

SUPPLEMENTARY INFORMATION

Table S1. Primers used in this study for screening of antimicrobial resistance genetic markers (ARG).

Target	Primer sequence (5' – 3')	Amplicon (bp)	Reference
<i>bla</i> _{CTX-M}	ATGTGCAGYACCAGTAAAG GGTCACCAGAAGGAGC	562	<i>Antimicrob Agents Chemother</i> 2009; 53 : 465-475
<i>bla</i> _{KPC}	ATGTCACTGTATGCCGTCT TTTCAGAGCCTTACTGCC	892	<i>Antimicrob Agents Chemother</i> 2009; 53 : 465-475
<i>bla</i> _{SHV}	CTTTACTCGCCTTATCGGC TTACCGACCGGCATCTTCC	982	<i>Antimicrob Agents Chemother</i> 2009; 53 : 465-475
<i>bla</i> _{TEM}	GTGCGCGGAACCCATT TTACCAATGCTTAATCAGTGAGGC	968	<i>Antimicrob Agents Chemother</i> 2009; 53 : 465-475
<i>bla</i> _{OXA-23}	GATCGGATTGGAGAACCGAGA ATTCTGACCGCATTTCCAT	501	<i>Int J Antimicrob Agents</i> 2006; 27 : 351-353
<i>bla</i> _{OXA-51}	TAATGCTTGATCGGCCTTG TGGATTGCACITCATCTTGG	353	<i>Int J Antimicrob Agents</i> 2006; 27 : 351-353
<i>bla</i> _{OXA-24}	GGTTAGTTGGCCCCCTTAAA AGTTGAGCGAAAAGGGGATT	246	<i>Int J Antimicrob Agents</i> 2006; 27 : 351-353
<i>bla</i> _{OXA-143}	TGGCACTTCAGCAGTTCC TAATCTTGAGGGGGCCAACC	149	<i>Int J Antimicrob Agents</i> 2010; 35 : 305
<i>bla</i> _{OXA-58}	AAGTATTGGGGCTTGTGCTG CCCCTCTGCGCTCTACATAC	599	<i>Int J Antimicrob Agents</i> 2006; 27 : 351-353
<i>bla</i> _{OXA-10}	TCTTTGAGTACGGCATTAGC CCAATGATGCCCTCACTTCC	760	<i>J Antimicrob Chemother</i> 2000; 46 : 703-711
<i>bla</i> _{OXA-2}	GCCAAAGGCACCGATAGTTG GCGTCGGAGTTGACTGCCGG	701	<i>Antimicrob Agents Chemother</i> 2002; 46 : 3031-3034
<i>bla</i> _Z	ACTTCAACACCTGCTGCTTTC TGACCACTTTATCAGCAACC	173	<i>Antimicrob Agents Chemother</i> 2000; 44 : 231-238
<i>cfxA/cfxA2</i>	CGTAGTTTGAGTATACTT GATGTTGCCCTATATATGTC	802	<i>J Antimicrob Chemother</i> 2003; 51 : 1293-1296
<i>ampC</i>	ATAACCACCCAGTCACGC CAGTAGCGAGACTGCGCA	630	<i>Clin Infect Dis</i> 2002; 35 : 140-145
<i>blasPM-1</i>	CCTACAATCTAACGGCGACC TCGCCGTGTCCAGGTATAAC	649	<i>J Antimicrob Chemother</i> 2003; 52 : 699-702
<i>cfa</i>	TCCATGCTTTCCCTGTCGAGTTAT GGGCTATGGCTTGAAGTGC	683	<i>J Med Microbiol</i> 2004; 53 : 413-419
<i>tet(A)</i>	GCTACATCCTGCTGCCTTC CATAGATGCCGTGAAGAGG	210	<i>Mol Cell Probes</i> 2001; 15 : 209-215
<i>tet(B)</i>	TTGGTTAGGGCCAAGTTTG GTAATGGGCCATAAACACCG	659	<i>Mol Cell Probes</i> 2001; 15 : 209-215
<i>tet(E)</i>	AAACCACATCCTCCATACCG AAA TAG GCC ACA ACC GTC AG	278	<i>Mol Cell Probes</i> 2001; 15 : 209-215
<i>tet(K)</i>	GTA GCG ACA ATA GGT AAT AGT GTA GTG ACA ATA AAC CTC CTA	360	<i>J Clin Microbiol</i> 2003; 41 : 4089-4094
<i>tet(L)</i>	TCG TTA GCG TGC TGT CAT TC GTA TCC CAC CAA TGT AGC CG	267	<i>Mol Cell Probes</i> 2001; 15 : 209-215
<i>tet(M)</i>	AGT GGA GCG ATT ACA GAA CAT ATG TCC TGG GGT GTC TA	158	<i>J Clin Microbiol</i> 2003; 41 : 4089-4094
<i>tet(O)</i>	AGC GTC AAA GGG GAA TCA CTA TCC CGG CGG GGT TGG CAA ATA	1723	<i>J Antimicrob Chemother</i> 2000; 45 : 763-770
<i>tet(Q)</i>	TTA TAC TTC CTC CGG CAT CG ATC GGT TCG AGA ATG TCC AC	904	<i>Mol Cell Probes</i> 2001; 15 : 209-215
<i>mrsA</i>	TCC AAT CAT AGC ACA AAA TC AAT TCC CTC TAT TTG GTG GT	163	<i>Jpn J Infect Dis</i> 2007; 60 : 183-187
<i>mecA</i>	GTA GAA ATG ACT GAA CGT CCG ATA A CCA ATT CCA CAT TGT TTC GGT CTA A	310	<i>J Clin Microbiol</i> 2004; 42 : 4947-4955
<i>mef</i>	AGT ATC ATT AAT CAC TAG TGC	348	<i>Jpn J Infect Dis</i>

	TTC TTC TGG TAC AAA AGT GG		2007; 60: 183-187
ereA	AAC ACC CTG AAC CCA AGG GAC G CTT CAC ATC CGG ATT CGC TCG A	420	<i>Antimicrob Agents Chemother</i> 1996; 40: 1817-1824
ereB	AGA AAT GGA GGT TCA TAC TTA CCA CAT ATA AAT CAT CAC CAC CAA TGG CA	546	<i>Antimicrob Agents Chemother</i> 1996; 40: 1817-1824
mphA	AAC TGT ACG CAC TTG C GGT ACT CTT CGT TAC C	837	<i>Antimicrob Agents Chemother</i> 1996; 40: 1817-1824
ermA	AAG CGG TAA ACC CCT CTG A TTC GCA AAT CCC TTC TCA AC	190	<i>J Clin Microbiol</i> 2003; 41: 4089-4094
ermB	CTA TCT GAT TGT TGA AGA AGG ATG AAA GTT TAC TCT TGG TTT AGG ATG AAA	142	<i>J Antimicrob Chemother</i> 2000; 46: 527-534
ermC	AAT CGT CAA TTC CTG CAT GT TAA TCG TGG AAT ACG GGT TTG	299	<i>J Clin Microbiol</i> 2003; 41: 4089-4094
qnrB	GAT CGT GAA AGC CAG AAA GG ATG AGC AAC GAT GCC TGG TA	476	<i>Antimicrob Agents Chemother</i> 2009; 53: 639-645
qnrS	GCA AGT TCA TTG AAC AGG GT TCT AAA CCG TCG AGT TCG GCG	428	<i>Antimicrob Agents Chemother</i> 2009; 53: 639-645
sul1	ATG GTG ACG GTG TTC GGC ATT CTG A CTA GGC ATG ATC TAA CCC TCG GTC T	815	<i>J Antimicrob Chemother</i> 2003; 52: 1022-1024
sul2	CCT GTT TCG TCC GAC ACA GA GAA GCG CAG CCG CAA TTC AT	396	<i>Int J Antimicrob Agents</i> 2011; 37: 230-234
sul3	GAG CAA GAT TTT TGG AAT CG CAT CTG CAG CTA ACC TAG GGC TTT GGA	396	<i>J Antimicrob Chemother</i> 2003; 52: 1022-1024
nim1	ATG TTC AGA GAA ATG CGG CGT AAG CG GCT TCC TTG CCT GTC ATG TGC TC	458	<i>J Clin Microbiol</i> 1996; 34: 2078-2084
nim2	ATG TTC AGA GAA ATG CGG CGT AAG CG GCT TCC TTG CCT GTC ATG TGC TC	458	<i>J Clin Microbiol</i> 1996; 34: 2078-2084
nim3	ATG TTC AGA GAA ATG CGG CGT AAG CG GCT TCC TTG CCT GTC ATG TGC TC	458	<i>J Clin Microbiol</i> 1996; 34: 2078-2084
nim4	ATG TTC AGA GAA ATG CGG CGT AAG CG GCT TCC TTG CCT GTC ATG TGC TC	458	<i>J Clin Microbiol</i> 1996; 34: 2078-2084
aacA-aphD	TAA TCC AAG AGC AAT AAG GGC GCC ACA CTA TCA TAA CCA CTA	227	<i>J Clin Microbiol</i> 2003; 41: 4089-4094
vatA	TGG TCC CGG AAC AAC ATT TAT TCC ACC GAC AAT AGA ATA GGG	268	<i>J Clin Microbiol</i> 2003; 41: 4089-4094
vatB	GCT GCG AAT TCA GTT GTT ACA CTG ACC AAT CCC ACC ATT TTA	136	<i>J Clin Microbiol</i> 2003; 41: 4089-4094
vatC	AAG GCC CCA ATC CAG AAG AA TCA ACG TTC TTT GTC ACA ACC	467	<i>J Clin Microbiol</i> 2003; 41: 4089-4094
vga	CCG AAC TGC TAT TAG CAG A AAG TTC GTT TCT CTT TTC GA	470	<i>Antimicrob Agents Chemother</i> 1999; 43: 1062-1066
vgb	ACT AAC CAA GAT ACA CAG GAC C TTA TTG CTT GTC AGC CTT CC	734	<i>Antimicrob Agents Chemother</i> 1999; 43: 1062-1066
linA	GGT GGC TGG GGG GTA GAT GT GCT TCT TTT GAA ATA CAT GG	323	<i>Antimicrob Agents Chemother</i> 1996; 40: 1817-1824
mrsB	TAT GAT ATC CAT AAT AAT TAT CCA ATC AAG TTTA TAT CAT GAA TAG ATT GTC CTG TT	334	<i>Antimicrob Agents Chemother</i> 1999; 43: 1062-1066
cepA	TTT CTG CTA TGT CCT GCC C ATC TTT CAC GAA GAC GGC	743	<i>Clinics</i> 2011; 66: 543-547
mexB	GTG TTC GGC TCG CAG TAC TC AAC CGT CGG GAT TGA CCT TG	244	<i>FEMS Microbiol Lett</i> 2005; 243: 125-131
mexD	CGA GCG CTA TTC GCT GC CGA GCG CTA TTC GCT GC	165	<i>BMC Microbiol</i> 2010; 10: 217
mexF	GGC AGT TGC ACC TCG A CGC CTG GTC ACC GAG GAA GAG T	255	<i>Apmis</i> 2005; 113: 187-196
mexY	TAG TCC ATG GCT TGC GGG AAG C CCG CTA CAA CGG CTA TCC CT	250	<i>FEMS Microbiol Lett</i> 2005; 243: 125-131

Table S2. Sequence of oligonucleotide probes used for fluorescence in situ hybridization.

Probe	Target	Sequence (5' – 3')	Formamide	Reference
Bfrag998	<i>Bacteroides fragilis</i>	GTTTCCACATCATTCCACTG	30%	<i>Syst Appl Microbiol</i> 2003; 26 : 110-118
Bfra602	<i>Bacteroides fragilis</i> group	GAGCCGCAAACCTTCACAA	30%	<i>Appl Environ Microbiol</i> 1998; 64 : 3336
ECO1167	<i>Escherichia coli</i>	GCATAAGCGTCGCTGCCG	40%	<i>Syst Appl Microbiol</i> 1995; 18 : 113-122
Enc1259	<i>Enterococcus</i> spp.	GAAGTCGCGAGGCTAAGC	35%	<i>Syst Appl Microbiol</i> 2000; 23 : 563-572
FUS664	<i>Fusobacterium</i> sp.	CTTGTAGTTCCGCYTACCTC	40%	<i>Eur J Oral Sci</i> 2004; 112 : 33-41
PINT649	<i>Prevotella intermedia</i>	GCCGCCRCTGAASTCAGGCC	40%	<i>Microbiology</i> 2002; 148 : 1379-1387
PNG675	<i>Prevotella nigrescens</i>	TCCGCCTGCGCTGCGTGTA	40%	<i>Microbiology</i> 2002; 148 : 1379-1387
Pae997	<i>Pseudomonas</i> sp.	TCTGGAAAGTTCTCAGCA	35%	<i>Syst Appl Microbiol</i> 1995; 18 : 113-122
KO218	<i>Staphylococcus</i> sp.	GAAGCAAGCTTCTCG TCCG	20%	<i>J Microbiol Methods</i> 2005; 62 : 37-56
Str	<i>Streptococcus</i> spp.	CACTCTCCCCTCTGCAC	30%	<i>Med Microbiol Immunol</i> 2000; 188 : 169-175
ACA652	<i>Acinetobacter</i> spp.	ATCCTCTCCCATACTCTA	35%	<i>Appl Environ Microbiol</i> 1994; 60 : 792-800

Table S3. Anthropometric and nutritional characteristics of the participants.

Characteristics (mean ± SD)	Groups of individuals			p < 0.05*
	Eutrophic (n = 24)	Overweight (n = 24)	Obese (n = 24)	
Gender (male/female)	05/19	12/12	09/15	
Average age (years ± SD)	37.91 ± 12	38.12 ± 13.38	42.79 ± 11.67	
Average Body Mass Index	22.81 ± 01.8	27.05 ± 01.3	36.92 ± 06.0	a,b,c
Abdominal circumference	88.05 ± 7.4	94.65 ± 5.5	114.98 ± 15.3	a,b,c
Waist circumference	80.95 ± 06.7	88.70 ± 4.5	111.99 ± 16.6	a,b,c
Hip circumference	99.60 ± 04.5	105.40 ± 4.30	117.45 ± 08.6	a,b,c
Waist-hip ratio	0.81± 0.55	0.84 ± 0.52	0.97 ± 0.12	a,b,c
Mean daily calorie intake (kcal)	1 891.58 ± 679	2 018.79 ± 845	2 191.33 ± 1,049	c
Mean daily carbohydrate intake (%)	53.60 ± 9.10	55.02 ± 7.78	57.43 ± 8.02	c
Mean daily lipids intake (%)	27.27 ± 8.79	29.87 ± 5.92	28.74 ± 5.60	-
Mean daily protein intake (%)	19.13 ± 4.30	15.11 ± 4.54	13.83 ± 3.27	-
Mean daily fibers intake (g)	23.75 ± 8.30	25.03 ± 13.30	30.49± 5.42	b,c

* Significant statistical analysis: *a*—comparison between eutrophic and overweight volunteers' values; *b*—comparison between overweight and obese volunteers' values; *c*—comparison between eutrophic and obese volunteers' values. SD = Standard Deviation

Table S4. Declaration of habitual intake of xenobiotic, other than antimicrobial drugs, as reported by the participants.

Xenobiotics	Habitual intake (%)		
	Eutrophic (n=24)	Overweight (n=24)	Obese (n= 24)
Antihypertensive	4.2	12.5	79.2
Antacids	0	0	8.4
Anti-inflammatories	0	4.2	8.4
Diuretic	0	0	4.2
Barbiturates	0	0	4.2
Antidepressant	0	0	4.2
Hormone Drugs (Tamoxifen)	0	4.2	0
Contraceptional	0	4.2	0
Whey protein	4.2	0	0
Nutritional Supplements	Glucosamine Sulfate	4.2	0
	Chondroitin sulfate	4.2	0
	Others	8.4	0
	Sucralose and Acesulfame K	8.4	8.4
Artificial sweeteners	Stevioside and cyclamate-saccharin	0	4.2
	cyclamate-saccharin	4.2	8.4
	Others	0	0
Indeterminate	0	4.2	0