

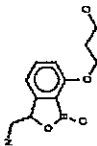
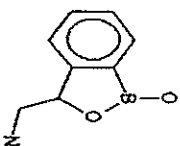
Reg #	Structure	
A46		
A1		
		$32 < = *$ $32 - 8.01 = **$ $8.00 - 2.00 = ***$ $2 > = ****$
		E. cloacae ATCC 13047 MIC
		E. coli ECM 1194 MIC
		E. coli ECM 1888 MIC
		E. faecalis ATCC 29212 MIC
		H. influenzae ATCC 49766 MIC
		K. pneumoniae ATCC 13883 MIC
		P. aeruginosa ATCC 27853 MIC
		P. aeruginosa PAO1 MIC
		P. aeruginosa PAO1d3 MIC
		S. aureus ATCC 29213 MIC
		S. pneumoniae ATCC 6301 MIC

FIG 1A

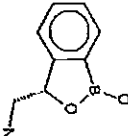
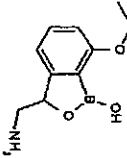
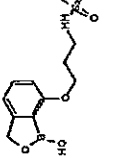
Reg #	Structure			
A2		****	****	<i>E. cloacae</i> ATCC 13047 MIC
A2		***	***	<i>E. coli</i> ECM 1194 MIC
A2		****	***	<i>E. coli</i> ECM 1888 MIC
A2		*	*	<i>E. faecalis</i> ATCC 29212 MIC
A2		***	***	<i>H. influenzae</i> ATCC 49766 MIC
A2		***	***	<i>K. pneumoniae</i> ATCC 13883 MIC
A2		**	*	<i>P. aeruginosa</i> ATCC 27853 MIC
A2		****	*	<i>P. aeruginosa</i> PAO1 MIC
A2		****	**	<i>P. aeruginosa</i> PAO1d3 MIC
A2		*	**	<i>S. aureus</i> ATCC 29213 MIC
A2		***	***	<i>S. pneumoniae</i> ATCC 6301 MIC
A52		****	****	
A52		***	***	
A52		***	***	
A52		*	*	
A52		***	***	
A52		***	***	
A52		**	*	
A52		****	**	
A52		****	***	
A15		**	**	
A15		**	***	
A15		***	***	
A15		*	*	
A15		**	***	
A15		**	***	
A15		*	*	
A15		***	***	

FIG 1B

Reg #	Structure	A49	A11	A16
E. cloacae ATCC 13047 MIC		*****	***	**
E. coli ECM 1194 MIC		***	***	**
E. coli ECM 1888 MIC		***	***	***
E. faecalis ATCC 29212 MIC		***	**	*
H. influenzae ATCC 49766 MIC		*****	***	**
K. pneumoniae ATCC 13883 MIC		*****	***	**
P. aeruginosa ATCC 27853 MIC		***	***	*
P. aeruginosa PAO1 MIC		****		
P. aeruginosa PAO1d3 MIC		*****		
S. aureus ATCC 29213 MIC		***	**	**
S. pneumoniae ATCC 6301 MIC		***	***	***

FIG 1C

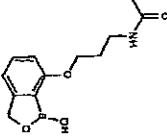
Reg #	Structure	A12	A13	A39
E. cloacae ATCC 13047 MIC		**	***	****
E. coli ECM 1194 MIC		**	***	***
E. coli ECM 1888 MIC		***	***	***
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		**	***	****
K. pneumoniae ATCC 13883 MIC		*	***	***
P. aeruginosa ATCC 27853 MIC		*	**	***
P. aeruginosa PAO1 MIC				****
P. aeruginosa PAO1d3 MIC				****
S. aureus ATCC 29213 MIC		**	**	*
S. pneumoniae ATCC 6301 MIC		***	****	****

FIG 1D

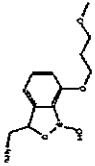
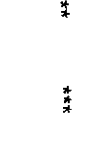
Reg #	Structure	A38	A3	A16
E. cloacae ATCC 13047 MIC		***	***	***
E. coli ECM 1194 MIC		***	***	***
E. coli ECM 1888 MIC		***	***	***
E. faecalis ATCC 29212 MIC		*	*	
H. influenzae ATCC 49766 MIC		***	***	
K. pneumoniae ATCC 13883 MIC		***	**	
P. aeruginosa ATCC 27853 MIC		***	*	
P. aeruginosa PAO1 MIC		***		
P. aeruginosa PAO1d3 MIC		***		
S. aureus ATCC 29213 MIC		*	*	
S. pneumoniae ATCC 6301 MIC		***	***	***

FIG 1E

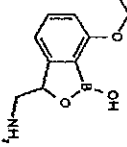
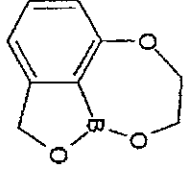
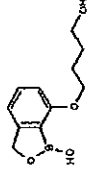
Reg #	Structure	
A52		
A61		
A22		
	E. cloacae ATCC 13047	
	MIC	***
	E. coli ECM 1194	
	MIC	***
	E. coli ECM 1888	
	MIC	***
	E. faecalis ATCC 29212	
	MIC	*
	H. influenzae ATCC 49766	
	MIC	***
	K. pneumoniae ATCC 13883	
	MIC	***
	P. aeruginosa ATCC 27853	
	MIC	**
	P. aeruginosa PAO1	
	MIC	*
	P. aeruginosa PAO1d3	
	MIC	*
	S. aureus ATCC 29213	
	MIC	**
	S. pneumoniae ATCC 6301	
	MIC	***

FIG 1F

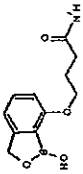
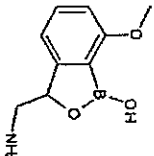
Reg #	Structure	
A10		
A14		
A53		
	<i>E. cloacae</i> ATCC 13047 MIC	****
	<i>E. coli</i> ECM 1194 MIC	***
	<i>E. coli</i> ECM 1888 MIC	***
	<i>E. faecalis</i> ATCC 29212 MIC	*
	<i>H. influenzae</i> ATCC 49766 MIC	***
	<i>K. pneumoniae</i> ATCC 13883 MIC	***
	<i>P. aeruginosa</i> ATCC 27853 MIC	***
	<i>P. aeruginosa</i> PAO1 MIC	*
	<i>P. aeruginosa</i> PAO1d3 MIC	**
	<i>S. aureus</i> ATCC 29213 MIC	*
	<i>S. pneumoniae</i> ATCC 6301 MIC	***

FIG 1G

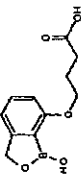
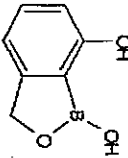
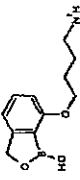
Reg #	Structure	A9	A26	A17
E. cloacae ATCC 13047 MIC				
E. coli ECM 1194 MIC		**	**	***
E. coli ECM 1888 MIC		***	**	**
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	***
K. pneumoniae ATCC 13883 MIC		**	**	***
P. aeruginosa ATCC 27853 MIC		*	*	**
P. aeruginosa PAO1 MIC		*		
P. aeruginosa PAO1d3 MIC		**		
S. aureus ATCC 29213 MIC		*	*	**
S. pneumoniae ATCC 6301 MIC		***	**	***

FIG. 1H

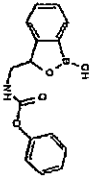
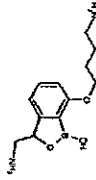
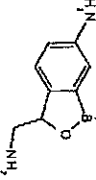
Reg #	Structure	
A29		
A41		
A32		
	E. cloacae ATCC 13047 MIC	***
	E. coli ECM 1194 MIC	**
	E. coli ECM 1888 MIC	**
	E. faecalis ATCC 29212 MIC	*
	H. influenzae ATCC 49766 MIC	***
	K. pneumoniae ATCC 13883 MIC	***
	P. aeruginosa ATCC 27853 MIC	***
	P. aeruginosa PAO1 MIC	
	P. aeruginosa PAO1d3 MIC	
	S. aureus ATCC 29213 MIC	*
	S. pneumoniae ATCC 6301 MIC	**

FIG 1J

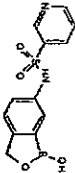
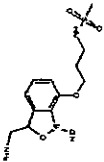
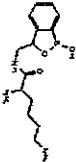
Reg #	Structure	
A4		
A44		
A27		
	<i>E. cloacae</i> ATCC 13047 MIC	*
	<i>E. coli</i> ECM 1194 MIC	**
	<i>E. coli</i> ECM 1888 MIC	**
	<i>E. faecalis</i> ATCC 29212 MIC	*
	<i>H. influenzae</i> ATCC 49766 MIC	***
	<i>K. pneumoniae</i> ATCC 13883 MIC	*
	<i>P. aeruginosa</i> ATCC 27853 MIC	*
	<i>P. aeruginosa</i> PAO1 MIC	**
	<i>P. aeruginosa</i> PAO1d3 MIC	***
	<i>S. aureus</i> ATCC 29213 MIC	*
	<i>S. pneumoniae</i> ATCC 6301 MIC	***

FIG 1K

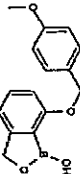
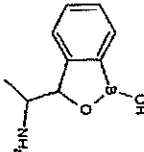
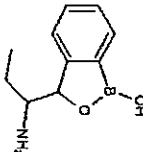
Reg #	Structure	A25	A54	A59
E. cloacae ATCC 13047 MIC				*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3 MIC				
S. aureus ATCC 29213 MIC		*	*	*
S. pneumoniae ATCC 6301 MIC		*	*	*

FIG. 1L

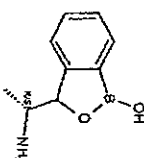
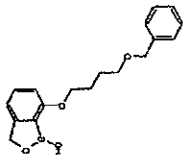
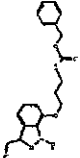
Reg #	Structure	A58	A21	A43
				
<i>E. cloacae</i> ATCC 13047 MIC	*	*	*	*
<i>E. coli</i> ECM 1194 MIC	*	*	*	*
<i>E. coli</i> ECM 1888 MIC	*	*	*	*
<i>E. faecalis</i> ATCC 29212 MIC	*	*	*	*
<i>H. influenzae</i> ATCC 49766 MIC	*	*	*	**
<i>K. pneumoniae</i> ATCC 13883 MIC	*	*	*	**
<i>P. aeruginosa</i> ATCC 27853 MIC	*	*	*	*
<i>P. aeruginosa</i> PAO1 MIC				
<i>P. aeruginosa</i> PAO1d3 MIC				
<i>S. aureus</i> ATCC 29213 MIC	*	*	**	*
<i>S. pneumoniae</i> ATCC 6301 MIC	*	*	**	***

FIG 1M

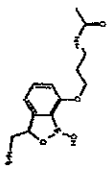
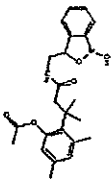
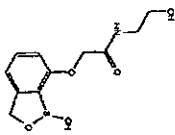
Reg #	Structure	A35	A27a	A7
E. cloacae ATCC 13047 MIC				+
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC		*	*	*
P. aeruginosa PAO1d3 MIC		***	*	*
S. aureus ATCC 29213 MIC		*	*	*
S. pneumoniae ATCC 6301 MIC		***	*	***

FIG. 1N

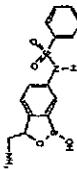
Reg #	Structure	A33	A34	A30
E. cloacae ATCC 13047 MIC		*		**
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*		*
H. influenzae ATCC 49766 MIC		**	**	***
K. pneumoniae ATCC 13883 MIC		*		*
P. aeruginosa ATCC 27853 MIC		*	**	**
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3 MIC				
S. aureus ATCC 29213 MIC		*	*	*
S. pneumoniae ATCC 6301 MIC		**	**	**

FIG 10

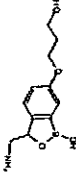
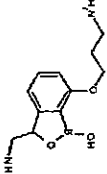
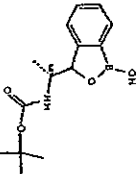
Reg #	Structure	A31	A36	A55
E. cloacae ATCC 13047 MIC				
E. coli ECM 1194 MIC	*	*	*	*
E. coli ECM 1888 MIC	*	*	*	*
E. faecalis ATCC 29212 MIC	*	*	*	*
H. influenzae ATCC 49766 MIC	***	**	*	*
K. pneumoniae ATCC 13883 MIC	*	*	*	*
P. aeruginosa ATCC 27853 MIC	*	*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3 MIC				
S. aureus ATCC 29213 MIC	*	*	*	*
S. pneumoniae ATCC 6301 MIC	*	**	*	*

FIG 1P

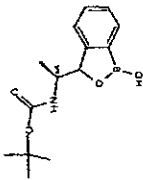
Reg #	Structure	A56	A40	A57
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	**	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3 MIC				
S. aureus ATCC 29213 MIC		*	**	*
S. pneumoniae ATCC 6301 MIC		*	**	*

FIG 10

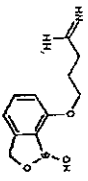
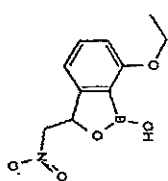
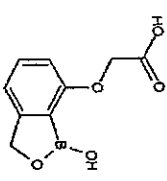
Reg #	Structure	A19	A51	A6
E. cloacae ATCC 13047 MIC				*
E. coli ECM 1194 MIC	*	*	*	*
E. coli ECM 1888 MIC	*	*	*	*
E. faecalis ATCC 29212 MIC	*	*	*	*
H. influenzae ATCC 49766 MIC	*	**	*	*
K. pneumoniae ATCC 13883 MIC	*	*	*	*
P. aeruginosa ATCC 27853 MIC	*	*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3 MIC				
S. aureus ATCC 29213 MIC	*	*	*	*
S. pneumoniae ATCC 6301 MIC	*	*	***	

FIG 1R

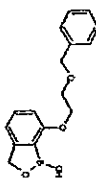
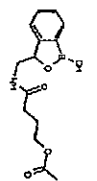
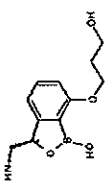
Reg #	Structure	A24	A28	A50
E. cloacae ATCC 13047 MIC				*
E. coli ECM 1194 MIC	*	*	*	*
E. coli ECM 1888 MIC	*	*	*	*
E. faecalis ATCC 29212 MIC	*	*	*	*
H. influenzae ATCC 49766 MIC	*	*	*	*
K. pneumoniae ATCC 13883 MIC	*	*	*	*
P. aeruginosa ATCC 27853 MIC	*	*	*	*
P. aeruginosa PAO1 MIC	*	*	*	*
P. aeruginosa PAO1d3 MIC	*	*	*	*
S. aureus ATCC 29213 MIC	*	*	*	*
S. pneumoniae ATCC 6301 MIC	*	*	*	**

FIG 1S

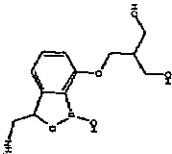
A45	Reg #	Structure
		
***	E. cloacae ATCC 13047	MIC
**	E. coli ECM 1194	MIC
*	E. coli ECM 1888	MIC
*	E. faecalis ATCC 29212	MIC
***	H. influenzae ATCC 49766	MIC
***	K. pneumoniae ATCC 13883	MIC
**	P. aeruginosa ATCC 27853	MIC
***	P. aeruginosa PAO1	MIC
***	P. aeruginosa PAO1d3	MIC
**	S. aureus ATCC 29213	MIC
*	S. pneumoniae ATCC 6301	MIC

FIG 17

FIG 2A

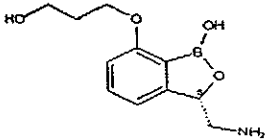
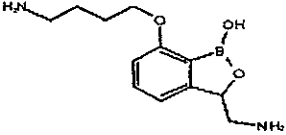
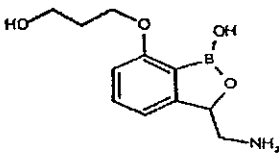
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
	400< = # 400 - 100.1 = ## 100 - 5 ### 5> = #####			
A49		####	####	
A41		####	####	####
A46		####	####	####

FIG 2B

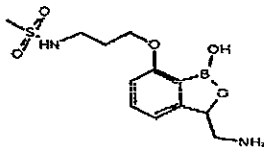
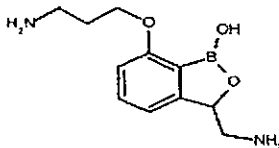
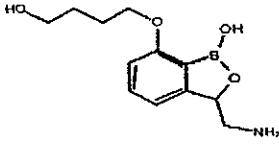
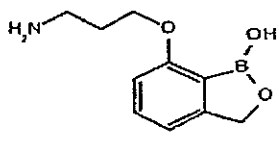
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A44		####	####	####
A36		####	####	####
A42		####	####	###
A11		####	####	###

FIG 2C

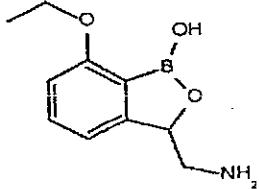
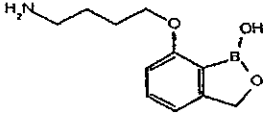
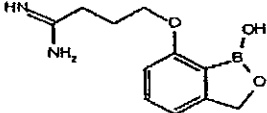
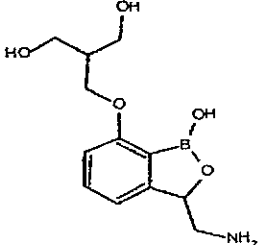
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A52		####	####	####
A17		####	####	###
A19		####	####	###
A45		####	####	####

FIG 2D

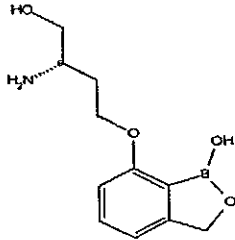
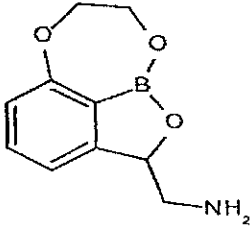
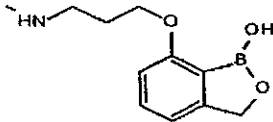
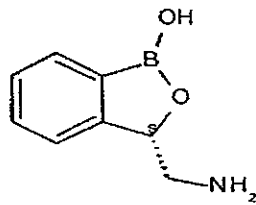
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1_LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC8325_LeuRS_wt IC50
A23		####	####	####
A39		####	####	####
A13		####	####	###
A2		####	####	####

FIG 2E

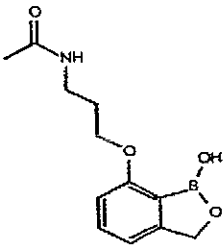
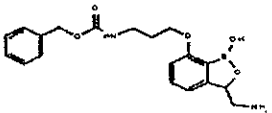
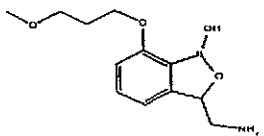
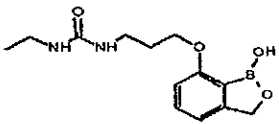
Compound ID	Structure	Escherichia_coll_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A12		####	####	###
A43		####	####	####
A38		####	###	###
A14		####	####	

FIG 2F

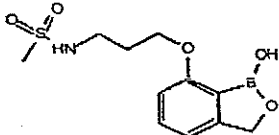
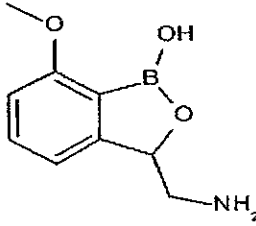
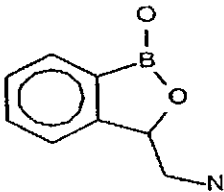
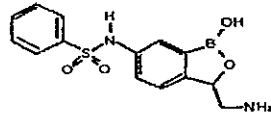
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A15		####	####	###
A53		####	####	####
A1		####	###	####
A33		####	###	###

FIG 2G

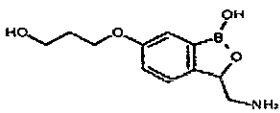
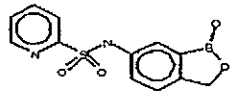
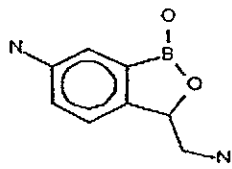
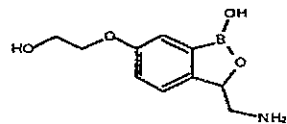
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A31		####	####	####
A3		####	####	###
A32		####	####	####
A30		####	###	####

FIG 2H

Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A34		####	###	###
A22		####	####	#
A58		###	###	#
A4		###	###	###

FIG 2I

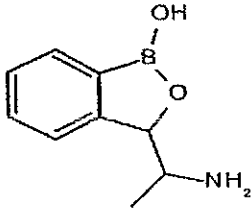
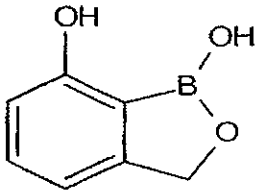
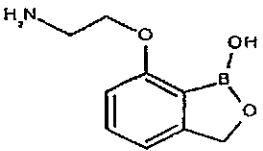
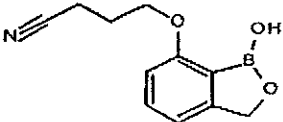
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1_LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC8325_LeuRS_wt IC50
A54		###	###	#
A26		###	#	###
A18		###	####	#
A16		###	####	#

FIG 2J

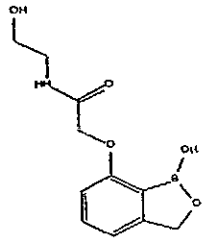
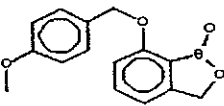
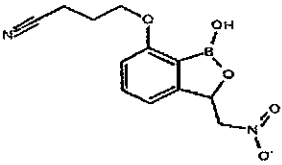
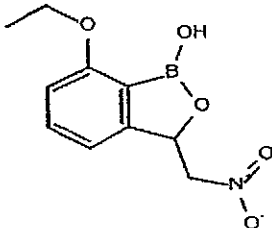
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1_LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC8325_LeuRS_wt IC50
A7		###	###	#
A25		###	###	#
A40		###	##	#
A51		###	##	#

FIG 2K

Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A6		##	###	
A56		##	##	#
A55		##	#	#
A59		#	#	#

FIG 2L

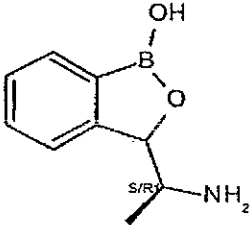
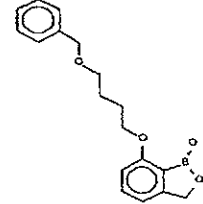
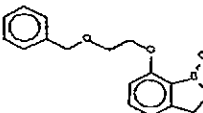
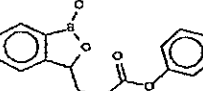
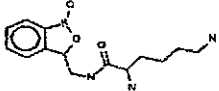
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A57		#	###	#
A21		#	#	#
A24		#	#	#
A29		#	#	#

FIG 2M

Compound ID	Structure	
A27		#
		#
		#

Escherichia_coli_K12_LeuRS_wt
IC50Pseudomonas_aeruginosa_PAO1
_LeuRS_wt
IC50Staphylococcus_aureus_NCTC83
25_LeuRS_wt
IC50