

## Synthesis, nanostructuring and *in silico* studies of a new imine bond containing macroheterocycle as a promising PBP-2a non- $\beta$ -lactam inhibitor

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## Supplementary Materials

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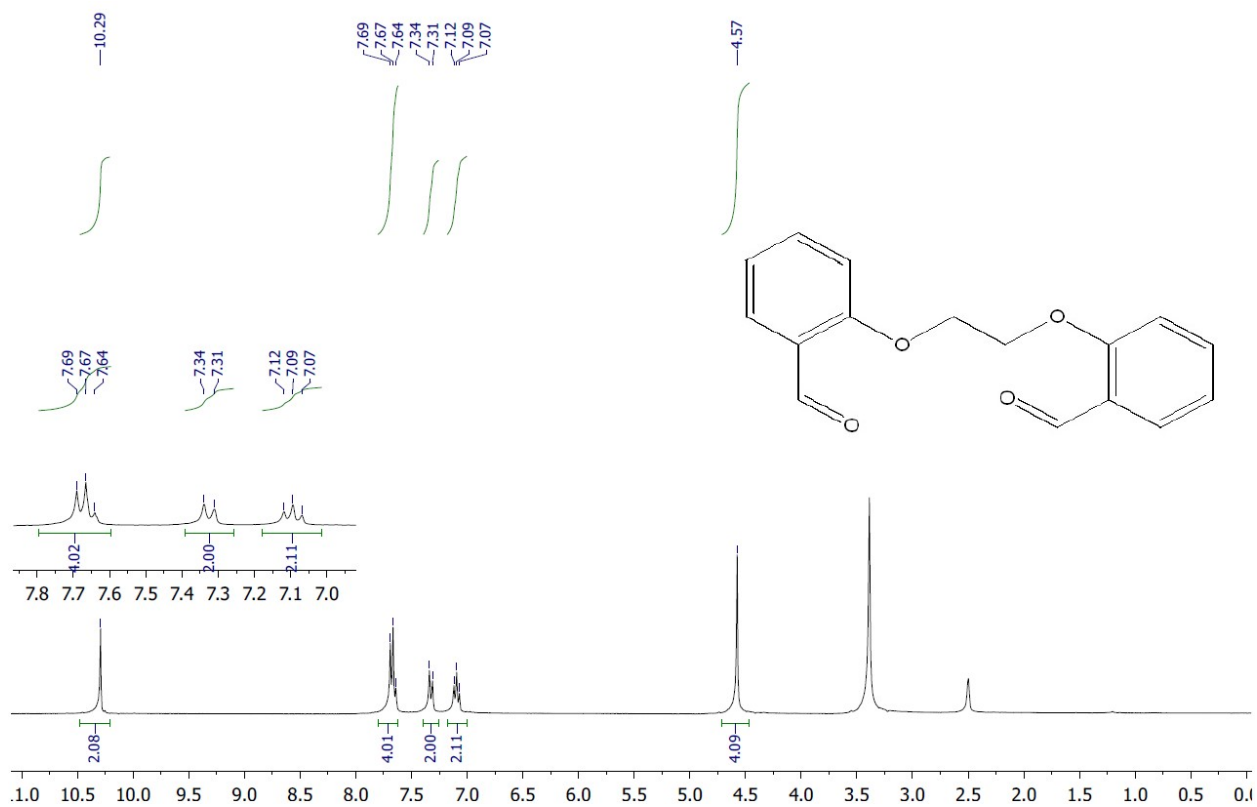


Figure 1S.  $^1\text{H}$  NMR spectrum of synthesized dialdehyde **1**

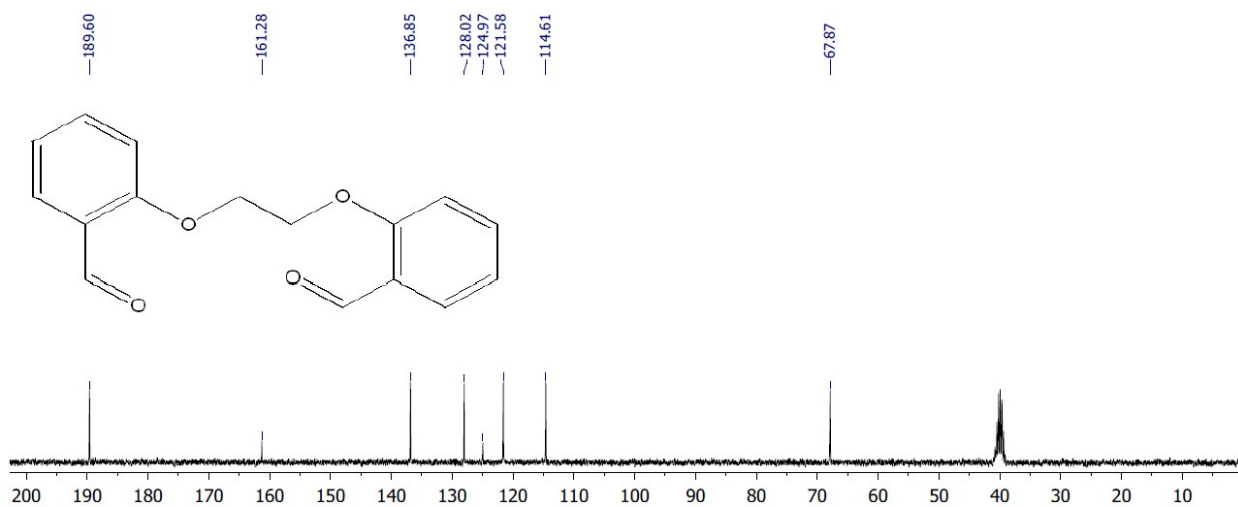
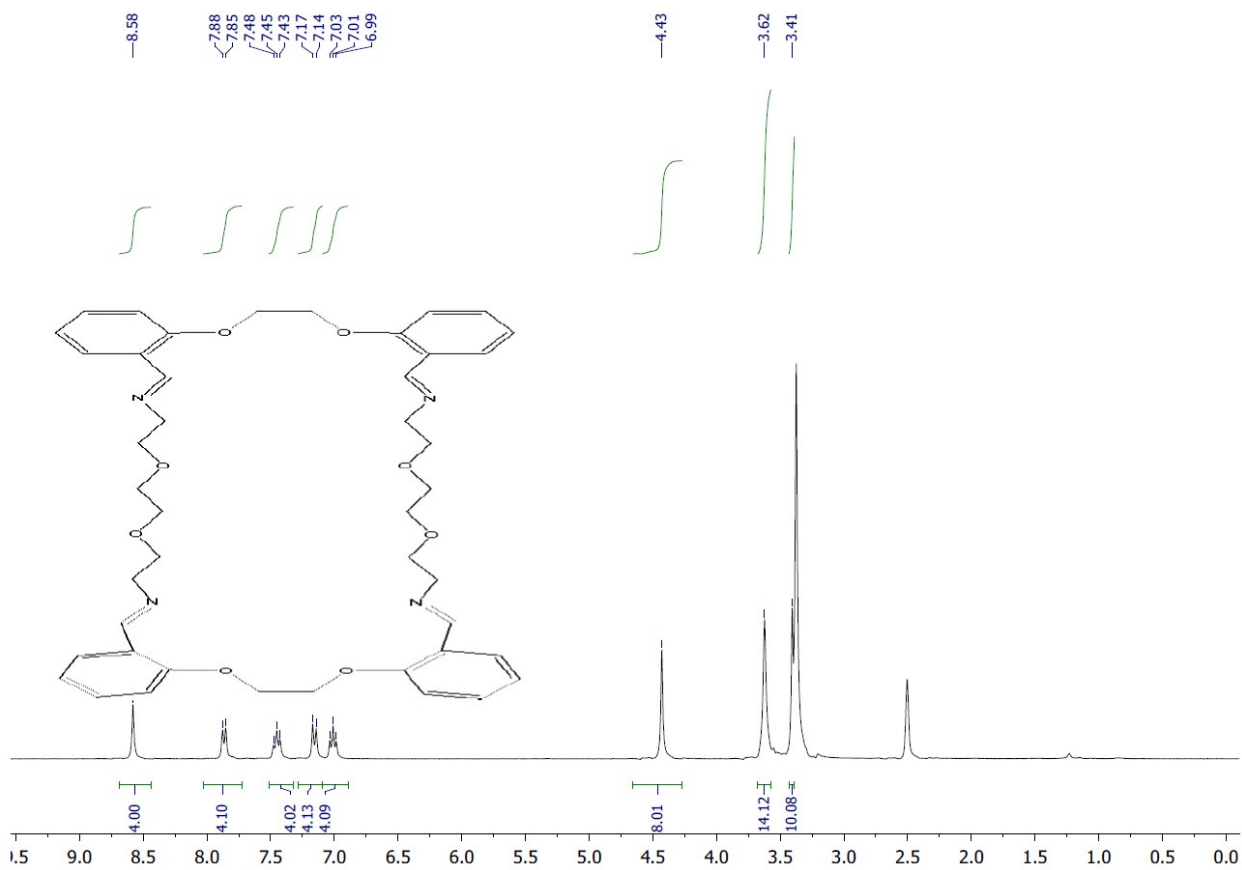
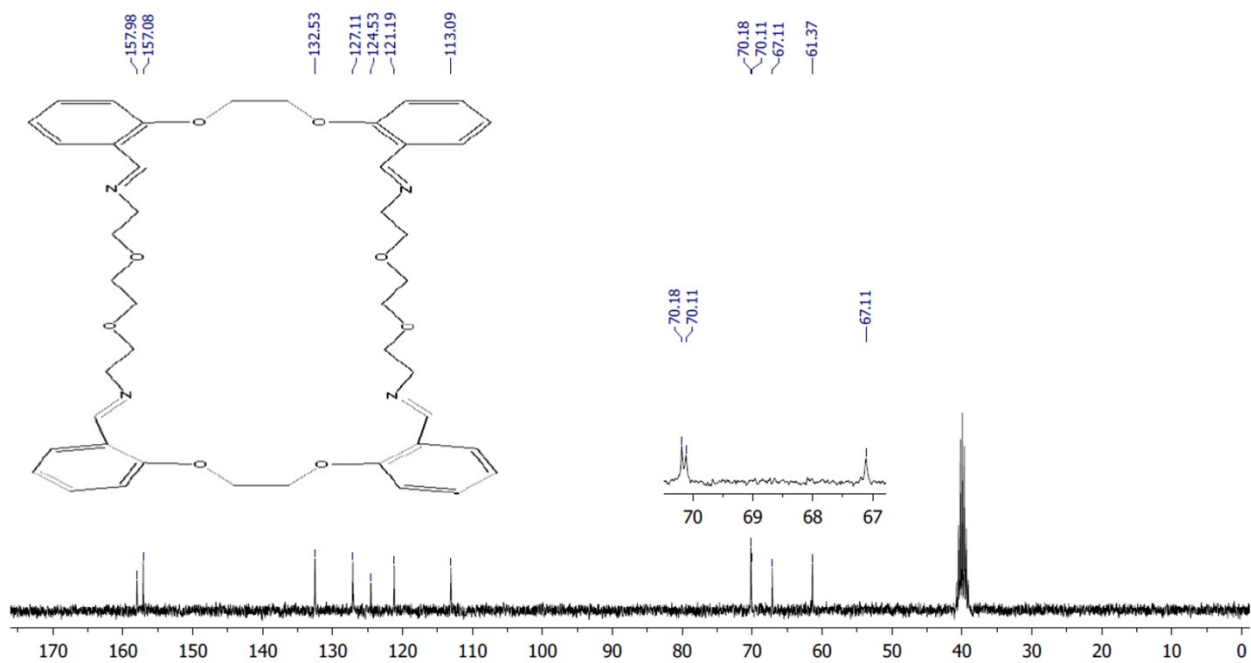


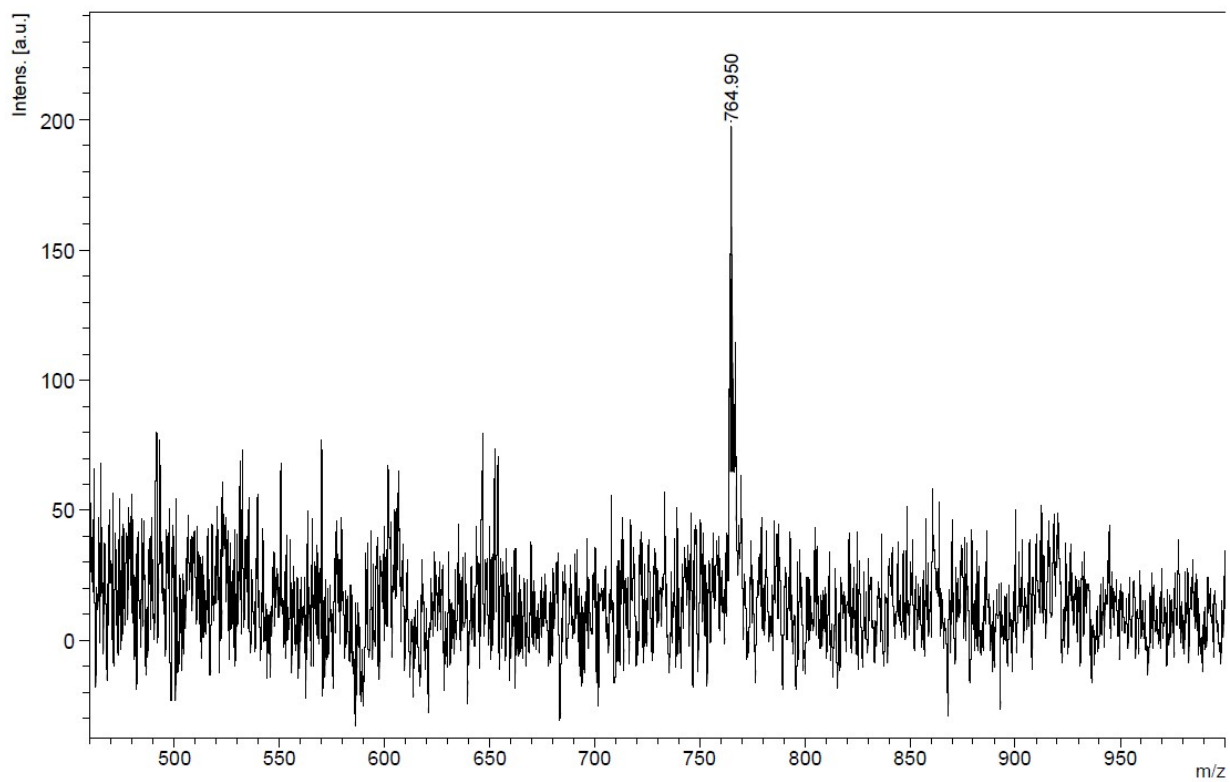
Figure 2S.  $^{13}\text{C}$  NMR spectrum of synthesized dialdehyde **1**



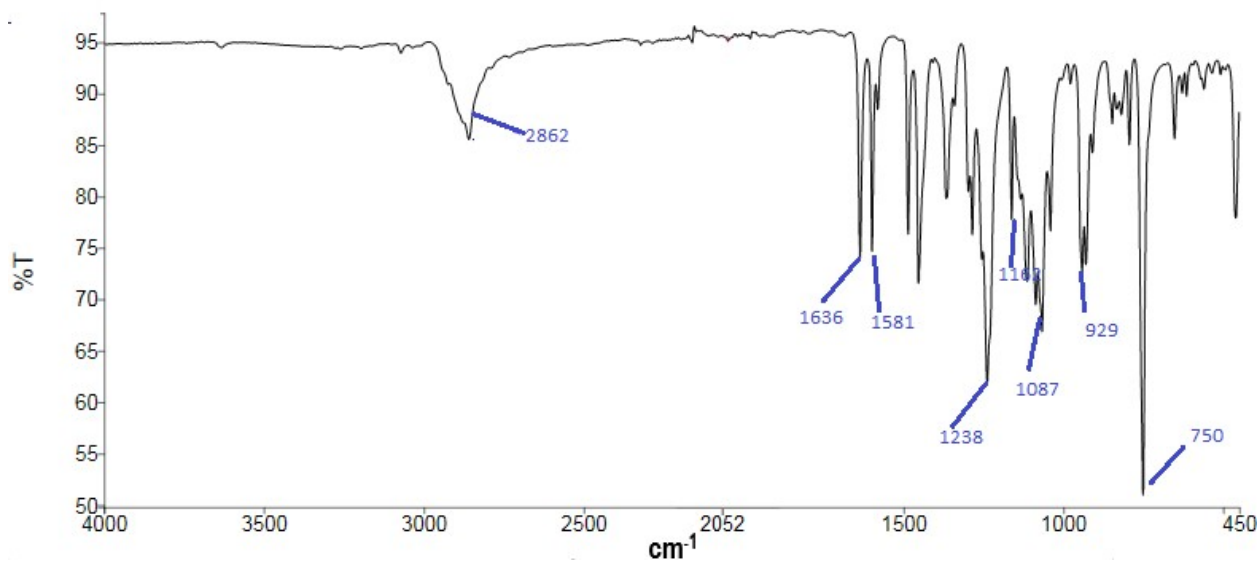
**Figure 3S.**  $^1\text{H}$  NMR spectrum of synthesized macrocycle **3**



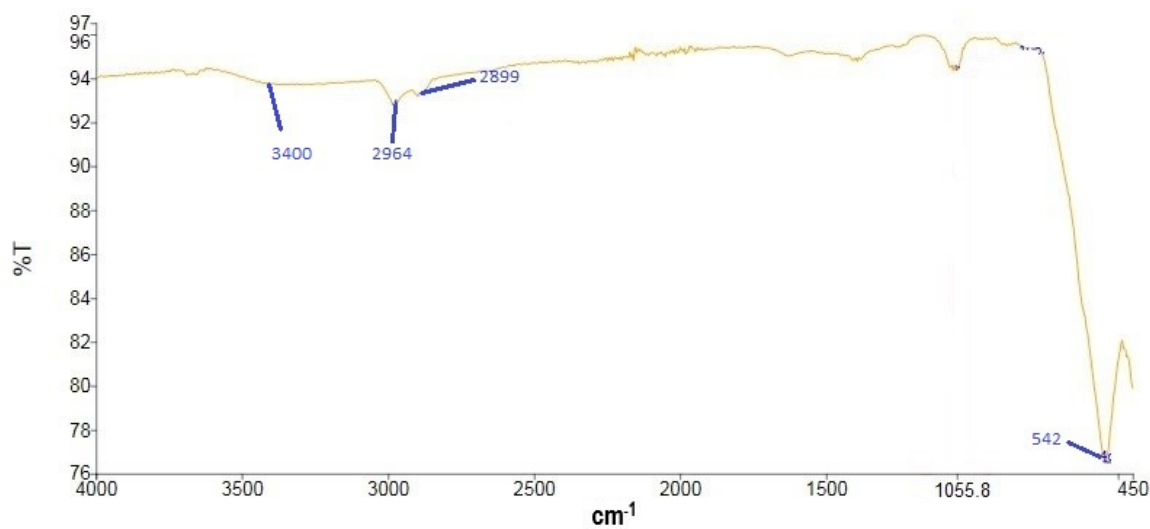
**Figure 4S.**  $^{13}\text{C}$  NMR spectrum of synthesized macrocycle **3**



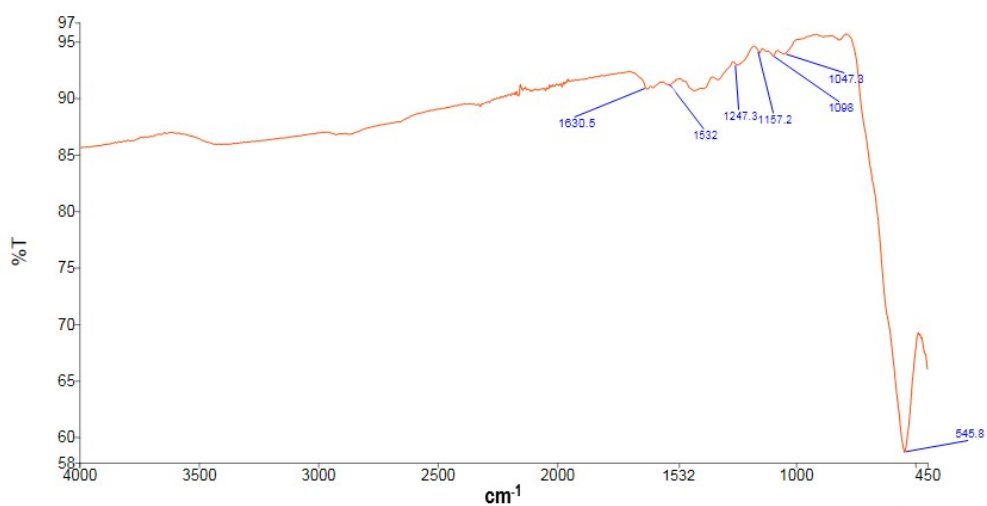
**Figure 5S.** MALDI-MS spectrum of synthesized macrocycle **3**



**Figure 6S.** FTIR spectrum of the MHC3



**Figure 7S.** FTIR spectrum of the magnetite nanoparticles



**Figure 8S.** FTIR spectrum of the MHC3@Fe<sub>3</sub>O<sub>4</sub>

**Table 1S.** Powder XRD qualitative results and peak list of the MHC3@Fe<sub>3</sub>O<sub>4</sub>

Phase name	Formula	Figure of merit	Phase reg. detail	DB card number
Magnetite	Fe <sub>3</sub> O <sub>4</sub>	0.789	ICDD (PDF-2/Release 2011 RDB)	00-001-1111
Phase name	Formula	Space group	Phase reg. detail	DB card number
Magnetite	Fe <sub>3</sub> O <sub>4</sub>	227 : Fd-3m,choice-2	ICDD (PDF-2/Release 2011 RDB)	00-001-1111

No.	2-theta(deg)	Interplanar spacing ( <i>d</i> spacing) (Å°)	Height(cps)	FWHM(deg)	Int. I(cps deg)	Int. W(deg)	Asym. factor	Miller indices (hkl)
1	12.6(3)	7.04(15)	246(20)	2.6(2)	691(97)	2.8(6)	0.9(4)	(111)
2	30.13(4)	2.964(4)	1221(45)	0.68(5)	1208(64)	0.99(9)	0.9(3)	(220)
3	35.536(13)	2.5242(9)	3873(80)	0.633(12)	3094(57)	0.80(3)	1.18(1)	(311)
4	43.17(6)	2.094(3)	949(40)	0.74(8)	1164(59)	1.23(11)	1.1(5)	(400)
5	57.07(4)	1.6126(11)	1193(45)	0.72(4)	911(63)	0.76(8)	0.9(2)	(333) (511)
6	62.75(3)	1.4796(7)	1738(54)	0.79(3)	1676(62)	0.96(7)	1.1(2)	(440)

**Table 2S.** Crystallite size of the MHC3@Fe<sub>3</sub>O<sub>4</sub>

Data set name	Crystallite size(A)	Phase name	Crystallite size(A)
MHC3@Fe <sub>3</sub> O <sub>4</sub>	91(12)	Magnetite	91(12)

All the MIC experiments were done in triplicate (Table 1). The box and whisker plot does not demonstrate any variations in the results, because the MIC test done by the application of resazurin disodium salt is considered a precise method, which if performed appropriately and for individual compounds provides identical values for each compound in triplicate (Figure 1).

Table 3S. Results of triplicate 96-well microtiter assay of tested compounds on bacterial strains

Bacterial strains	MIC, $\mu\text{g/mL}$							Ampicillin	Ampicillin
	MHC3	MHC3	MHC3	MHC3	MHC3	MHC3	$\text{Fe}_3\text{O}_4$ NPs		
				@ $\text{Fe}_3\text{O}_4$	@ $\text{Fe}_3\text{O}_4$	@ $\text{Fe}_3\text{O}_4$			
<i>K. pneumoniae</i>	512	512	512	>1024	>1024	>1024	-	512	512
<i>P. aeruginosa</i>	256	256	256	1024	1024	1024	-	512	512
<i>S. aureus</i>	32	32	32	0.5	0.5	0.5	-	16	16

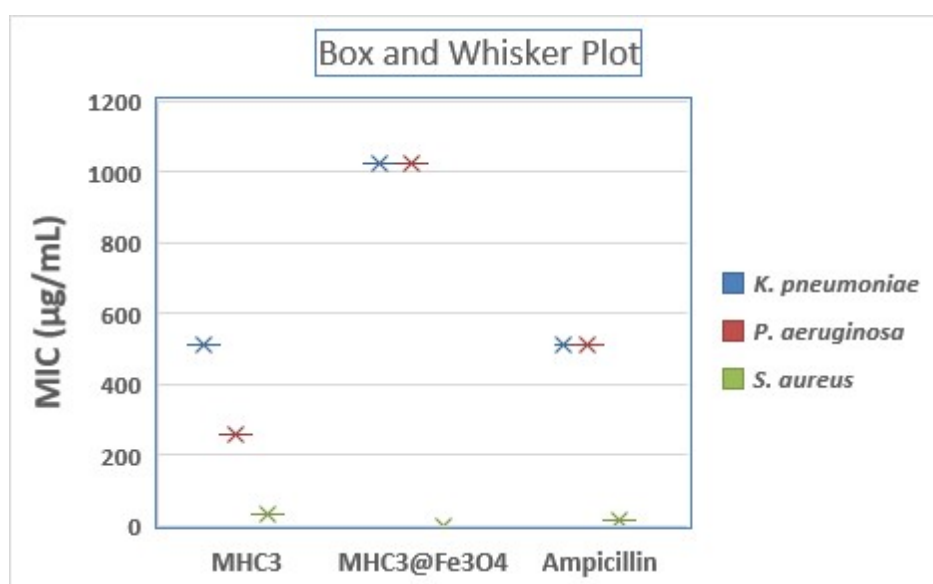


Figure 9S. The Box and Whisker plot for the results of the 96-well microtiter assay

The two-way and post hoc ANOVA were applied to the obtained results by using GraphPad Prism (version 9 for Windows, GraphPad Software, San Diego, CA, USA, www.graphpad.com). Post hoc analysis by two-way ANOVA was applied as multiple comparisons in three families (Figure 2, Table 2). The antibacterial activity of tested compounds on three bacterial strains was compared with each other and ampicillin as a control. The adjusted P value is less than 5%, so the obtained results can be considered significantly different. The only not significant difference has been observed for ampicillin and MHC3 activity towards *Klebsiella pneumoniae* because values of MIC for both compounds were equal to 512  $\mu\text{g/mL}$ .

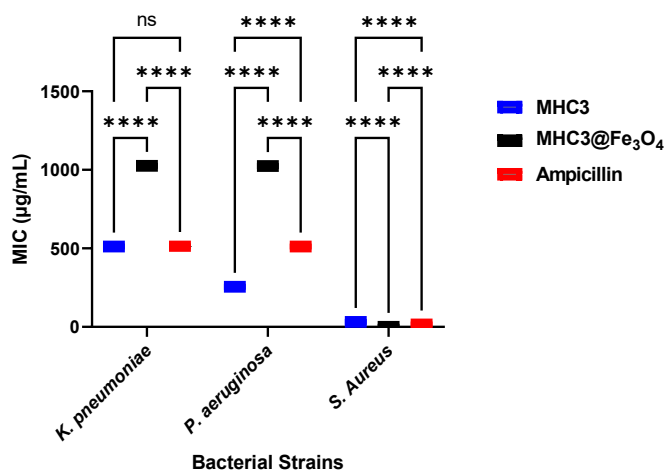


Figure 10S. ANOVA: Tukey's multiple comparisons results

Table 4S. ANOVA: Tukey's multiple comparisons results

Number of comparisons per family	3							
Alpha	0.05							
Tukey's multiple comparisons test	Mean Diff.	95.00% CI of diff.	Below threshold?	Summary	Adjusted P Value			
<b><i>K. pneumoniae</i></b>								
MHC3 vs. MHC3@Fe <sub>3</sub> O <sub>4</sub>	-513.0	-514.2 to -511.8	Yes	****	<0.0001			
MHC3 vs. Ampicillin	-1.000	-2.203 to 0.2031	No	ns	0.1135			
MHC3@Fe <sub>3</sub> O <sub>4</sub> vs. Ampicillin	512.0	510.8 to 513.2	Yes	****	<0.0001			
<b><i>P. aeruginosa</i></b>								
MHC3 vs. MHC3@Fe <sub>3</sub> O <sub>4</sub>	-768.0	-769.2 to -766.8	Yes	****	<0.0001			
MHC3 vs. Ampicillin	-256.0	-257.2 to -254.8	Yes	****	<0.0001			
MHC3@Fe <sub>3</sub> O <sub>4</sub> vs. Ampicillin	512.0	510.8 to 513.2	Yes	****	<0.0001			
<b><i>S. Aureus</i></b>								
MHC3 vs. MHC3@Fe <sub>3</sub> O <sub>4</sub>	31.50	30.30 to 32.70	Yes	****	<0.0001			
MHC3 vs. Ampicillin	16.00	14.80 to 17.20	Yes	****	<0.0001			
MHC3@Fe <sub>3</sub> O <sub>4</sub> vs. Ampicillin	-15.50	-16.70 to -14.30	Yes	****	<0.0001			
<b>Test details</b>								
	Mean 1	Mean 2	Mean Diff.	SE of diff.	N1	N2	q	DF
<b><i>K. pneumoniae</i></b>								
MHC3 vs. MHC3@Fe <sub>3</sub> O <sub>4</sub>	512.0	1025	-513.0	0.4714	3	3	153	18.



							9	00
MHC3 vs. Ampicillin	512.0	513.0	-1.000	0.4714	3	3	3.0 00	18. 00
MHC3@Fe <sub>3</sub> O <sub>4</sub> vs. Ampicillin	1025	513.0	512.0	0.4714	3	3	153 6	18. 00
<b><i>P. aeruginosa</i></b>								
MHC3 vs. MHC3@Fe <sub>3</sub> O <sub>4</sub>	256.0	1024	-768.0	0.4714	3	3	230 4	18. 00
MHC3 vs. Ampicillin	256.0	512.0	-256.0	0.4714	3	3	768 .0	18. 00
MHC3@Fe <sub>3</sub> O <sub>4</sub> vs. Ampicillin	1024	512.0	512.0	0.4714	3	3	153 6	18. 00
<b><i>S. Aureus</i></b>								
MHC3 vs. MHC3@Fe <sub>3</sub> O <sub>4</sub>	32.00	0.5000	31.50	0.4714	3	3	94. 50	18. 00
MHC3 vs. Ampicillin	32.00	16.00	16.00	0.4714	3	3	48. 00	18. 00
MHC3@Fe <sub>3</sub> O <sub>4</sub> vs. Ampicillin	0.5000	16.00	-15.50	0.4714	3	3	46. 50	18. 00