

Target-response relationship of bevacizumab may be more relevant than exposure-response: a Target-Mediated Drug Disposition model

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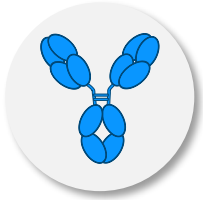
²Tours University Hospital, Department of Gastroenterology and Digestive Oncology, France

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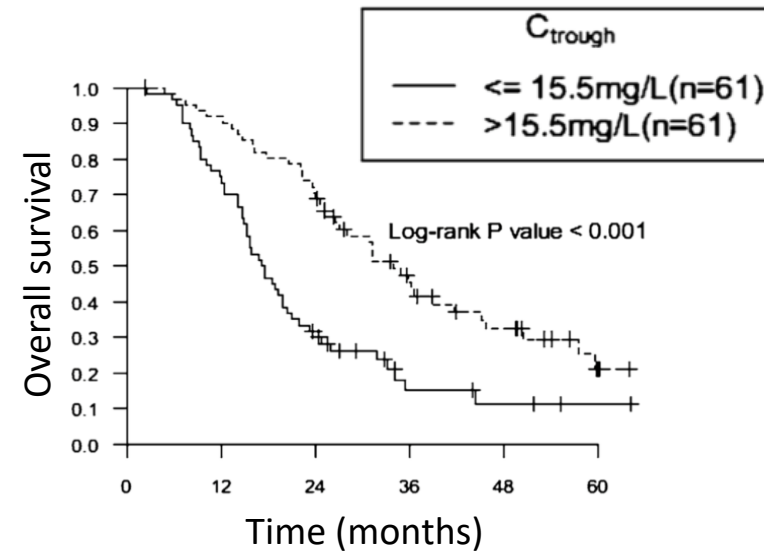
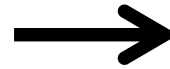
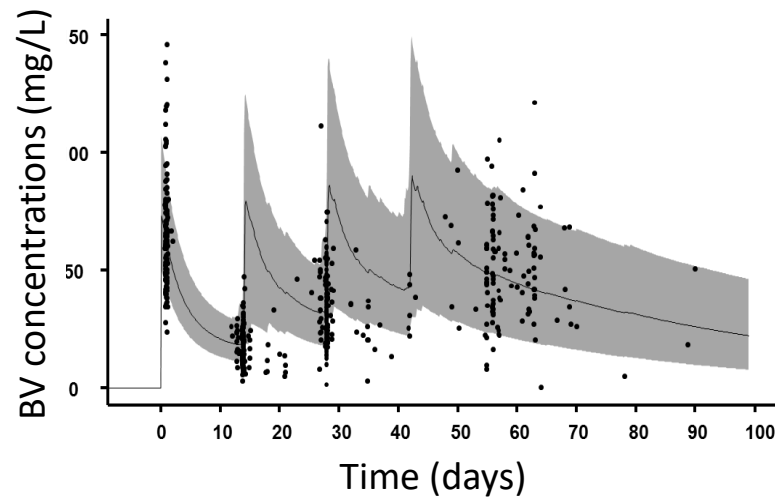
⁴Tours University Hospital, Department of clinical pharmacology, France

⁵Tours University, EA4245 Transplantation, Immunologie, Inflammation (T2i), France

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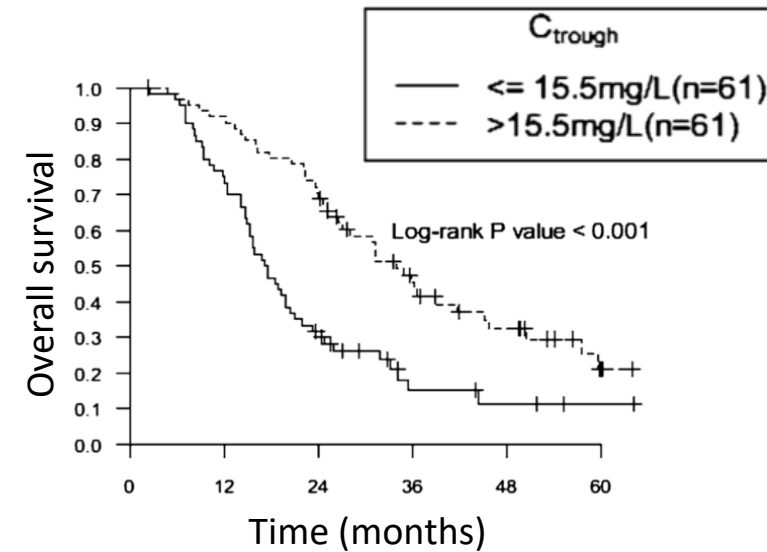
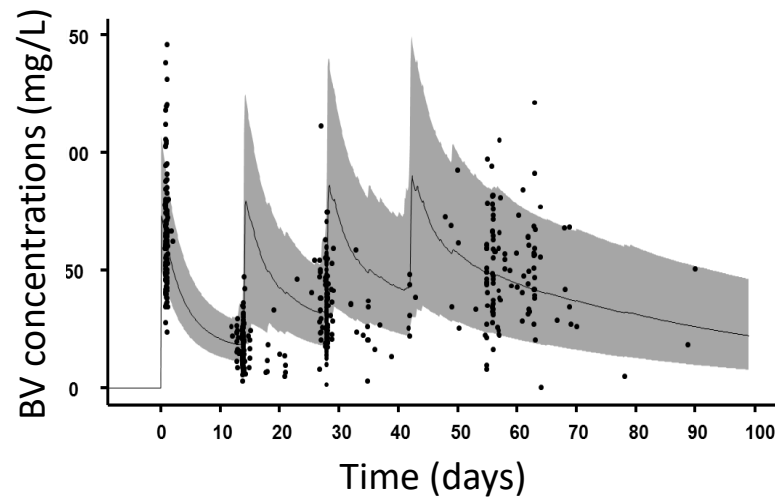
- anti-VEGF monoclonal antibody
- first-line treatment in metastatic colorectal cancer



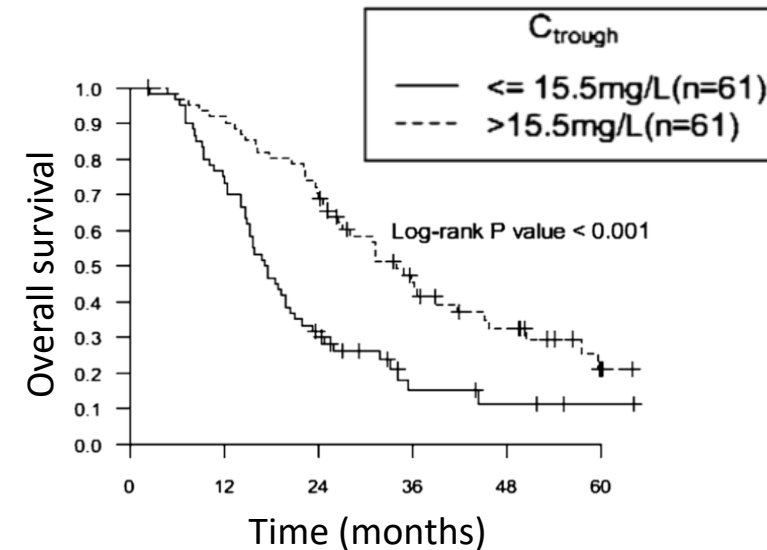
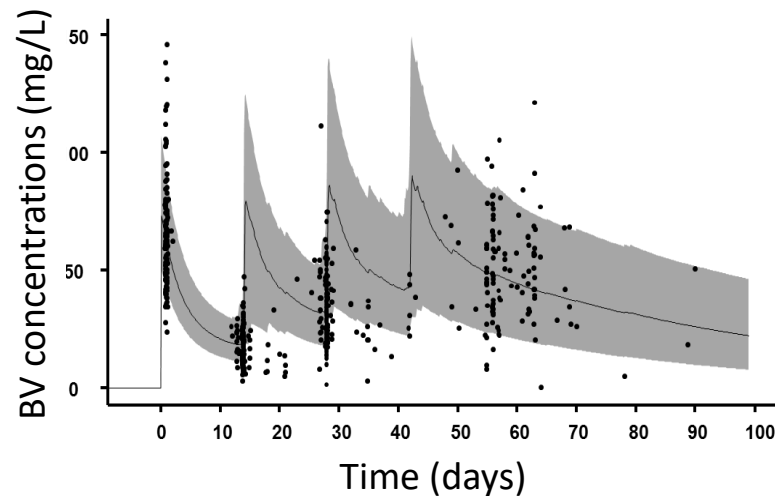
Caulet, 2016

High PK interindividual variability

↗ BV Concentration ⇨ ↗ Survival



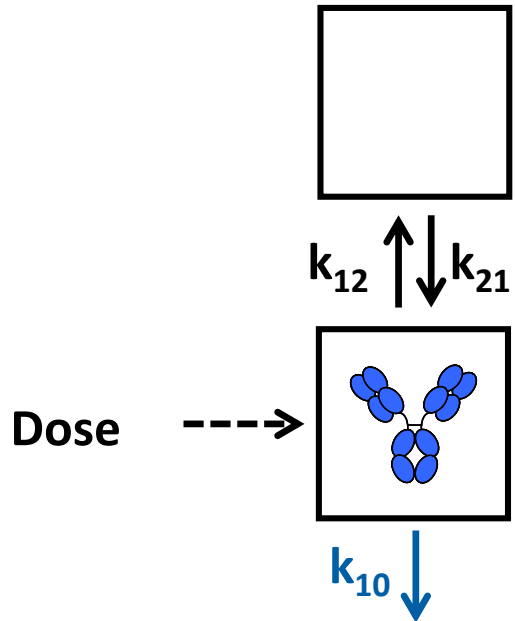
Ref.	Patients	Cancer	Relationship
<i>Lu, 2008</i>	491	Various	✗
<i>Caulet, 2016</i>	130	mCRC	✓
<i>Akbulut, 2018</i>	88	mCRC	✓
<i>Papachristos, 2020</i>	46	mCRC	✓
<i>Peña-Cabia, 2021</i>	28	mCRC	✗



Several confounding factors

- Tumor burden ↗
 - Protein consumption (cachexia)
- } ⇒ CL ↗
} ⇒ mAb «consumption» ↗

Endogenous elimination

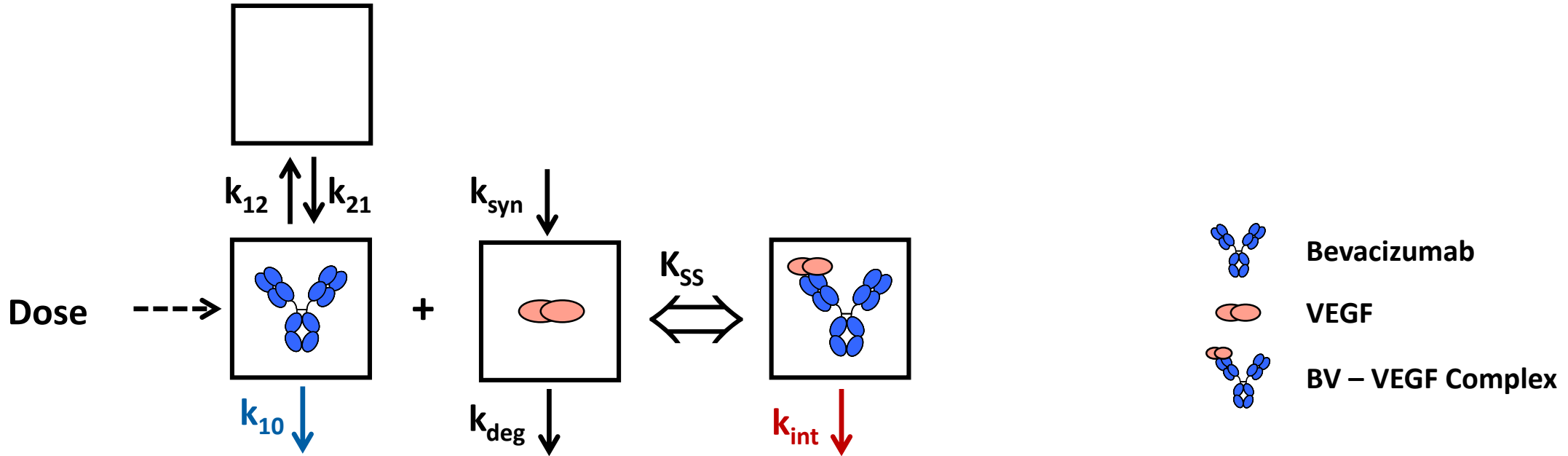


Target-mediated elimination



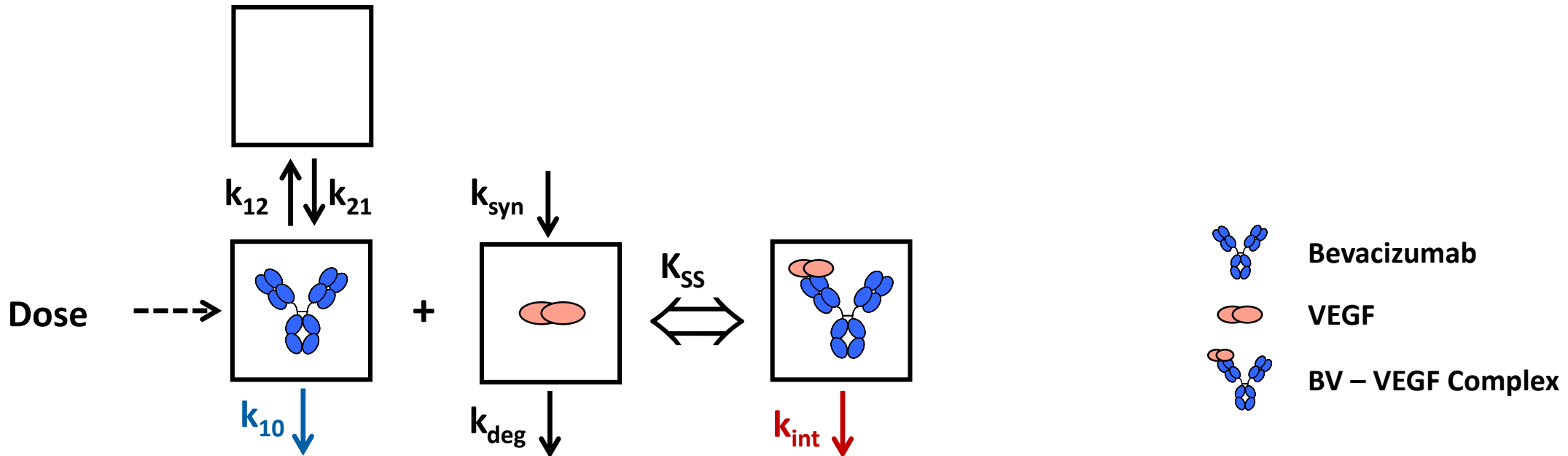
Endogenous elimination

Target-mediated elimination



Endogenous elimination

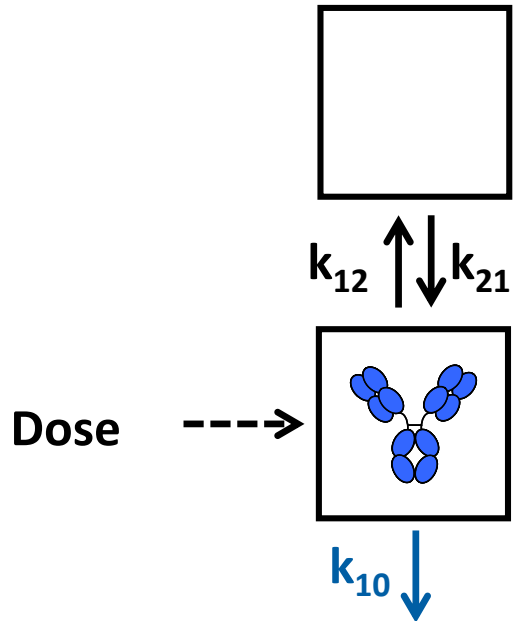
Target-mediated elimination



↗ Target-mediated elimination:

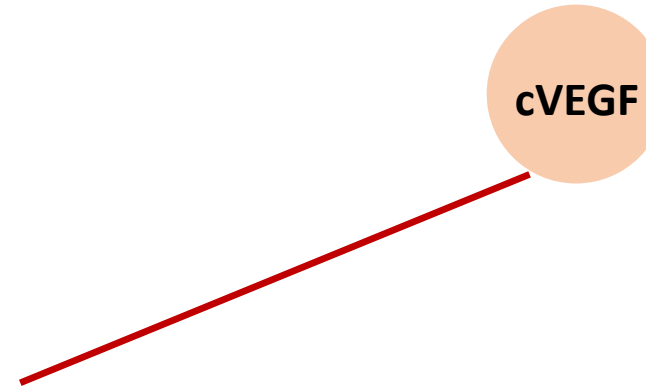
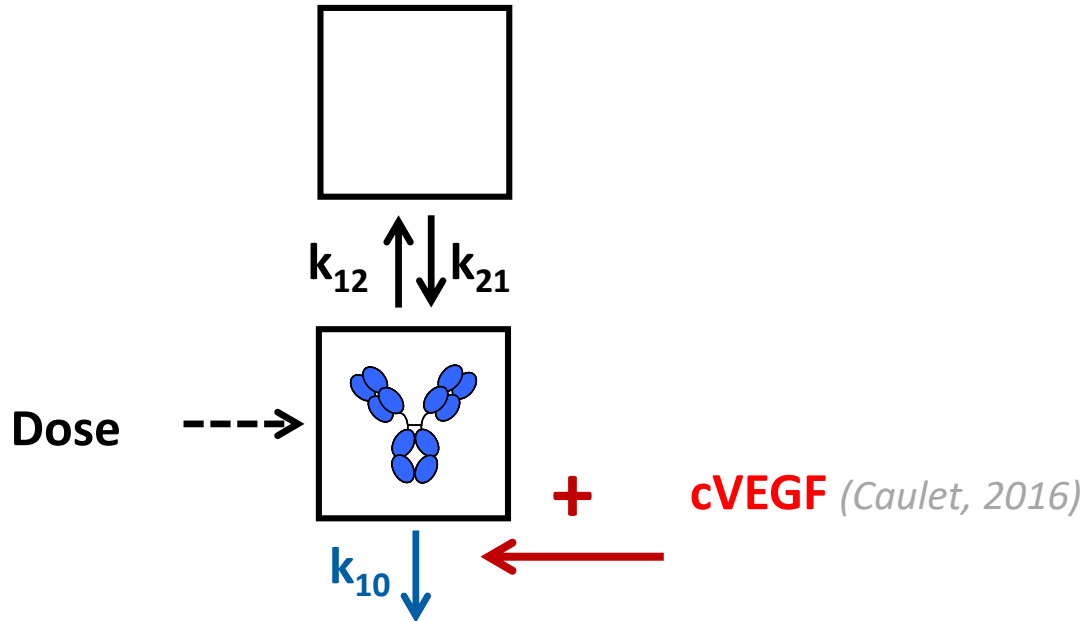
- ↗ target level
- ↘ mAb concentration

} ⇒ Non-linear elimination

Endogenous elimination**Target-mediated elimination**

Endogenous elimination

Target-mediated elimination

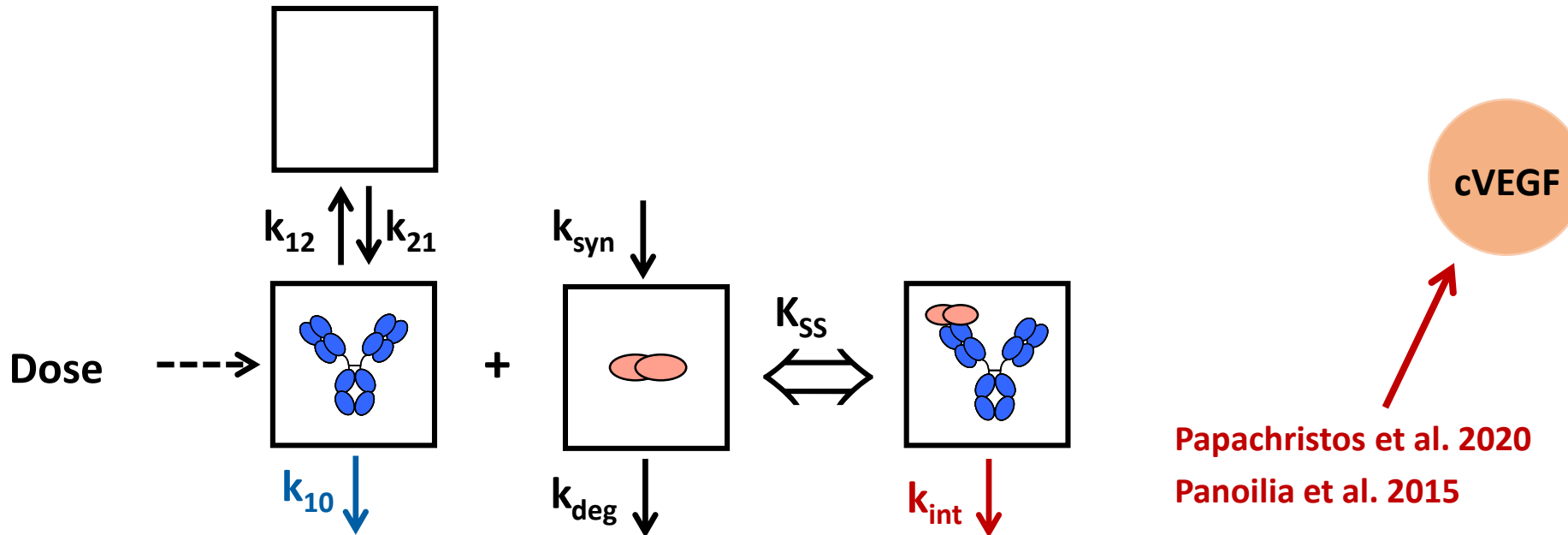


cVEGF = circulating VEGF

↗ cVEGF \Rightarrow ↗ k_{10} \rightarrow Target-mediated elimination

Endogenous elimination

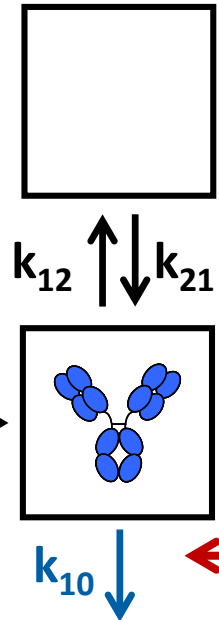
Target-mediated elimination



2 publications with TMDD models – measurements of

- Bevacizumab concentrations
- cVEGF levels

Endogenous elimination



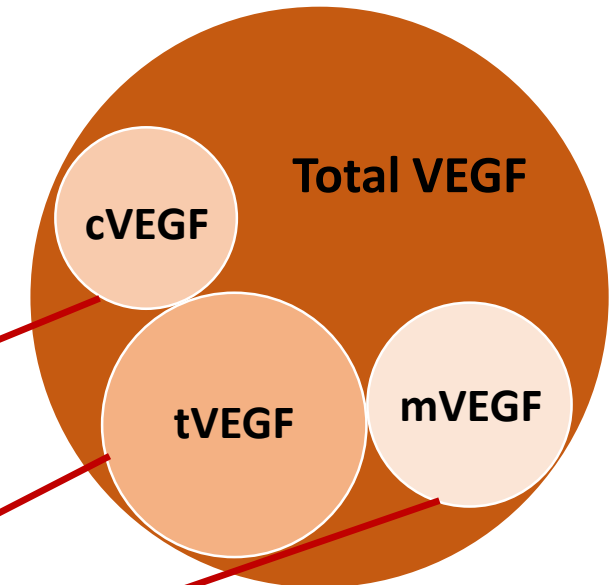
Dose



+

cVEGF (Caulet, 2016)**Tumor type** (Li, 2013; Han, 2015)**Baseline CEA** (Caulet, 2016)**EH metastases** (Caulet, 2016)

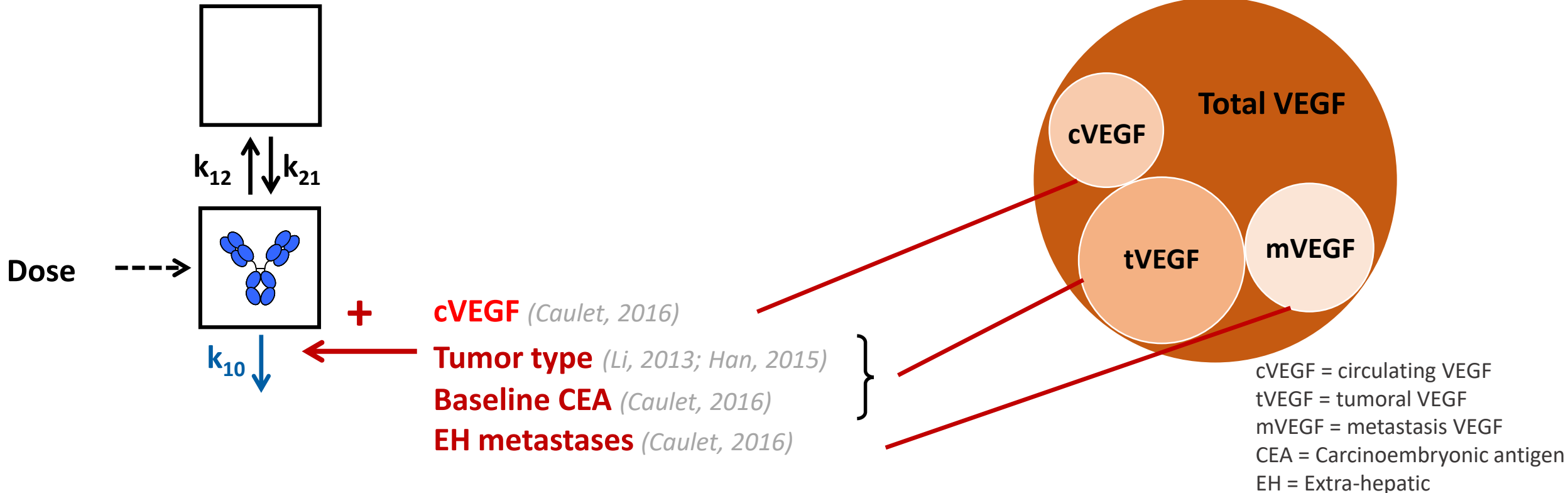
Target-mediated elimination



cVEGF = circulating VEGF
 tVEGF = tumoral VEGF
 mVEGF = metastasis VEGF
 CEA = Carcinoembryonic antigen
 EH = Extra-hepatic

Endogenous elimination

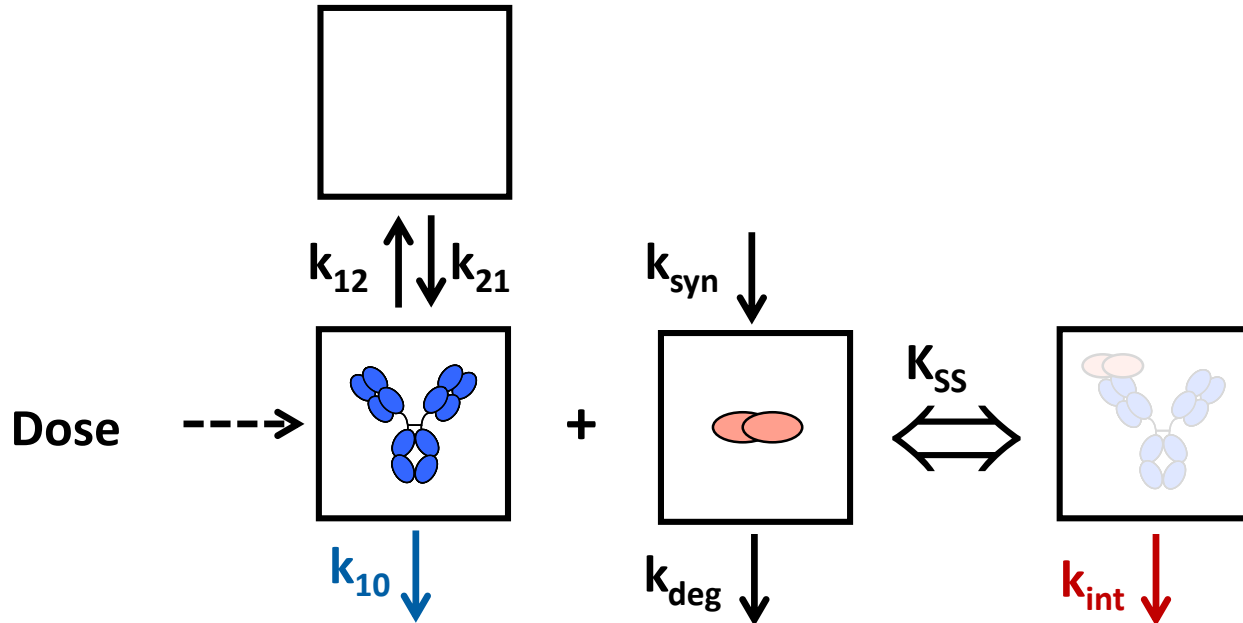
Target-mediated elimination



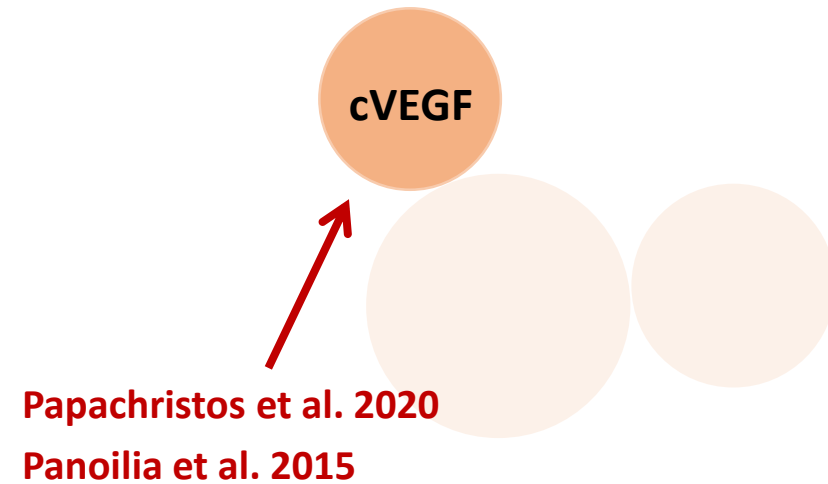
Tumor VEGF > circulating VEGF (Kut, 2007; Martins, 2011)

↗ Tumor burden \Rightarrow ↗ k_{10} \rightarrow Target-mediated elimination

Endogenous elimination

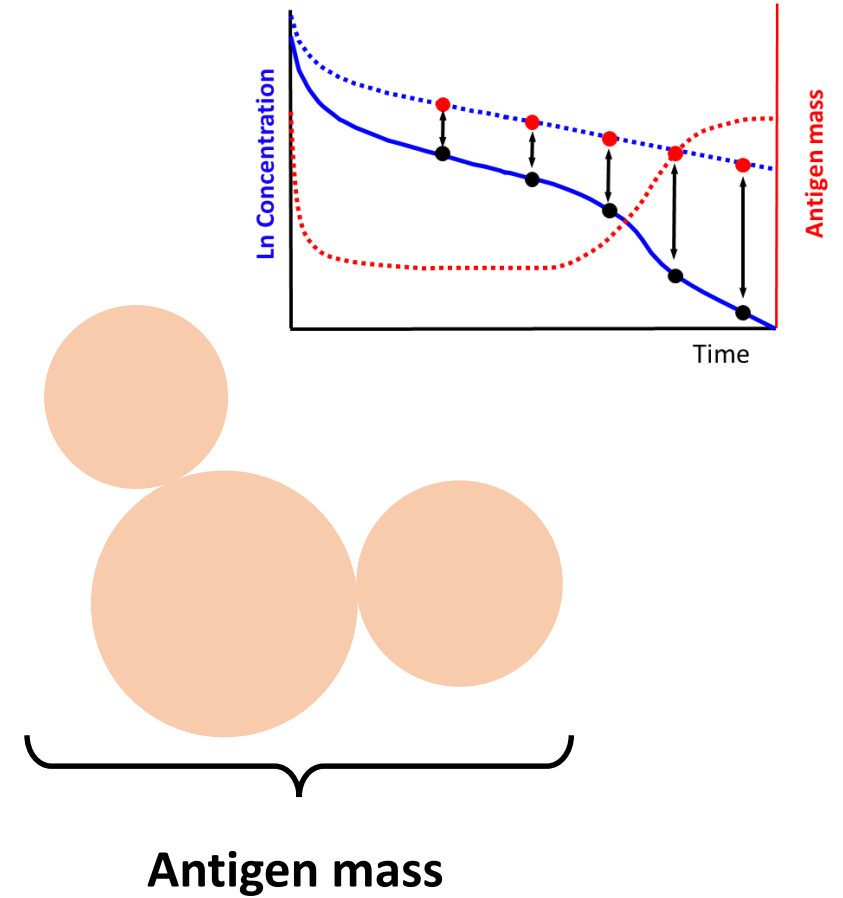
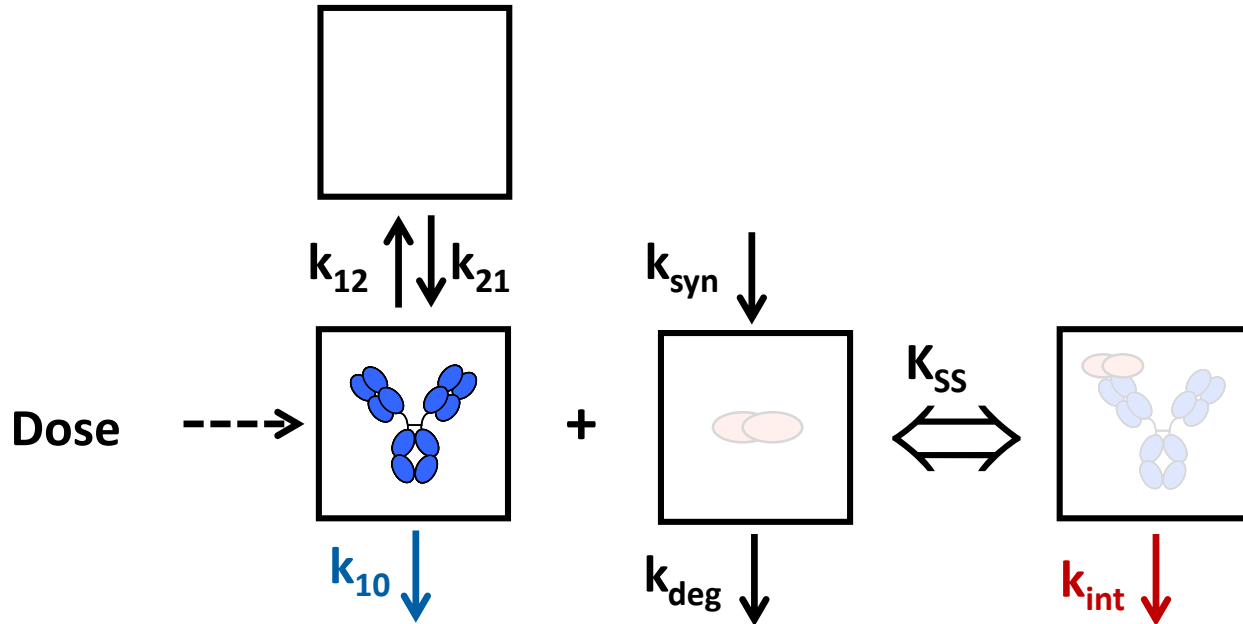


Target-mediated elimination



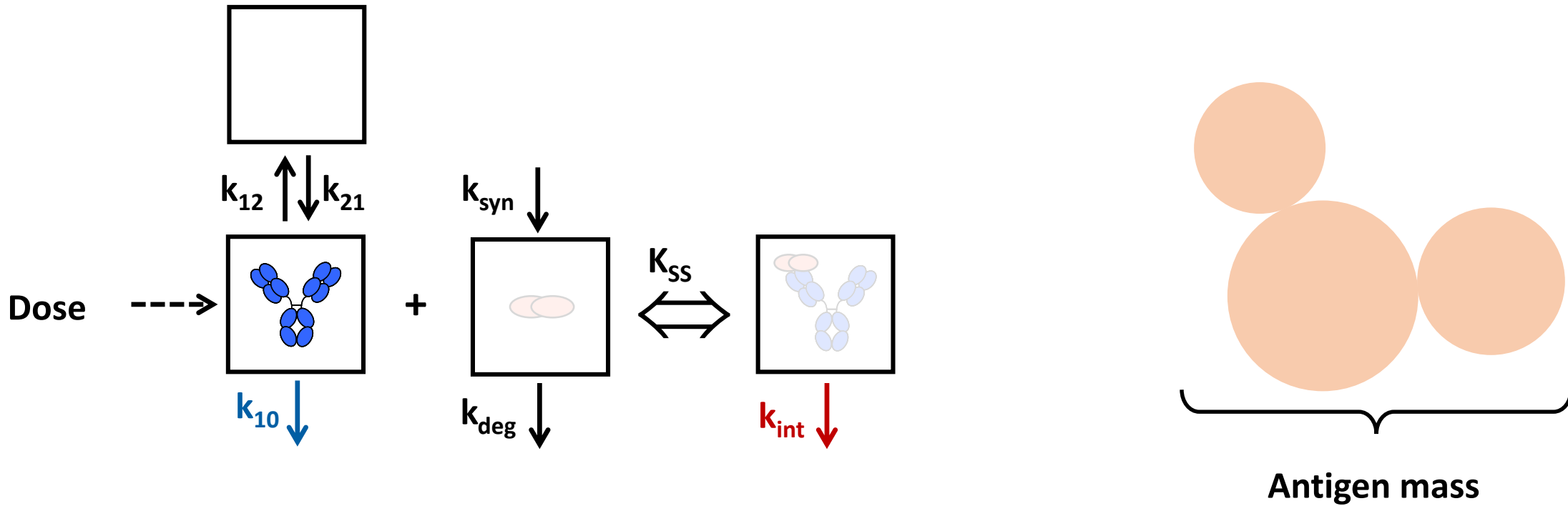
2 publications with TMDD models – measurements of

- Bevacizumab concentrations
- cVEGF levels \Rightarrow ***only part of target amount levels***



Antigen mass

- Total target amount bound to bevacizumab
- Estimated with TMDD model (latent)



- Question 1: is antigen mass related to cVEGF and tumor burden?
- Question 2: is antigen mass associated with survival?

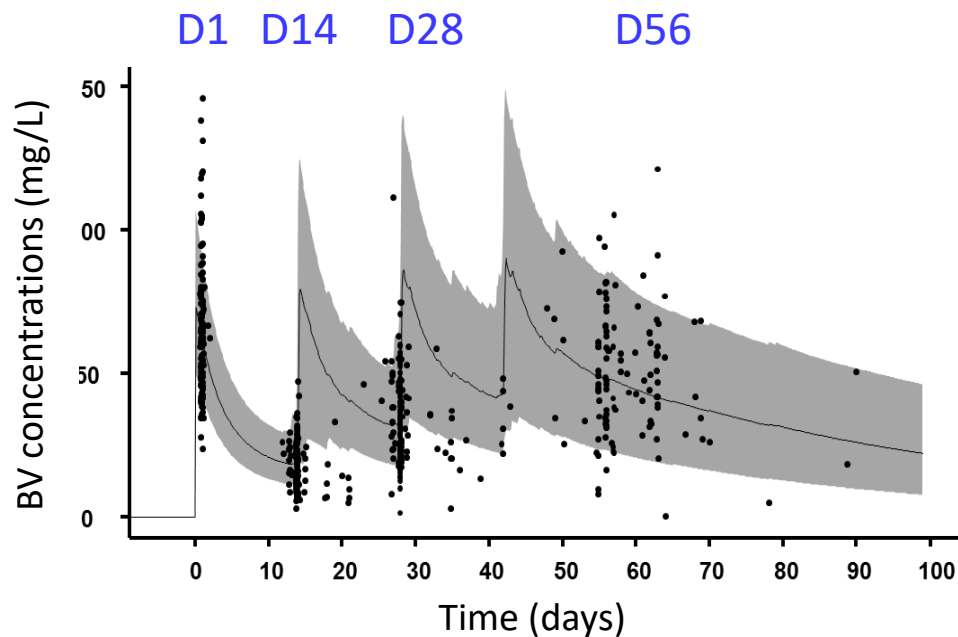
Non-comparative Phase II multicenter study (NCT00489697): *PIs* : Profs. T Lecomte, G Paintaud

137 mCRC patients with liver metastasis

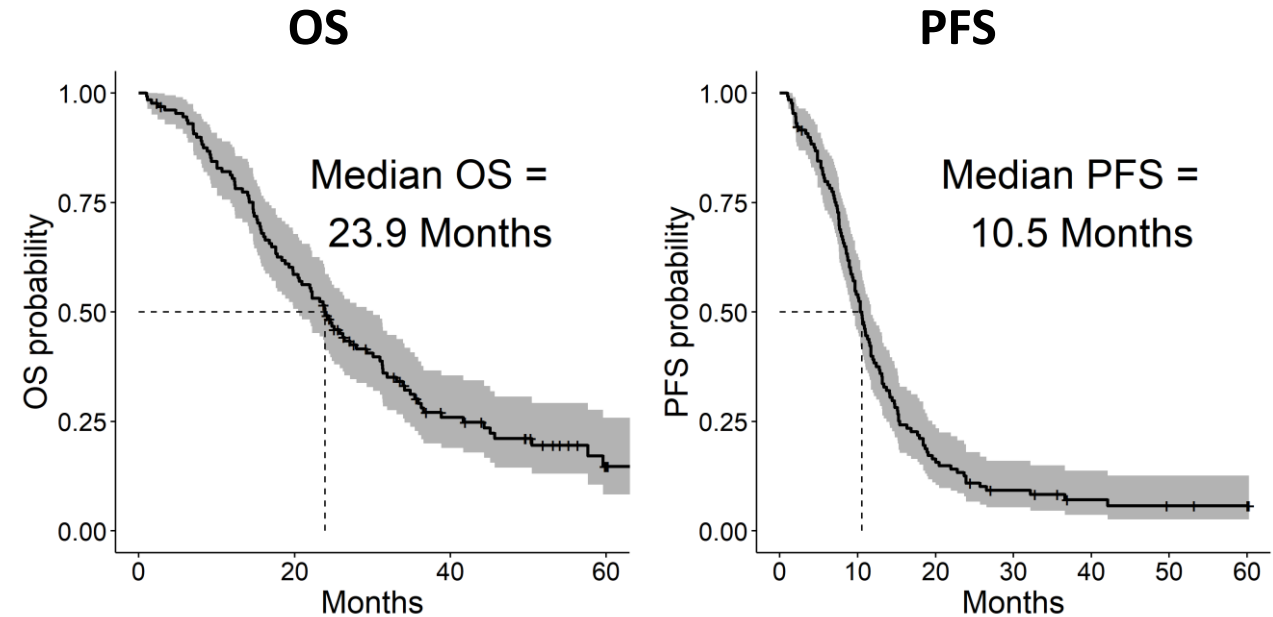
First-line treatment

Bevacizumab + CHEMO
4 cycles 5 mg/kg Q2W

BV concentrations (N=130; n=486)



Survival (N=130)

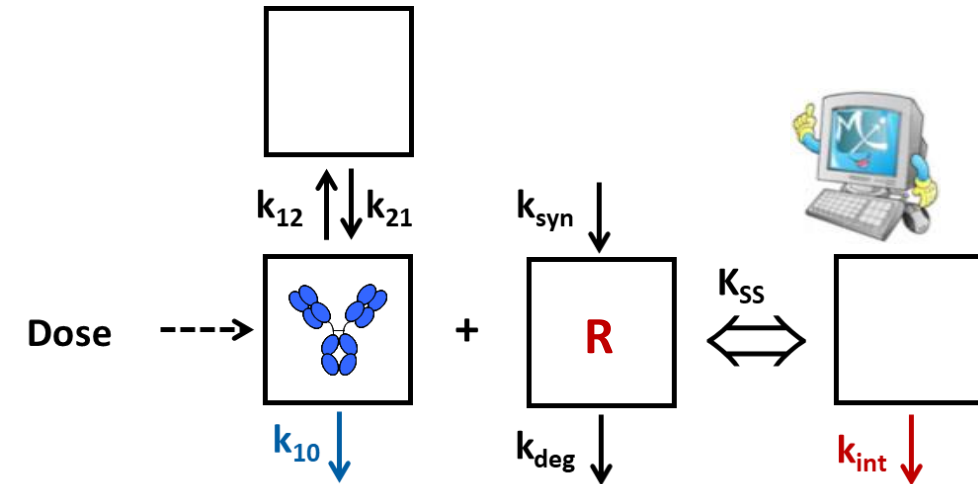


Parameters	Estimates (RSE %)	$\omega_{\%}$ (RSE %)	P*
V_1 , L	4.1 (3.8)	32 (10)	
HT on V_1	3.6 (20)		4.1×10^{-7}
CL, L/day	0.16 (7.8)	30 (18)	
HT on CL	4.0 (26)		6.5×10^{-5}
V_2 , L	4.7 (6.8)	-	-
Q, L/day	1.0 (0.28)	-	-
R_0 , nM	8.4 (14)	29 (19)	
cVEGF on R_0	0.34 (40)		1.2×10^{-2}
CEA on R_0	0.072 (37)		6.6×10^{-3}
mEH on R_0	0.20 (50)		4.7×10^{-2}
k_{int} , day ⁻¹	0.56 (5.8)	-	-
k_{deg} , day ⁻¹	0.96 (0.57)	-	-
K_{SS} , nM	12 (24)	-	-
σ_{prop} , %	21 (4,3)		

*based on Wald test

R_0 = initial target amount

$$R_0 = \frac{k_{syn}}{k_{deg}}$$



TMDD model, QSS approximation

$$\frac{dC_T}{dt} = In(t) - \frac{CL}{V_1} \cdot C - \frac{Q}{V_1} \cdot C + \frac{Q}{V_2} \cdot C_P - k_{int} \cdot (C_T - C)$$

$$\frac{dC_P}{dt} = \frac{Q}{V_1} \cdot C - \frac{Q}{V_2} \cdot C_P$$

$$\frac{dR_T}{dt} = k_{syn} - k_{deg} \cdot (R_T - C_T + C) - k_{int} \cdot (C_T - C)$$

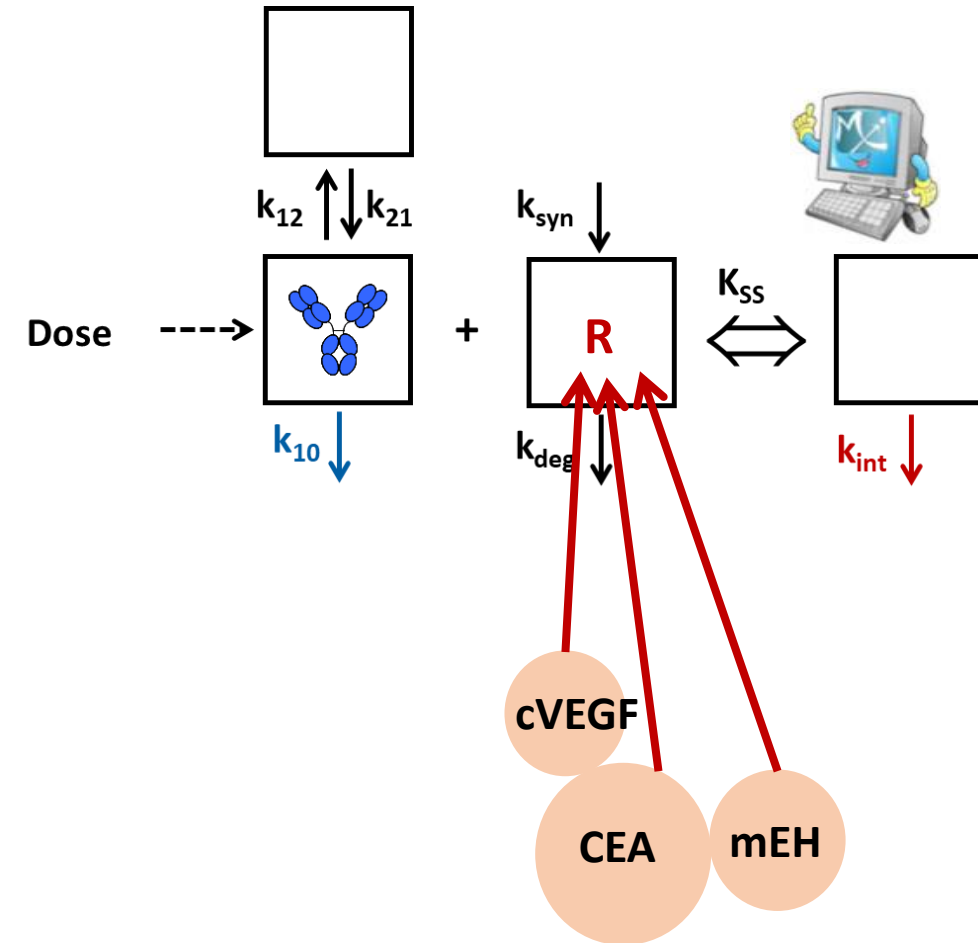
$$C = \frac{1}{2} \left[(C_T - R_T - K_{SS}) + \sqrt{(C_T - R_T - K_{SS})^2 - 4 \cdot K_{SS} \cdot C_T} \right]$$

Parameters	Estimates (RSE %)	$\omega_{\%}$ (RSE %)	P*
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k_{deg} , day ⁻¹	0.96 (0.57)	-	-
K_{SS} , nM	12 (24)	-	-
σ_{prop} , %	21 (4,3)		

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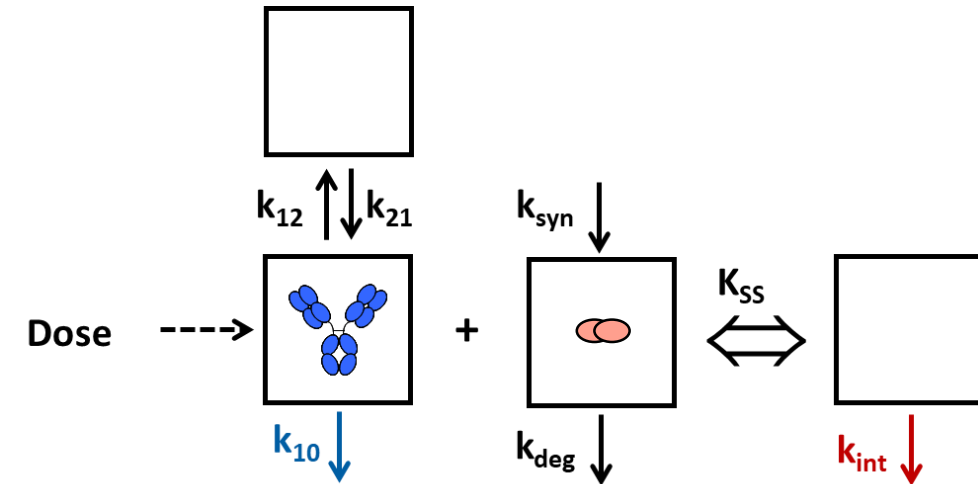


Parameters	Estimates	$\omega_{\%}$	Panoilia, 2015		Papachristos, 2020	
			Estimates	$\omega_{\%}$	Estimates	$\omega_{\%}$
V_1 , L	4.1	32	3.2	22	5.8	17
HT on V_1	3.6					
CL, L/day	0.16	30	0.18	20	0.34	31
HT on CL	4.0					
V_2 , L	4.7	-	3.1	-	3.2	56
Q, L/day	1.0	-	1.4	-	0.14	20
R_0, nM	8.4	29	0.0053	33	0.014	24
cVEGF on R_0	0.34					
CEA on R_0	0.072					
mEH on R_0	0.20					
k_{int} , day ⁻¹	0.56	-	0.056*	-	0.059*	-
k_{deg} , day ⁻¹	0.96	-	0.401	-	0.12	-
K_{SS} , nM	12	-	267	-	135	-
$\sigma_{prop, BV}$, %	21		28		25	
$\sigma_{prop, VEGF}$, %			32		29	

R_0 = initial target amount

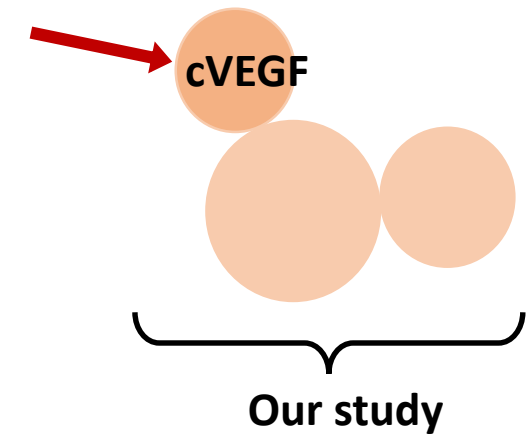
$$R_0 = \frac{k_{syn}}{k_{deg}}$$

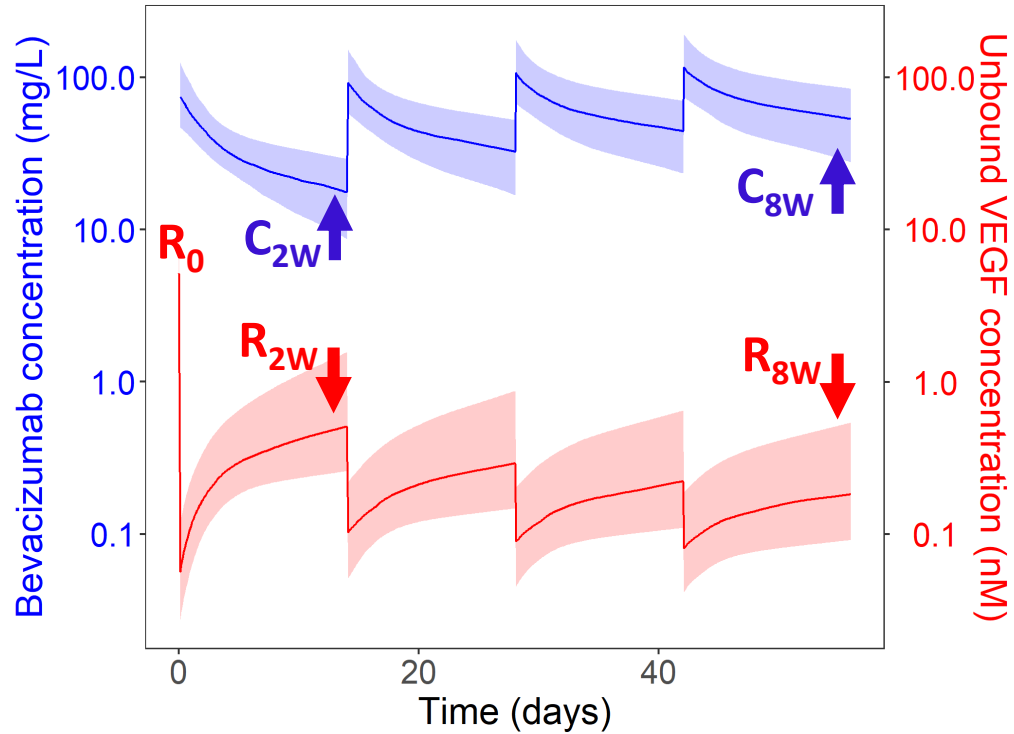
*fixed to k_{10} value



Papachristos, 2020

Panoilia, 2015

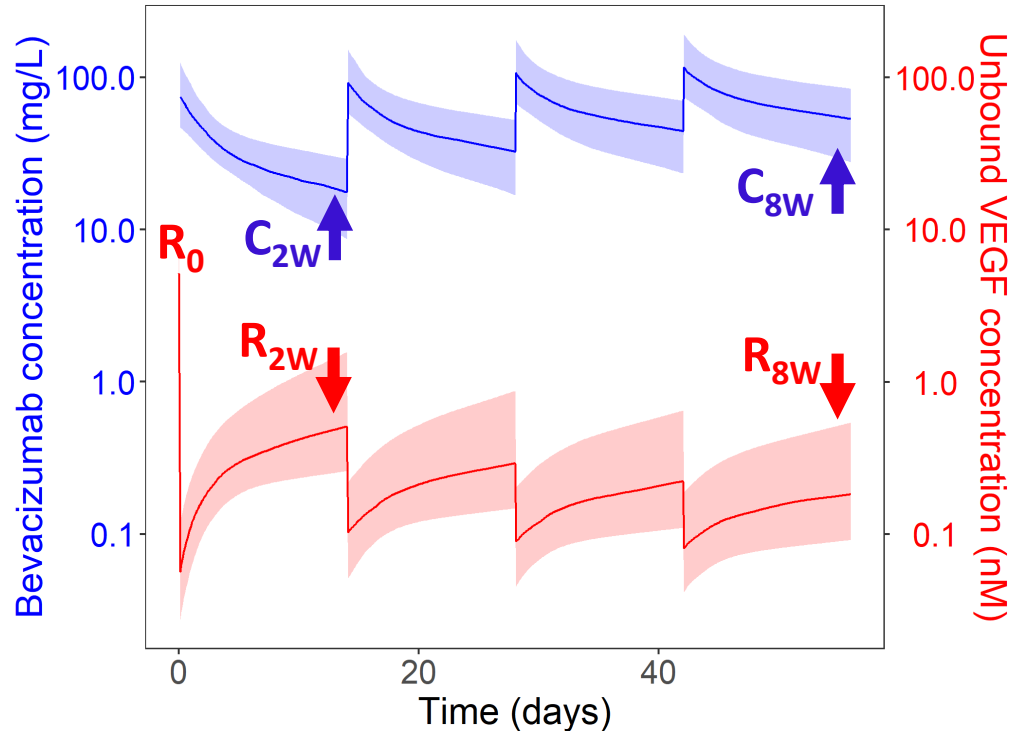


Bevacizumab, mCRC, 5 mg/kg Q2W

R_0 = initial target amount

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Bevacizumab, mCRC, 5 mg/kg Q2W



Cox Analysis (n = 130)

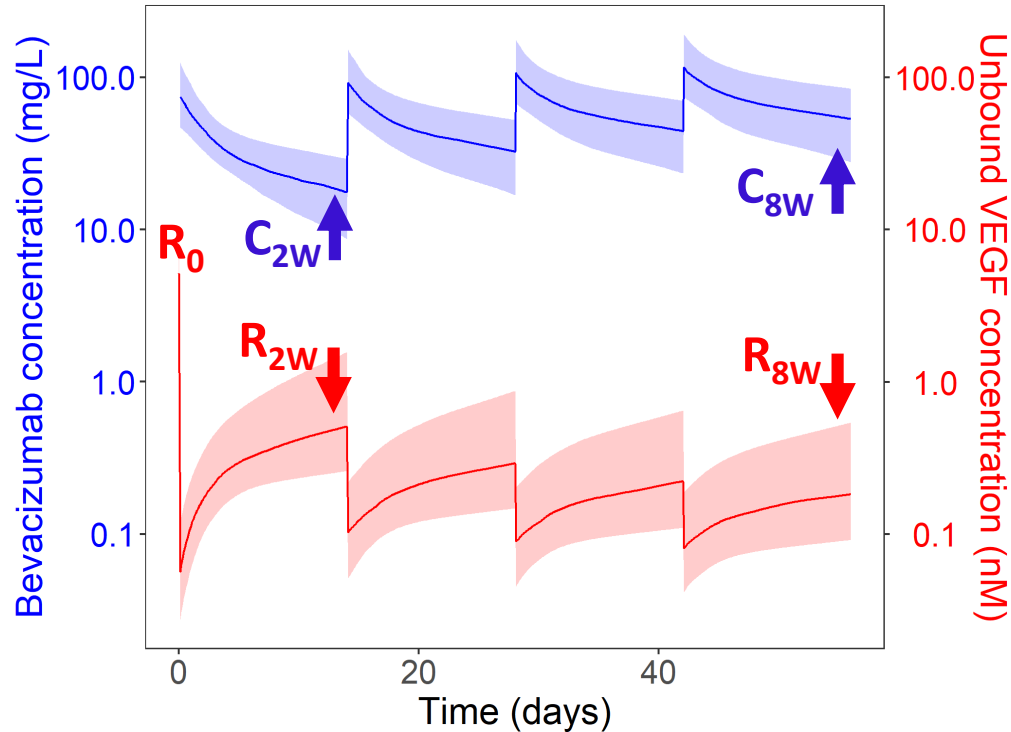
	Progression free survival			Overall survival		
	HR	95% CI	<i>P</i>	HR	95% CI	<i>P</i>
R _{2w}	1.8	1.2-2.6	0.0022	2.3	1.5-3.5	9.2x10 ⁻⁵
R ₀	1.8	1.2-2.6	0.0028	2.1	1.4-3.1	5.6x10 ⁻⁴
R _{8w}	1.7	1.2-2.5	0.005	2.1	1.4-3.2	0.0007
C _{2w}	0.70	0.49-1.0	0.06	0.57	0.37-0.85	0.0068
C _{8w}	0.75	0.51-1.1	0.13	0.68	0.45-1.0	0.073

**all parameters were dichotomized around the median value*

PFS and OS are most strongly associated with

- Target involvement (R_{2w})
- Baseline antigen mass (R₀)

Bevacizumab, mCRC, 5 mg/kg Q2W



Cox Analysis (n = 130)

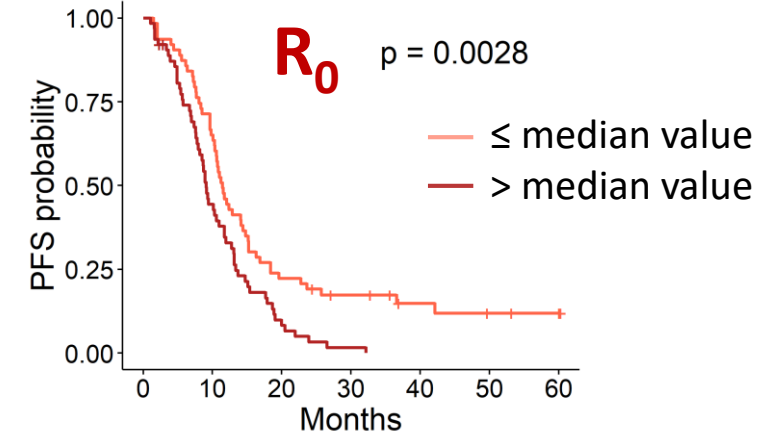
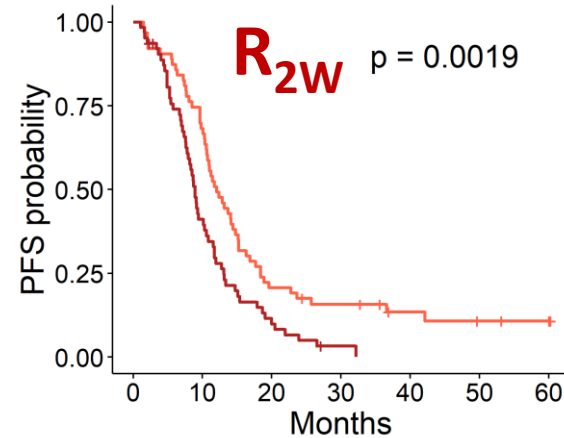
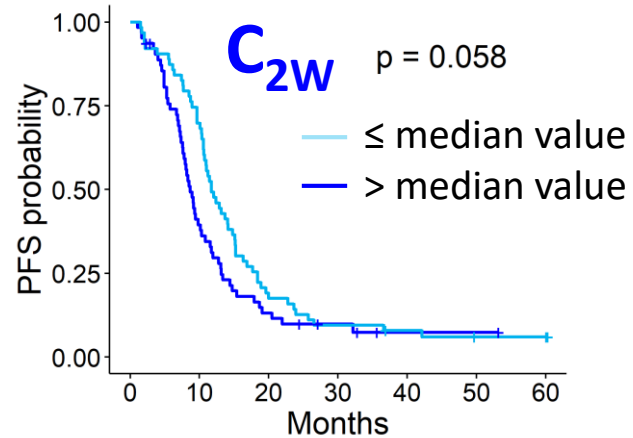
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R_0	1.8	1.2-2.6	0.0028	2.1	1.4-3.1	5.6×10^{-4}
R_{8w}	1.7	1.2-2.5	0.005	2.1	1.4-3.2	0.0007
C_{2w}	0.70	0.49-1.0	0.06	0.57	0.37-0.85	0.0068
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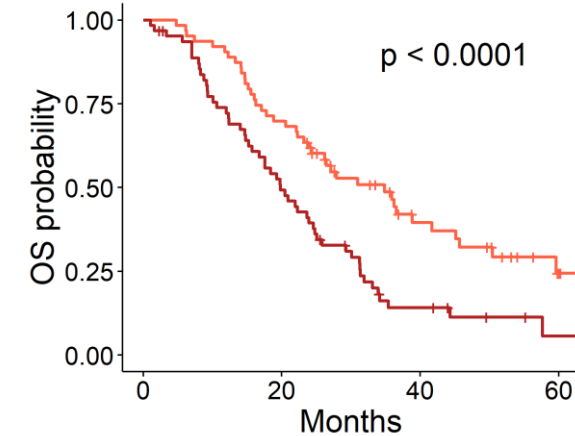
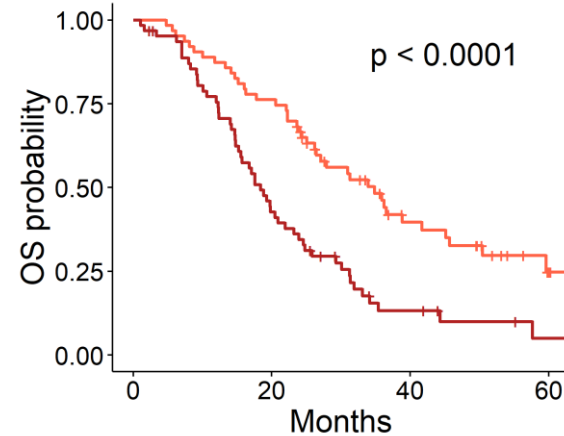
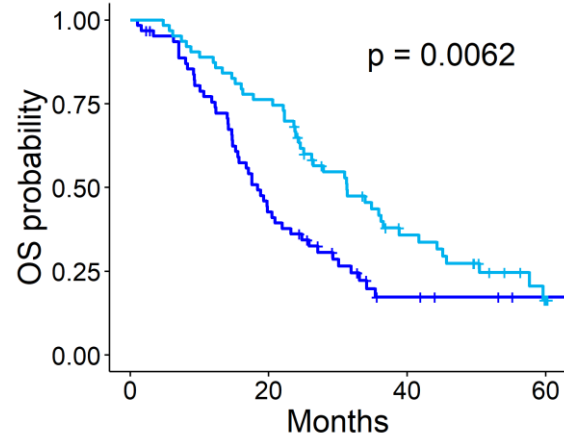
PFS and OS are most strongly associated with

- Target involvement (R_{2w})
- Baseline antigen mass (R_0)

PFS

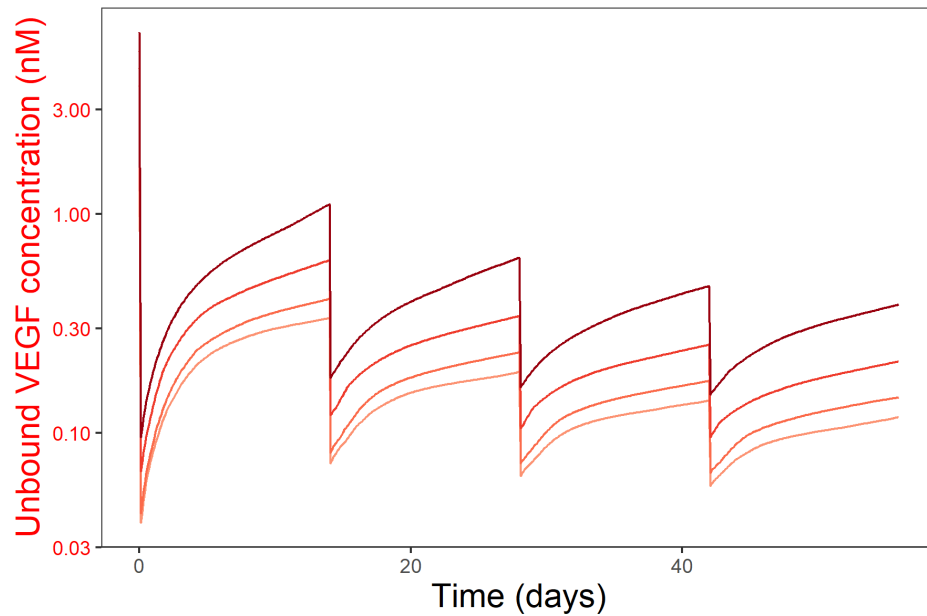
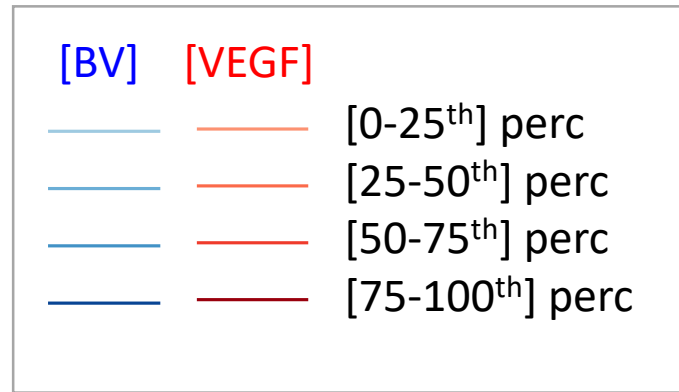
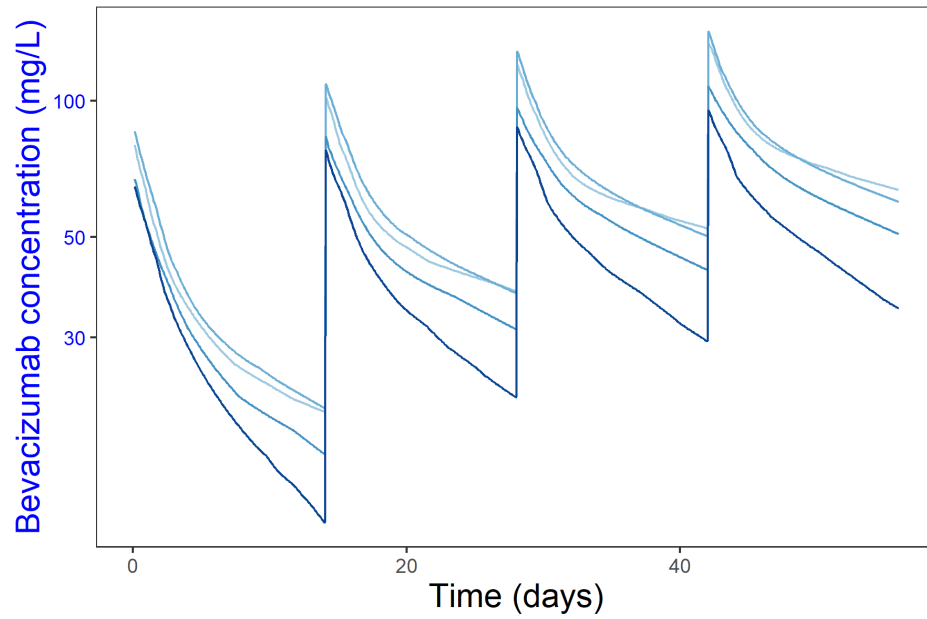


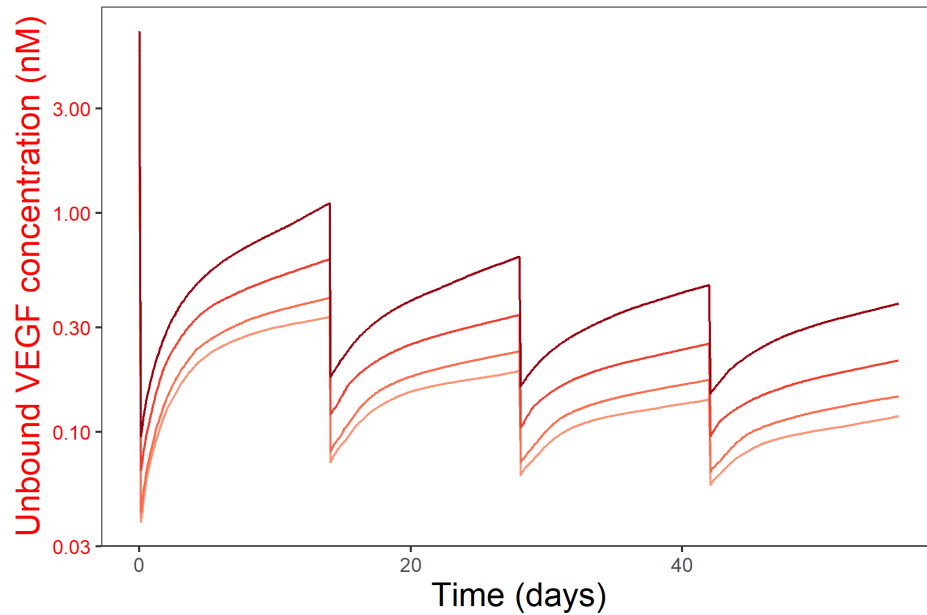
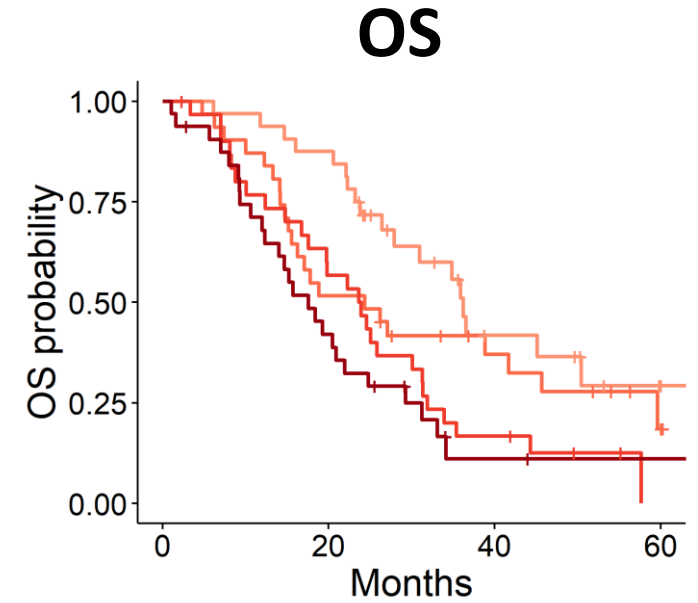
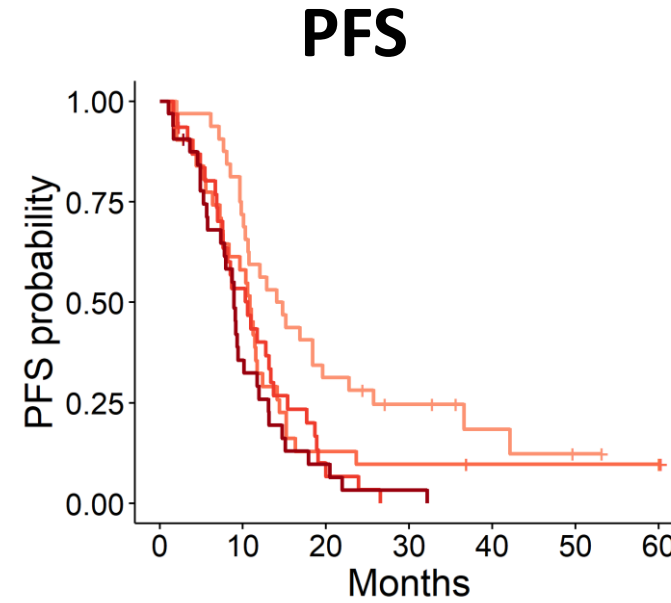
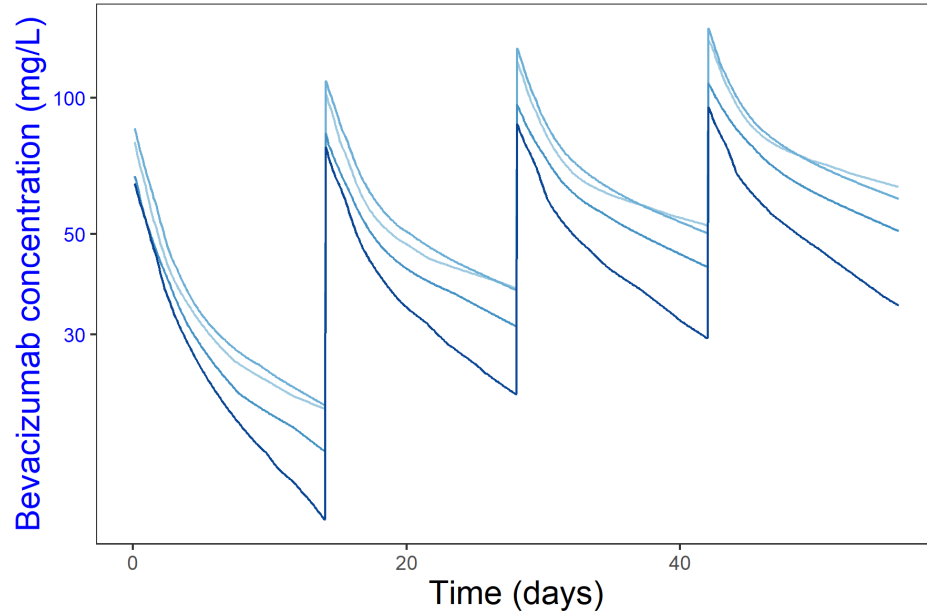
OS



Short PFS and OS are associated with

- Low C_{2W} \Rightarrow High BV « consumption » *(as found by Caulet, 2016)*
- High R_{2W} \Rightarrow Low target involvement ?
- High R_0 \Rightarrow High target expression





$R_{0\ cat}$	Median PFS	Median OS
[0-25 th] perc	14.4	36.3
[25-50 th] perc	10.9	24.4
[50-75 th] perc	10.6	23.8
[75-100 th] perc	8.9	17.6

Latent antigen mass estimated using TMDD modeling

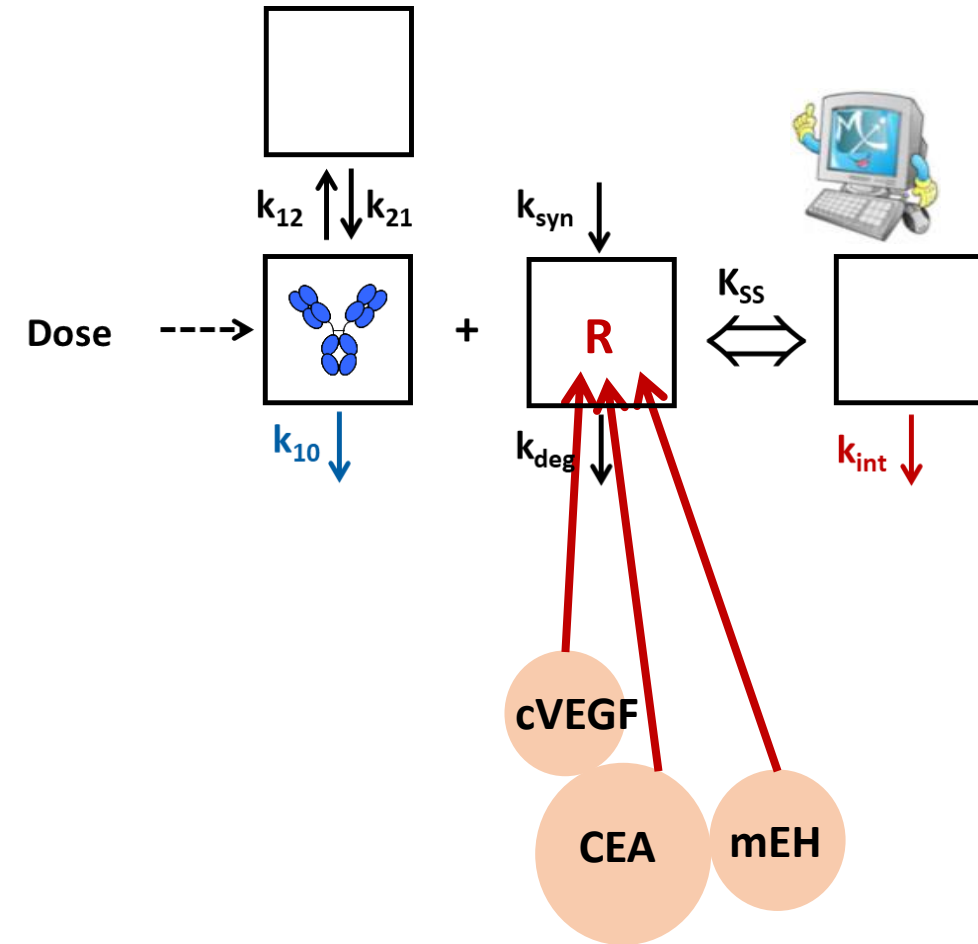
- Using unbound BV concentrations
- Accounts for total target amount bound to BV

Latent antigen mass computed associated with

- Tumor burden and cVEGF
- Progression-free survival and overall survival
 ⇒ Relevant predictor of bevacizumab efficacy ?

Further work needed

- ⇒ Refine relationships of both target involvement and sarcopenia/cachexia with clinical efficacy
- ⇒ Demonstrate benefit of dosing intensification in patient with poor target involvement



Acknowledgments



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Dr. Stéphane Servais
Dr. William Raoul



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Dr. Nicolas Azzopardi



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Dr. David Ternant
Prof. Theodora Bejan-Angoulvant
Dr. Céline Desvignes



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Caroline Guerineau



Thanks for your attention