

A novel method for visual microarray detection of Antibiotic resistance genes

Shonglong Ma^{a, b}, Baowei Zhao^{a, *}, Rui Zhang^{b, *}, Hui Liu^a, Yunxia Li^b

¹ *School of Environmental and Municipal Engineering , Lanzhou Jiaotong University , Lanzhou 730070 , China;* ² *Key Laboratory of Sensor and Sensing Technology of Gansu Province , Institute of Sensor Technology , Gansu Academy of Sciences , Lanzhou 730000 , China*

Keywords: Microplastics; Antibiotic; Antibiotic Resistance Genes; Biofilm

Corresponding authors

Baowei Zhao, School of Environmental and Municipal Engineering, Lanzhou Jiaotong University, Lanzhou 730000, China

E-mail address: baoweizhao@mail.lzjtu.cn

Rui Zhang, Institute of Sensor Technology, Gansu Academy of Sciences, Lanzhou 730000, China

E-mail address: ruizhanglzu@sina.com

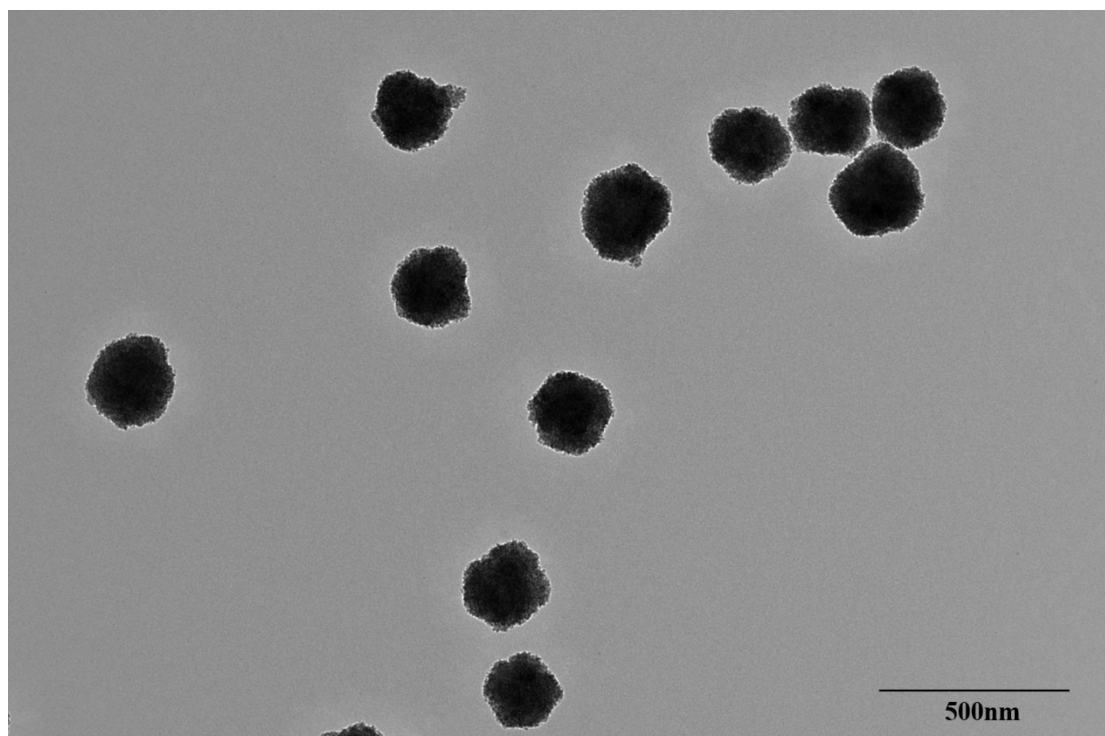


Fig. S1 TEM characterization of magnetic bead

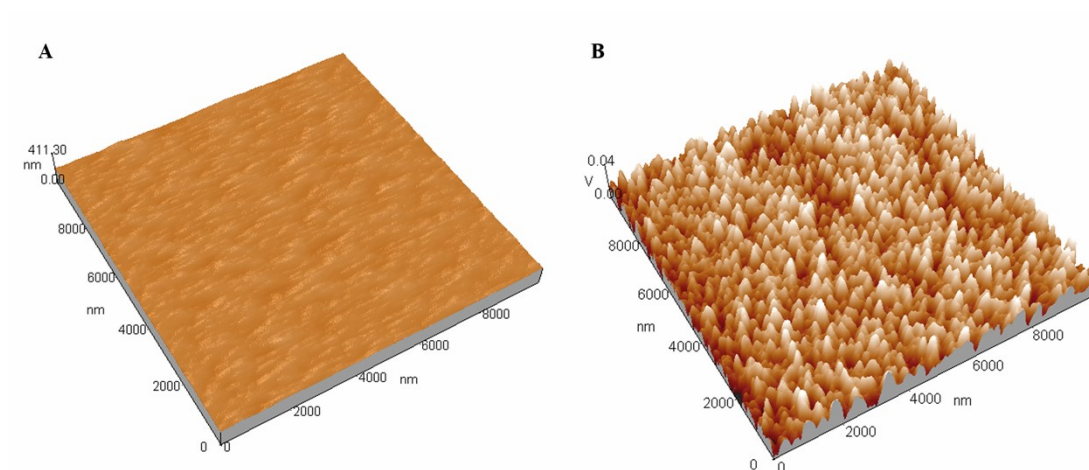


Fig. S2 Production of AFM images of aminated slides, represents untreated slides (A), represents aminated slides (B).

Table.S1 PCR Primers

	Forward Primer (5'-3')	Reverse Primer (5'-3')
<i>SulI</i>	TCCTTCTTCGATGAGAGCCG	TGCTGTCGATTGAAACACGGT
<i>macB</i>	TCAGCGAGCTGCATTACATCGA	GGGTGTTGTGGTCAATCACCA
<i>QepA</i>	TTCCTGCCCCGAGTATCGTGAT	TAGGCGATGTGGCCCTGG
<i>VanR</i>	ACAAGCGGCCTTACTATCTGTC	GATAACTGCTTCTCGTTCAGAT
<i>TetA</i>	ACTCATGCTCGGAATGATTG	CGTTGTTATAGAAGCCGCATAG

Table.S2 Target and probe sequences (Se is the sense chain target sequence, while Ase is the antisense chain target sequence.)

	Name	Sequence (5'-3')
<i>QepA</i>	Q-Se (target sequence)	CGAGTATCGTGATCCGGACGCGGGGCACCTGGACCTG.....CTC GATGGCTGCGCTGCTGGCCGGGCTGGCG
	Q-ASe	CGCCAGCCCGGCCAGCAGCGCAGCCATCGAG.....CAGGTCCAG GTGCCCCGCGTCCGGATCACGATACTCG
	Q-Se-Target probe 1	COOH- TTTCAGGTCCAGGTGCCCCGCGTCCGGATCACGATACTCG
	Q-Ase-Target probe 2	COOH-TTTTTTCTCGATGGCTGCGCTGCTGGCCGGGCTGGCG
	Q-Se-Detection probe 1	SH-TTTTTTCGCCAGCCCGGCCAGCAGCGCAGCCATCGAG
	Q-Ase-Detection probe 2	SH- TTTTTCGAGTATCGTGATCCGGACGCGGGGCACCTGGACCTG
<i>macB</i>	m-Se	CTGCATTACATCGACAGCGCCACGCCAACG.....CCAGGACGA TGACACGCGCCAGGCCCAAGTG
	m-ASe	CACTTGGGCCTGGCGCGTGTTCATCGTCCTGG.....CGTTGGGCGT GGCGCTGTCGATGTAATGCAG
	m-Se-Target probe 1	COOH-TTTTTTCGTTGGGCGTGGCGCTGTCGATGTAATGCAG
	m-Ase-Target probe 2	COOH-TTTTTTCCAGGACGATGACACGCGCCAGGCCCAAGTG
	m-se-Detection probe 1	SH-TTTTTTCACTTGGGCCTGGCGCGTGTTCATCGTCCTGG
	m-Ase-Detection probe 2	SH-TTTTTTCTGCATTACATCGACAGCGCCACGCCAACG
<i>vanR</i>	v-Se	AGGGACAAGCACACCTATCCGATTATCATG.....TAATTGCTCGG GTAAAGGCCCAGTTGCGCC
	v-ASe	GGCGCAACTGGGCCTTTACCCGAGCAATTA.....CATGATAATCG GATAGGTGTGCTTGTCCCT
	v-Se-Target probe 1	COOH-TTTTTTCATGATAATCGGATAGGTGTGCTTGTCCCT
	v-Ase-Target probe 2	COOH-TTTTTTTAATTGCTCGGGTAAAGGCCCAGTTGCGCC

	v-se-Detection probe 1	SH-TTTTTTGGCGCAACTGGGCCTTTACCCGAGCAATTA
	v-Ase-Detection probe 2	SH-TTTTTTAGGGACAAGCACACCTATCCGATTATCATG
<i>tetA</i>	t-Se	TTGCCTTCGCGACACGGGGATGGATGGCGTTCCC.....AGGCAG GTGGATGAGGAACGTCAGGGGCAGC
	t-ASe	GCTGCCCCTGACGTTTCCTCATCCACCTGCCT.....GGGAACGCCA TCCATCCCCGTGTCGCGAAGGCAA
	t-Se-Target probe 1	COOH-TTTTTGGGAACGCCATCCATCCCCGTGTCGCGAAGGCAA
	t-Ase-Target probe 2	COOH-TTTTTTAGGCAGGTGGATGAGGAACGTCAGGGGCAGC
	t-Se-Detection probe 1	SH-TTTTTTGCTGCCCCTGACGTTTCCTCATCCACCTGCCT
	t-Ase-Detection probe 2	SH-TTTTTTTTGCCTTCGCGACACGGGGATGGATGGCGTTCCC
<i>sulI</i>	s-Se	CCGGCGCTGTCACCGCGGCGATCGAAATGCT.....CCGCTCTTAG ACGCCCTGTCCGATCAGATGC
	s-ASe	GCATCTGATCGGACAGGGCGTCTAAGAGCGG.....AGCATTTTCG ATCGCCGCGGTGACAGCGCCGG
	s-Se-Target probe 1	COOH-TTTTTTAGCATTTTCGATCGCCGCGGTGACAGCGCCGG
	s-Ase-Target probe 2	COOH-TTTTTTCCGCTCTTAGACGCCCTGTCCGATCAGATGC
	s-Se-Detection probe 1	SH-TTTTTTGATCTGATCGGACAGGGCGTCTAAGAGCGG
	s-Ase-Detection probe 2	SH-TTTTTTCCGGCGCTGTCACCGCGGCGATCGAAATGCT