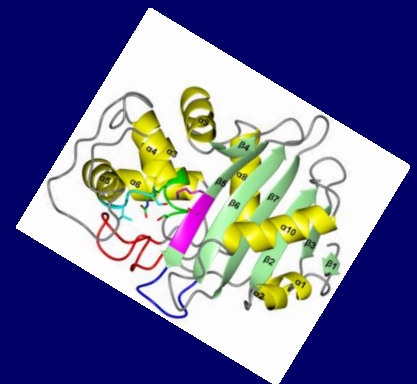
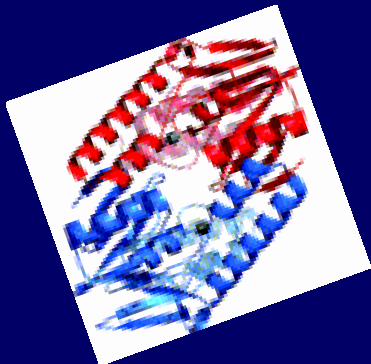


# ***CARBAPENEMASES WHAT THEY ARE AND WHAT THEY DO***

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# Financial Disclosures

## Retirement income

Johnson & Johnson  
Bristol-Myers Squibb  
Wyeth (Pfizer)

## Consultant and research income received (June 2012 to June 2013)

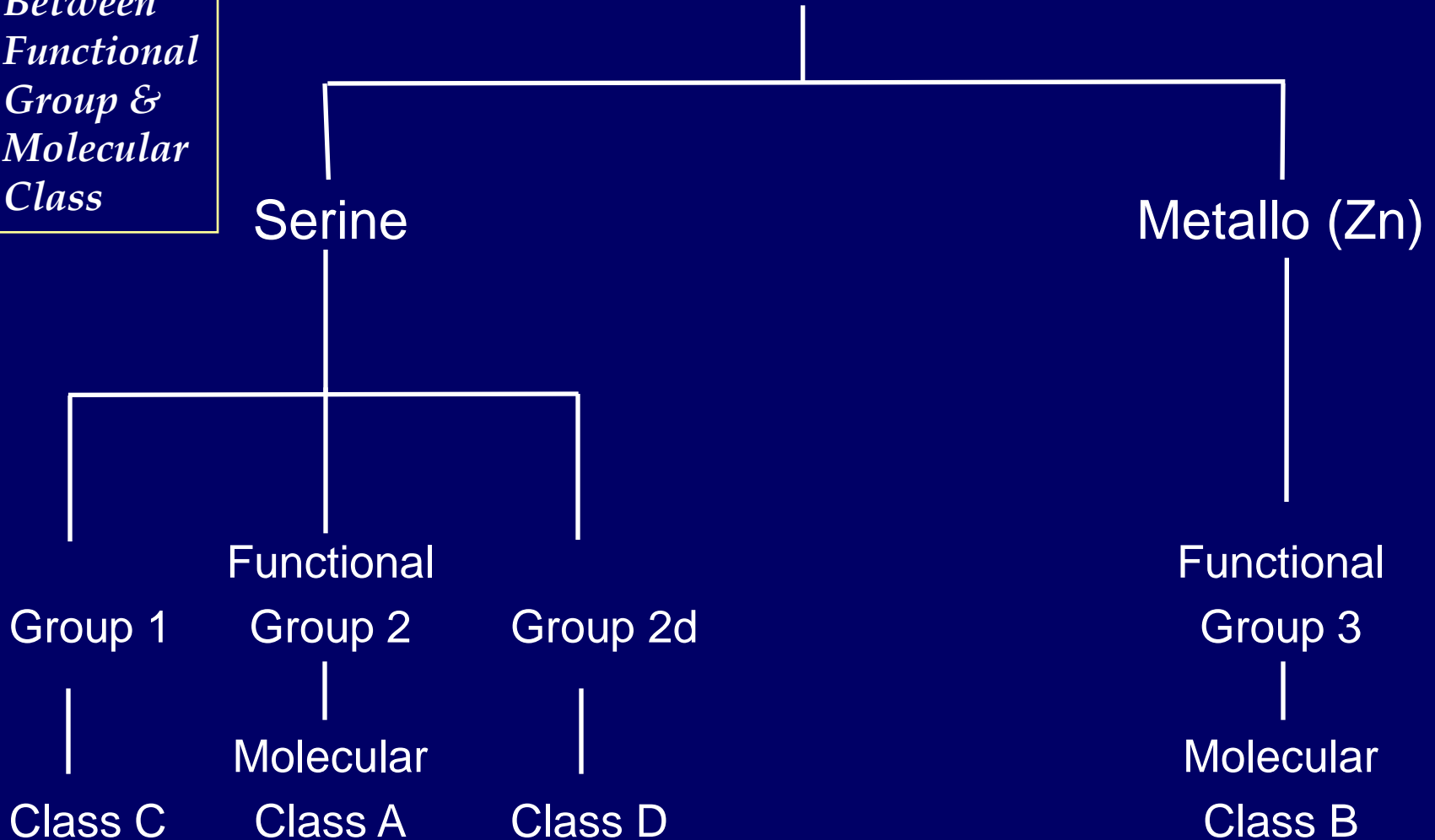
AstraZeneca  
Cubist  
Fedora  
Forest  
Medivir  
Merck

# Carbapenemases

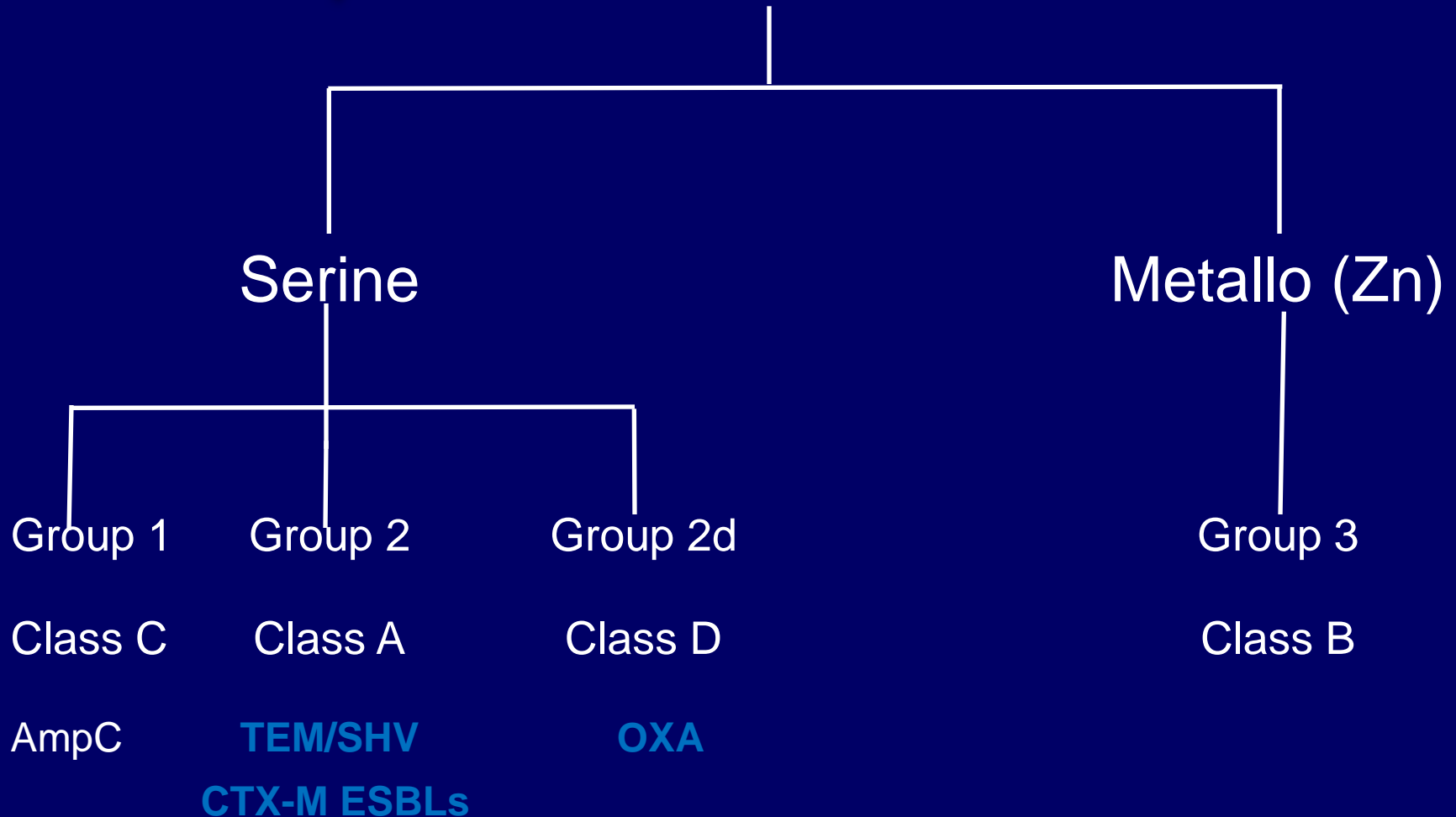
- ▣ What are they?
- ▣ What are the most common types?
- ▣ What molecular characteristics are responsible for their phenotypic behavior?

# $\beta$ -Lactamases

*Relations  
Between  
Functional  
Group &  
Molecular  
Class*

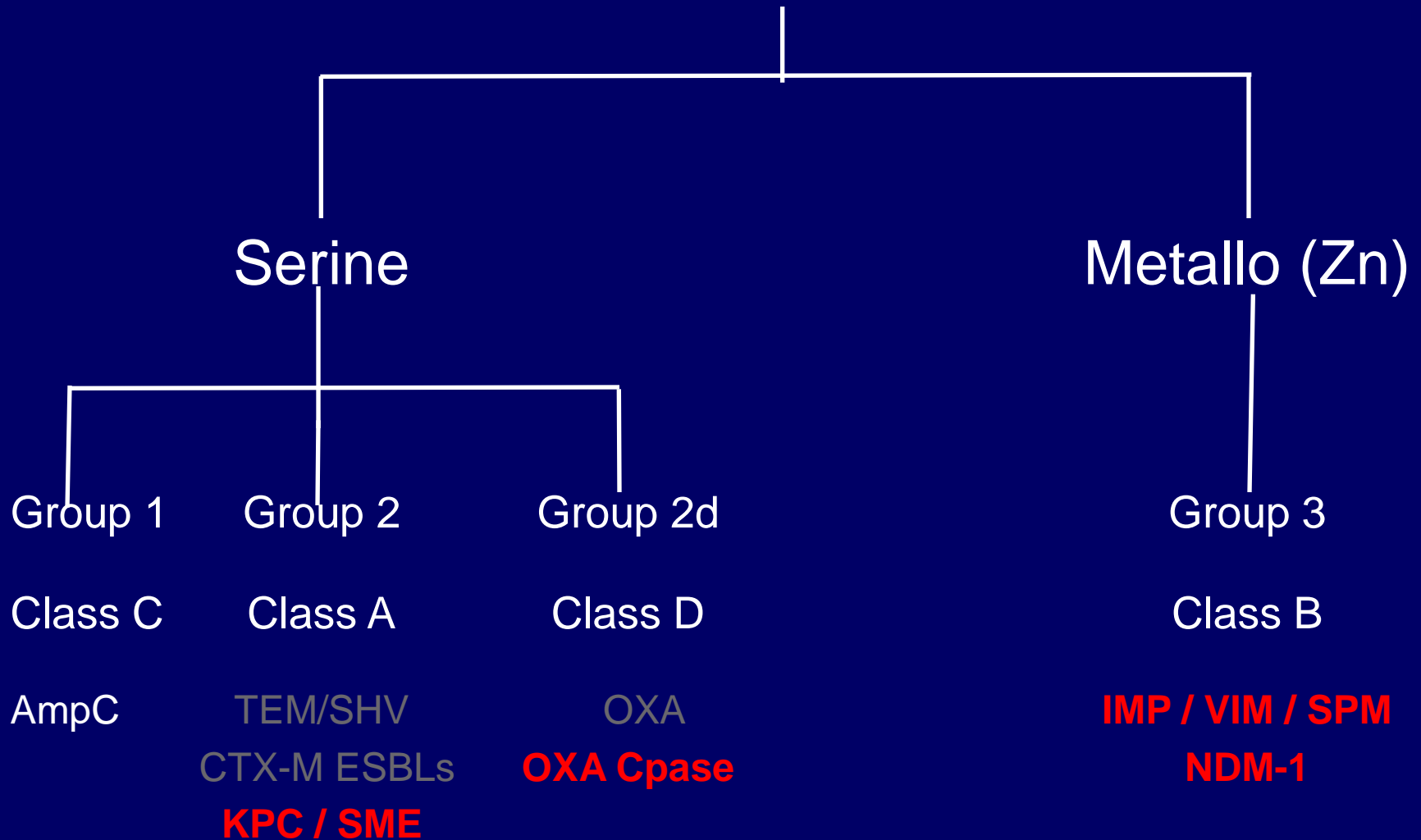


# $\beta$ -Lactamases / ESBLs



Based on Bush, Jacoby & Medeiros AAC:39:1211 (1995)

# $\beta$ -Lactamases / Carbapenemases



Based on Bush, Jacoby & Medeiros AAC:39:1211 (1995); Bush & Jacoby, AAC 54:969 (2010)

# Distinguishing Features of Carbapenem-Hydrolyzing $\beta$ -Lactamases

## ▣ Serine carbapenemases

- Hydrolyze cephalosporins, carbapenems, monobactams
- Inhibited (*in vitro*) by clavulanic acid, tazobactam and boronic acid

## ▣ Metallo- $\beta$ -lactamases

- Most can hydrolyze everything except monobactams
- Not inhibited by clavulanic acid
- Inhibited by metal chelators such as EDTA
- Diagnostic Etest with carbapenem/EDTA not always predictable
  - ▣ EDTA may permeabilize membranes to allow carbapenem easier access to AmpC hydrolysis, especially in derepressed mutants

# Increase in Major Carbapenemase Families

<u>Enzyme type</u>		<u>Number in class</u>	
		<u>1995</u>	<u>2010</u>
Serine	SME	1	3
	IMI	1	2
	NMC	1	1
	KPC	0	10
	GES	0	16
	OXA	0	37
Metallo- (Plasmid)	IMP	1	28
	VIM	0	26

<http://www.lahey.org/Studies> ; Bush, Jacoby & Medeiros AAC:39:1211 (1995);  
Bush & Jacoby, AAC 54:969 (2010)

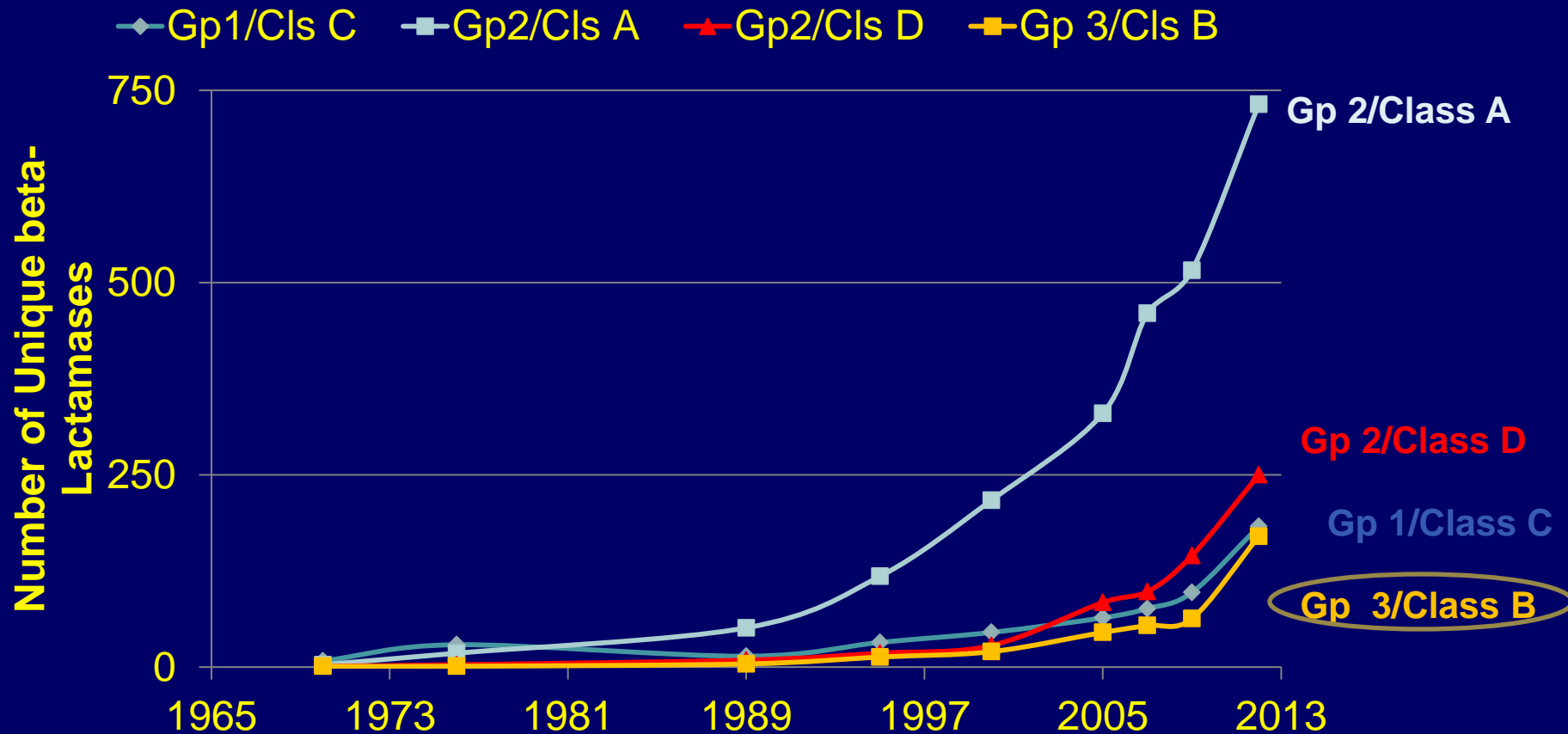


# Major Plasmid-Encoded Carbapenemase Families

<u>Enzyme type</u>	<u>Number in Class</u>			
	<u>1961</u>	<u>2000</u>	<u>2011</u>	<u>June 2013</u>
KPC	0	1	10	14
OXA	0	3	40	>60*
IMP	0	3	31	44
VIM	0	2	32	39
NDM	0	0	2	9

\*Total number of OXAs = 347, many from *Acinetobacter* spp.

# Increasing Numbers of $\beta$ -Lactamases by Class



Compilation of Unique  $\beta$ -Lactamase Sequences from Natural Isolates

Based on Bush & Fisher Annu. Rev. Microbiol. 2011. 65:455; Bush, Ann. NY Acad. Sc. (2013)  
Jacoby & Bush: <http://www.lahey.org/Studies/>

# Serine Carbapenemases

- ▣ Functional group 2f / Molecular class A
- ▣ Chromosomal often from environmental sources
  - Species-specific
  - IMI and NMC isolated from *Enterobacter* spp.
  - Most important are SME-1, -2, -3 from *S. marcescens*
- ▣ Plasmid-encoded
  - Most common are the KPC enzymes
    - ▣ *Klebsiella pneumoniae* carbapenemase
  - Global dissemination
  - May occur in clonal outbreaks (ST258)

Yang et al. AAC 34:755 (1990); Queenan et al. AAC 44:3035 (2000); Yigit et al. AAC 47:3881 (2003)

# Resistance Profiles Associated with SME and KPC Serine Carbapenemases

<u>Antibiotic</u>	MIC ( $\mu\text{g/mL}$ )	
	<u><i>S. marcescens</i> (SME-1)</u>	<u><i>K. oxytoca</i> (KPC-2)</u>
Ampicillin	>16	>64
Cefuroxime	>16	Not tested
Cefoxitin	>16	>64
Ceftazidime	<=1	>64
Cefotaxime	<=2	>64
Cefepime	<=2	>32
Aztreonam	>16	>64
Imipenem	>8	32
Meropenem	>8	32
Piperacillin-tazobactam	Not tested	>128/4
Tigecycline	<=1	Not tested
Gentamicin	<=1	8

Bush et al. IJAA 41:1-4 (2013); Yigit et al. AAC 47:3881 (2003)

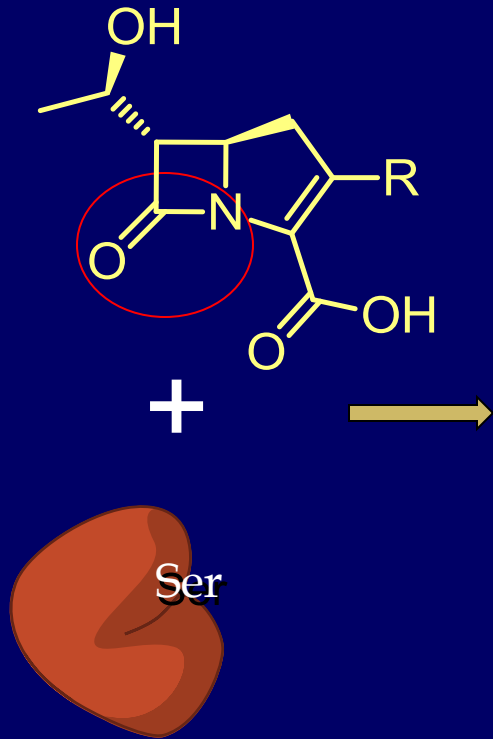
# Hydrolysis of Key Substrates by Serine $\beta$ -Lactamases

Antibiotic	$k_{cat}$ (sec <sup>-1</sup> ) *		
	TEM-1	SME-3	KPC-2
Penicillin G	1600	6.3	530
Cephaloridine	1500	1370	Not tested
Cefotaxime	9	1.5	22
Ceftazidime	0.3	0.14	$\leq 0.12$
Aztreonam	<1	148	30
Imipenem	<1	322	15
Meropenem	Not tested	3.2	4.0

\*Molecules of substrate hydrolyzed per sec per molecule of enzyme

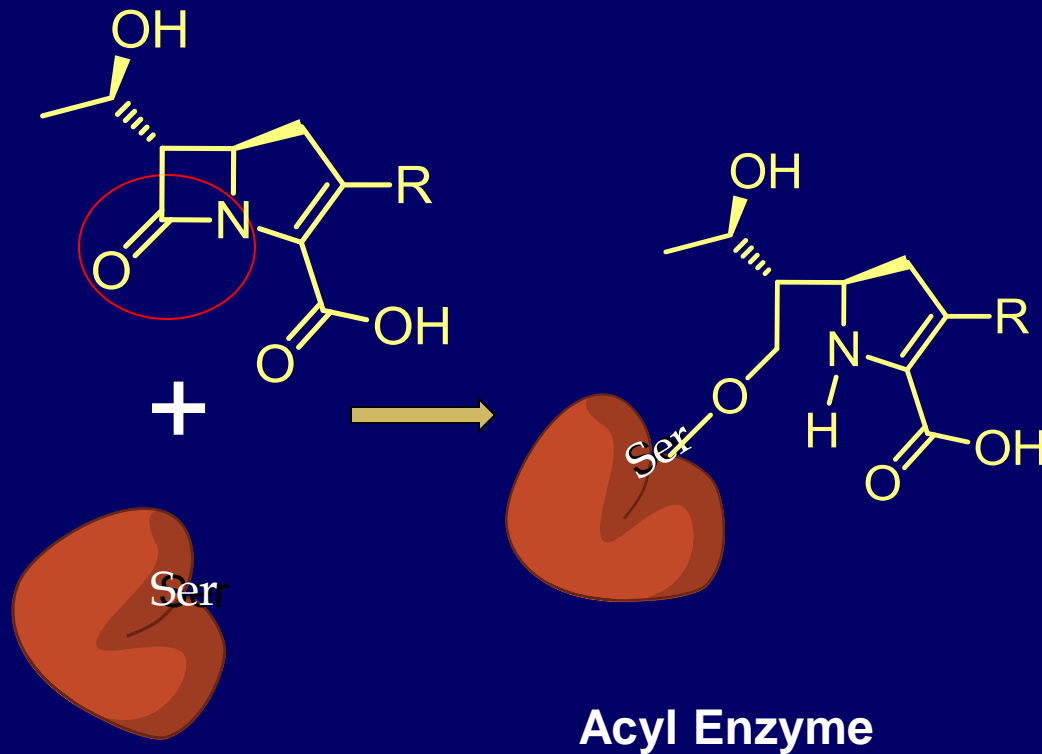
Raquet et al. J. Mol. Biol. 244:625 (1994); Queenan et al. AAC (2006);  
Yigit et al. AAC 47:3881 (2003)

# Hydrolysis of a Carbapenem by a Serine $\beta$ -Lactamase

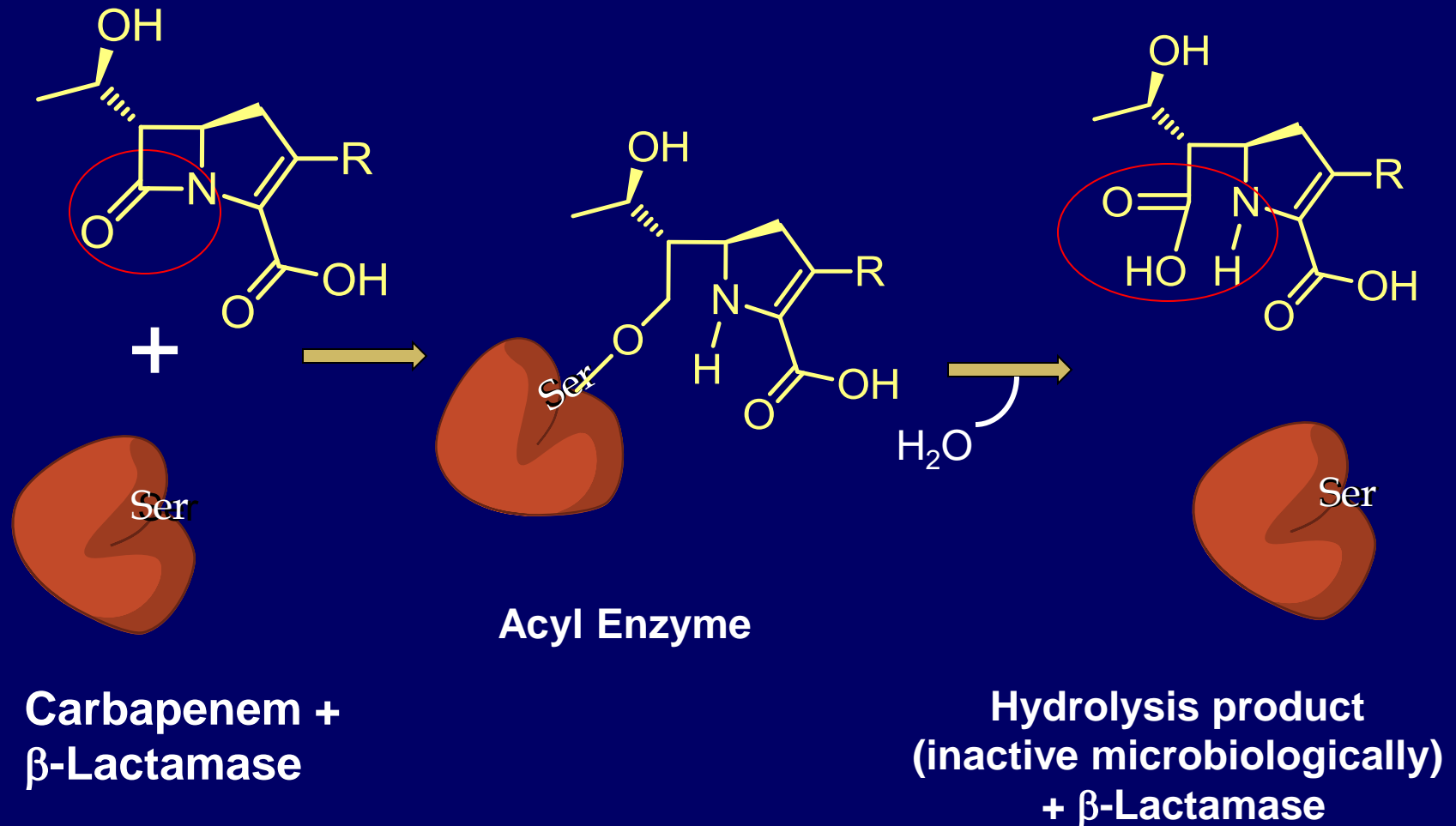


Carbapenem +  
 $\beta$ -Lactamase

# Hydrolysis of a Carbapenem by a Serine $\beta$ -Lactamase



# Hydrolysis of a Carbapenem by a Serine $\beta$ -Lactamase





# KPC Carbapenemases (Early History)

- KPC- family of serine carbapenemases (transferable)
  - High level of enzyme production encoded in integrons on plasmids
  - Major resistance problem in Brooklyn/New York City (*K. pneumoniae*)
- Initial KPC-1 isolate from *K. pneumoniae* in North Carolina (1997) now known to be KPC-2 (Yigit et al., AAC45:1151, 2001, + Erratum))
- KPC-3 identified in New Jersey and New York in early 2000s
- Each enzyme differs by one amino acid
- Generally appear with multiple enzymes per isolate
- Marketed  $\beta$ -lactamase inhibitors are not effective *in vivo*
  - Promising inhibitory activity by avibactam and RPX7009
  - Boronic acid inhibition used as a diagnostic test

# Inhibitor Resistance of KPC-2

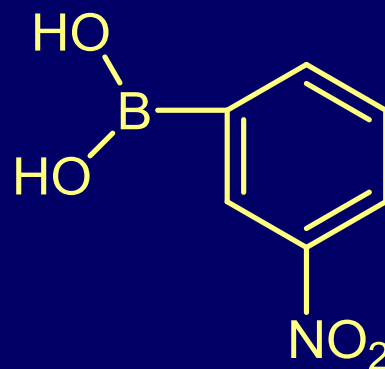
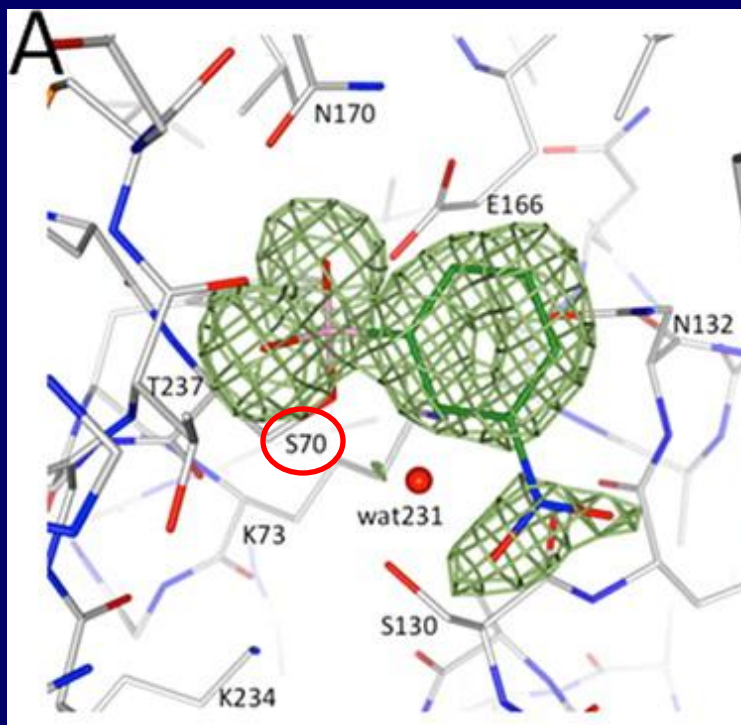
- KPC-2 is considered to be poorly susceptible to current inhibitors
- Inhibitor hydrolysis occurs before inactivation
- Tazobactam is the most effective of the three inhibitors tested

<u>Enzyme</u>	<u>Partition ratios (<math>k_{cat}/k_{inact}</math>)</u>		
	<u>Clavulanic acid</u>	<u>Sulbactam</u>	<u>Tazobactam</u>
KPC-2	2,500	1,000	500
TEM-2*	115	7,000	125

- Turnover of numbers for KPC-2 before inactivation were higher than for TEM-2 for clavulanic acid and tazobactam.

\*Turnover numbers before inactivation: Fisher et al. Biochemistry 17: 2180 (1978). Fisher et al. Biochemistry 20: 2726 (1981). Bush et al. AAC 37:851 (1993).

# Electron Density Map of the Serine $\beta$ -Lactamase Inhibitor 3-Nitrophenyl Boronic Acid in the Active Site of KPC-2



- Reversible covalent interaction with active site serine
- $K_m = 1.0 \mu\text{M}$

# Identification of KPC Carbapenemases

KPC	Species	Year	Location
1 / 2	<i>K. pneumoniae</i> / <i>K. oxytoca</i>	1996, 1998	East coast USA
3	<i>K. pneumoniae</i>	2000	New York
4	<i>Enterobacter cancerogenus</i>	2003	Scotland
5	<i>P. aeruginosa</i>	2006	Puerto Rico
6	<i>K. pneumoniae</i>	2003	Puerto Rico
7	<i>K. pneumoniae</i>	2007	Ohio
8	<i>K. pneumoniae</i>	2008	Puerto Rico
9	<i>E. coli</i>	2009	Israel
10	<i>A. baumannii</i>	2009	Puerto Rico
11	<i>K. pneumoniae</i>	2010	Greece
12-15	No data	2011-2013	No data

# OXA-Carbapenemases

- Most diverse family of carbapenemases
  - 9 remotely related sub-families (36-50% homology common)
- Non-fermenters are frequent hosts
- Plasmid-encoded enzymes
  - Initial OXA carbapenemase in *Acinetobacter baumannii* (1985 Scottish isolate) = ARI-1 or OXA-23
    - Now in England, Brazil, Singapore, Polynesia, Korea and China
  - OXA-58 from central and southern Europe, Argentina, Kuwait
- Chromosomal OXA-51/69-like enzymes in all *A. baumannii*
- OXA-23 outbreak in MDR - *A. baumannii* clones in Rio de Janeiro
  - January 2006-September 2007: 110 isolates; 5 genotypes
- Genomic sequencing of *Acinetobacter* spp. from various labs has added >85 new variants in the past 6 months

Perez et al. AAC 51:3471 (2007); Queenan & Bush, Clin Micro Rev 20:440 (2008); Carvalho, Carvalho-Assef, Peirano, Santos, Pereira, Asensi. Intl. J. Antimicrob. Agents 34:25 (2009).

# Metallo- $\beta$ -lactamases

- Functional group 3, Molecular class B
- Chromosomal
  - *Bacteroides* spp., *Bacillus* spp., *Stenotrophomonas maltophilia*
- Plasmid-encoded MBLs first described in Japan in 1990
  - VIM, IMP, and NDM are major families
  - NDM has become one of the most wide-spread carbapenemases globally
- Multiple  $\beta$ -lactamases of different classes usually found in carbapenemase-producing bacteria

# Chromosomal Metallo- $\beta$ -Lactamases

- All are co-produced with a more efficient  $\beta$ -lactamase
- *Bacillus cereus* –prototypical MBL in the literature
- *Bacillus anthracis* Bla2 [Materon et al. AAC 47:2040, 2003]
  - 92% identity with *B. cereus* 569H MBL
  - Poor affinity for standard substrates
  - Catalytic efficiency ( $k_{cat}/K_m$ ) 4-fold lower for imipenem than for cefotaxime; at least one order of magnitude lower than for Bla1 penicillin hydrolysis
- *Stenotrophomonas maltophilia* L1
  - The only tetrameric  $\beta$ -lactamase
- *Aeromonas sobria*
  - Asb1, group 1 cephalosporinase [Rasmussen et al. AAC 38:2078, 1994]
  - Asb2, group 2d oxacillinase
  - AsbM1, group 3 MBL [Yang & Bush, FEMS Microbiol. Lett. 127:193, 1996]

# Occurrence of Transferable Metallo- $\beta$ -Lactamases

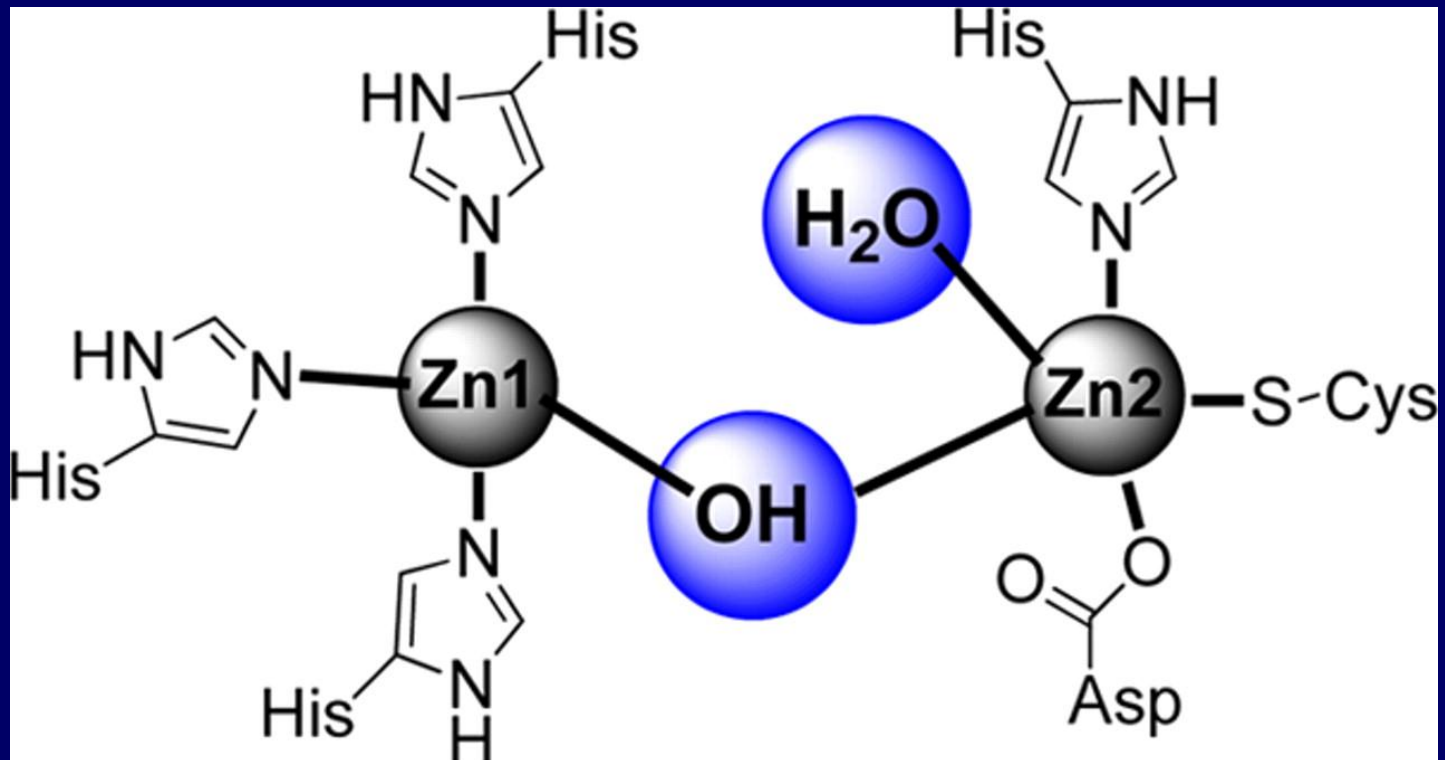
- IMP on transferable element in 1991 in Japan (Bandoh et al., AAC 35:371. 1991 )
- VIM appeared in Italy, France, Greece (1997)
- Eventually, IMP in Europe and VIM in Japan
- SPM-1 appeared to be confined to Brazil or patients associated with Brazil
- Rare in the United States
- MBLs usually are produced in combination with at least one other  $\beta$ -lactamase
  - Plasmids for IMP enzymes may be lost when antibiotic selection pressure removed
  - Metallo- $\beta$ -lactamases often have lower catalytic efficiencies than other common  $\beta$ -lactamases



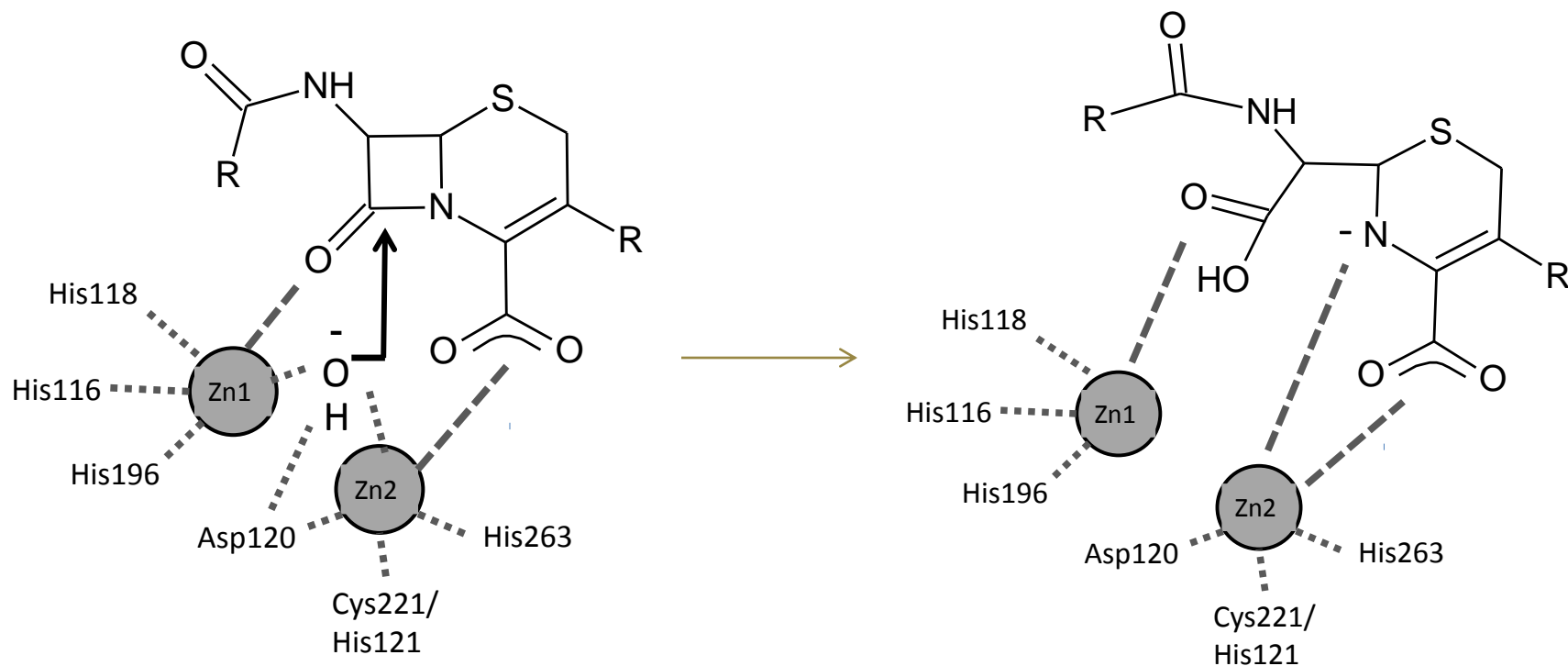
# Molecular Diversity of MBLs

Molecular Subclasses	Typical enzymes	Substrate Preference	Number of active site Zn <sup>2+</sup> atoms
B1a	BCII, IMP family, VIM family, SPM-1, CcrA	All $\beta$ -lactams except monobactams	2
B1b	NDM family	All $\beta$ -lactams except monobactams	2
B2	CphA, Sfh-1	Carbapenems	1
B3	L1, FEZ-1, CAU-1	All $\beta$ -lactams except monobactams	2

# Schematic representation of the Zn<sup>2+</sup>-binding site of dinuclear subclass B1 metallo- $\beta$ -lactamases such as *B. cereus* BclI.

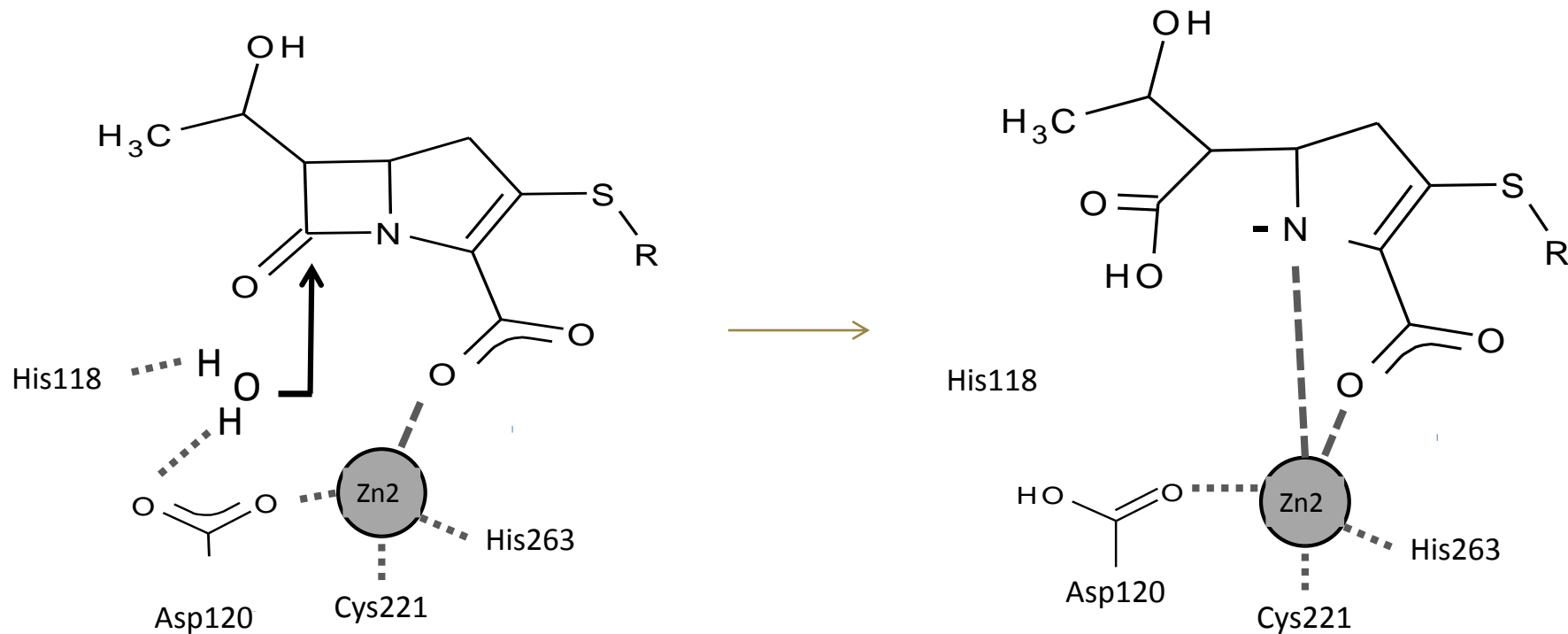


# Schematic Illustration of Cephalosporin Binding to Di-zinc Metallo- $\beta$ -Lactamase Active Site (subclasses B1 and B3)



Adapted from T. Palzkill *Annals NY Acad Sci.* 1277:91-104 (2013)

# Carbapenem Substrate and Anionic Intermediate Binding to Mono-zinc Metallo- $\beta$ -lactamase Active Site (subclass B2).



Adapted from T. Palzkill *Annals NY Acad Sci.* 1277:91-104 (2013)

# Microbiological Impact of Plasmid-Encoded Metallo- $\beta$ -Lactamases

<u>Organism</u>	<u>Bla</u>	MIC in $\mu\text{g/ml}$				
		<u>CAZ</u>	<u>CTX</u>	<u>FEP</u>	<u>IMP</u>	<u>MER</u>
<i>P. aeruginosa</i>	None	8	2	2	2	0.5
<i>P. aeruginosa</i>	VIM-1	>128	>128	>128	>128	128
<i>P. aeruginosa</i>	VIM-2	16	256	16	16	8
<i>P. aeruginosa</i>	VIM-4	>512	128	256	256	256
<i>P. aeruginosa</i>	IMP-1	--	--	8	16	--
<i>P. aeruginosa</i>	IMP-9	256	256	--	32	8
<i>P. aeruginosa</i>	IMP-16	--	--	>32	16	
<i>P. aeruginosa</i>	SPM-1	--	--	>32	>32	--

Franceschini et al. AAC 44:3003 (2000); Xiong et al. AAC 50:355 (2006); Marra et al. AAC 50:368 (2006); Fiett et al. AAC 50:880 (2006)

# NDM – 1 : The Newest Star in the $\beta$ -Lactamase Arena

- Swedish patient returning from India where he had been in a surgical ward
- Urinary isolate grew a carbapenem-R *Klebsiella pneumoniae* strain – not responsible for infection
  - ST14 single variant locus of ST15
  - ST15 associated with CTX-M and KPC *Klebsiellae*
- MBL named NDM-1 (New Delhi Metallo- $\beta$ -lactamase)
- Low identity with other MBLs
  - Most similar MBLs to VIM-1/VIM-2, with 32.4% identity
- Susceptibility profile
  - MICs >32  $\mu$ g/ml for all  $\beta$ -lactams including carbapenems, also ciprofloxacin
  - Colistin MIC 0.75  $\mu$ g/ml

# Unimpressive Hydrolysis Profile of NDM-1

- Hydrolysis profile similar to IMP-1 & VIM-2
  - Lower hydrolysis rates for most substrates
    - NDM-1 : Penicillin /cephalosporin  $k_{cat}$  values 1 - 15 sec<sup>-1</sup>
    - IMP/VIM: Penicillin/cephalosporin  $k_{cat}$  values 3 - 950 sec<sup>-1</sup>
  - Less efficient hydrolysis of imipenem than IMP-1 and VIM-2 ; MEM similar

	Imipenem			Meropenem		
	$K_m$ (μM)	$k_{cat}$ (s <sup>-1</sup> )	$k_{cat}/K_m$	$K_m$ (μM)	$k_{cat}$ (s <sup>-1</sup> )	$k_{cat}/K_m$
NDM-1	94	20	0.21	49	12	0.25
IMP-1	39	46	1.2	10	50	0.12
VIM-2	10	10	1.0	5	1	0.28

# Multiple $\beta$ -Lactamases are Common in Enterobacteriaceae

- KPC enzymes are frequently co-produced with weak ESBLs or strong penicillinases
  - SHV-11 or SHV-12; TEM-1; OXA-9
- Metallo-  $\beta$ -lactamases, chromosomal and plasmid-encoded, are always produced with at least one other  $\beta$ -lactamase
- Greek isolate of *K. pneumoniae*
  - Plasmid-encoded TEM-1 and CMY-2 cephalosporinase
  - ESBL CTXM-15
  - Two carbapenemases
    - VIM-19 and the serine carbapenemase KPC-2



# Two *Enterobacter cloacae* Isolates Produced KPC-3 and VIM-1 in Indianapolis

- Two patients admitted 11 months apart
  - Isolates collected 33 days apart in same ward in same hospital (2011)

MICs ( $\mu\text{g/ml}$ )	<u>Patient #1</u>	<u>Patient #2</u>
Imipenem	4	8
Meropenem	4	1

- Full sequencing confirmed KPC-3 and VIM-1 in both isolates
- MLST and plasmid profiles indicated non-clonal origins
- In 2012 an additional 41 CRE isolates were found positive for *bla*<sub>KPC</sub> by PCR.
  - 7 isolates were also positive for *bla*<sub>VIM</sub>

# Summary

- Carbapenemase production is increasing in Gram-negative bacteria
- Serine carbapenemases of clinical significance include the KPC enzymes and the OXA family of enzymes
- The most important metallo- $\beta$ -lactamases are in the IMP, VIM and NDM families
- Carbapenemases are often less efficient at hydrolyzing  $\beta$ -lactams than other  $\beta$ -lactamases
- Multiple  $\beta$ -lactamase production in a single organism must be considered in the development of new agents