

ABP 215 – Bevacizumab Biosimilar Candidate

FDA Advisory Committee

13 July 2017

Introduction

Richard Markus, MD, PhD

VP Global Development, Amgen

Agenda

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Richard Markus, MD, PhD
Global Development, Amgen

Analytical Similarity

Simon Hotchin
Regulatory Affairs, Amgen

Non-Clinical and Clinical Similarity and Extrapolation to All Indications

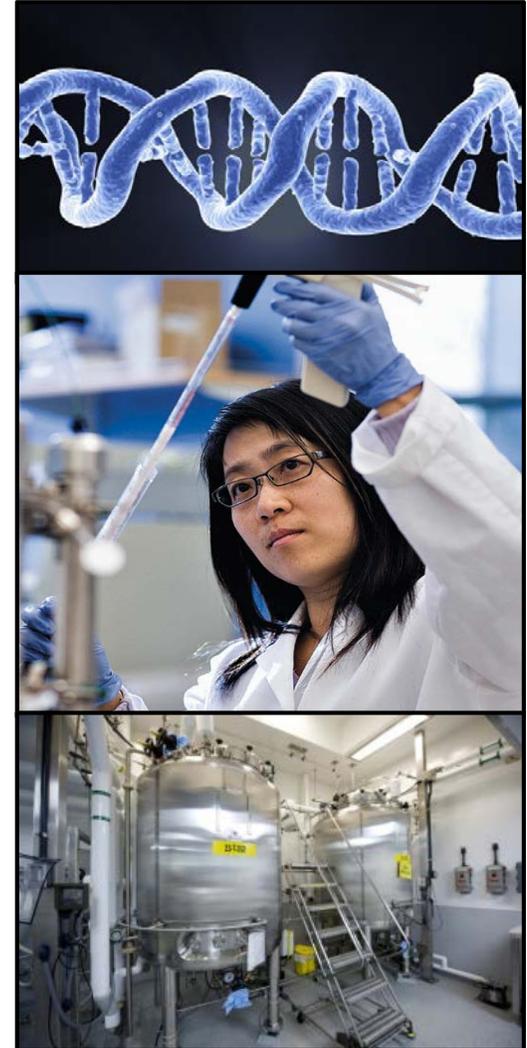
Richard Markus, MD, PhD
Global Development, Amgen

Conclusion

Lisa Bollinger, MD
Regulatory Affairs and Safety, Amgen

Amgen: A Biotechnology Pioneer

- ◆ More than 35 years as innovative leader
- ◆ Capability to discover, develop, and manufacture complex biologics
- ◆ Broad pipeline of innovative medicines, and also biosimilars
- ◆ Same scientists and laboratories to develop our biosimilars
- ◆ Same manufacturing network and quality systems to produce our biosimilars



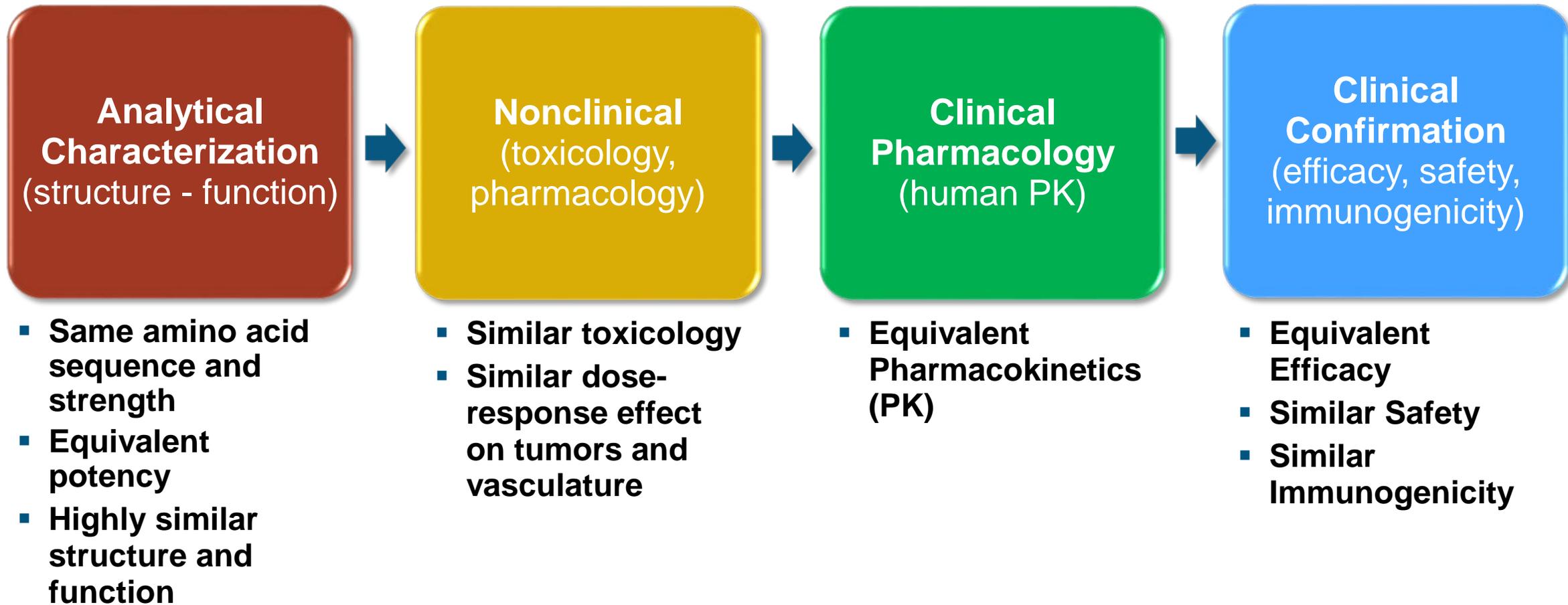
Bevacizumab and ABP 215 Mechanism of Action

- ◆ **VEGF is secreted by tumors**
- ◆ **Products bind and neutralize VEGF**

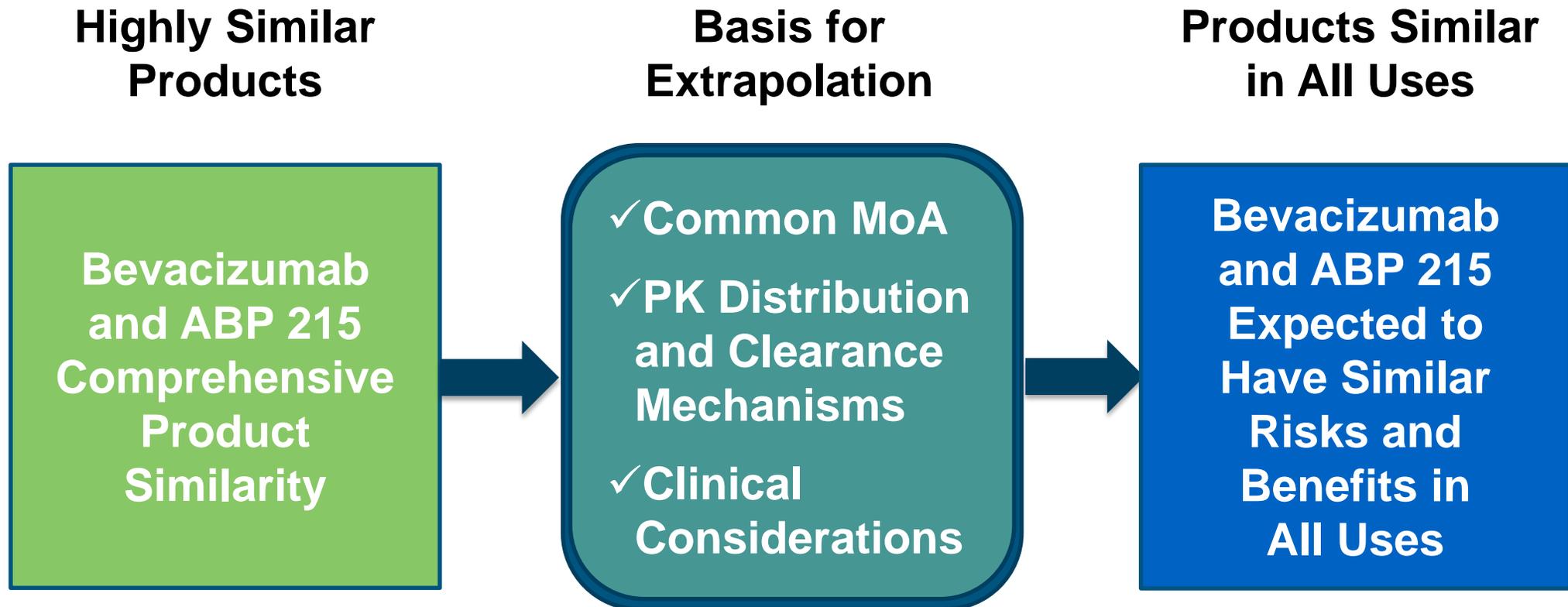
ABP 215

| A proposed biosimilar
of bevacizumab

Biosimilar Development and Approval Is Based on the Totality of Evidence



Biosimilar Extrapolation of Indications



ABP 215 Proposed Indications of Use

- ◆ **Nonsquamous Non-small Cell Lung Cancer**
- ◆ **Metastatic Colorectal Cancer**
- ◆ **Metastatic Renal Cell Carcinoma**
- ◆ **Glioblastoma**
- ◆ **Cervical Cancer**

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Analytical Similarity to Bevacizumab

Simon Hotchin

Executive Director, Regulatory Affairs, Amgen

Overview

- ◆ **ABP 215 development**
- ◆ **Design of the analytical similarity assessment**
- ◆ **Analytical similarity results and conclusions**
 - Structural and purity attributes
 - Functional activities

ABP 215 Development

ABP 215 Product and Process Design

◆ Cell Line Development

- Amgen carefully designed the ABP 215 cell line, screening a large number of clones
- Matched amino acid sequence and critical attributes of the reference product

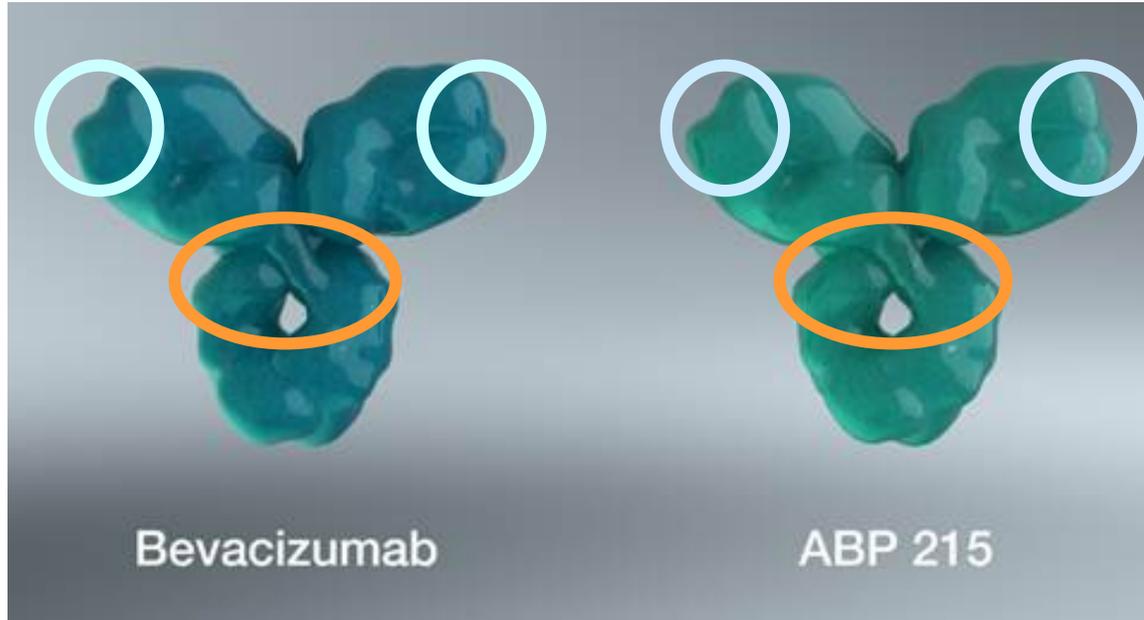
◆ Manufacturing Process and Formulation

- Process developed with multiple controls to ensure similarity
- Commercial manufacturing process established at the outset of development
- Drug product presentations and formulation match the reference product



Design of the Analytical Similarity Assessment

Design Considerations: Fab and Fc Domain Characterization



- ◆ The mechanism of action for ABP 215 and bevacizumab in all indications is to bind and neutralize VEGF
- ◆ The Fab domain binds all isoforms of VEGF

- ◆ The Fc domain binds Fc receptors, as well as complement component C1q in-vitro
- ◆ Fc-mediated effector functions do not occur for these products
- ◆ Tested for confirmation of structural similarity

Design Considerations: Analytical Similarity Assessment Criteria

Considerations and Criteria

Attributes with the highest risk to clinical outcomes and includes assay(s) that evaluate clinically relevant primary mechanism(s) of action

Similarity if 90% CI around difference in means is within $\pm 1.5 \times \text{SD}$ of US reference product

Attributes with relatively lower risk to clinical outcomes

Similarity if at least 90% of the ABP 215 lots are within mean $\pm 3 \times \text{SD}$ of the US reference product

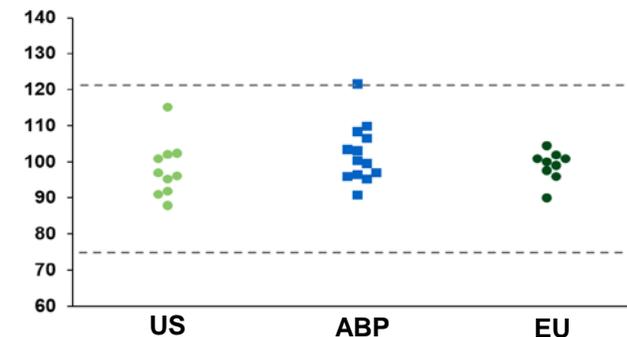
Attributes with the lowest risk to clinical outcomes

Qualitative comparisons

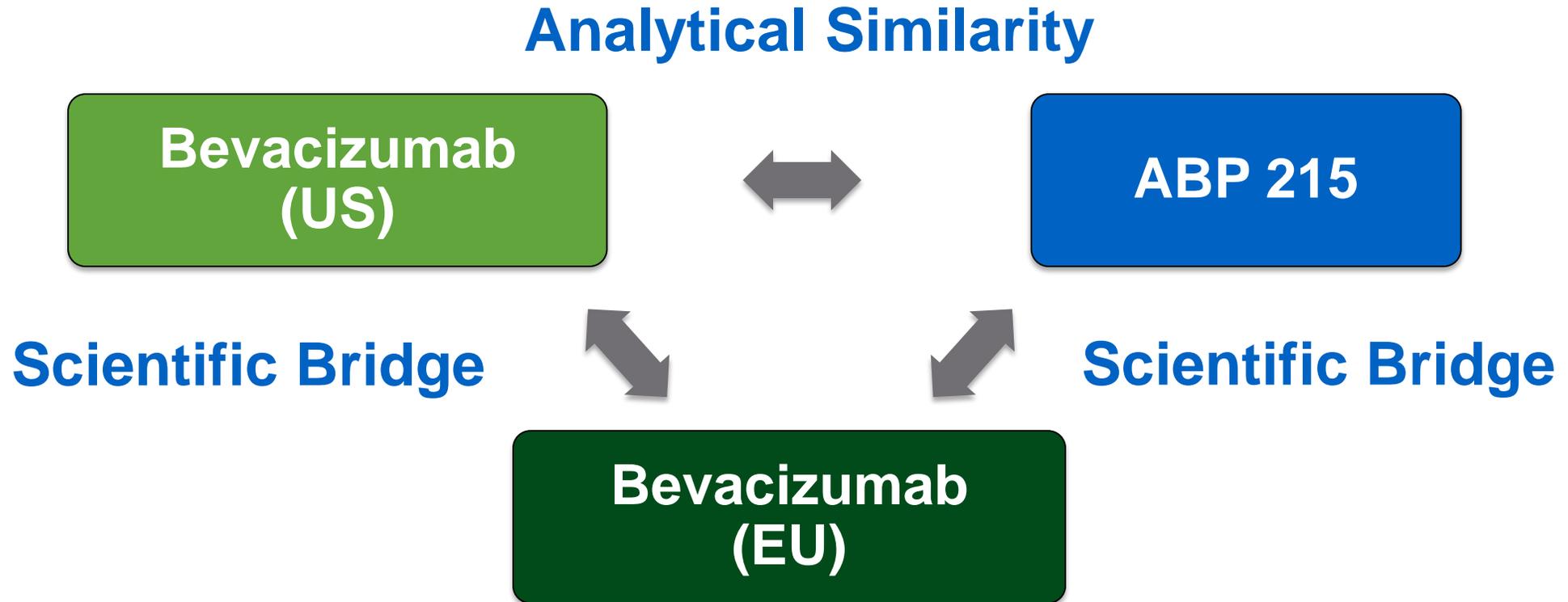
Results



EAC = equivalence acceptance criteria



Design Considerations: Establishing the Scientific Bridge



**Analytical and PK Similarity Data Established
the Required Scientific Bridge**

Analytical Similarity of ABP 215 to Bevacizumab Was Demonstrated for ~ 100 Attributes/Assays

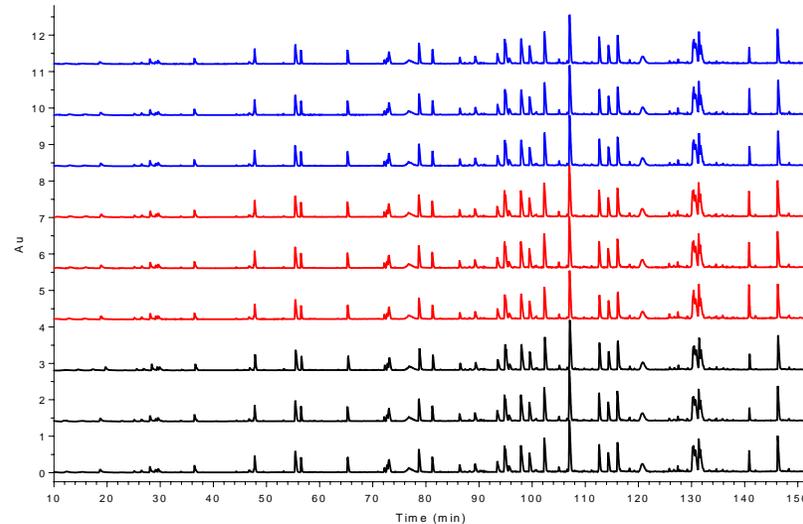
	Analytical Testing/Attributes		Analytical Testing/Attributes		Analytical Testing/Attributes				
Biological Activity	Inhibition of proliferation in HUVEC	Higher Order Structure	FTIR: Spectral similarity	Thermal Degradation @ 50°C	SE-HPLC: HMW				
	Binding to VEGF by ELISA		FTIR: Profile		rCE-SDS: HC + LC				
	On and off rates (VEGF)		Near UV CD: Spectral similarity		rCE-SDS: LMW + MMW				
	Binding to VEGF isoforms		Near UV CD: Profile		CEX-HPLC: Main peak				
	Inhibition of VEGFR-2 RTK autophosphorylation		DSC: T _m 1		CEX-HPLC: Acidic peaks				
	Specificity for VEGF by VEGFR-2 RTK autophosphorylation		DSC: T _m 2		CEX-HPLC: Basic peaks				
	FcRn binding		DSC: Profile		Proliferation inhibition bioassay				
	FcγRIa binding		Particles and Aggregates		SE-HPLC: HMW	Thermal Stability @ 40°C	SE-HPLC: HMW		
	FcγRIIa (131H) binding				HIAC: ≥ 2 μm particles		rCE-SDS: HC + LC		
	FcγRIIb binding				HIAC: ≥ 5 μm particles		rCE-SDS: LMW + MMW		
	FcγRIIIa (158V) binding				HIAC: ≥ 10 μm particles		CEX-HPLC: Main peak		
	FcγRIIIa (158F) binding				HIAC: ≥ 25 μm particles		CEX-HPLC: Acidic peaks		
	FcγRIIIb binding				MFI: ≥ 5 μm particles		CEX-HPLC: Basic peaks		
	C1q binding				MFI: ≥ 5 μm non-spherical particles		Proliferation inhibition bioassay		
	Lack of ADCC				FFF: Submicron particles		SE-HPLC: HMW	Thermal Stability @ 25°C	SE-HPLC: HMW
	Lack of CDC				DLS (Hydrodynamic Radius)		rCE-SDS: HC + LC		
	Primary Structures				Intact molecular mass		AUC-SV: % Monomer		rCE-SDS: LMW + MMW
Intact molecular mass; profiles		AUC-SV: Profile		CEX-HPLC: Main peak	CEX-HPLC: Acidic peaks				
Reduced and deglycosylated molecular masses of HC and LC		SLS: Molar mass		CEX-HPLC: Basic peaks	CEX-HPLC: Basic peaks				
Reduced and deglycosylated molecular masses of HC and LC; profiles		Product-related Substance and Impurities		Proliferation inhibition bioassay	General Properties		Proliferation inhibition bioassay		
Reduced peptide map				SE-HPLC: HMW			Protein Concentration		
Reduced peptide map; profiles				SE-HPLC: Profile			Volume		
Non-reduced peptide map				rCE-SDS: HC+LC			Osmolality		
Non-reduced peptide map; profiles				rCE-SDS: NGHC			pH		
Glycan Map; % high mannose			rCE-SDS: LMW + MMW	Appearance					
Glycan Map; % galactosylation			rCE-SDS: Profile	Color					
Glycan Map; % afucosylation			nrCE-SDS: Main peak	Clarity					
Glycan Map; % sialylation			nrCE-SDS: Pre-peaks	Process-related Impurities		HCP ELISA			
Glycan Map; profiles			nrCE-SDS: Profile			HCP LC-MS			
Capillary isoelectric focusing: Isoelectric point			CEX-HPLC: Acidic peaks			HCP 2D-DIGE			
Capillary isoelectric focusing: Profile			CEX-HPLC: Main peak			Residual Protein A ELISA			
Amino acid analysis/UV spectroscopy			CEX-HPLC: Basic peaks			Residual DNA qPCR			
Identity by ELISA			CEX-HPLC: Profile						

Evaluation of Structural and Purity Attributes

Primary Structure Similarity Results

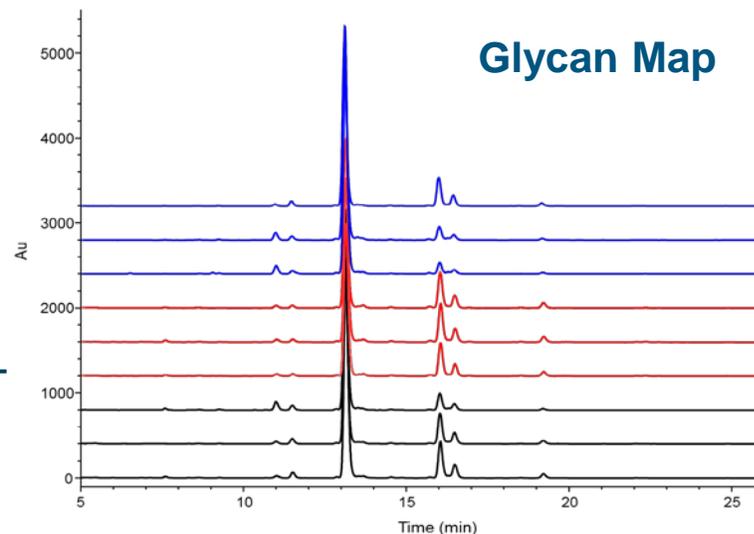
Primary Structure	Similarity Outcome
Intact whole mass	✓
Reduced deglycosylated molecular mass	✓
Reduced peptide map	✓
Nonreduced peptide map	✓
Glycan map	Minor quantitative differences
cIEF	✓
Extinction coefficient	✓
Identity by ELISA	✓

Reduced Peptide Map



Blue = Bevacizumab (EU) lots
 Red = ABP 215 lots
 Black = Bevacizumab (US) lots

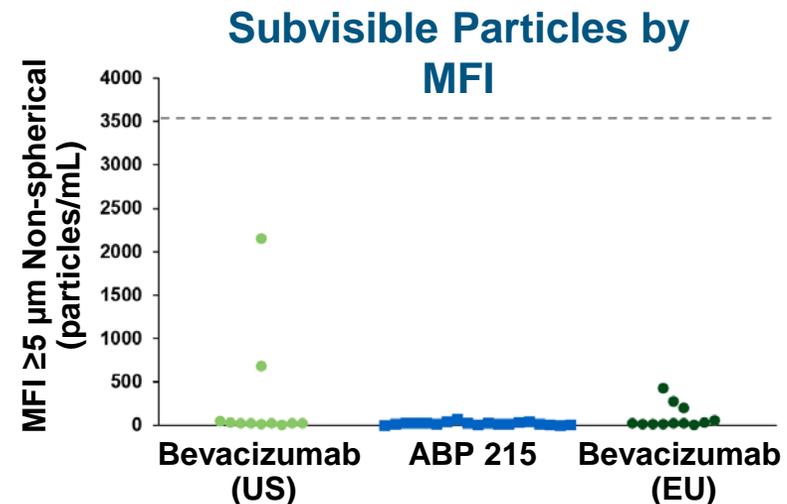
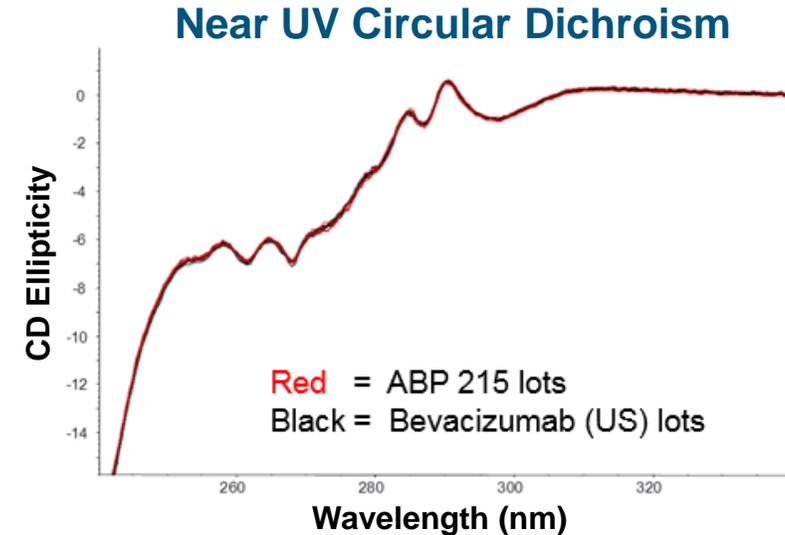
Glycan Map



Higher Order Structure and Particles and Aggregates Similarity Results

Higher Order Structure	Similarity Outcome
FTIR	✓
Near UV circular dichroism	✓
DSC	✓

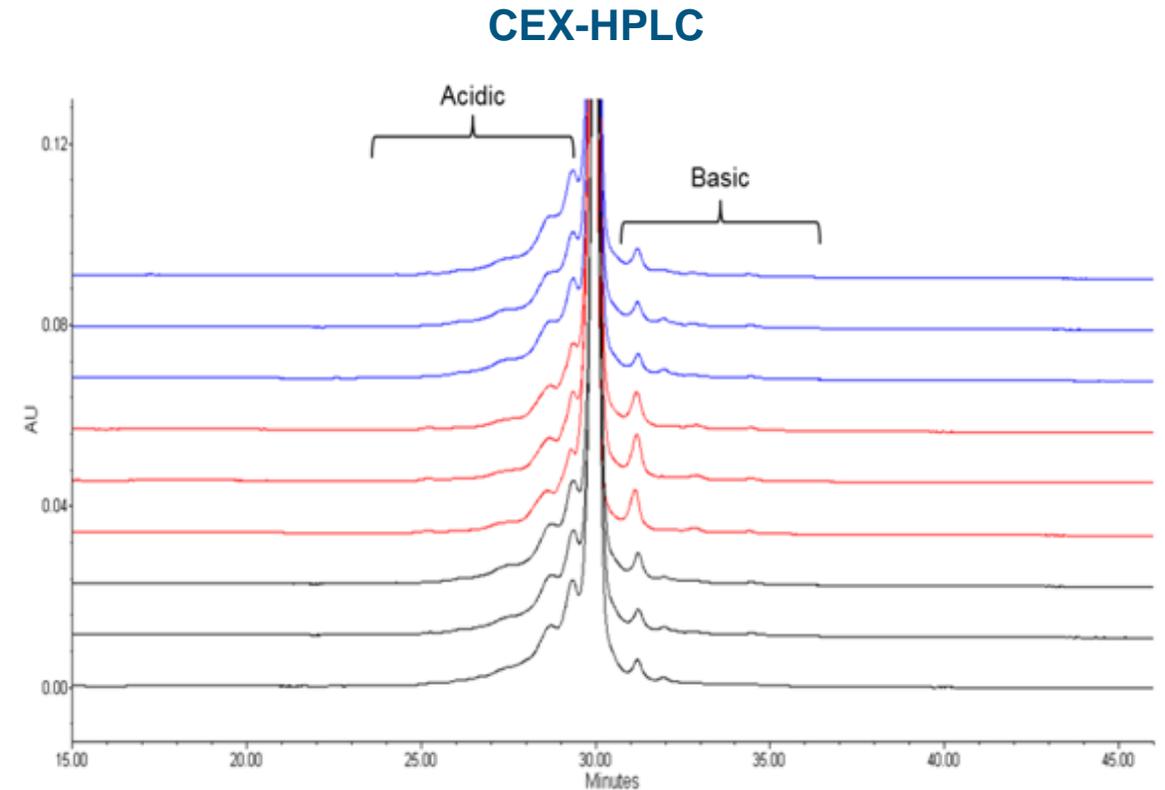
Particles and Aggregates	Similarity Outcome
Microflow imaging (MFI)	✓
Subvisible particle counts by HIAC	✓
Field flow fractionation	✓
Dynamic light scattering	✓
AUC sedimentation velocity	✓
SE-HPLC with light-scattering	✓



Product-related Substances and Impurities Similarity Results

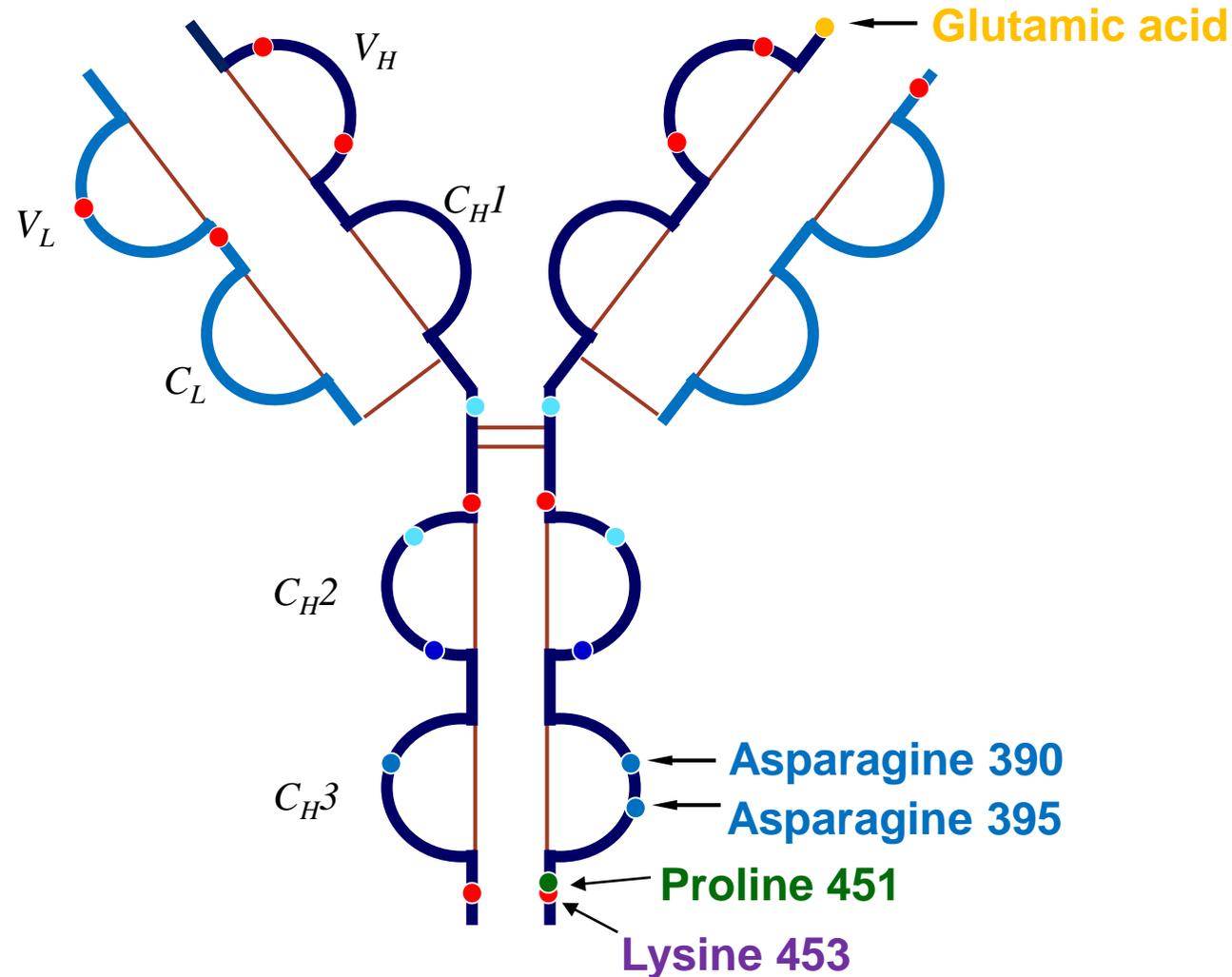
Product-related Substances and Impurities	Similarity Outcome
Size Variants SE-HPLC rCE-SDS nrCE-SDS	Minor quantitative differences, all less than 2%
Charge Variants CEX-HPLC	Quantitative differences

- ◆ No new species observed
- ◆ No impact on functional activities



Blue = Bevacizumab (EU) lots
 Red = ABP 215 lots
 Black = Bevacizumab (US) lots

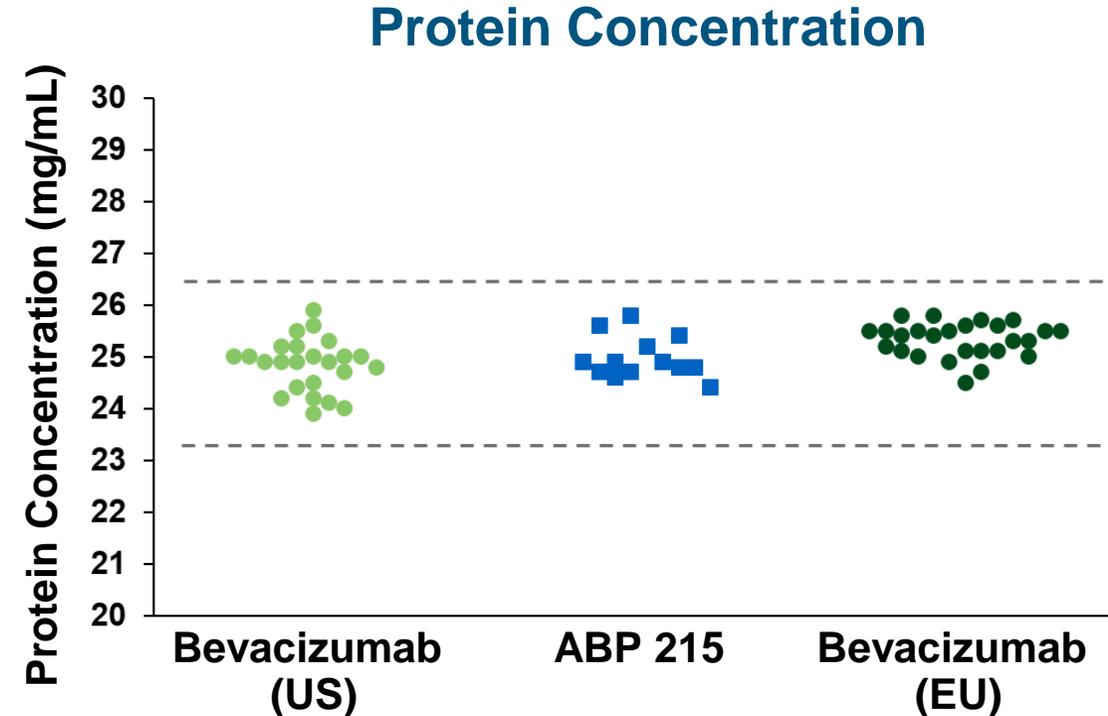
Assessment of Charge Profile Differences



- ◆ Same charge variants present in the reference product
- ◆ Charge variants are not within the Fab or Fc binding domains
- ◆ No observed impact to PK, efficacy, safety, or immunogenicity

General Properties Similarity Results

General Properties	Similarity Outcome
Appearance	✓
Color	✓
Clarity	✓
pH	✓
Osmolality	✓
Volume	✓
Protein concentration	✓

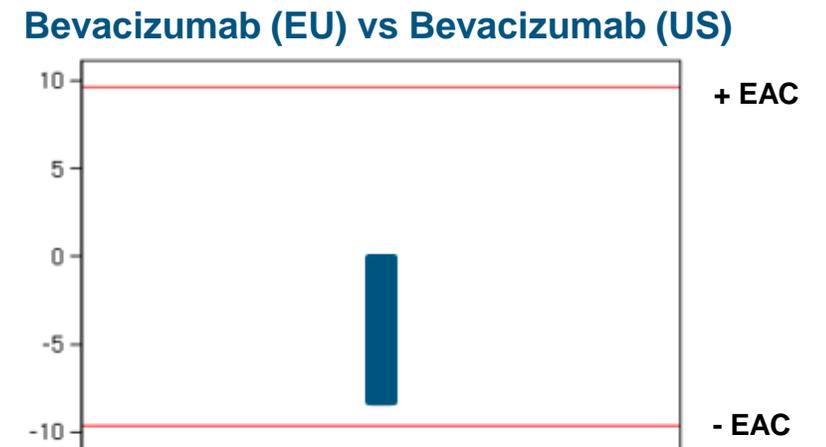
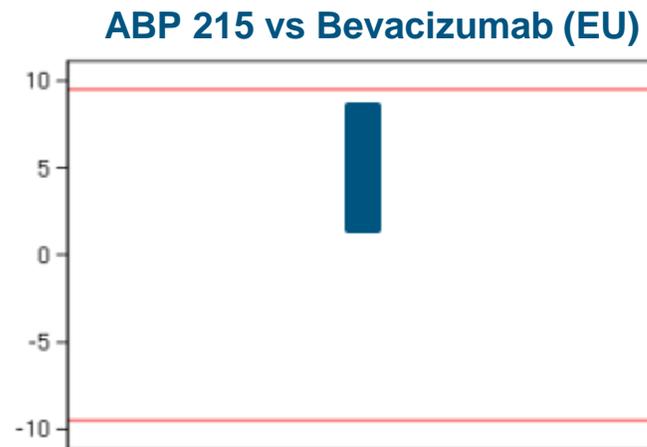
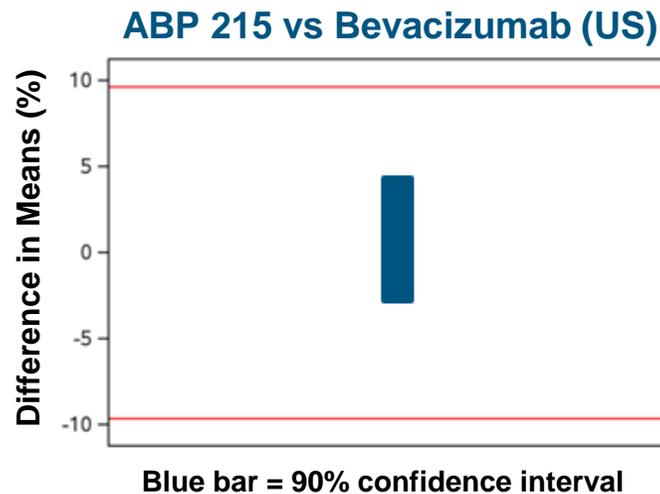
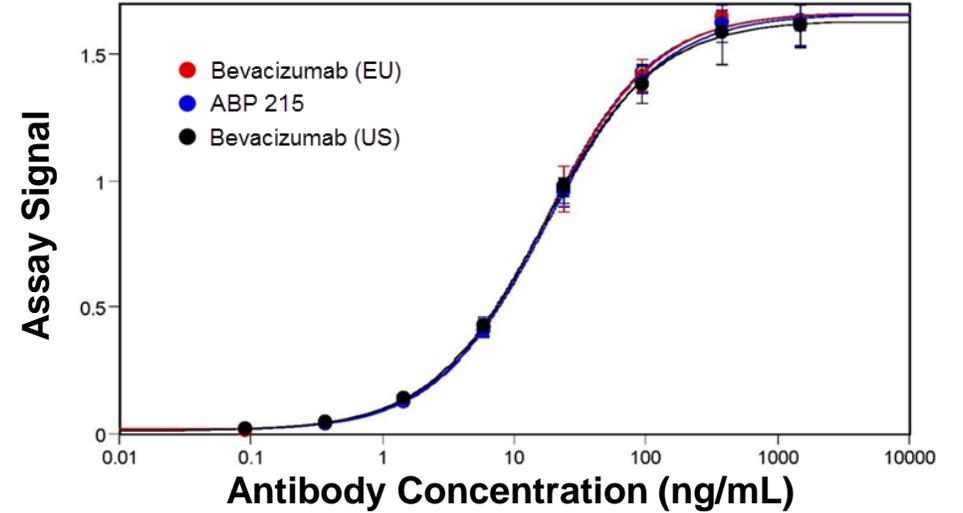
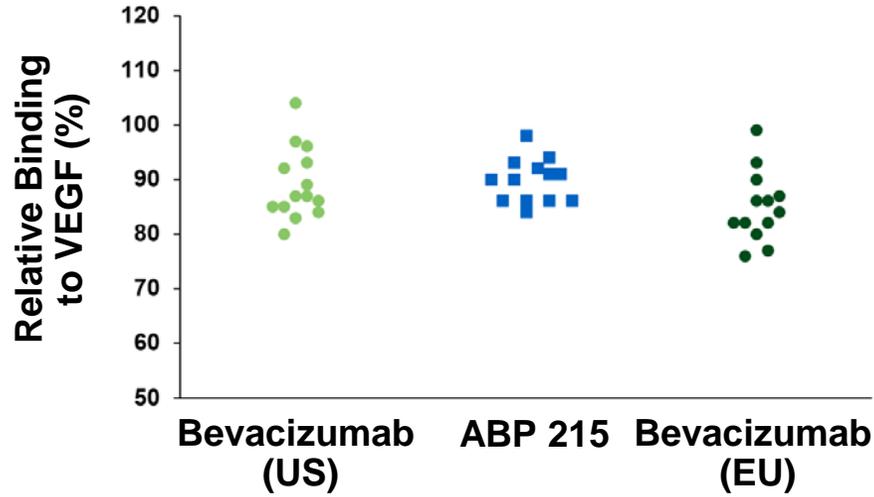


Evaluation of Functional Activities

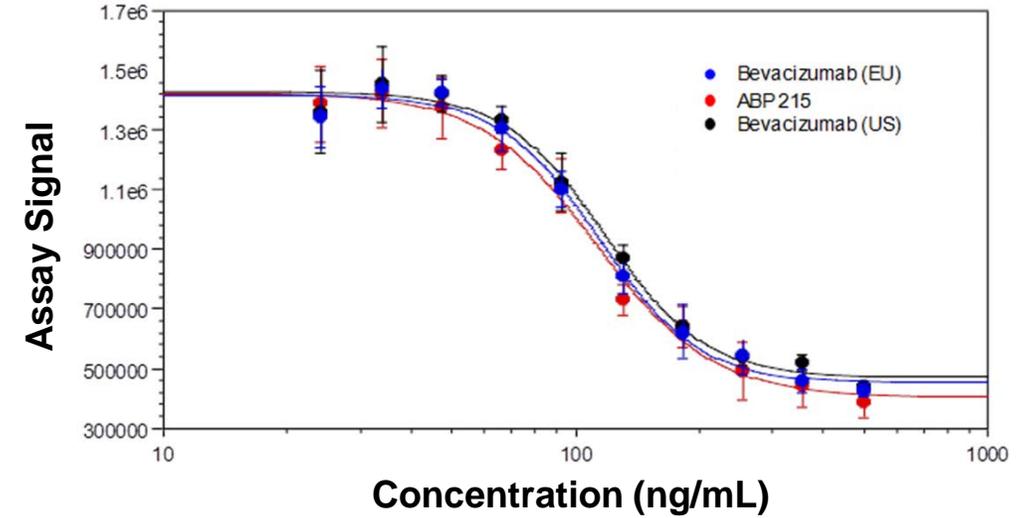
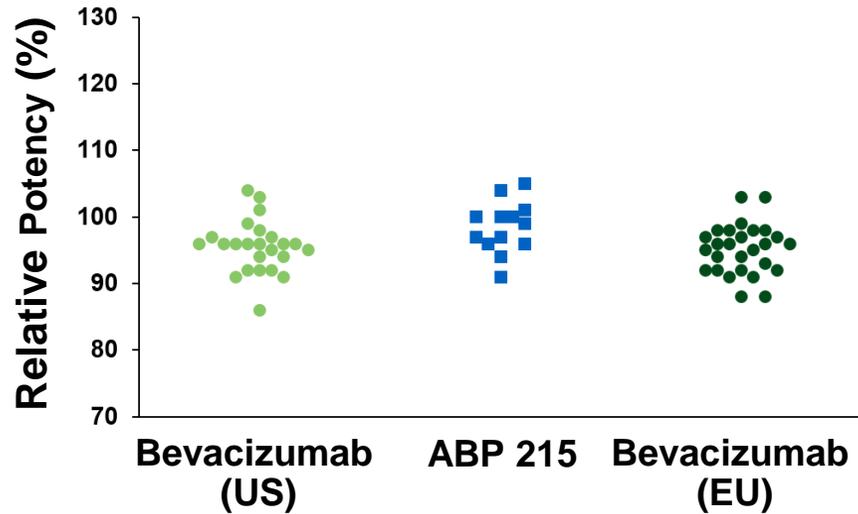
Similarity was Demonstrated in Functional Activities Evaluated

Fab-mediated Activities		Fc-mediated Characterization	
Binding to VEGF	✓	FcRn binding	✓
Inhibition of proliferation in HUVEC (potency)	✓	FcγRIa binding	✓
On and off rates (VEGF)	✓	FcγRIIa (131H) binding	✓
Binding to VEGF isoforms	✓	FcγRIIb binding	✓
Inhibition of VEGFR-2 RTK autophosphorylation	✓	FcγRIIIa (158V) binding	Minor difference unrelated to MoA
Specificity for VEGF by VEGFR-2 RTK autophosphorylation	✓	FcγRIIIa (158F) binding	✓
		FcγRIIIb binding	✓
		C1q binding	✓
Fab- and Fc-mediated Characterization			
		Lack of ADCC	✓
		Lack of CDC	✓

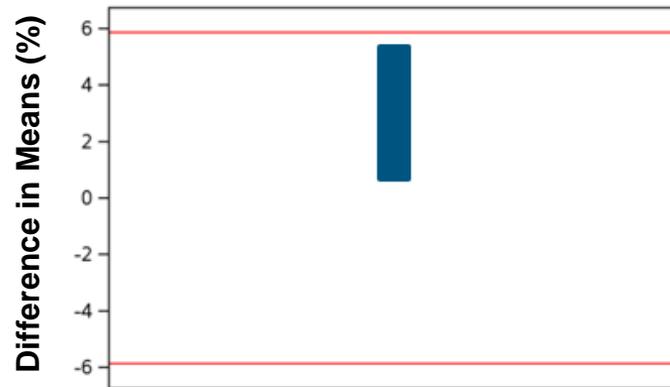
Binding to VEGF Meets Assessment Criteria



Potency via Inhibition of Proliferation Meets Assessment Criteria



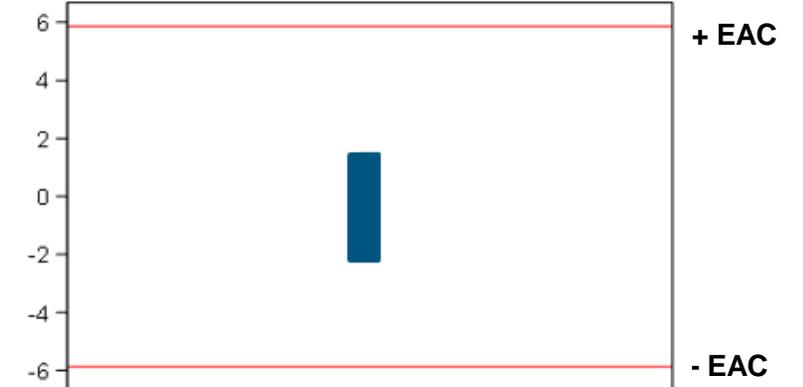
ABP 215 vs Bevacizumab (US)



ABP 215 vs Bevacizumab (EU)



Bevacizumab (EU) vs Bevacizumab (US)

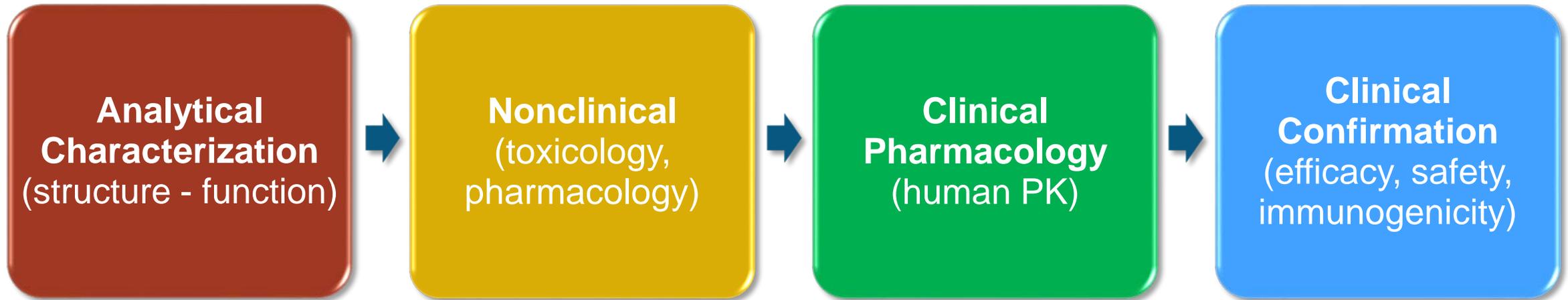


Blue bar = 90% confidence interval

Analytical Similarity of ABP 215 to Bevacizumab Was Demonstrated for ~ 100 Attributes/Assays

	Analytical Testing/Attributes	Results		Analytical Testing/Attributes	Results		Analytical Testing/Attributes	Results
Biological Activity	Inhibition of proliferation in HUVEC	Similar	Higher Order Structure	FTIR: Spectral similarity	Similar	Thermal Degradation @ 50°C	SE-HPLC: HMW	Similar
	Binding to VEGF by ELISA	Similar		FTIR: Profile	Similar		rCE-SDS: HC + LC	Similar
	On and off rates (VEGF)	Similar		Near UV CD: Spectral similarity	Similar		rCE-SDS: LMW + MMW	Similar
	Binding to VEGF isoforms	Similar		Near UV CD: Profile	Similar		CEX-HPLC: Main peak	Similar
	Inhibition of VEGFR-2 RTK autophosphorylation	Similar		DSC: T _m 1	Similar		CEX-HPLC: Acidic peaks	Similar
	Specificity for VEGF by VEGFR-2 RTK autophosphorylation	Similar		DSC: T _m 2	Similar		CEX-HPLC: Basic peaks	Similar
	FcRn binding	Similar		DSC: Profile	Similar		Proliferation inhibition bioassay	Similar
	FcγRIa binding	Similar	Particles and Aggregates	HIAC: ≥ 2 μm particles	Similar	Thermal Stability @ 40°C	SE-HPLC: HMW	Similar
	FcγRIIa (131H) binding	Similar		HIAC: ≥ 5 μm particles	Similar		rCE-SDS: HC + LC	Similar
	FcγRIIb binding	Similar		HIAC: ≥ 10 μm particles	Similar		rCE-SDS: LMW + MMW	Similar
	FcγRIIIa (158V) binding	Minor Diff		HIAC: ≥ 25 μm particles	Similar		CEX-HPLC: Main peak	Similar
	FcγRIIIa (158F) binding	Similar		MFI: ≥ 5 μm particles	Similar		CEX-HPLC: Acidic peaks	Similar
	FcγRIIIb binding	Similar		MFI: ≥ 5 μm non-spherical particles	Similar		CEX-HPLC: Basic peaks	Similar
	C1q binding	Similar		FFF: Submicron particles	Similar	Proliferation inhibition bioassay	Similar	
	Lack of ADCC	Similar		DLS (Hydrodynamic Radius)	Similar	Thermal Stability @ 25°C	SE-HPLC: HMW	Similar
	Lack of CDC	Similar		AUC-SV: % Monomer	Similar		rCE-SDS: HC + LC	Similar
	Primary Structures	Intact molecular mass		Similar	AUC-SV: Profile		Similar	rCE-SDS: LMW + MMW
Intact molecular mass; profiles		Similar	SLS: Molar mass	Similar	CEX-HPLC: Main peak		Similar	
Reduced and deglycosylated molecular masses of HC and LC		Similar	Product-related Substance and Impurities	SE-HPLC: HMW	Minor Diff		CEX-HPLC: Acidic peaks	Similar
Reduced and deglycosylated molecular masses of HC and LC; profiles		Similar		SE-HPLC: Profile	Similar		CEX-HPLC: Basic peaks	Similar
Reduced peptide map		Similar		rCE-SDS: HC+LC	Minor Diff	General Properties	Protein Concentration	Similar
Reduced peptide map; profiles		Similar		rCE-SDS: NGHC			Volume	Similar
Non-reduced peptide map		Similar		rCE-SDS: LMW + MMW			Osmolality	Similar
Non-reduced peptide map; profiles		Similar		rCE-SDS: Profile	Similar		pH	Similar
Glycan Map; % high mannose		Minor Diff		nrCE-SDS: Main peak	Minor Diff		Appearance	Similar
Glycan Map; % galactosylation				nrCE-SDS: Pre-peaks			Color	Similar
Glycan Map; % afucosylation		Similar		nrCE-SDS: Profile	Similar	Clarity	Similar	
Glycan Map; % sialylation		Similar		CEX-HPLC: Acidic peaks	Minor Diff	Process-related Impurities	HCP ELISA	Similar
Glycan Map; profiles		Similar	CEX-HPLC: Main peak		HCP LC-MS		Similar	
Capillary isoelectric focusing: Isoelectric point		Similar	CEX-HPLC: Basic peaks		HCP 2D-DIGE		Similar	
Capillary isoelectric focusing: Profile		Similar	CEX-HPLC: Profile	Similar	Residual Protein A ELISA		Similar	
Amino acid analysis/UV spectroscopy		Similar			Residual DNA qPCR	Similar		
Identity by ELISA		Similar						

Analytical Characterization Supports Biosimilarity



- ◆ **Comprehensive analytical characterization was conducted**
- ◆ **Results demonstrate that ABP 215 is highly analytically similar to bevacizumab**
- ◆ **Analytical similarity results form the foundation of the scientific justification for extrapolation**

Agenda

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Analytical Similarity

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**Non-Clinical and Clinical Similarity and
Extrapolation to All Indications**

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Global Development, Amgen

Conclusion

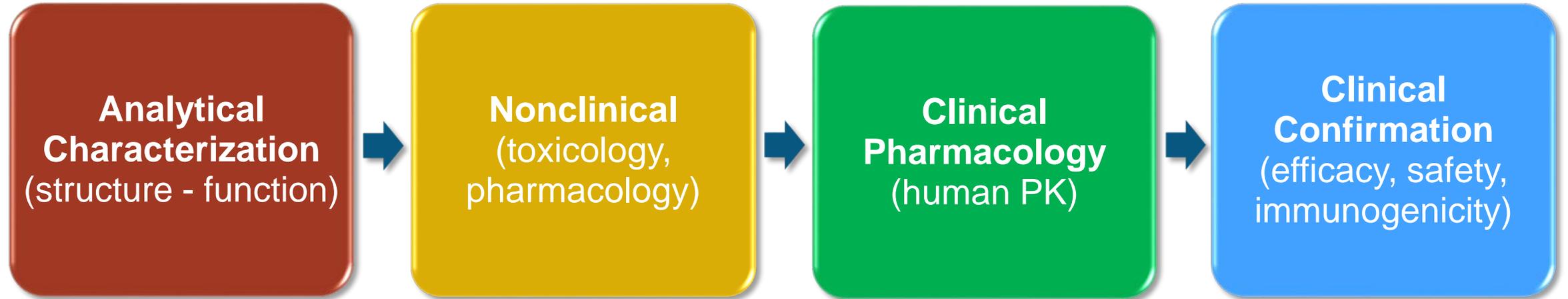
Lisa Bollinger, MD
Regulatory Affairs and Safety, Amgen

Non-Clinical and Clinical Similarity and Extrapolation to All Indications

Richard Markus, MD, PhD

VP Global Development, Amgen

Nonclinical Assessment is Next in Stepwise Development



- Same amino acid sequence and strength
- Equivalent potency
- Highly similar structure and function

Nonclinical: Toxicology

Study Design:

- ◆ **ABP 215 vs bevacizumab**
- ◆ **Four weeks: 50 mg/kg IV twice a week**
- ◆ **3 male and 3 female cynomolgus monkeys/group**

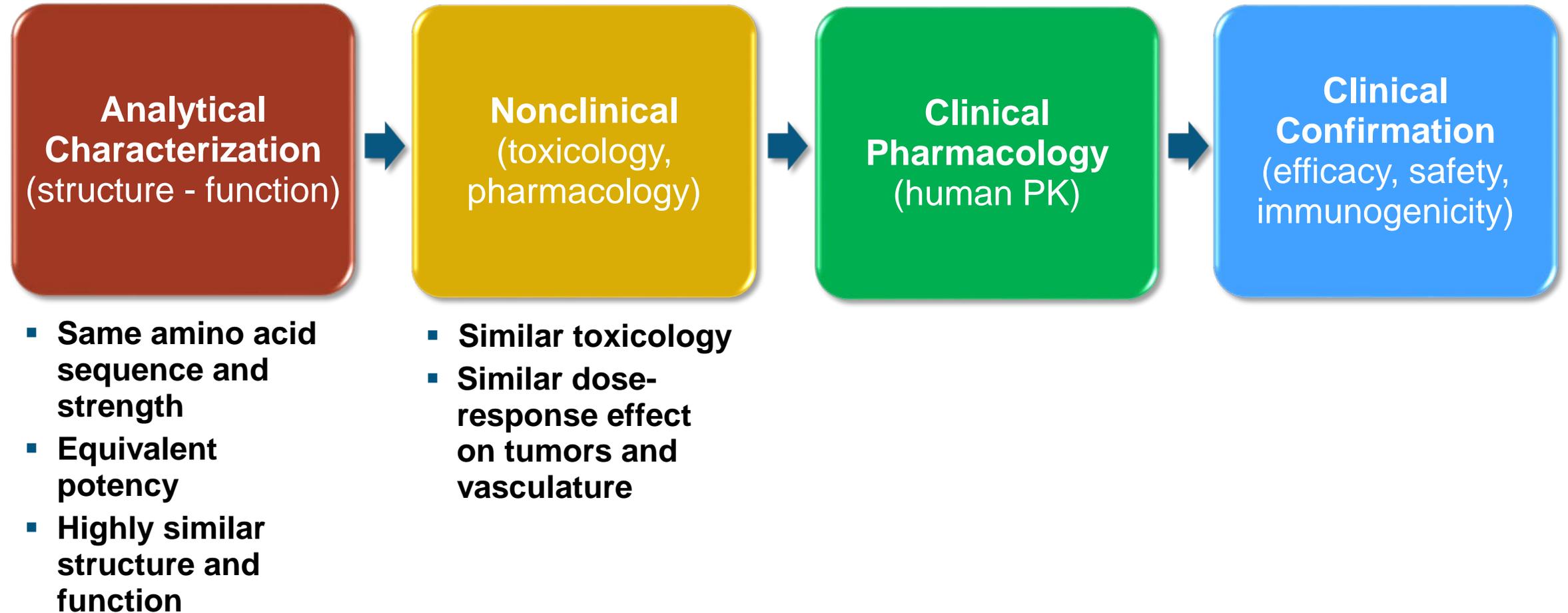
Findings:

- ◆ **Anticipated finding of physeal dysplasia of the femur similar in incidence and severity in both groups**
- ◆ **No unexpected toxicities**

Nonclinical Pharmacology Studies

Evaluation	Treatment Arms	Results
A431 Xenograft (Epidermoid Cancer Model)	Bevacizumab & ABP 215 10 and 100 µg; IgG1 control 100 µg	ABP 215 inhibited tumor growth and vasculature similar to bevacizumab
Colo205 Xenograft (Colon Cancer Model)	Bevacizumab & ABP 215 10 and 100 µg; IgG1 control 100 µg	ABP 215 inhibited tumor growth and vasculature similar to bevacizumab
Vascular Permeability induced by huVEGF expressing cells	Bevacizumab & ABP 215 3, 10, 30, and 100 µg; IgG1 control 100 µg	ABP 215 inhibited vascular permeability similar to bevacizumab

Clinical Pharmacology is Next in Stepwise Development



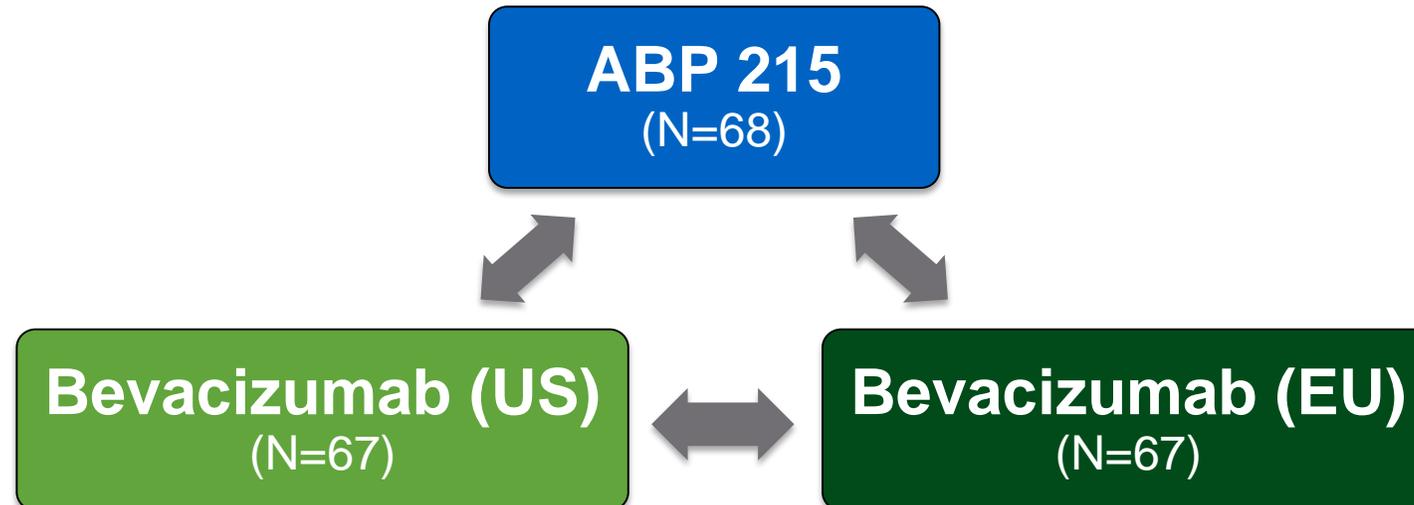
Clinical Pharmacology – PK Similarity

◆ Design:

- Healthy male volunteers
- Single 3 mg/kg IV dose
- 85 days follow-up

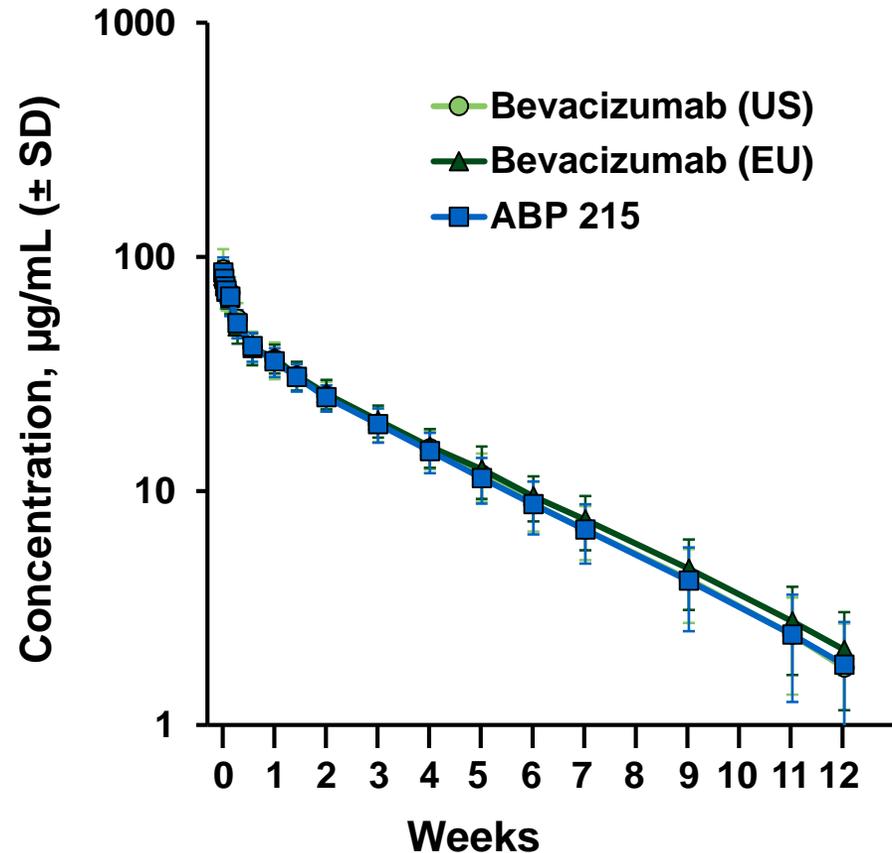
◆ Endpoints:

- C_{max} , AUC_{inf} , AUC_{last}
(Margin 80-125%)
- Safety
- Immunogenicity



PK Similarity Results

Mean (\pm SD) Serum
Concentration-Time Profiles



Ratio of Adjusted LS Geometric
Means with 90% CI

C_{max} ($\mu\text{g/mL}$)

ABP 215 vs. bev (US)

ABP 215 vs. bev (EU)

Bev (US) vs. (EU)

AUC_{inf} ($\text{h}\cdot\mu\text{g/mL}$)

ABP 215 vs. bev (US)

ABP 215 vs. bev (EU)

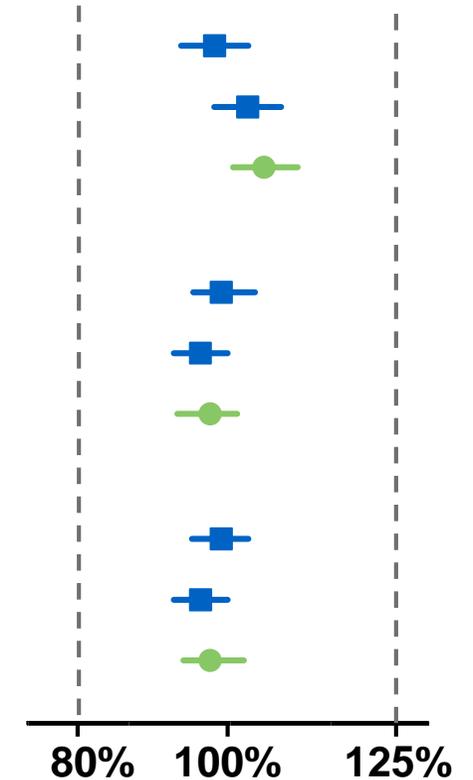
Bev (US) vs. (EU)

AUC_{last} ($\text{h}\cdot\mu\text{g/mL}$)

ABP 215 vs. bev (US)

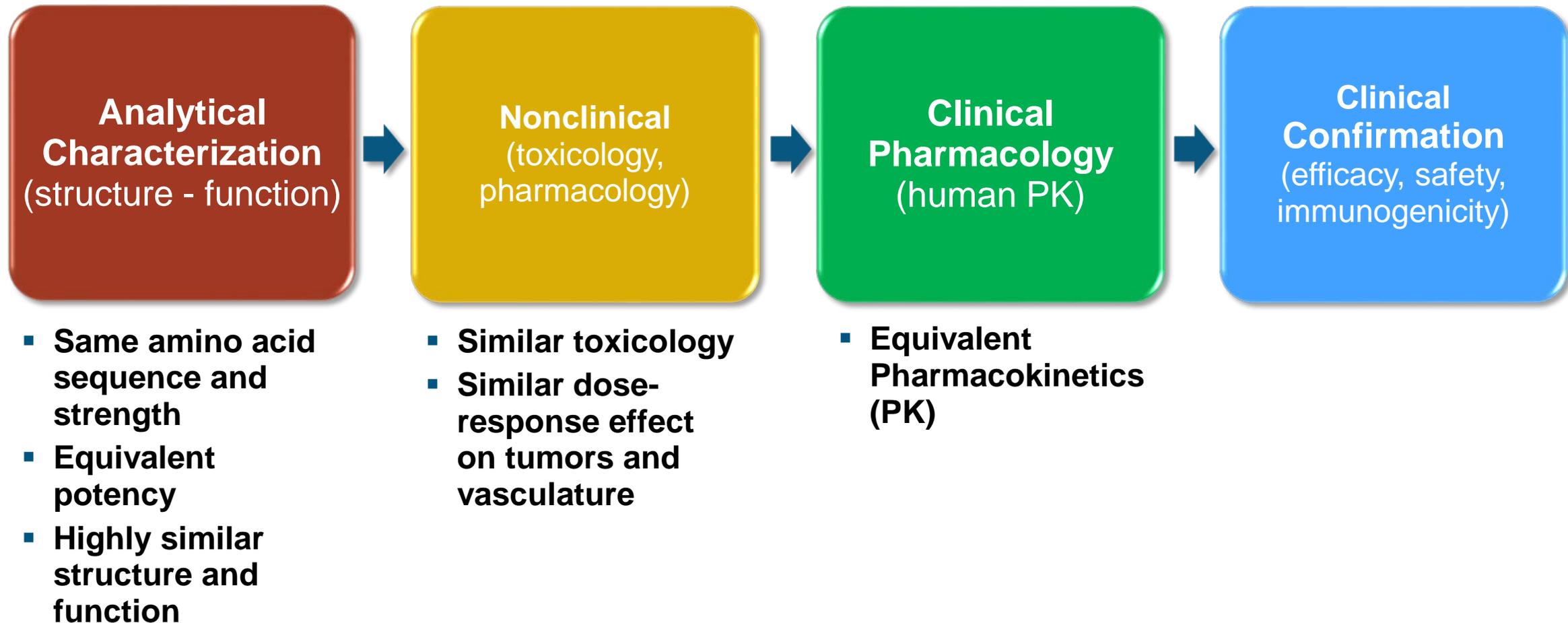
ABP 215 vs. bev (EU)

Bev (US) vs. (EU)



LS = least squares.

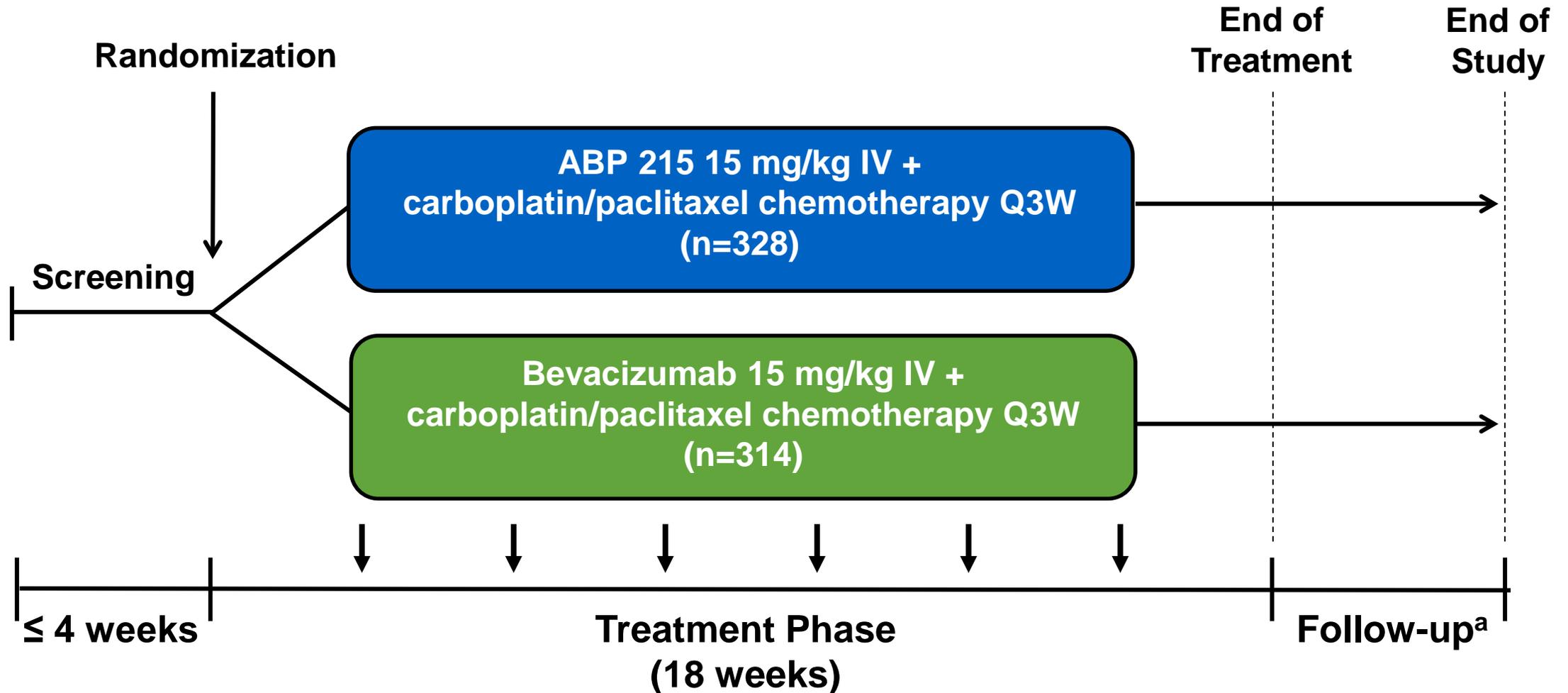
Clinical Confirmation is Next in Stepwise Development



Clinical Study Considerations

- ◆ **Objective: Head-to-head comparison of anti-tumor efficacy, safety, and immunogenicity**
 - Confirm similarity with no clinically meaningful differences
- ◆ **Advanced NSCLC with tumor response primary endpoint**
 - Objective response rate (ORR) is a measurement of anti-tumor activity due to treatment
 - ORR of Bevacizumab in NSCLC has relatively large magnitude of response

Study Schema – Advanced NSCLC



a. Maintenance monotherapy not included.

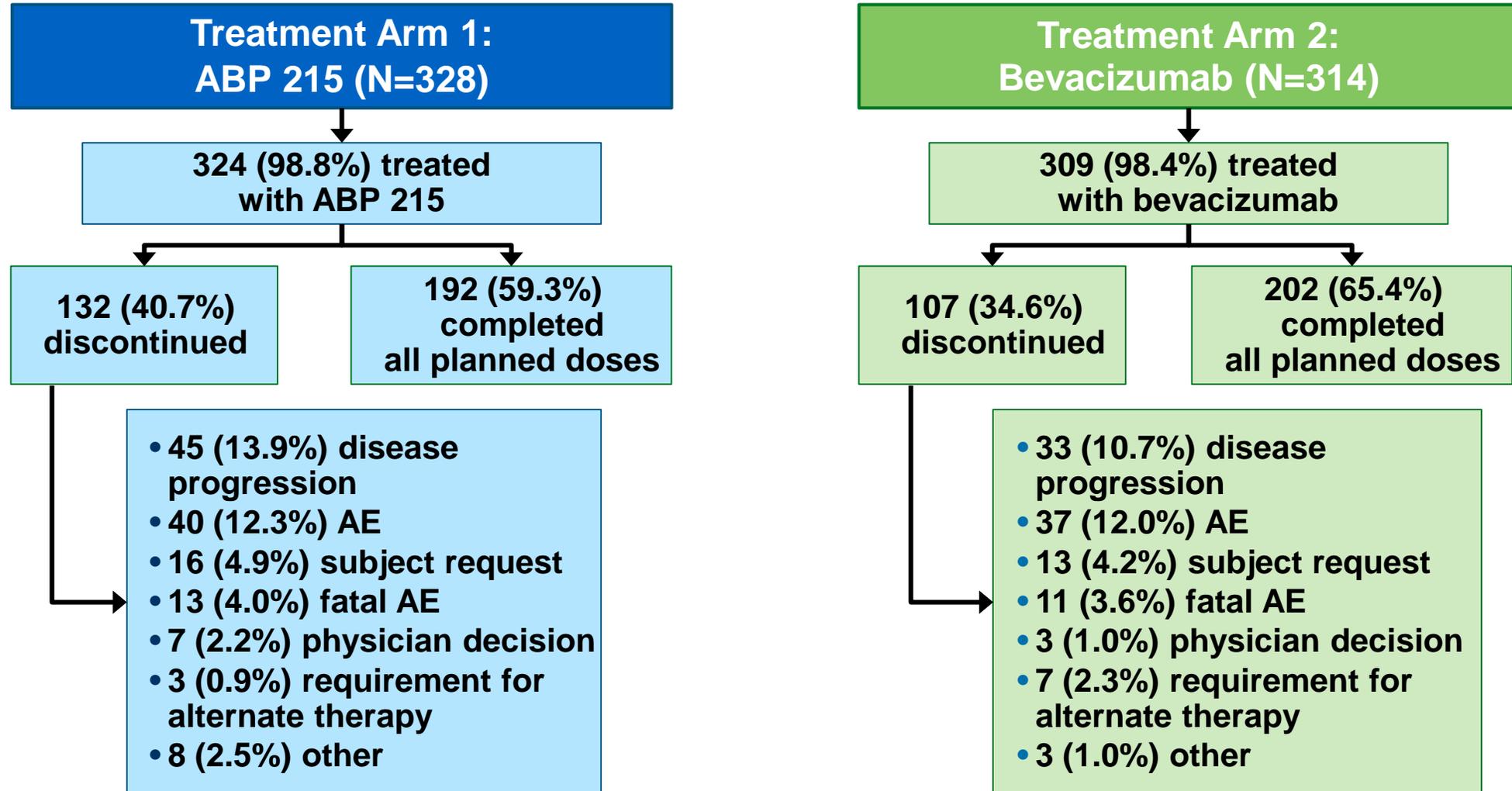
Endpoints

- ◆ **Primary endpoint**
 - Risk Ratio (RR) of objective response rate (ORR)
- ◆ **Secondary endpoints**
 - Risk difference of ORR
 - Progression-free survival
 - Duration of response
- ◆ **Safety endpoints**
 - AE's and SAE's
 - Overall survival
 - Development of anti-drug antibodies

Primary Analysis

- ◆ **Objective Response Rate (ORR): Best overall response rate (complete response or partial response)**
 - ORR based on independent central radiology evaluation
- ◆ **Pre-specified similarity margin for RR = (0.67, 1.50)**
 - FDA revised margin RR = (0.73, 1.36)
- ◆ **90% confidence interval for RR must be entirely within the pre-specified similarity margin**
 - Equates to ORR risk difference within 6%

Subject Disposition



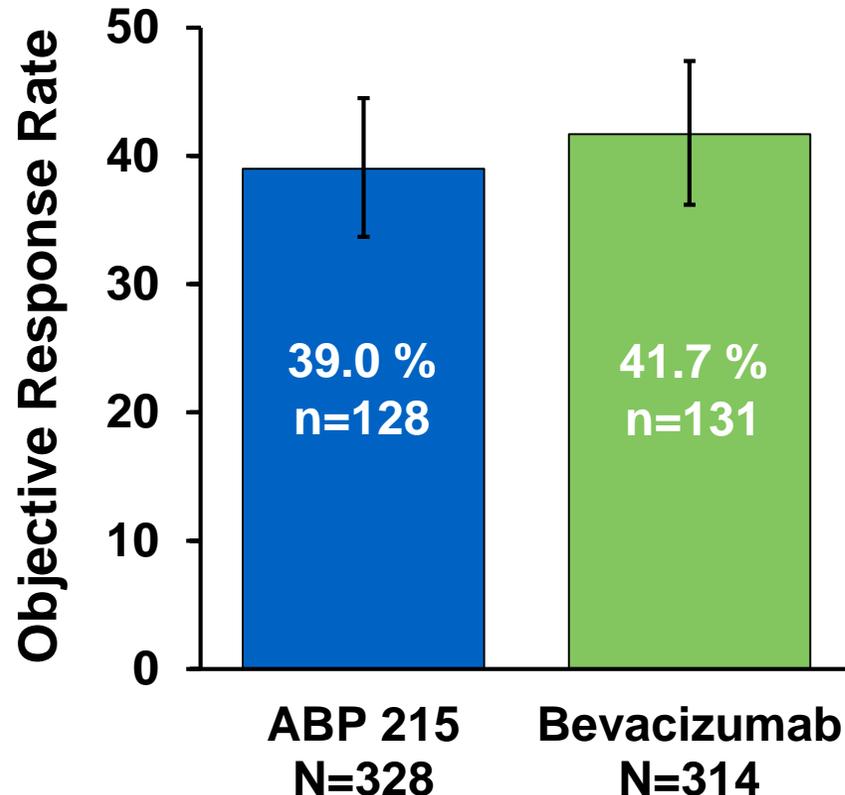
Baseline Demographics

Characteristic	ABP 215 N=328	Bevacizumab N=314
Age, years, mean (SD)	61.6 (9.1)	61.6 (8.9)
< 65 years, n (%)	199 (60.7)	191 (60.8)
≥ 65 years, n (%)	129 (39.3)	123 (39.2)
Weight, kg, mean (SD)	71.2 (14.7)	73.5 (15.3)
White, n (%)	315 (96.0)	300 (95.5)
Male, n (%)	196 (59.8)	188 (59.9)
Smoking status, n (%)		
Never	65 (19.8)	76 (24.2)
Former	163 (49.7)	158 (50.3)
Current	100 (30.5)	80 (25.5)

Baseline Disease Characteristics

Characteristic, n (%)	ABP 215 N=328	Bevacizumab N=314
Staging of original diagnosis		
≤ Stage IIIA	23 (7.0)	25 (8.0)
Stage IIIB	2 (0.6)	7 (2.2)
Stage IV	303 (92.4)	281 (89.5)
Stage IV/recurrent disease at baseline		
Stage IV	309 (94.2)	290 (92.4)
Recurrent disease	19 (5.8)	24 (7.6)
Weight loss in last 6 months		
0%–5%	289 (88.1)	276 (87.9)
> 5%–10%	39 (11.9)	37 (11.8)
ECOG PS		
Grade 0	127 (38.7)	117 (37.3)
Grade 1	201 (61.3)	197 (62.7)

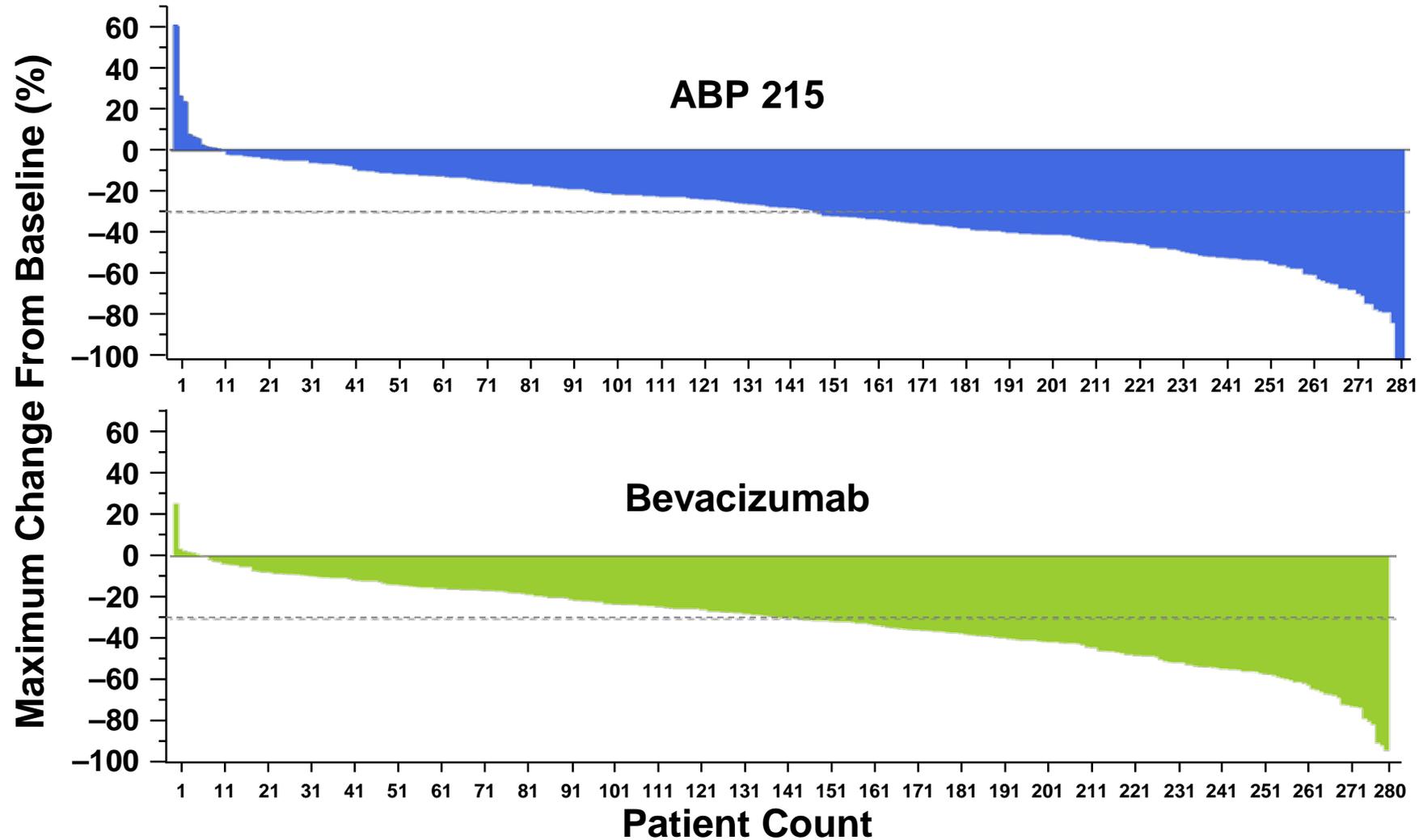
Primary Endpoint Results



Primary Analysis	ABP 215 vs Bevacizumab
RR (90% CI)	0.93 (0.80, 1.09)
Pre-specified equivalence margin	(0.67, 1.50)
FDA revised margin	(0.73, 1.36)

Primary analysis of ORR using RECIST 1.1 by central radiology review of ITT population.
 Generalized linear model adjusted for randomization stratification factors; geographic region, ECOG PS, and sex.
 Whiskers in the bar chart represent 95% CI of ORR.

Similar Magnitude of Tumor Response



Based on independent central tumor review using RECIST v1.1.
Figures exclude patients with a missing maximum change from baseline.

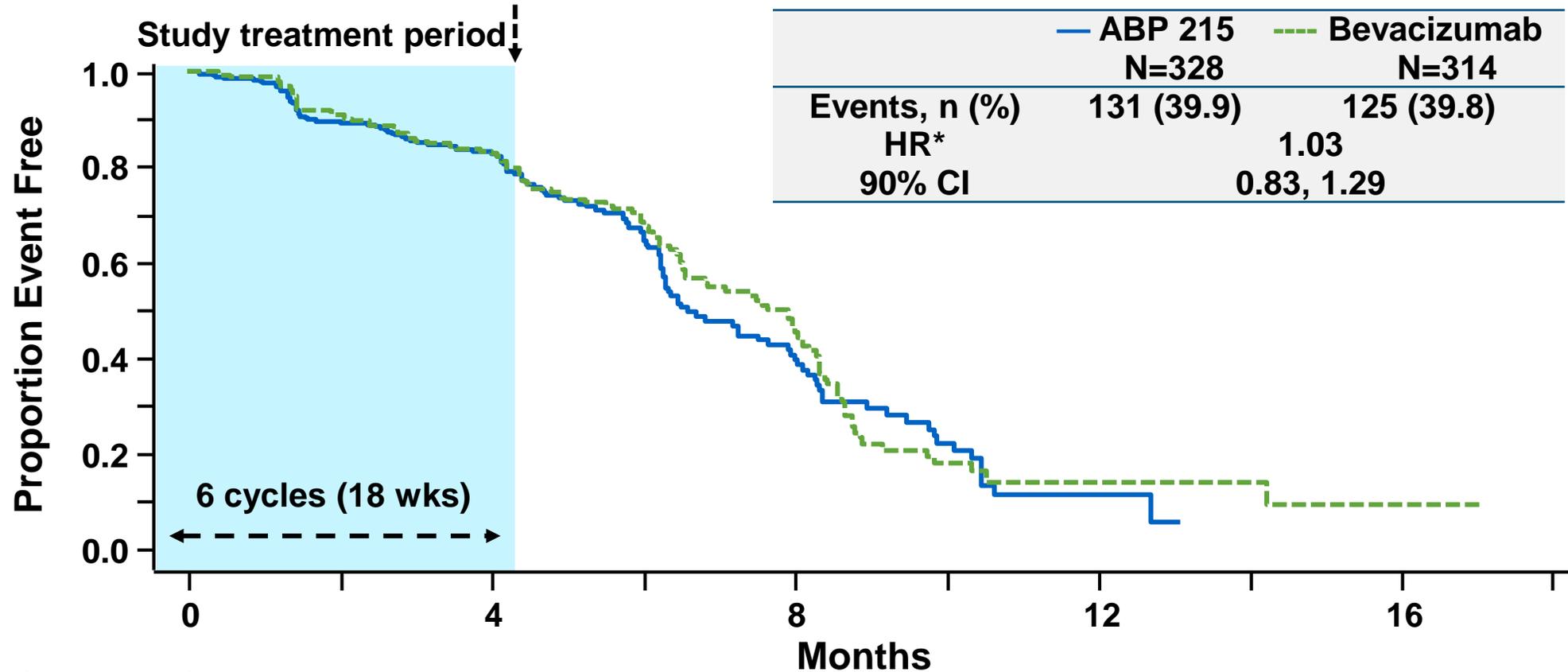
Secondary Efficacy Endpoints

Endpoint	ABP 215 N=328	Bevacizumab N=314
Objective Response Rate (ORR)		
Responders, n	128	131
ORR % (95% CI)	39.0 (33.7, 44.5)	41.7 (36.2, 47.4)
Risk Difference % (90% CI) ^a	-2.90 (-9.26, 3.45)	
Progression Free Survival (PFS)		
Progression or death, n (%)	131 (39.9)	125 (39.8)
HR (90% CI) ^b	1.03 (0.83, 1.29)	
Duration of Response (DOR)		
Responders, n (%)	128 (39.0)	131 (41.7)
Progression among responders	43 (33.6)	45 (34.4)
Median Time, months (95% CI)	5.8 (4.9, 7.7)	5.6 (5.1, 6.3)

^a Based on generalized linear model adjusted for randomization stratification factors: geographic region, ECOG PS, and sex.

^b Based on a Cox proportional hazards model stratified by geographic region, ECOG PS, and sex.

NSCLC: Progression Free Survival



Subjects at risk:

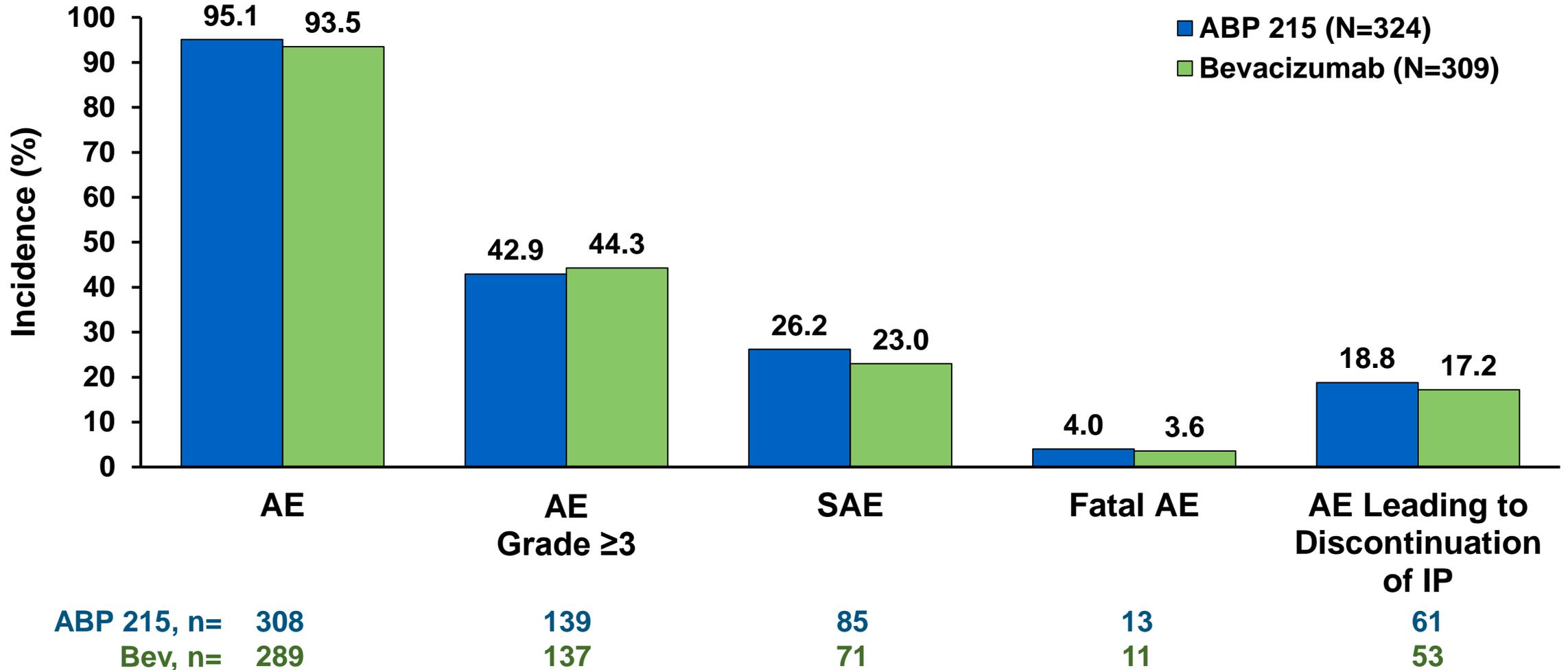
ABP 215, n= 328	254	202	98	39	15	4	0	0	0
Bev, n= 314	253	201	92	49	12	5	3	1	0

ITT population. Independent central radiology review.

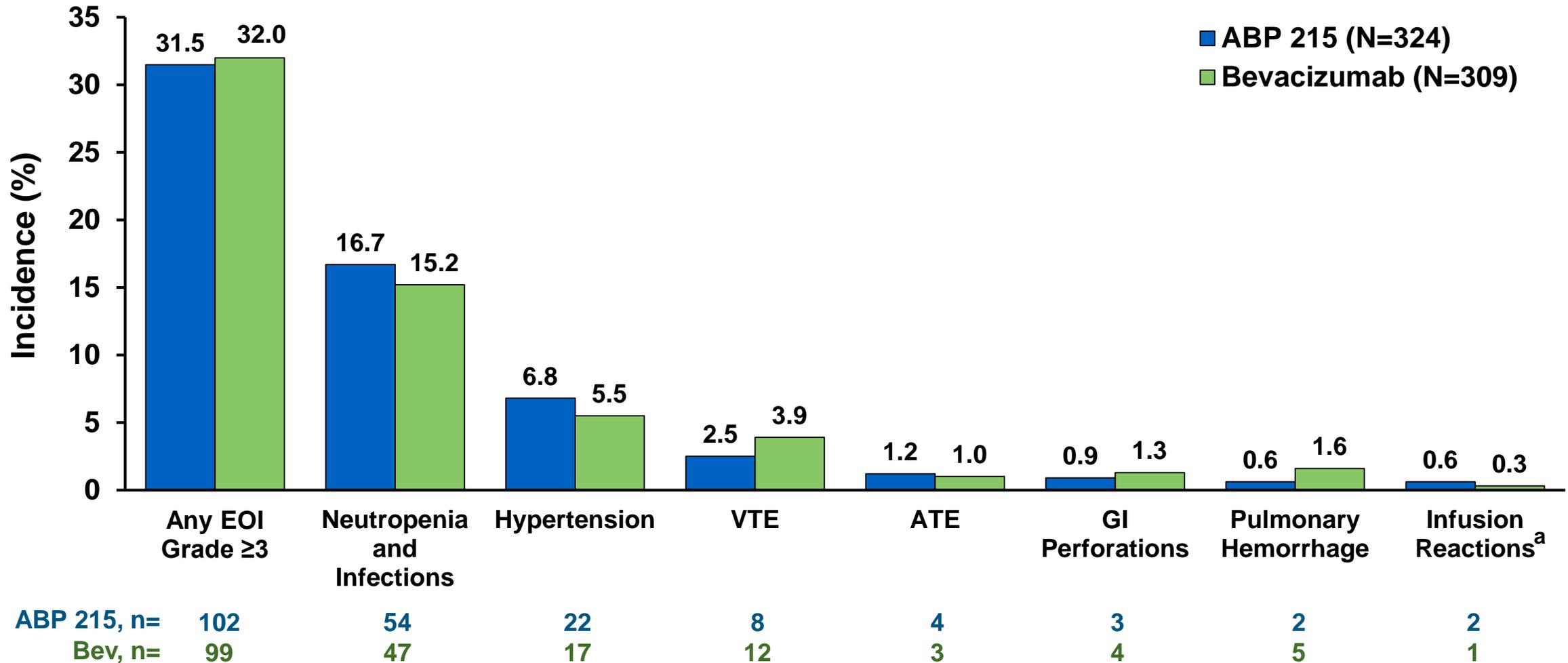
*Hazard ratio (HR) for ABP 215 relative to bevacizumab, based on a Cox proportional hazards model stratified by randomization factors.

Clinical Safety and Immunogenicity

Summary of Adverse Events



Key Events of Interest Grade ≥ 3



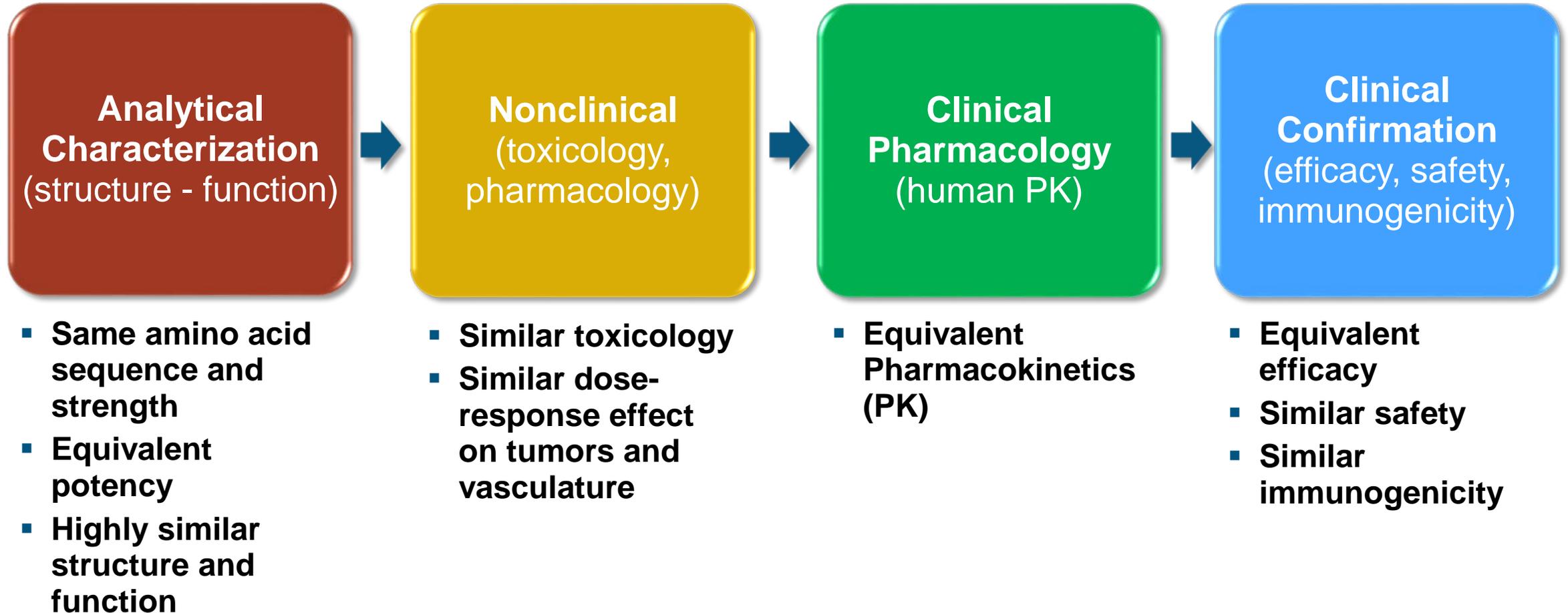
a. Infusion reactions = events within 2 days of first dose; VTE = Venous Thromboembolic Events; ATE = Arterial Thromboembolic Events; GI=Gastrointestinal.

Immunogenicity: Anti-drug Antibodies

Number of Subjects	ABP 215 N=324	Bevacizumab N=309
Baseline	315	303
Pre-existing binding antibody incidence, n (%)	0	3 (1.0)
Post-baseline	294	284
Binding antibody positive post-baseline with a negative or no result at baseline, n (%)	4 (1.4)	7 (2.5)

- ◆ No neutralizing antibodies in either group

Totality of Evidence Supports Biosimilarity

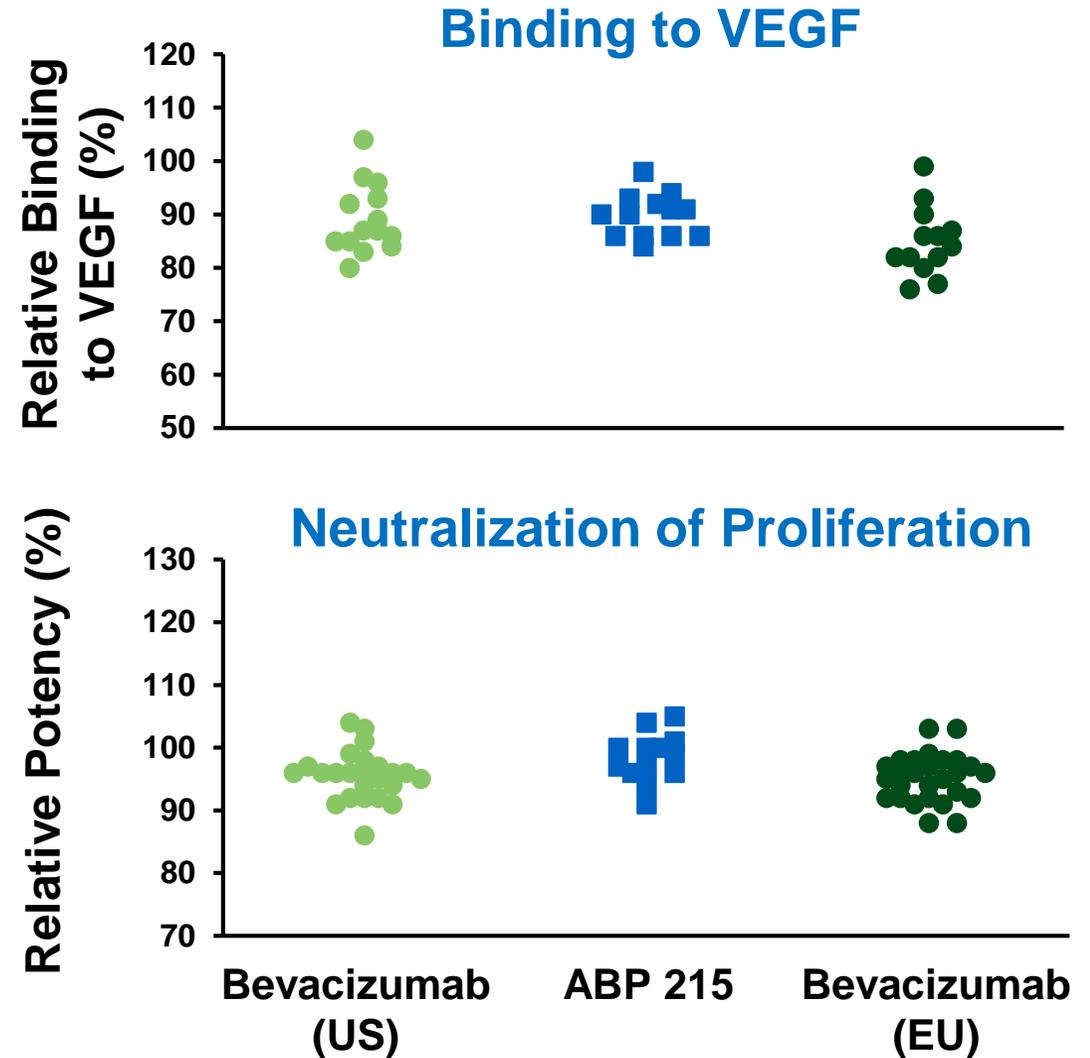


Extrapolation Considerations

- ◆ **Product similarity (ABP 215 and bevacizumab)**
 - Analytical similarity
 - Nonclinical
 - Clinical pharmacology
 - Safety, efficacy, and immunogenicity
- ◆ **Considerations across indications (bevacizumab)**
 - Mechanism of action
 - PK distribution and clearance
 - Clinical considerations

Extrapolation is applying the knowledge of efficacy and safety of the reference product to the highly similar biosimilar

Mechanism of Action in All Uses



Similar Mechanism of Action

Key Molecular Mechanisms

- Binding to all splice isoforms of VEGF-A
- Neutralization of all downstream VEGF-A signaling outcomes through VEGFR1 and VEGFR2—irrespective of cell type

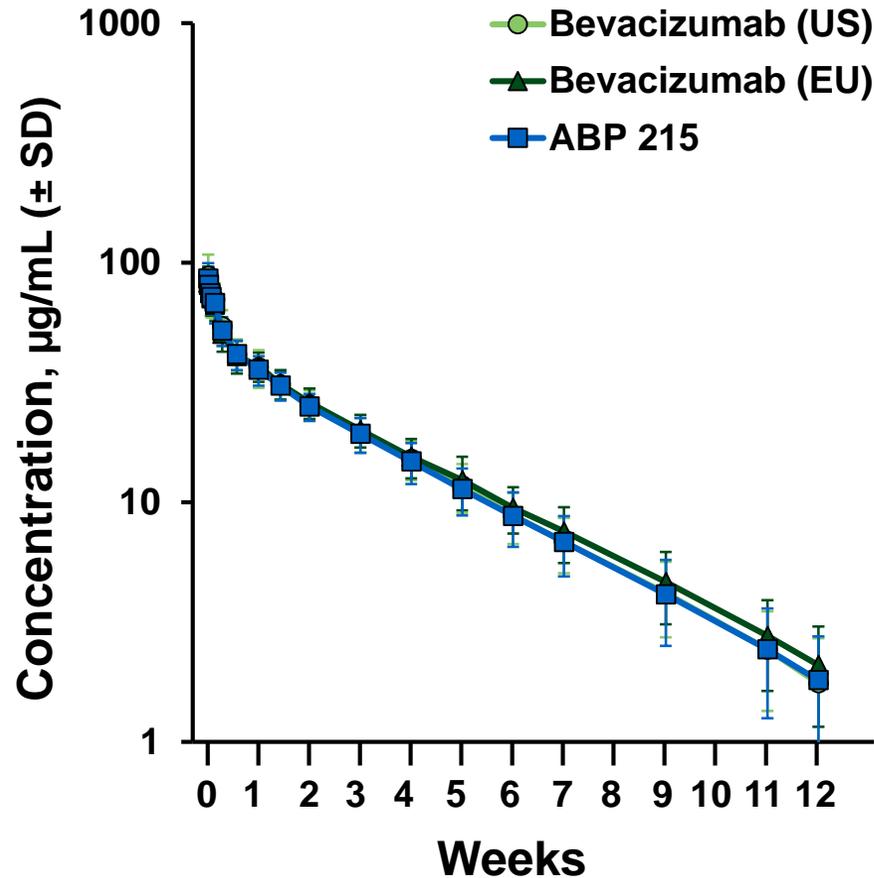
✓ **Common
MOA**

Proposed Indications

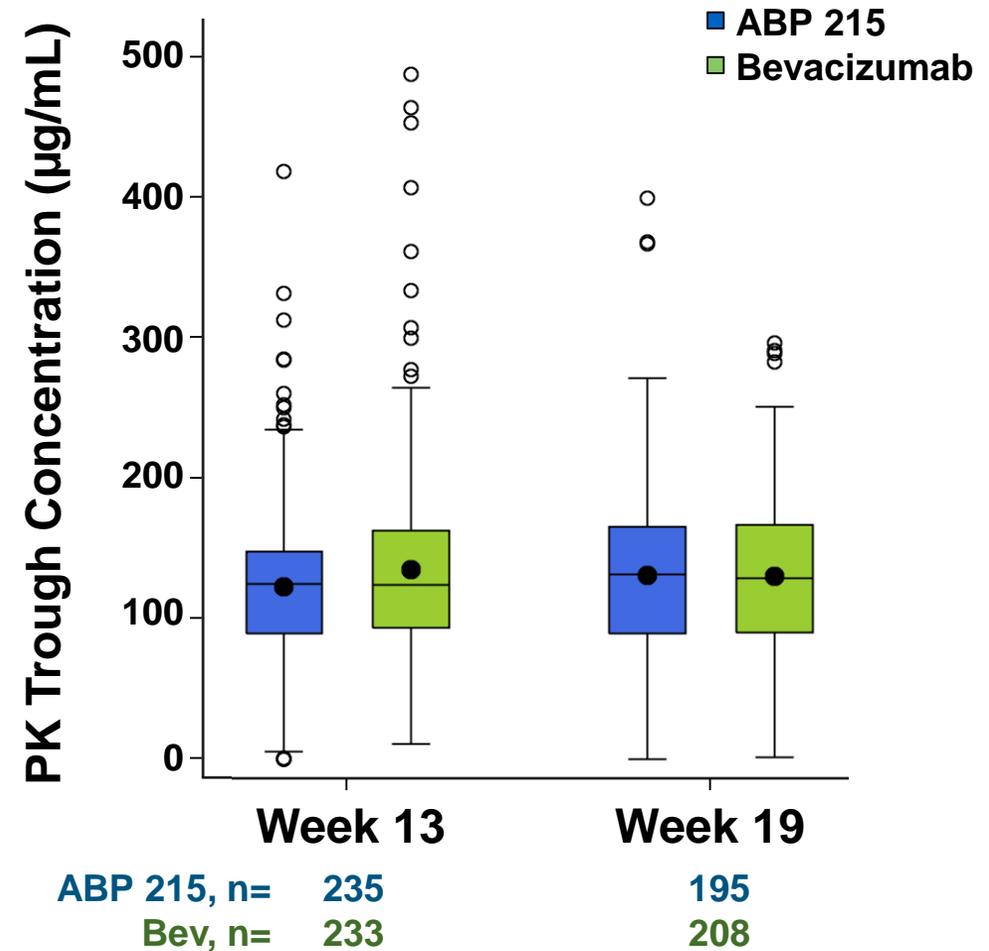
- Nonsquamous Non-small Cell Lung Cancer
- Metastatic Colorectal Cancer
- Metastatic Renal Cell Carcinoma
- Glioblastoma
- Cervical Cancer

ABP 215 and Bevacizumab Pharmacokinetics

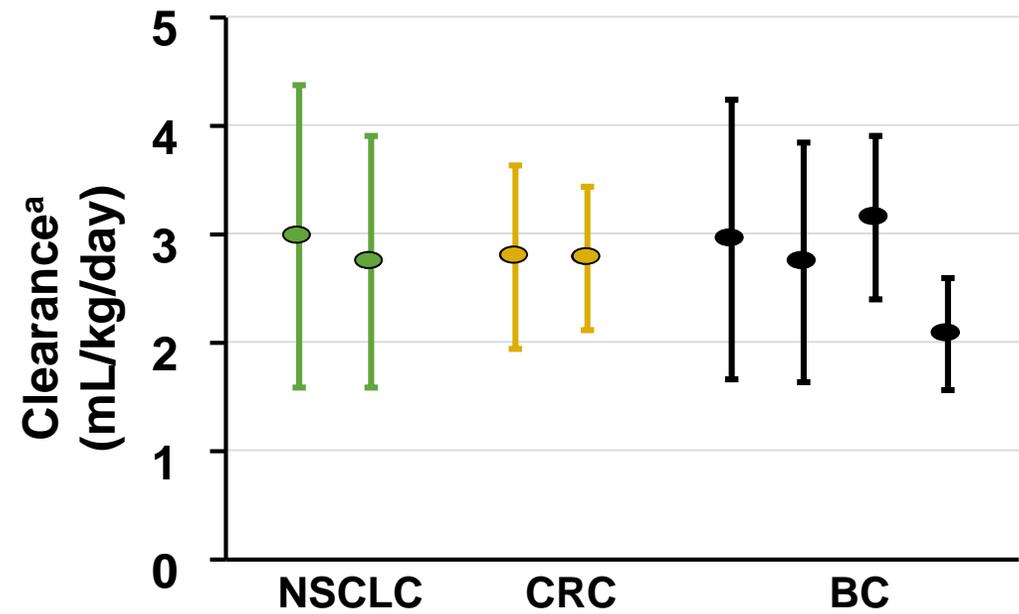
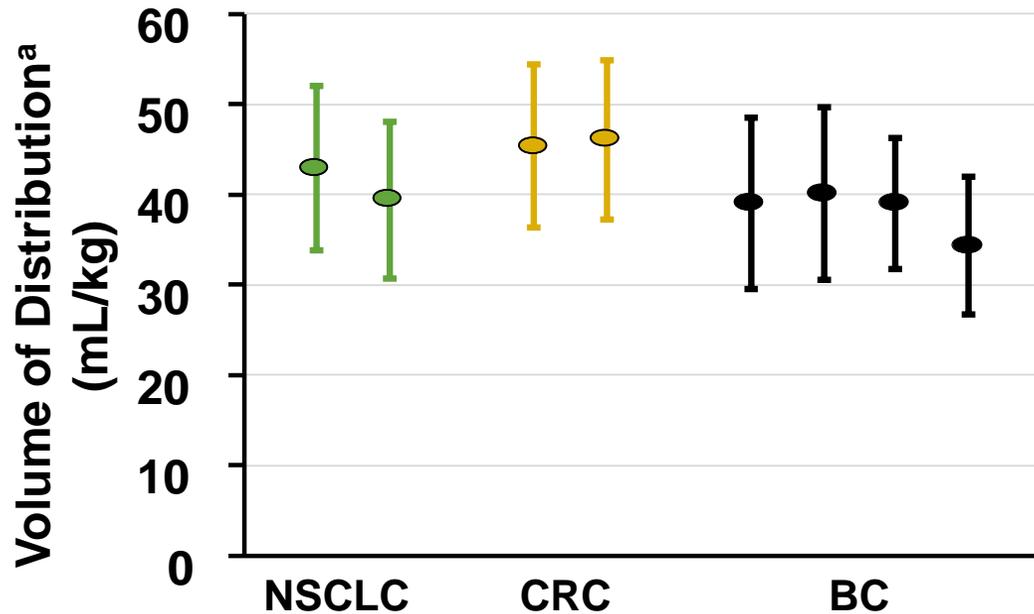
Healthy Volunteers (3 mg/kg)



Advanced NSCLC (15 mg/kg)



Similar PK Distribution and Clearance of Bevacizumab Across Different Tumor Types

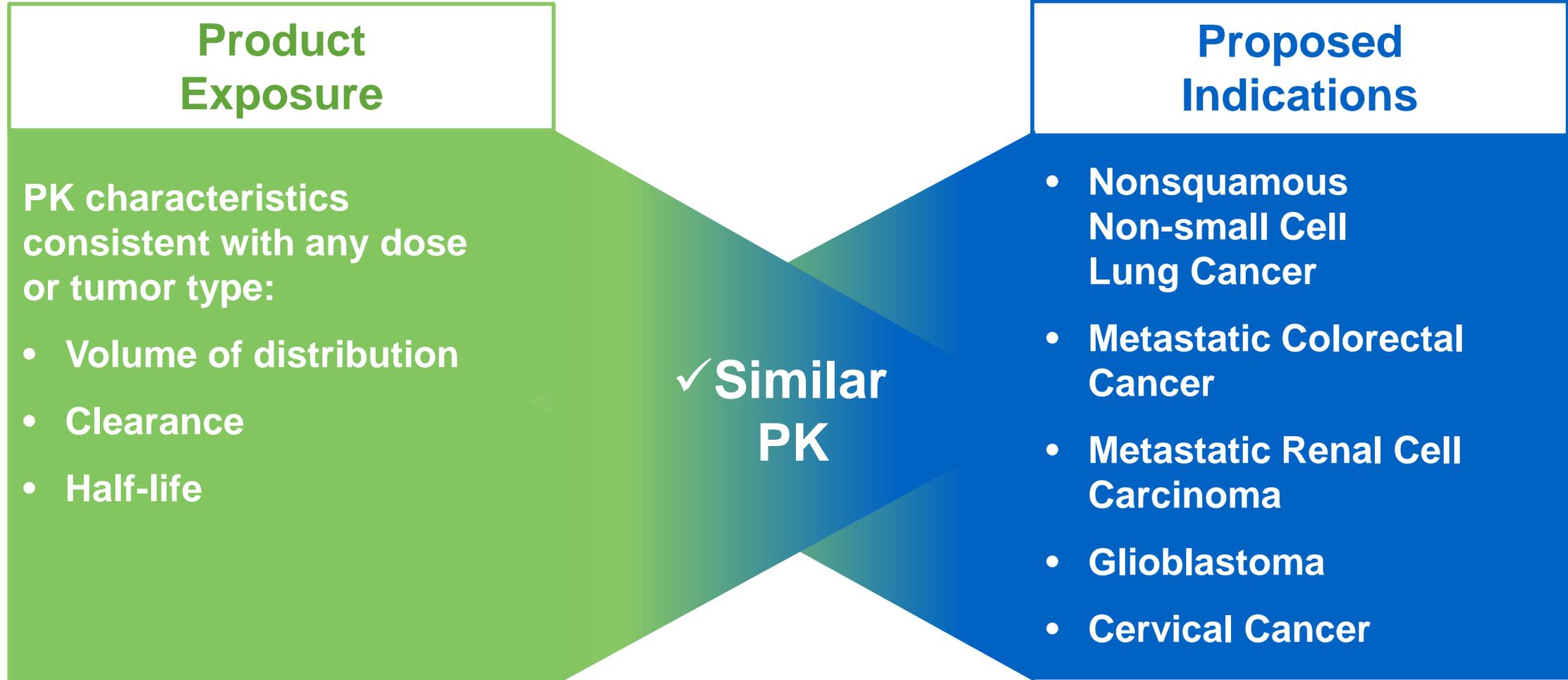


- ◆ Consistent PK properties also demonstrated in population PK analysis of bevacizumab (in NSCLC, CRC, GBM, BC, RCC, other tumors)^b

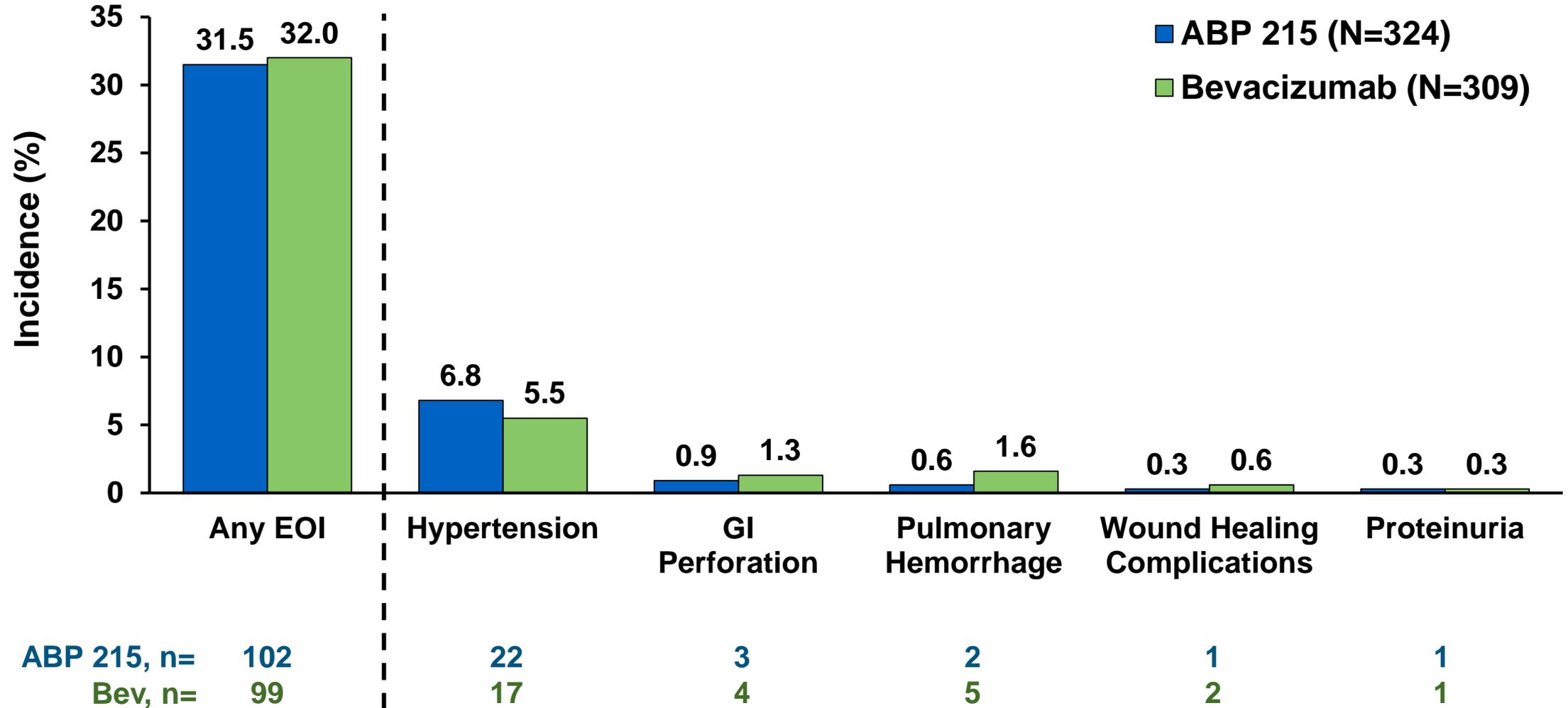
a. US and EMA prescribing information, Mean \pm SD, BC approved in EU only.

b. Han et al, *Cancer Chemother Pharmacol* 2016. CRC=Colorectal Carcinoma, GBM=Glioblastoma Multiforme, BC=Breast Cancer, RCC=Renal Cell Carcinoma.

Similar Pharmacokinetics



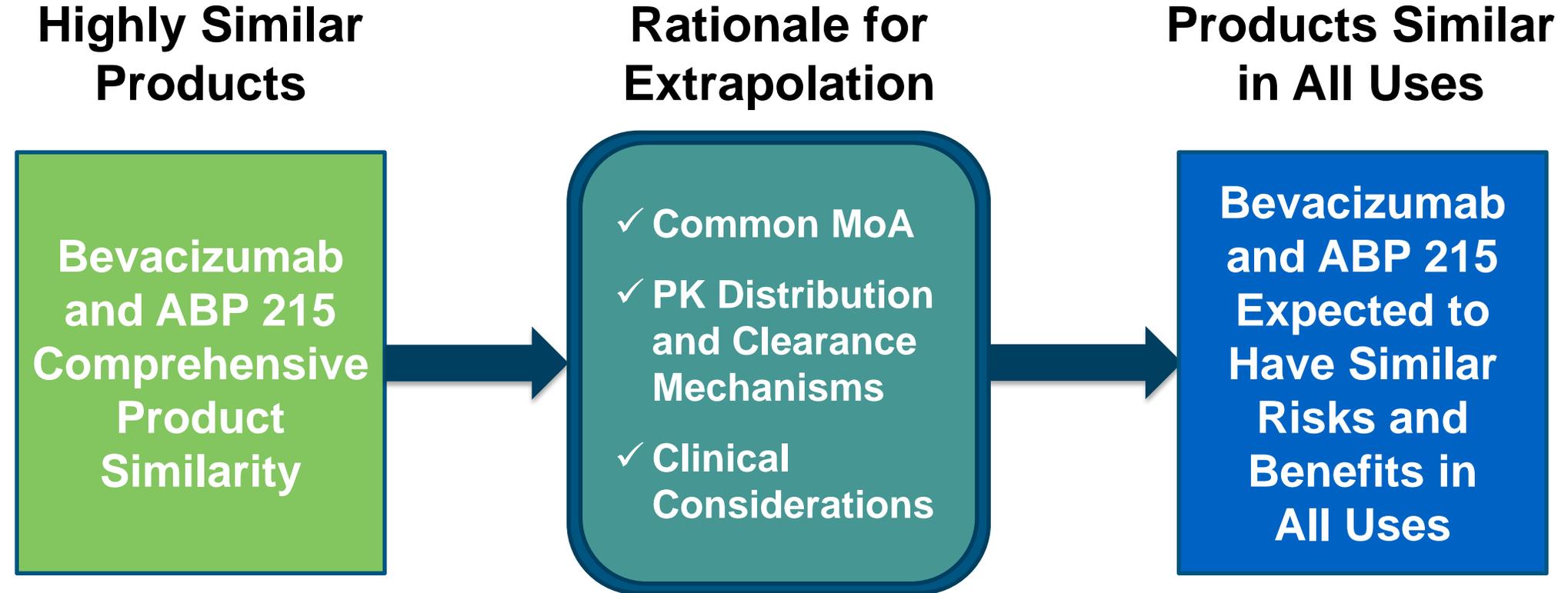
Anti-VEGF Toxicities: AEs Grade ≥ 3 in NSCLC Study



Similar Safety Profiles



Extrapolation of Indications



ABP 215 Proposed Indications of Use

- ◆ **Nonsquamous Non-small Cell Lung Cancer**
- ◆ **Metastatic Colorectal Cancer**
- ◆ **Metastatic Renal Cell Carcinoma**
- ◆ **Glioblastoma**
- ◆ **Cervical Cancer**

Agenda

Introduction

Richard Markus, MD, PhD
Global Development, Amgen

Analytical Similarity

Simon Hotchin
Regulatory Affairs, Amgen

Non-Clinical and Clinical Similarity and
Extrapolation to All Indications

Richard Markus, MD, PhD
Global Development, Amgen

Conclusion

Lisa Bollinger, MD
Regulatory Affairs and Safety, Amgen

Conclusions

Lisa Bollinger, MD

VP Regulatory Affairs & Safety

Statutory Requirements: Biosimilarity

Section 351(i) of the PHS Act defines biosimilarity to mean

- ◆ “that the biological product is **highly similar** to the reference product notwithstanding minor differences in clinically inactive components” and
 - √ *Structural and functional similarity demonstrated*
- ◆ that “there are **no clinically meaningful differences** between the biological product and the reference product in terms of the safety, purity, and potency of the product.”
 - √ *Equivalent PK, efficacy, similar safety and immunogenicity demonstrated*

Statutory Requirements: Extrapolation

PHS Act allows biosimilar sponsor to seek licensure for one or more conditions of use if:

- ◆ Biosimilarity supported by data from one or more clinical study in an appropriate condition of use
 - √ *Double blind, randomized study in patients with NSCLC (n=642)*

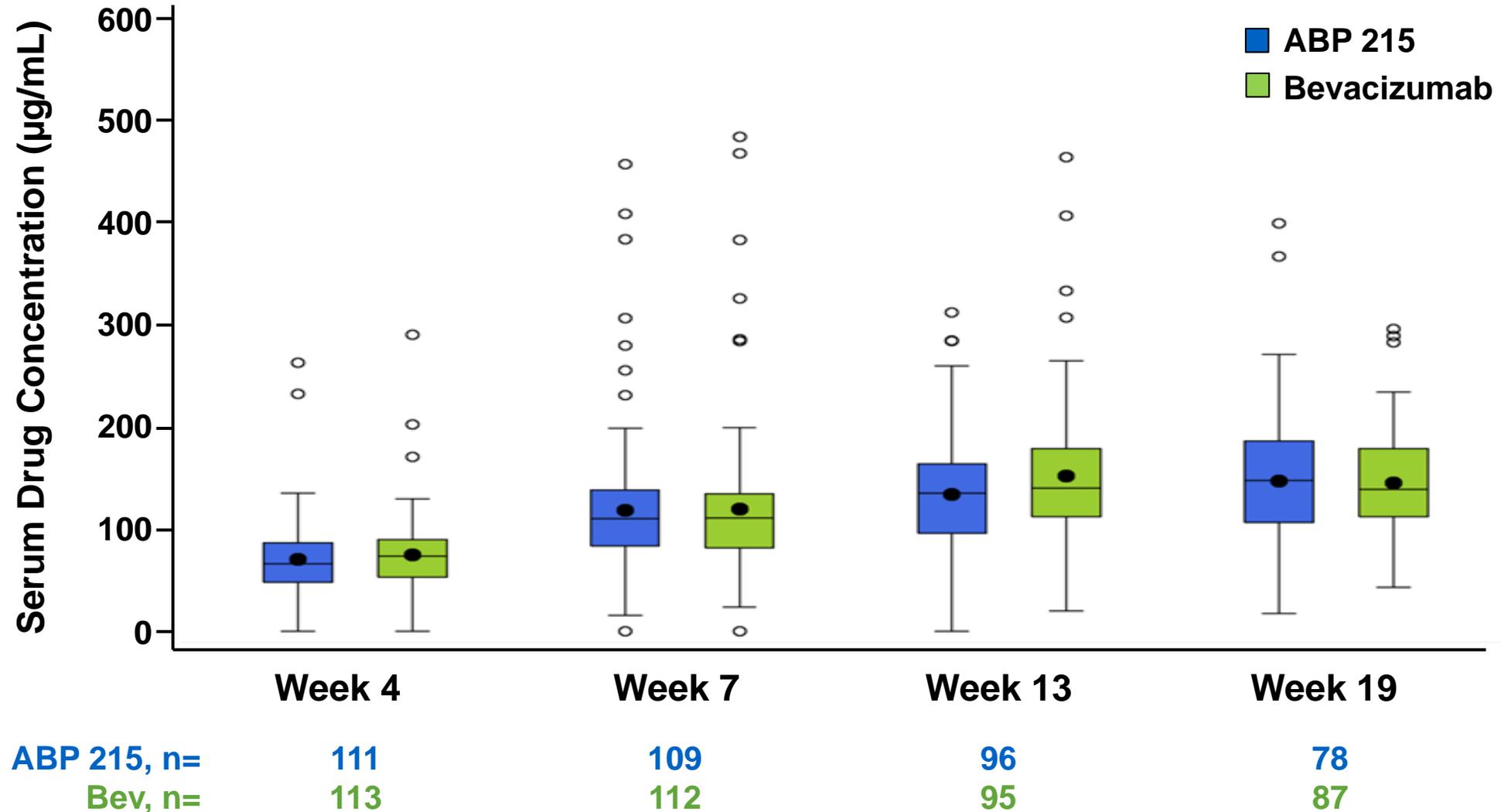
- ◆ **Scientific justification for extrapolation of biosimilarity to each condition of use is provided**
 - √ *Mechanism of Action (VEGF neutralization)*
 - √ *PK distribution and clearance*
 - √ *Clinical considerations*

Amgen's Commitment

- ◆ **Increased access to an important oncology therapy**
- ◆ **Same pharmacovigilance system and manufacturing as other Amgen products**
- ◆ **High quality biosimilar option for patients**

Supporting Slides

NSCLC: PK Trough Concentrations by Visit (Female)



NSCLC: Exposure Summary – IP

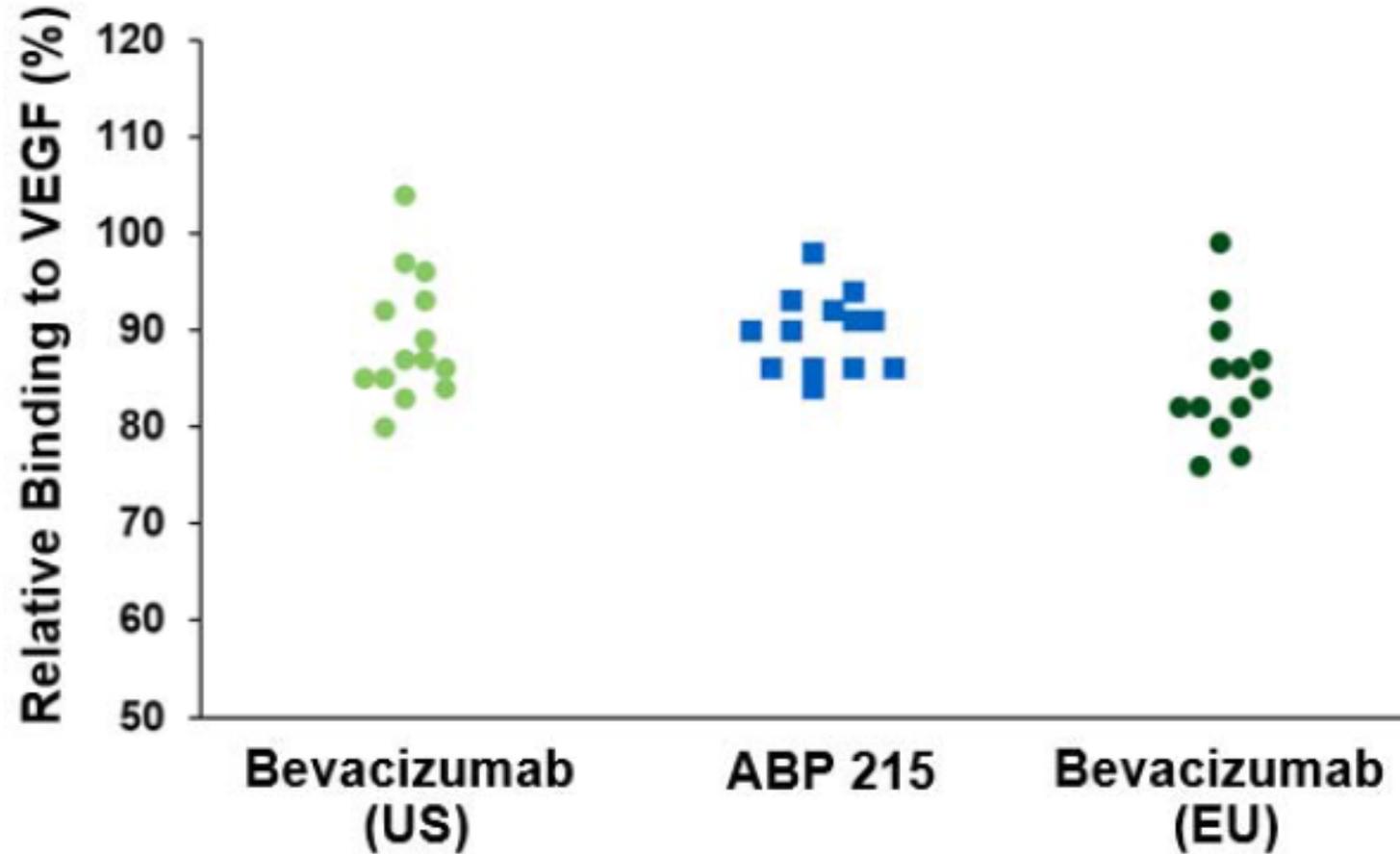
	ABP 215 N=324	Bevacizumab N=309
Total number of IP doses administered		
Mean (SD)	4.8 (1.76)	5.0 (1.61)
Median	6.0	6.0
Q1, Q3	4.0, 6.0	4.0, 6.0
Min, Max	1, 7	1, 6
Cumulative total dose of IP administered (mg/kg)^a		
Mean (SD)	71.3 (26.32)	74.8 (24.22)
Median	90.0	90.0
Q1, Q3	60.0, 90.0	60.0, 90.0
Min, Max	15, 105	15, 90

a. Four subjects received partial dose and were therefore excluded.

NSCLC: Exposure Summary – Chemotherapy

	ABP 215 N=324	Bevacizumab N=309
Subjects receiving ≥ 1 dose of paclitaxel, n	323	309
Total number of doses administered		
Mean (SD)	4.5 (1.69)	4.7 (1.56)
Median	5.0	5.0
Q1, Q3	4.0, 6.0	4.0, 6.0
Min, Max	1, 7	1, 6
Subjects receiving ≥ 1 dose of carboplatin, n	320	309
Total number of doses administered		
Mean (SD)	4.6 (1.67)	4.7 (1.57)
Median	5.0	5.0
Q1, Q3	4.0, 6.0	4.0, 6.0
Min, Max	1, 7	1, 6

F2. pg 9. Relative Binding to VEGF



VEGF = vascular endothelial growth factor.

Binding was calculated relative to the ABP 215 reference standard.