

# **Soil Contamination in China: Current Status and Mitigation Strategies**

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**Supporting information: 1 Table**

**Table S1.** Quantitative trait loci (QTLs) for grain Cd and As concentrations in rice

Traits	Population	QTLs	Chromosome	Nearest marker	Marker type	R <sup>2</sup> (%)	Reference
Grain Cd concentration	CSSLs (Koshihikari×Kasalath)	CdG3	3	S1513	RFLP	-	1
		CdG6	6	R2171	RFLP	-	
		CdG8	8	C390	RFLP	-	
	BILs (Nipponbare×Kasalath)	qcd4-1	4	R93	RFLP	8.2	2
		qcd4-2	4	R514	RFLP	7.9	
		qcd11	11	C189	RFLP	10.5	
	BILs (Sasanishiki×Habataki)	qGCd2	2	R2511	RFLP	11.8	3
		qGCd7	7	C383	RFLP	35.5	
	RILs (Bala×Azucena)	qCd1	1	C949	RFLP	18.9	4
		qCd2	2	RM6	SSR	9.7	
		qCd3	3	RG191	RFLP	8.7	
		qCd7	7	G338	RFLP	20.2	
		qCd9	9	C506	RFLP	12.2	
		qCd10	10	AB1001	Undefined	16.8	
	RILs (Fukuhibiki×LAC23)F6	qLCdG3-1	3	RM426	SSR	13.86	5
		qLCdG11	11	NBLAC61	CAPS	9.41	
	RILs (Fukuhibiki×LAC23)F7	qLCdG3-2	3	RM1940	SSR	8.29	
qLCdG11		11	NBLAC61	CAPS	12.91		
BILs (Koshihikari×Jarjan)	qCdp7	7	RM3859	SSR	46	6	
DH (JX17×ZYQ8)	qCdc3	3	G249	RFLP	18.21	7	

		qCdc4	4	C513	RFLP	22.44		
		qCdc6	6	G122	RFLP	13.56		
	CSSLs (Koshihikari×LAC23)	qlGCd3	3	RM16153	SSR	-	8	
	RILs (Suwon490×SNU-SG1)	gcc3	3	RM545	SSR	16.7	9	
		gcc9	9	RM1328	SSR	17.8		
		gcc11	11	RM286	SSR	16.1		
	LT-RILs (Lemont×TeQing)	Cd1	1	RG236	RFLP	6.3	10	
		Cd9	9	RG570	RFLP	5.8		
	BILs (TeQing-into-Lemont ) unflooded	Cd2.1	2	RM6840	SSR	-		
		Cd2.2	2	RM166	SSR	-		
	BILs (TeQing-into-Lemont ) flooded	Cd3	3	RM132	SSR	-		
		Cd5	5	RM13	SSR	-		
		Cd6	6	OSR21	SSR	-		
		Cd8	8	RM149	SSR	-		
Grain As concentration	DH (CJ06×TN1)	AsSe1	6	RM3	SSR	26.3		11
		AsSe2	8	RM1111	SSR	35.2		
	RILs (Bala×Azucena)	qAs8	8	R902	RFLP	22.8	12	
		qAs10	10	a12372	Genomic DNA	8.5		
	LT-RILs (Lemont×TeQing)	As5	5	RG13	RFLP	3.7	10	
	BILs (TeQing-into-Lemont ) flooded	As2	2	RM341	SSR	-		
		As9	9	RM1189	SSR	-		
As11		11	RM167	SSR	-			

	BILs (TeQing-into-Lemont ) unflooded	As3	3	RM16	SSR	-	
Grain DMA concentration*	F2 (Padi Perak×Koshihikari)	qDMAs6.1	6	AE06000003	SNP	36	13
		qDMAs6.2	6	AE06001721	SNP	25	
		qDMAs8	8	AE08000162	SNP	12	

\* DMA, dimethylarsinic acid.

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