

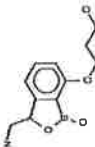
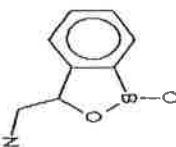
Reg #	Structure	
	$32 < = *$ $32 - 8.01 = **$ $8.00 - 2.00 = ***$ $2 > = ****$	
A46		
A1		
		E. cloacae ATCC 13047 MIC
		E. coli ECM 1194 MIC
		E. coli ECM 1888 MIC
		E. faecalis ATCC 29212 MIC
		H. influenzae ATCC 49766 MIC
		K. pneumoniae ATCC 13883 MIC
		P. aeruginosa ATCC 27853 MIC
		P. aeruginosa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

FIG 1A

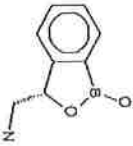
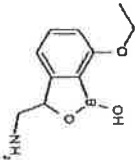
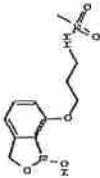
Reg #	Structure	A2	A52	A15
E. cloacae ATCC 13047 MIC				***
E. coli ECM 1194 MIC	***	***	***	**
E. coli ECM 1888 MIC	***	***	***	***
E. faecalis ATCC 29212 MIC	*	*	*	*
H. influenzae ATCC 49766 MIC	***	***	***	**
K. pneumoniae ATCC 13883 MIC	***	***	***	**
P. aeruginosa ATCC 27853 MIC	*	**	*	*
P. aeruginosa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

FIG 1B

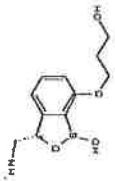
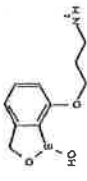
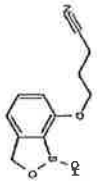
Reg #	Structure	A49	A11	A16
E. cloacae ATCC 13047 MIC				*****
E. coli ECM 1194 MIC	*****	***	***	***
E. coli ECM 1888 MIC	*****	***	***	***
E. faecalis ATCC 29212 MIC	***	**	*	*
H. influenzae ATCC 49766 MIC	*****	***	***	**
K. pneumoniae ATCC 13883 MIC	*****	***	**	**
P. aeruginosa ATCC 27853 MIC	***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

FIG 1C

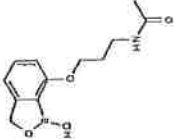
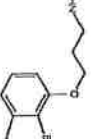
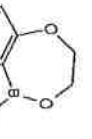
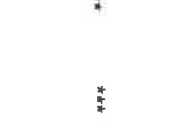
Reg #	Structure	A12	A13	A39
<i>E. cloacae</i> ATCC 13047 MIC		**	****	*****
<i>E. coli</i> ECM 1194 MIC		**	****	****
<i>E. coli</i> ECM 1888 MIC		****	****	****
<i>E. faecalis</i> ATCC 29212 MIC		*	*	*
<i>H. influenzae</i> ATCC 49766 MIC		***	****	*****
<i>K. pneumoniae</i> ATCC 13883 MIC		*	****	****
<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 1D

Reg #	Structure	A16	A3	A38
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

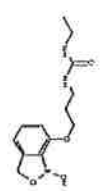
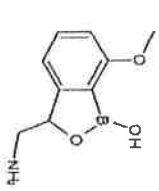
Reg #	Structure								
A10		*	**	***	*	*	*	*	***
A14		*	**	***	*	*	*	*	***
A53		****	***	***	*	***	***	***	****
					E. cloacae ATCC 13047 MIC				
					E. coli ECM 1194 MIC				
					E. coli ECM 1888 MIC				
					E. faecalis ATCC 29212 MIC				
					H. influenzae ATCC 49766 MIC				
					K. pneumoniae 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 1G

Reg #	Structure	A17	A26	A9
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

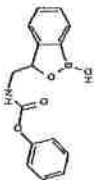
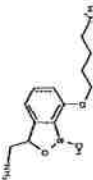
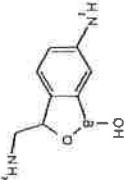
Reg #	Structure	
A29		
A41		
A32		
	<i>E. cloacae</i> ATCC 13047 MIC	***
	<i>E. coli</i> ECM 1194 MIC	**
	<i>E. coli</i> ECM 1888 MIC	**
	<i>E. faecalis</i> ATCC 29212 MIC	*
	<i>H. influenzae</i> ATCC 49766 MIC	**
	<i>K. pneumoniae</i> ATCC 13883 MIC	**
	<i>P. aeruginosa</i> ATCC 27853 MIC	**
	<i>P. 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 1J

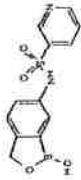
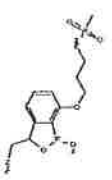
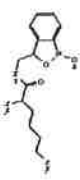
Reg #	Structure	A4	A44	A27
E. cloacae ATCC 13047 MIC		*	**	*
E. coli ECM 1194 MIC		*	*	**
E. coli ECM 1888 MIC		*	**	**
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		***	**	**
K. pneumoniae ATCC 13883 MIC		*	**	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC			**	
P. aeruginosa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

FIG 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

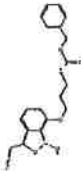
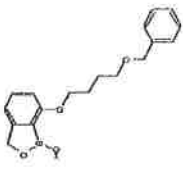
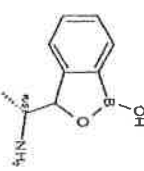
Reg #	Structure	A43	A21	A58
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		**	*	*
K. pneumoniae ATCC 13883 MIC		**	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa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

FIG 1M

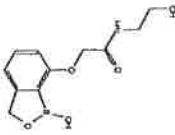
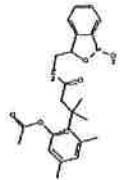
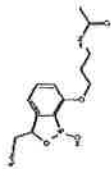
Reg #	Structure	A7	A27a	A35
				
E. cloacae ATCC 13047 MIC	*	*	††	
E. coli ECM 1194 MIC	*	*	*	
E. coli ECM 1888 MIC	*	*	*	
E. faecalis ATCC 29212 MIC	*	*	*	
H. influenzae ATCC 49766 MIC	*	*	*	
K. pneumoniae ATCC 13883 MIC	*	*	*	
P. aeruginosa ATCC 27853 MIC	*	*	*	
P. aeruginosa PAO1 MIC	*	*	*	
P. aeruginosa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

FIG IN

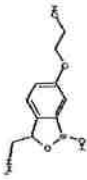
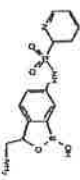
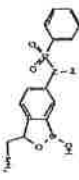
Reg #	Structure	
A30		
A34		
A33		
	E. cloacae ATCC 13047 MIC	**
	E. coli ECM 1194 MIC	*
	E. coli ECM 1888 MIC	*
	E. faecalis ATCC 29212 MIC	*
	H. influenzae ATCC 49766 MIC	***
	K. pneumoniae ATCC 13883 MIC	*
	P. aeruginosa ATCC 27853 MIC	**
	P. aeruginosa PAO1 MIC	**
	P. aeruginosa PAO1d3 MIC	*
	S.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

FIG 10

Reg #	Structure	
A55		
A36		
A31		
	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

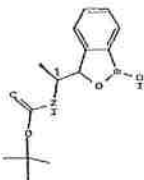
Reg #	Structure	A56	A40	A57
E. cloacae ATCC 13047		*	*	*
MIC				
E. coli ECM 1194		*	*	*
MIC				
E. coli ECM 1888		*	*	*
MIC				
E. faecalis ATCC 29212		*	*	*
MIC				
H. influenzae ATCC 49766		*	**	*
MIC				
K. pneumoniae ATCC 13883		*	*	*
MIC				
P. aeruginosa ATCC 27853		*	*	*
MIC				
P. aeruginosa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

FIG 1Q

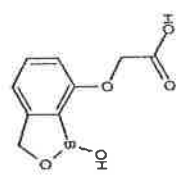
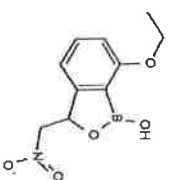
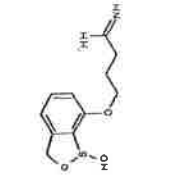
Reg #	Structure	A6	A51	A19
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3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

FIG 1R

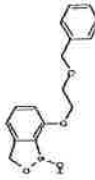
Reg #	Structure	A50	A28	A24
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC		*	*	*
P. aeruginosa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

FIG 1S

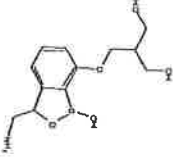
Reg #	Structure
A45	
***	E. cloacae ATCC 13047 MIC
**	E. coli ECM 1194 MIC
*	E. coli ECM 1888 MIC
*	E. faecalis ATCC 29212 MIC
***	H. influenzae ATCC 49766 MIC
***	K. pneumoniae ATCC 13883 MIC
**	P. aeruginosa ATCC 27853 MIC
***	P. aeruginosa PAO1 MIC
***	P. aeruginosa PAO1d3 MIC
**	S. aureus ATCC 29213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

FIG 1T

FIG 2A

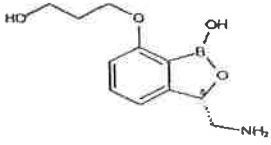
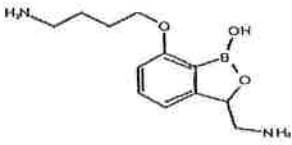
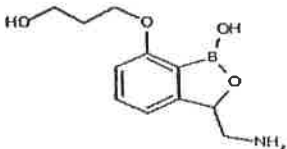
Compound ID	Structure	Escherichia_coll_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
	400< = # 400 - 100.1 = ## 100 - 5 ### 5> = #####			
A49		####	####	
A41		####	####	####
A46		####	####	####

FIG 2B

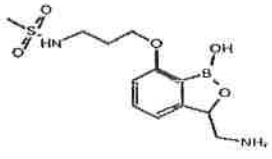
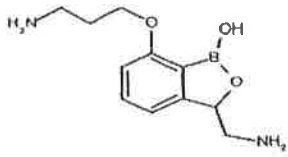
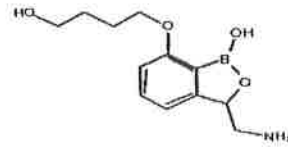
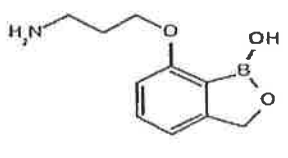
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A44		####	####	####
A36		####	####	####
A42		####	####	###
A11		####	####	###

FIG 2C

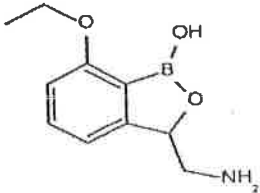
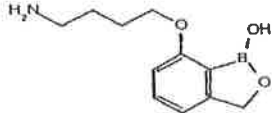
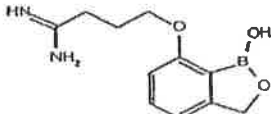
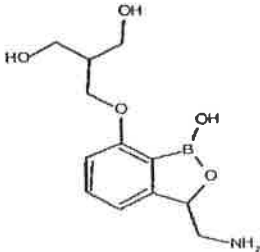
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A52		####	####	####
A17		####	####	###
A19		####	####	###
A45		####	####	####

FIG 2D

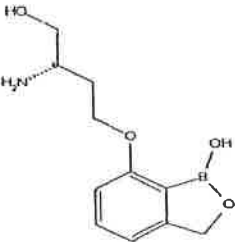
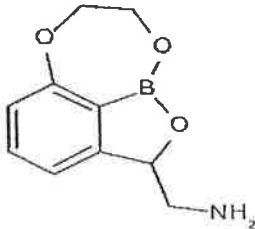
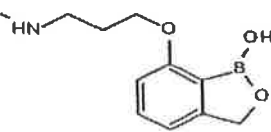
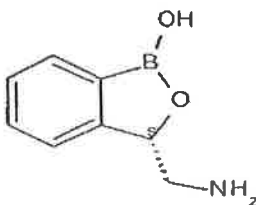
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1_LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC8325_LeuRS_wt IC50
A23		####	####	####
A39		####	####	####
A13		####	####	###
A2		####	####	####

FIG 2E

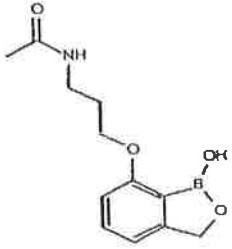
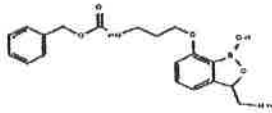
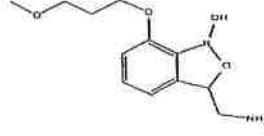
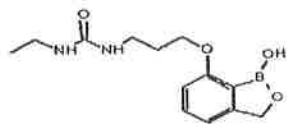
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A12		####	####	###
A43		####	####	####
A38		####	###	###
A14		####	####	

FIG 2F

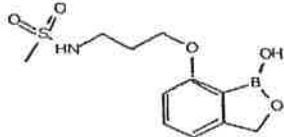
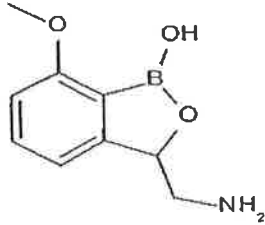
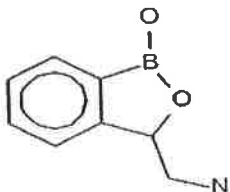
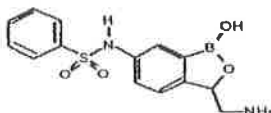
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A15		####	####	###
A53		####	####	####
A1		####	###	####
A33		####	###	###

FIG 2G

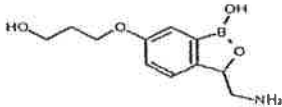
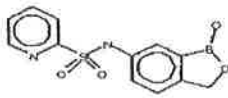
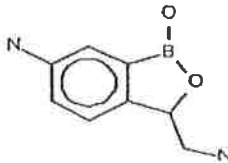
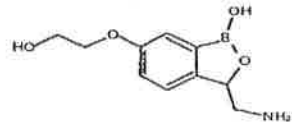
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A31		####	####	####
A3		####	####	###
A32		####	####	####
A30		####	###	####

FIG 2H

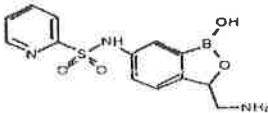
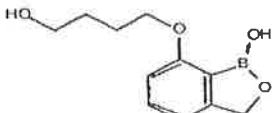
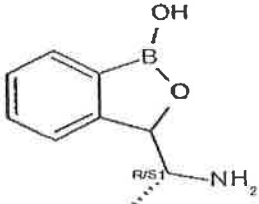
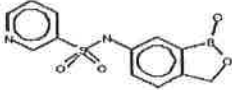
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A34		####	###	###
A22		####	####	#
A58		###	###	#
A4		###	###	###

FIG 2I

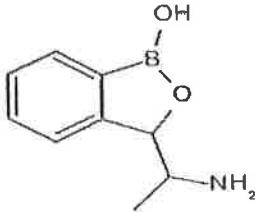
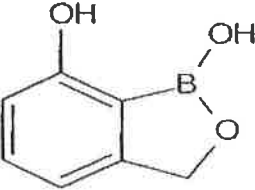
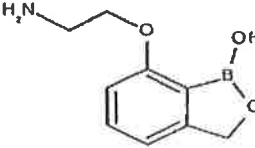
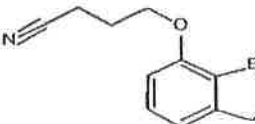
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1_LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC8325_LeuRS_wt IC50
A54		###	###	#
A26		###	#	###
A18		###	####	#
A16		###	####	#

FIG 2J

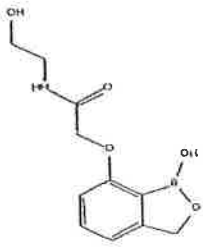
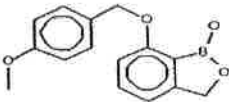
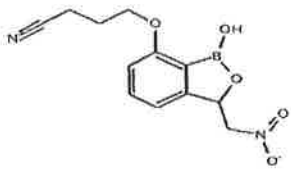
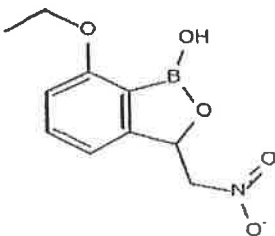
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A7		###	###	#
A25		###	###	#
A40		###	##	#
A51		###	##	#

FIG 2K

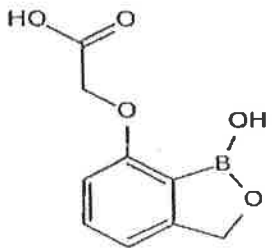
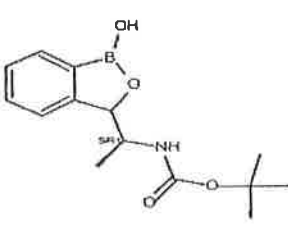
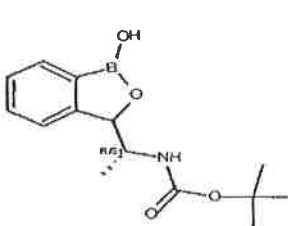
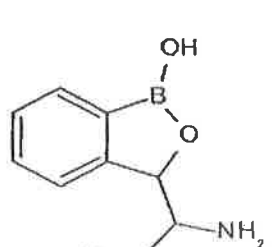
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A6		##	###	
A56		##	##	#
A55		##	#	#
A59		#	#	#

FIG 2L

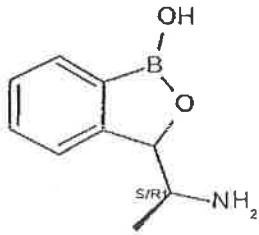
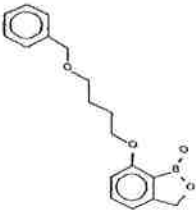
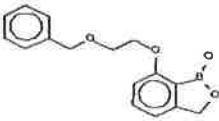
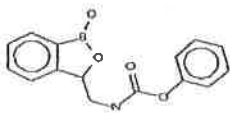
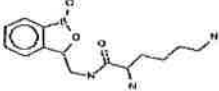
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A57		#	###	#
A21		#	#	#
A24		#	#	#
A29		#	#	#

FIG 2M

Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A27		#	#	#