

S1 Table. List of primers used for screening metal tolerant genes in *Chryseobacterium* sp. PMSZPI

Primer	Oligo sequences 5'-3'	Target Genes (Organisms)	Annealing Tm (°C)
czcAF1	CGACCTGCGCACCNENCARGAYTG	<i>czcA</i> gene (<i>Chryseobacterium</i> sp., <i>Serratia</i> sp., <i>Pseudomonas</i> sp.)	46
czcAR1	CGGGTGGAACATCTTGCCYTCNWSNCC		
czcDF2	CGATGCCGGCCATATGYTNWSNGA	<i>czcD</i> gene (<i>Arthrobacter</i> sp.)	49
czcDR2	CGGATGTAATTGTCCAAATATGNARRTCRTG		
81JC	GGATGTCCTTGTGCTYTART	<i>cadA</i> gene (<i>Bacillus</i> , <i>Acinobacter</i> , <i>Pseudomonas</i>)(Martinez et al.2006)	49
84JC	GGAGCATCGTTAATDCCRDCC		

Table 2: List of primers for qRT-PCR studies

Primer	Oligo sequences 5'-3'	Amplicon Size (bp)	Target Gene
ZP1Fc	GGTGGATAGAGATGGGCACG	107	<i>16S rRNA</i>
ZP1Rc	AGTACCAGTGTGGGGGATCA		
ZP1F1a	TGATGTTTTTCGCAGCATTGG	103	<i>czcA</i>
ZP1R1a	TCCCAAAGTCCCTCACTCC		
PMZF2b	CCCGGTTGCTACTGCAATTC	110	<i>cadA</i>
PMZR2b	TTCCATTCACCGTTGCTTTCA		
ZP1F1d	GGAGTCATGATTGCCGGAGTT	120	<i>czcD</i>
ZP1R1d	AGCCTCCATCAACAGTTTCCA		

Table 3: BLASTX analysis of metal transporters from *Chryseobacterium* sp. (PMSZPI)

Gene Name (Accession. No)	The closest match of metal tolerant genes from PMSZPI with those in NCBI Database after BLASTX	Similarity Percentage
<i>czcA</i> (MH423682)	CusA/CzcA family heavy metal efflux RND transporter [<i>Chryseobacterium</i> sp. UNC8MFCoI] (WP_027372322.1)	98
<i>czcD</i> (MH394701)	Cation diffusion facilitator family transporter [<i>Chryseobacterium</i> sp. ISE14] (WP109713444.1)	97
<i>cadA</i> (JN034431)	Cadmium-translocating P-type ATPase [<i>Chryseobacterium</i> sp. ISE14] (WP_103246897.1)	98