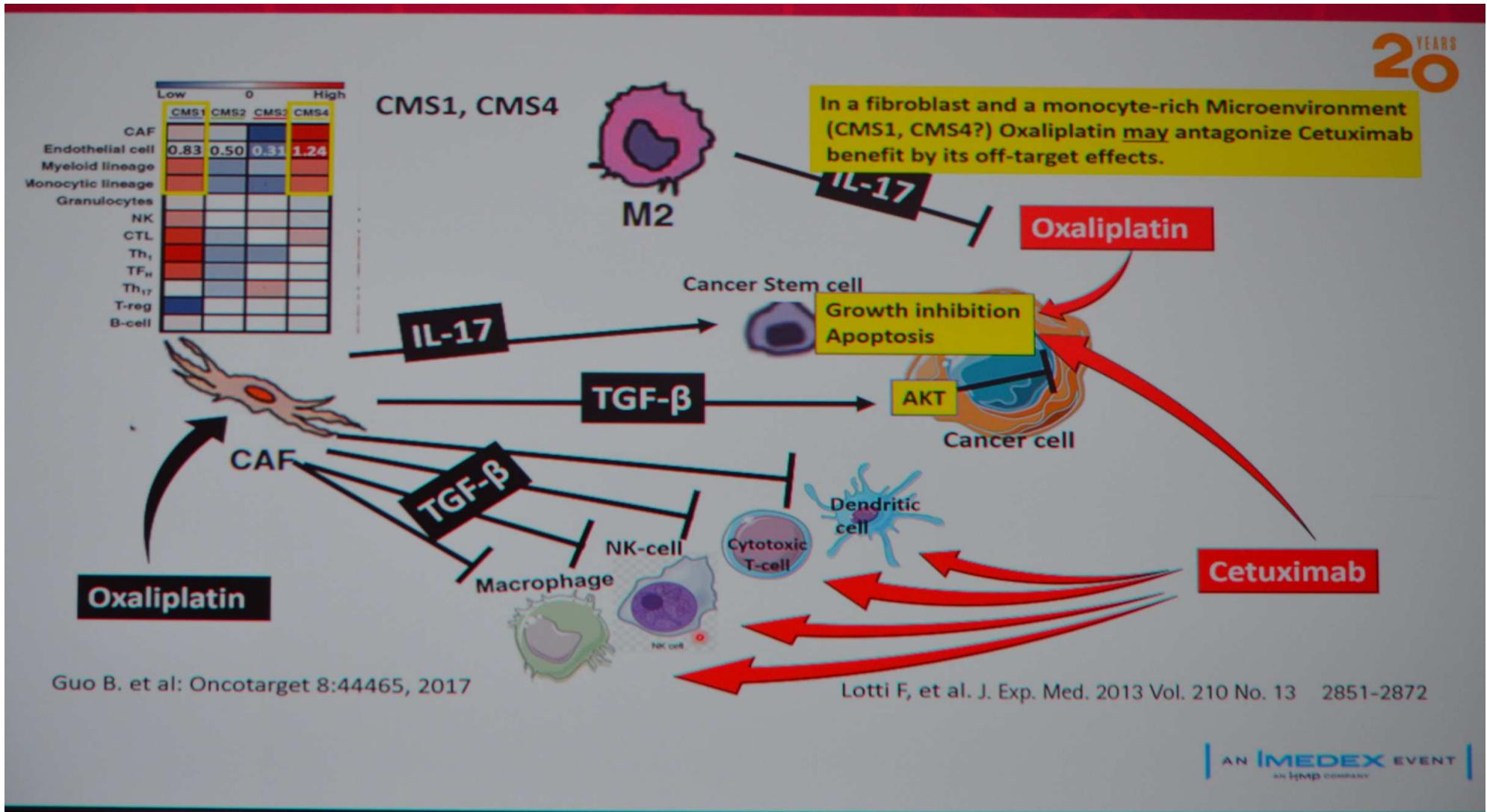
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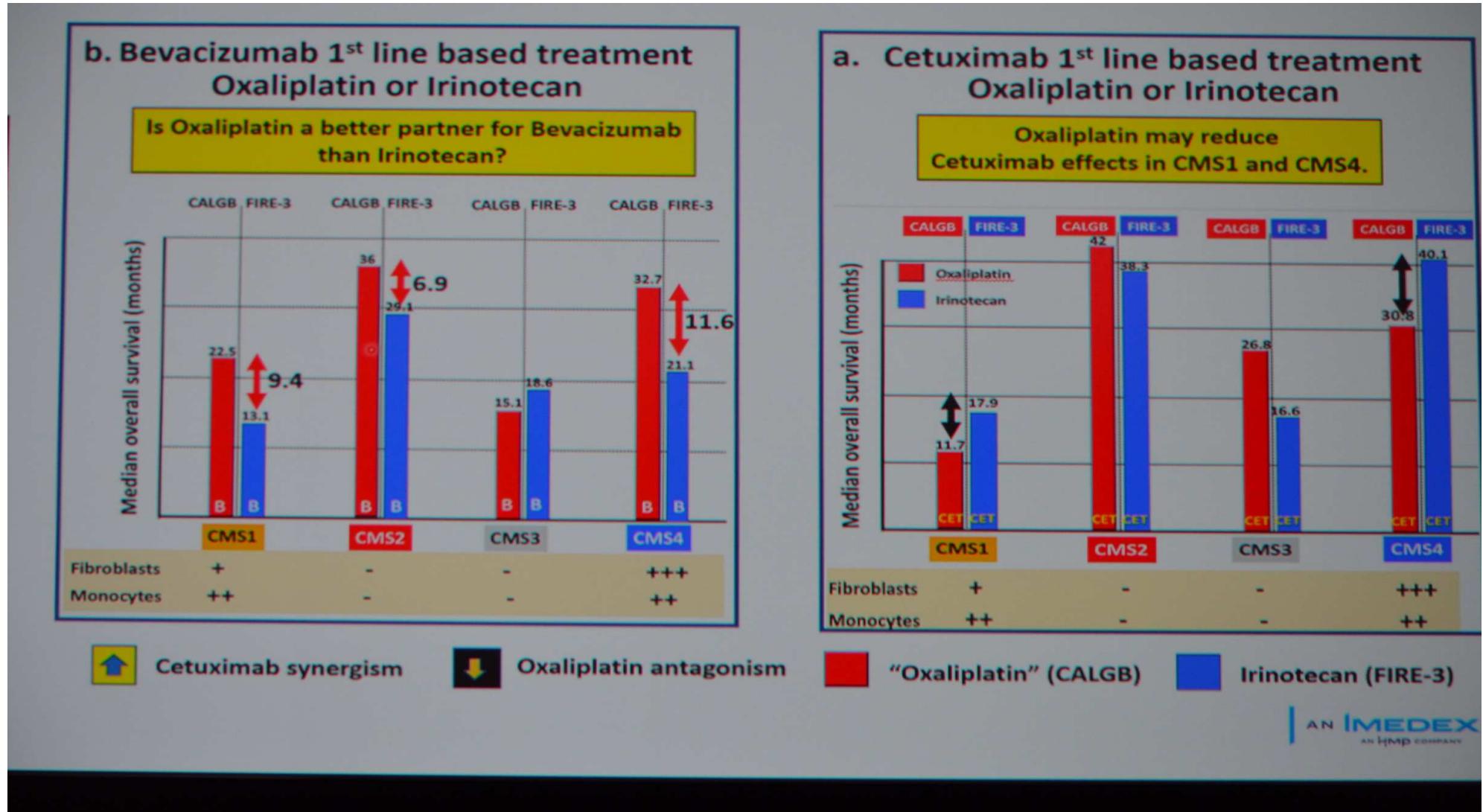
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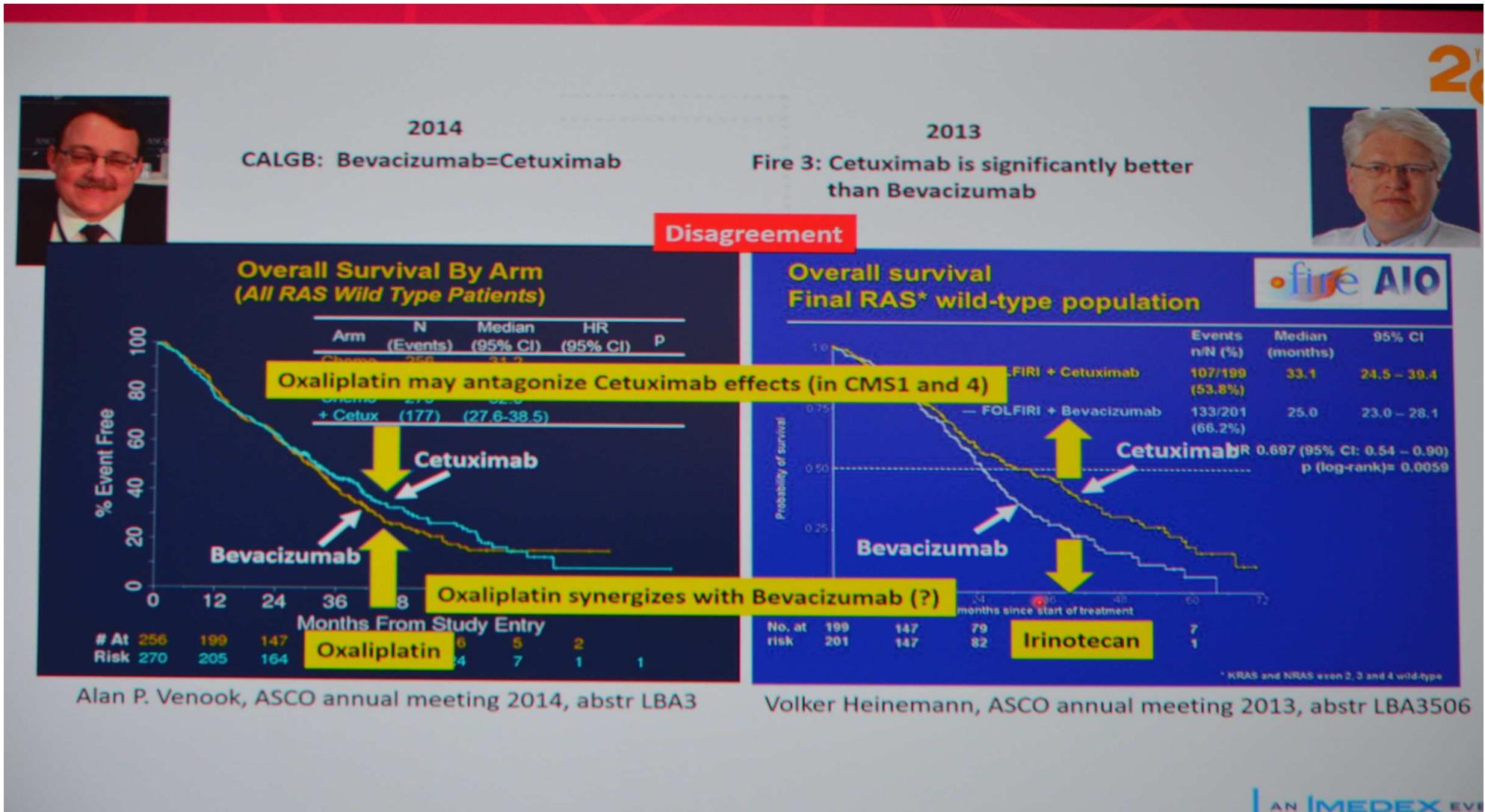
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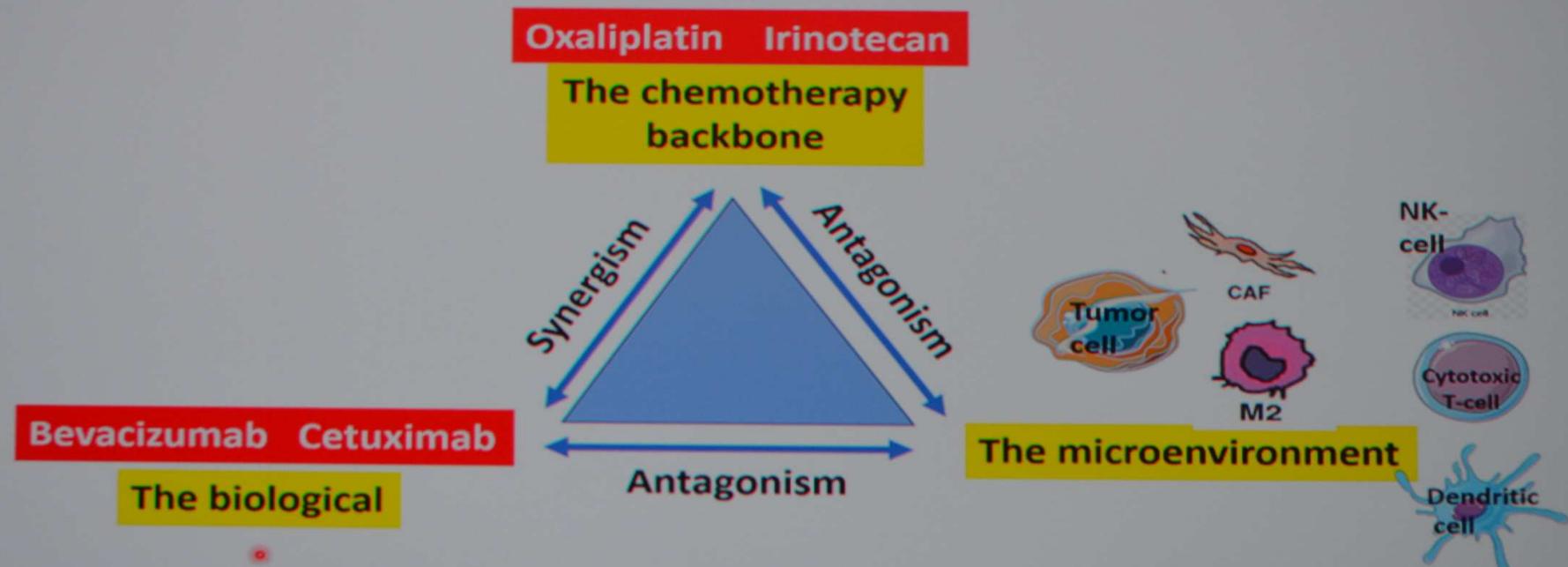
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Conclusions:

1. The response to treatment is not dependent solely on a single variable such as a biological but on a complex synergistic-antagonistic interaction between the biologicals, the chemotherapy backbone and the specific tumor microenvironment.



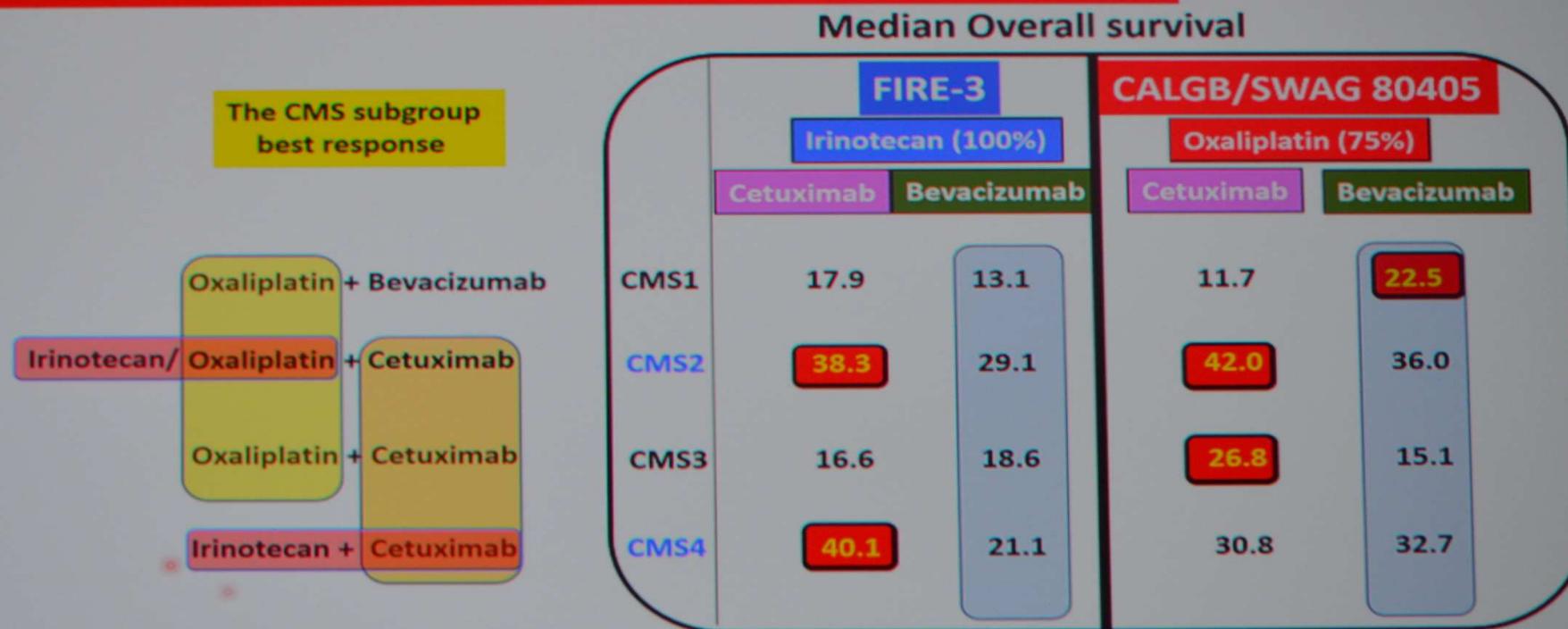
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CMS-dependent of explaining the "unexplainable" – analysis

2

Conclusions:

4. The best chemotherapy +biological combination for each colon cancer subtype:



Conclusions

- Molecular subtyping is a key mechanism to improve a patient outcome and opens new avenues for optimization of the personalized treatment in the different mCRC subtypes
- Current molecular subtypes with proven clinical activity:
 - *RAS/BRAF^{V600E}wild type* – cetuximab, panitumumab
 - *RAS mutation* – bevacizumab, afibbercept
 - *BRAF^{V600E} mutation* – triple EGFR + BRAF + MEK inhibition
 - *HER-2 amplification* – trastuzumab + lapatinib or pertuzumab
 - *MSI high* – nivolumab/ipilimumab
- If the presented analysis will be further validated, future personalized therapies must incorporate CMS based

