

Table S2. Primers used in the study.

Primer	Sequence 5' - 3'	Purpose of the primer
2S_APH(6)_clon_F 2S_APH(6)_clon_R	AGCGAACCCTACCTGAGCCG TCACTTCTCCGCCAGCGCCT	Cloning of <i>S. maltophilia</i> APH(6) family gene
2S_APH(3)_clon_F 2S_APH(3)_clon_R	GGAGCACCCGATCCCTTCA TCAGAAGAAGCTCGTCGAGCA	Cloning of <i>S. maltophilia</i> APH(3') family gene
2C_aden_trans_clon_F 2C_aden_trans_clon_R	AAAATAAGAGAAGATAAGCTTGA ACACATCAGGAATTCTGGTTG	Cloning of <i>Chryseobacterium</i> sp. ANT(6) family gene
2C_TETR_MFS_clon_F 2C_TETR_MFS_clon_R	GAAAATTCAAGGAAAAAAGCTG CTATTTTTTAATATCCTTCTTTTTT	Cloning of <i>Chryseobacterium</i> sp. tetracycline MFS efflux pump gene
2C_IND_clon_F 2C_IND_clon_R	AGAAAAAGTATTCGATTTTTTAATTA TAGCCTCGAGCTATTTTTTATTTTCA TTTAGAAGT	Cloning of <i>Chryseobacterium</i> sp. IND-like MBL gene
2C_Met_beta_lact_clon_F 2C_Met_beta_lact_clon_R	AAAACCTACCCTTACCCATATAG TAGCCTCGAGTTATTGAGTTTTAAA TAATGTAATG	Cloning of <i>Chryseobacterium</i> sp. putative MBL gene
Steno_kanAPH(3)_F Steno_kanAPH(3)_R	CSTTCSTGAARTCGGAAGTGATCG CCRCARTCRATGAAACCRCTGAA	Detection of <i>S. maltophilia</i> APH(3') family gene
Steno_strepAPH(6)_F Steno_strepAPH(6)_R	NCCRCASAGRTCCGGRITG GGYGACCTGCAYCAYGACAA	Detection of <i>S. maltophilia</i> APH(6) family gene
Chrbct_Ind_F Chrbct_Ind_R	GCAACACATTCGCATGATGATA TGKCCRCKCCTTYCCATT	Detection of <i>Chryseobacterium</i> sp. IND-like MBL gene
Chrbct_StrR_F Chrbct_StrR_R	TCCYYATGCWCKGTAGATGA TCCCACCARAAATCATTNAKC	Detection of <i>Chryseobacterium</i> sp. ANT(6) family gene
Chrbct_TetrMFS_F Chrbct_TetrMFS_R	GYTDGGWTTTGCSTAYGC RCRAACATCATCCADCCYT	Detection of <i>Chryseobacterium</i> sp. tetracycline MFS efflux pump gene
blaL1_Sm_F blaL1_Sm_R	ACCACACCTGGCAGATCGG TCGCCATCCATGATGATGCG	Detection of <i>S. maltophilia</i> blaL1
blaL2_Sm_F blaL2_Sm_R	GAGCGCTTCCCGATGTGCA CATTGCTGCCGGTCTTGTC	Detection of <i>S. maltophilia</i> blaL2
IND_sig_F IND_R	CAGGTAAGAGATTTTGTGATC CTATTTTTTATTTTCATTTAGAAGT	IND-like MBL purification
MBL_sig_F MBL_R	TGTGGCTCTCAACATAAAG TTATTGAGTTTTAAATAATGTAATG	Putative MBL purification