

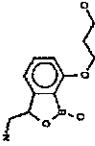
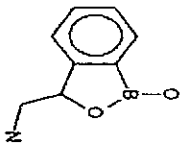
A46	A1	Reg #	Structure
		32 < = *	
		32 - 8.01 = **	
		8.00 - 2.00 = ***	
		2 > = ****	
****	****	E. cloacae ATCC 13047 MIC	
***	***	E. coli ECM 1194 MIC	
****	****	E. coli ECM 1888 MIC	
**	*	E. faecalis ATCC 29212 MIC	
****	***	H. influenzae ATCC 49766 MIC	
***	***	K. pneumoniae ATCC 13883 MIC	
**	***	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

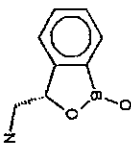
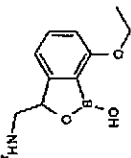
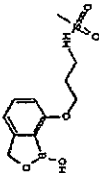
Reg #	Structure	
A2		
A52		
A15		
		E. cloacae ATCC 13047 MIC
		E. coli ECM 1194 MIC
		E. coli ECM 1888 MIC
		E. faecalis ATCC 29212 MIC
		H. influenzae ATCC 49766 MIC
		K. pneumoniae ATCC 13883 MIC
		P. aeruginosa ATCC 27853 MIC
		P. aeruginosa PAO1 MIC
		P. aeruginosa PAO1d3 MIC
		S. aureus ATCC 29213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

FIG 1B

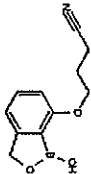
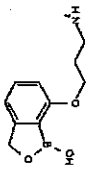
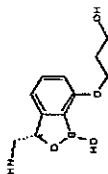
Reg #	Structure	
A16		**
A11		***
A49		****
	<i>E. cloacae</i> ATCC 13047 MIC	***
	<i>E. coli</i> ECM 1194 MIC	***
	<i>E. coli</i> ECM 1888 MIC	***
	<i>E. faecalis</i> ATCC 29212 MIC	*
	<i>H. influenzae</i> ATCC 49766 MIC	**
	<i>K. pneumoniae</i> ATCC 13883 MIC	**
	<i>P. aeruginosa</i> ATCC 27853 MIC	*
	<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 1C

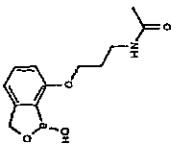
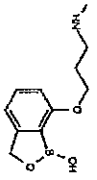
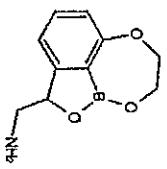
Reg #	Structure	A12	A13	A39
E. cloacae ATCC 13047 MIC				****
E. coli ECM 1194 MIC		**	***	***
E. coli ECM 1888 MIC		***	***	***
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		**	***	****
K. pneumoniae ATCC 13883 MIC		*	***	***
P. aeruginosa ATCC 27853 MIC		*	**	***
P. aeruginosa PAO1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

FIG 1D

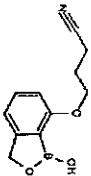
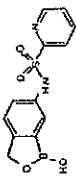
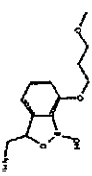
Reg #	Structure	
A16		
A3		
A38		
	E. cloacae ATCC 13047	***
	E. coli ECM 1194	***
	E. coli ECM 1888	***
	E. faecalis ATCC 29212	*
	H. influenzae ATCC 49766	***
	K. pneumoniae ATCC 13883	**
	P. aeruginosa ATCC 27853	*
	P. aeruginosa PAO1	***
	P. aeruginosa PAO1d3	***
	S. aureus ATCC 29213	*
	S. pneumoniae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

FIG 1E

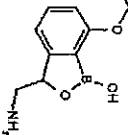
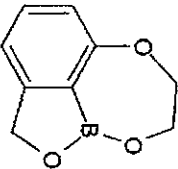
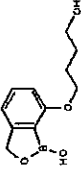
Reg #	Structure	
A52		
A61		
A22		
		E. cloacae ATCC 13047 MIC
		E. coli ECM 1194 MIC
		E. coli ECM 1888 MIC
		E. faecalis ATCC 29212 MIC
		H. influenzae ATCC 49766 MIC
		K. pneumoniae ATCC 13883 MIC
		P. aeruginosa ATCC 27853 MIC
		P. aeruginosa PAO1 MIC
		P. aeruginosa PAO1d3 MIC
		S. aureus ATCC 29213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

FIG 1F

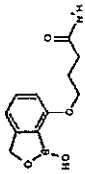
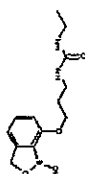
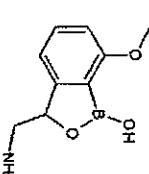
Reg #	Structure	
A10		
A14		
A53		
	E. cloacae ATCC 13047	****
	E. coli ECM 1194	***
	E. coli ECM 1888	***
	E. faecalis ATCC 29212	*
	H. influenzae ATCC 49766	***
	K. pneumoniae ATCC 13883	***
	P. aeruginosa ATCC 27853	***
	P. aeruginosa PAO1	*
	P. aeruginosa PAO1d3	**
	S. aureus ATCC 29213	*
	S. pneumoniae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

FIG 1G

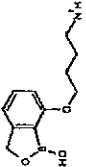
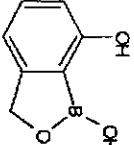
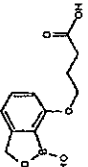
Reg #	Structure	
A17		
A26		
A9		
	E. cloacae ATCC 13047	
	MIC	***
	E. coli ECM 1194	
	MIC	***
	E. coli ECM 1888	
	MIC	**
	E. faecalis ATCC 29212	
	MIC	*
	H. influenzae ATCC 49766	
	MIC	***
	K. pneumoniae ATCC 13883	
	MIC	***
	P. aeruginosa ATCC 27853	
	MIC	**
	P. aeruginosa PAO1	
	MIC	*
	P. aeruginosa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

FIG. 1H

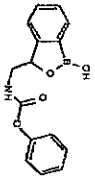
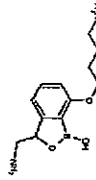
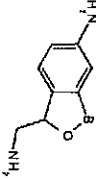
Reg #	Structure	
A29		
A41		
A32		
	E. cloacae ATCC 13047 MIC	***
	E. coli ECM 1194 MIC	**
	E. coli ECM 1888 MIC	**
	E. faecalis ATCC 29212 MIC	*
	H. influenzae ATCC 49766 MIC	***
	K. pneumoniae ATCC 13883 MIC	***
	P. aeruginosa ATCC 27853 MIC	***
	P. aeruginosa PAO1 MIC	
	P. aeruginosa PAO1d3 MIC	
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

FIG 1J

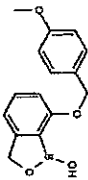
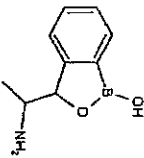
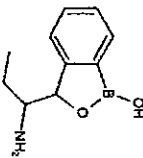
Reg #	Structure	A25	A54	A59
<i>E. cloacae</i> ATCC 13047 MIC				*
<i>E. coli</i> ECM 1194 MIC		*	*	*
<i>E. coli</i> ECM 1888 MIC		*	*	*
<i>E. faecalis</i> ATCC 29212 MIC		*	*	*
<i>H. influenzae</i> ATCC 49766 MIC		*	*	*
<i>K. pneumoniae</i> ATCC 13883 MIC		**	*	*
<i>P. aeruginosa</i> ATCC 27853 MIC		*	*	*
<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 IL

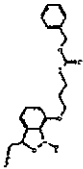
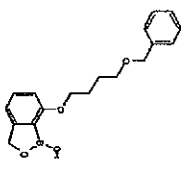
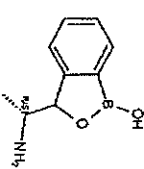
Reg #	Structure	A43	A21	A58
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		**	*	*
K. pneumoniae ATCC 13883 MIC		**	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa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

FIG 1M

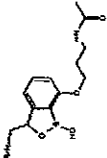
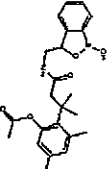
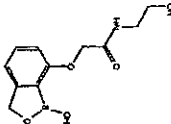
Reg #	Structure	A35	A27a	A7
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC		*	*	*
P. aeruginosa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

FIG IN

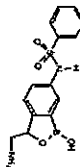
Reg #	Structure	A33	A34	A30
E. cloacae ATCC 13047 MIC		*		**
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*		*
H. influenzae ATCC 49766 MIC		**	**	***
K. pneumoniae ATCC 13883 MIC		*		*
P. aeruginosa ATCC 27853 MIC		*	**	**
P. aeruginosa PAO1 MIC				
P. aeruginosa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

FIG 10

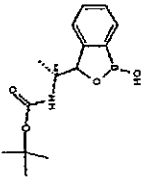
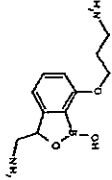
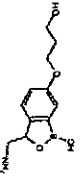
Reg #	Structure	
A55		
A36		
A31		
	E. cloacae ATCC 13047 MIC	**
	E. coli ECM 1194 MIC	*
	E. coli ECM 1888 MIC	*
	E. faecalis ATCC 29212 MIC	*
	H. influenzae ATCC 49766 MIC	***
	K. pneumoniae ATCC 13883 MIC	*
	P. aeruginosa ATCC 27853 MIC	*
	P. aeruginosa PAO1 MIC	
	P. aeruginosa PAO1d3 MIC	
	S. aureus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

FIG 1P

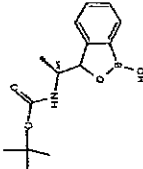
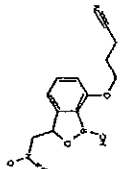
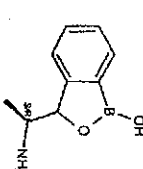
Reg #	Structure	A56	A40	A57
E. cloacae ATCC 13047 MIC				*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	**	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa PAO1d3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

FIG 1Q

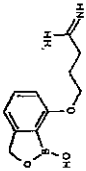
Reg #	Structure	A19	A51	A6
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		**	**	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC				
P. aeruginosa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

FIG 1R

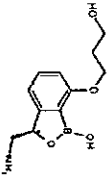
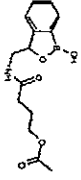
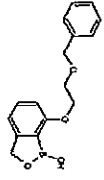
Reg #	Structure	A50	A28	A24
E. cloacae ATCC 13047 MIC		*	*	*
E. coli ECM 1194 MIC		*	*	*
E. coli ECM 1888 MIC		*	*	*
E. faecalis ATCC 29212 MIC		*	*	*
H. influenzae ATCC 49766 MIC		*	*	*
K. pneumoniae ATCC 13883 MIC		*	*	*
P. aeruginosa ATCC 27853 MIC		*	*	*
P. aeruginosa PAO1 MIC		*	*	*
P. aeruginosa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

FIG 1S

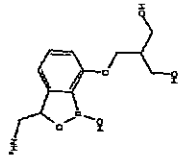
Reg #	Structure
A45	
***	E. cloacae ATCC 13047 MIC
**	E. coli ECM 1194 MIC
*	E. coli ECM 1888 MIC
*	E. faecalis ATCC 29212 MIC
***	H. influenzae ATCC 49766 MIC
***	K. pneumoniae ATCC 13883 MIC
**	P. aeruginosa ATCC 27853 MIC
***	P. aeruginosa PAO1 MIC
***	P. aeruginosa PAO1d3 MIC
**	S. aureus ATCC 29213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

FIG 11

FIG 2A

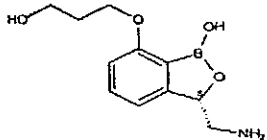
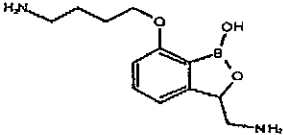
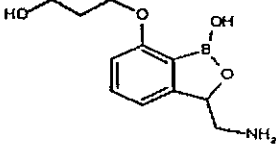
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
	400<=# 400-100.1= ##100-5### 5>=####			
A49		####	####	
A41		####	####	####
A46		####	####	####

FIG 2B

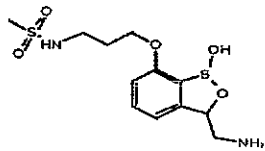
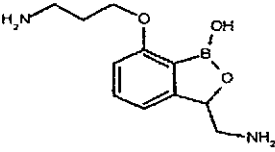
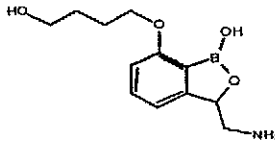
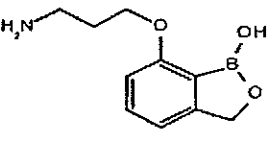
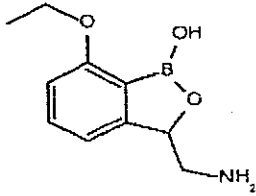
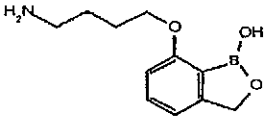
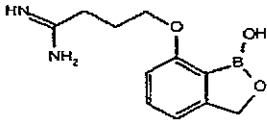
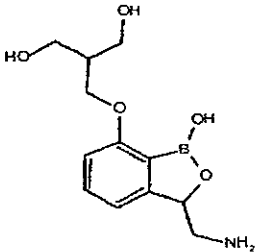
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A44		####	####	####
A36		####	####	####
A42		####	####	###
A11		####	####	###

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		####	####	####

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
A23		####	####	####
A39		####	####	####
A13		####	####	###
A2		####	####	####

FIG 2E

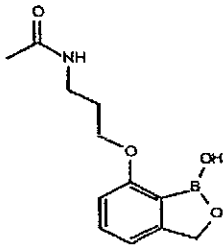
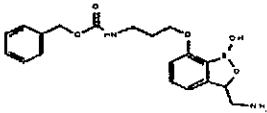
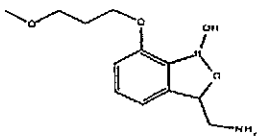
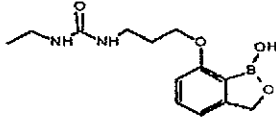
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A12		####	####	###
A43		####	####	####
A38		####	###	###
A14		####	####	

FIG 2F

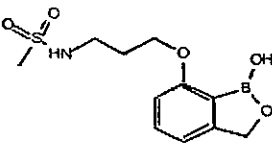
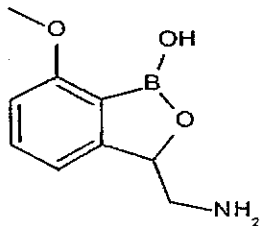
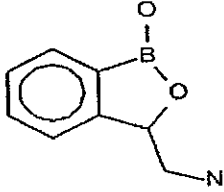
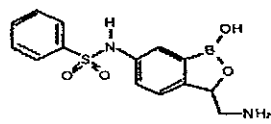
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A15		####	####	###
A53		####	####	####
A1		####	###	####
A33		####	###	###

FIG 2G

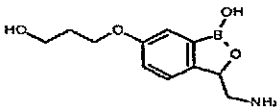
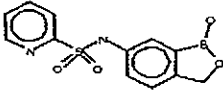
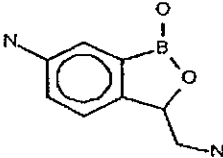
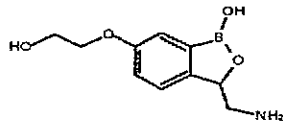
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A31		####	####	####
A3		####	####	###
A32		####	####	####
A30		####	###	####

FIG 2H

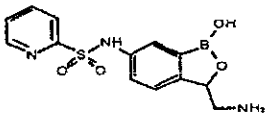
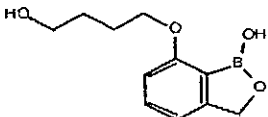
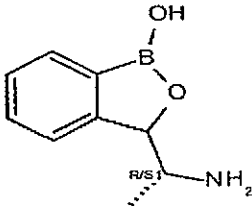
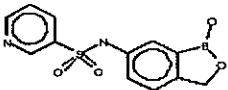
Compound ID	Structure	Escherichia_coli_K12_LeuRS_wt IC50	Pseudomonas_aeruginosa_PAO1 _LeuRS_wt IC50	Staphylococcus_aureus_NCTC83 25_LeuRS_wt IC50
A34		####	###	###
A22		####	####	#
A58		###	###	#
A4		###	###	###

FIG 2I

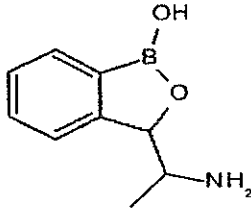
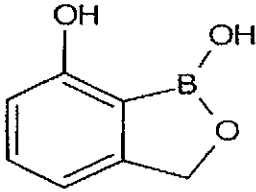
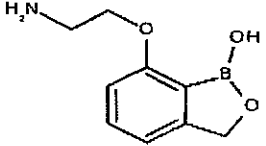
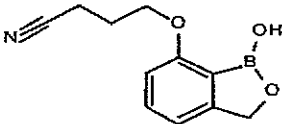
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas_aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus_aureus</i> _NCTC83 25_LeuRS_wt IC50
A54		###	###	#
A26		###	#	###
A18		###	####	#
A16		###	####	#

FIG 2J

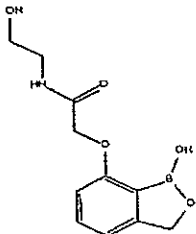
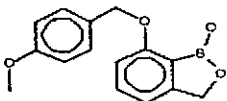
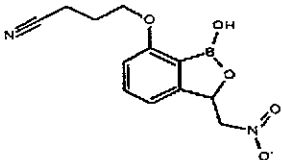
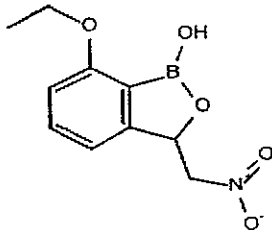
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A7		###	###	#
A25		###	###	#
A40		###	##	#
A51		###	##	#

FIG 2K

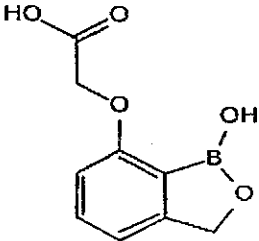
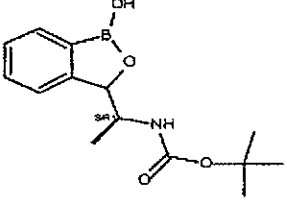
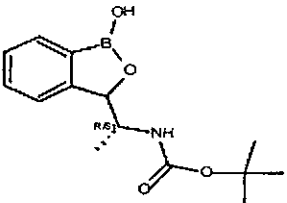
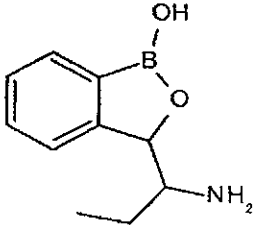
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A6		##	###	
A56		##	##	#
A55		##	#	#
A59		#	#	#

FIG 2L

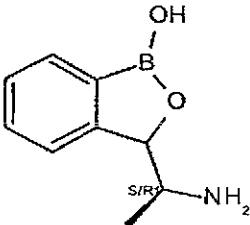
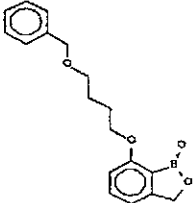
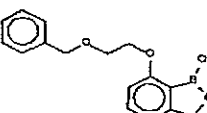
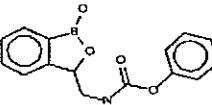
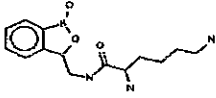
Compound ID	Structure	<i>Escherichia coli</i> _K12_LeuRS_wt IC50	<i>Pseudomonas aeruginosa</i> _PAO1 _LeuRS_wt IC50	<i>Staphylococcus aureus</i> _NCTC83 25_LeuRS_wt IC50
A57		#	###	#
A21		#	#	#
A24		#	#	#
A29		#	#	#

FIG 2M

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