

Table S1. Primers used for qRT-PCR in this study.

Gene Name	Forward Primers	Reverse Primers
<i>ACTIN1</i>	TCCATCTTGGCATCTCTCAG	GTACCCTCATCAGGCATCTG
<i>OsNramp1</i>	CATCGCATACCTTGATCCTAGT	GGAGTACCCATAGCAACGAATA
<i>OsNramp5</i>	TTCGTTTATATTTGTGCGGTCC	CACCTCCCCTCAAATGCTTATA
<i>OsIRT1</i>	GCAATTCGCTGCATTGTTAGAT	GAGAAGTCACAGTCACTGTACA
<i>OsIRT2</i>	CTTCCACCAGATGTTCGAGG	GGTGGAGAAGAAGAAGACCAG
<i>OsHMA2</i>	ATACTCATGCTGATTGCTGGTA	CAAGCCAAAATGCATGCATTAG
<i>OsHMA3</i>	CAATGGTGTGGTTCGTTGC	CTCCCATTTCTGCAGTCTTTC
<i>OsLCT1</i>	AGCACATCTCTGGCTTCCAC	CGGCTCATTGCATTCTGCTC
<i>OsNAP</i>	CAAGAAGCCGAACGGTTC	GTTAGAGTGGAGCAGCAT
<i>SGR</i>	AGGGGTGGTACAACAAGCTG	GCTCCTTGC GGAAGATGTAG
<i>RCCR1</i>	CGCATTTCTCATGGAATTT	CTTCTCACGCTGTTTGTCCA
<i>NYC1</i>	CATGCAACACCAACAAAAGG	GACCATTCCAGGAGAAGCAG
<i>NYC3</i>	TGTCGTTGCCATGTGAAGAT	TTGGTCACGCCACAAATCTA
<i>Osh69</i>	CCACAACACGGATAACTT	GGTGAACACTATGGAACA
<i>Osh36</i>	GCACGGAGGCGAACGA	TTGAGCGGTAGCACCCATT
<i>OsI57</i>	ACCCTAAAGTAAATGAAGTC	CCTGCTCTTGTCTTGTTA

Table S2. P value corresponding to Figure S10. P1, P value of CK versus +Cd; P2, P value of +Cd versus +Cd&CA; P3, P value of +Cd versus +Cd&M50; P4, P value of +Cd&M50 versus +Cd&M50-AuNPs; P5, P value of +Cd&M100 versus +Cd&M100-AuNPs; P6, P value of +Cd&M200 versus +Cd&M200-AuNPs. Student *t*'tests were used to generate P value. P < 0.05 was marked in red.

Gene	P-value	0d	5d	10d	Gene	P-value	0d	5d	10d
<i>OsNramp1</i> in roots	P1	0.39917	0.00168	0.00055	<i>OsNramp1</i> in leaves	P1	0.21445	0.00015	0.00017
	P2	0.40585	0.46044	0.08265		P2	0.08008	0.13819	0.05496
	P3	0.30028	0.28642	0.44985		P3	0.13619	0.10344	0.12347
	P4	0.31075	0.00394	0.03718		P4	0.19013	0.04356	0.43464
	P5	0.37621	0.01564	0.02399		P5	0.48898	0.02520	0.00893
	P6	0.49805	0.00425	0.02259		P6	0.21204	0.01049	0.00563
<i>OsNramp5</i> in roots	P1	0.47417	0.00034	0.00404	<i>OsNramp5</i> in leaves	P1	0.46831	0.00023	0.00020
	P2	0.35977	0.13978	0.15235		P2	0.16033	0.15943	0.41021
	P3	0.12855	0.43998	0.33198		P3	0.19057	0.47426	0.09335
	P4	0.26345	0.00018	0.00103		P4	0.08057	0.00540	0.03440
	P5	0.18091	0.00424	0.00043		P5	0.40106	0.00681	0.02434
	P6	0.21035	0.00361	0.00246		P6	0.12949	0.02801	0.00032
<i>OsIRT1</i> in roots	P1	0.22088	0.00005	0.00142	<i>OsIRT1</i> in leaves	P1	0.16501	0.00172	0.29538
	P2	0.06756	0.40174	0.37752		P2	0.24298	0.48296	0.15080
	P3	0.46318	0.20199	0.28821		P3	0.45281	0.09252	0.07243
	P4	0.25661	0.00280	0.03175		P4	0.08298	0.00123	0.07636
	P5	0.07214	0.00019	0.01525		P5	0.13910	0.04237	0.18450
	P6	0.05657	0.00029	0.00029		P6	0.49634	0.01929	0.33353
<i>OsIRT2</i> in roots	P1	0.25921	0.00036	0.00677	<i>OsIRT2</i> in leaves	P1	0.49792	0.00531	0.10411
	P2	0.06064	0.48351	0.19100		P2	0.10749	0.48317	0.19916
	P3	0.23843	0.13262	0.42825		P3	0.16332	0.23578	0.34102
	P4	0.25661	0.00135	0.00238		P4	0.33495	0.02386	0.38527
	P5	0.30064	0.00136	0.02021		P5	0.36216	0.00773	0.43732
	P6	0.31495	0.00037	0.00032		P6	0.30113	0.01051	0.09666
<i>OsHMA2</i> in roots	P1	0.00761	0.00128	0.00214	<i>OsHMA2</i> in leaves	P1	0.27665	0.00189	0.46963
	P2	0.29231	0.10656	0.42997		P2	0.37036	0.23606	0.15279
	P3	0.42854	0.11063	0.28935		P3	0.43231	0.11172	0.14842
	P4	0.35838	0.00447	0.02749		P4	0.28283	0.01048	0.36487
	P5	0.37283	0.00071	0.00698		P5	0.28670	0.00707	0.11446

	P6	0.49223	0.00010	0.00180		P6	0.41183	0.02520	0.32877
	P1	0.27086	0.00025	0.00148		P1	0.43786	0.00010	0.40838
	P2	0.35962	0.43063	0.48347		P2	0.38334	0.47339	0.23915
<i>OsHMA3</i>	P3	0.33794	0.34040	0.44742	<i>OsHMA3</i>	P3	0.27129	0.18493	0.45693
in roots	P4	0.42865	0.00489	0.01781	in leaves	P4	0.24401	0.01849	0.10806
	P5	0.43979	0.00083	0.00284		P5	0.18505	0.00345	0.27264
	P6	0.49475	0.00332	0.00000		P6	0.26716	0.02093	0.40869
	P1	0.19456	0.00001	0.35815		P1	0.21219	0.00051	0.29007
	P2	0.30302	0.19927	0.38987		P2	0.09464	0.11208	0.49092
<i>OsLCT1</i> in	P3	0.30566	0.00366	0.21671	<i>OsLCT1</i> in	P3	0.46281	0.17528	0.30280
roots	P4	0.40205	0.00014	0.22445	leaves	P4	0.38069	0.03614	0.24697
	P5	0.31025	0.00592	0.36463		P5	0.16530	0.02587	0.40672
	P6	0.31578	0.00089	0.43968		P6	0.40160	0.00427	0.14660

Table S3. P value corresponding to Figure S9. P1, P value of CK versus +Cd; P2, P value of +Cd versus +Cd&CA; P3, P value of +Cd versus +Cd&M50; P4, P value of +Cd&M50 versus +Cd&M50-AuNPs; P5, P value of +Cd&M100 versus +Cd&M100-AuNPs; P6, P value of +Cd&M200 versus +Cd&M200-AuNPs. Student *t*'tests were used to generate P value. P < 0.05 was marked in red.

Gene	P-value	0d	5d	10d	Gene	P-value	0d	5d	10d
<i>SGR</i>	P1	0.44335	0.00031	0.00027	<i>RCCR1</i>	P1	0.19319	0.00022	0.00015
	P2	0.06973	0.19966	0.09556		P2	0.34371	0.21132	0.43621
	P3	0.09495	0.01383	0.02604		P3	0.16290	0.01074	0.00462
	P4	0.14037	0.01190	0.00885		P4	0.26137	0.03337	0.02480
	P5	0.24098	0.00228	0.00029		P5	0.38496	0.00155	0.00693
	P6	0.30002	0.00482	0.00260		P6	0.08681	0.00039	0.00179
<i>NYCI</i>	P1	0.22319	0.00044	0.00083	<i>NYC3</i>	P1	0.45831	0.00024	0.00026
	P2	0.13441	0.21595	0.23295		P2	0.12588	0.12343	0.09556
	P3	0.09828	0.03493	0.00980		P3	0.26922	0.00681	0.04188
	P4	0.38986	0.00916	0.02051		P4	0.40420	0.00679	0.00963
	P5	0.19149	0.00093	0.00373		P5	0.16211	0.00146	0.00154
	P6	0.08008	0.00017	0.00222		P6	0.02585	0.00001	0.00068
<i>OsNAP</i>	P1	0.13429	0.00113	0.00036	<i>Osh69</i>	P1	0.17539	0.00074	0.00163
	P2	0.19017	0.05175	0.07088		P2	0.12893	0.14465	0.11419
	P3	0.14014	0.00617	0.01094		P3	0.05986	0.00875	0.03010
	P4	0.03893	0.00012	0.00065		P4	0.41421	0.01190	0.00188
	P5	0.34606	0.00018	0.01900		P5	0.44249	0.00029	0.00341
	P6	0.09335	0.00656	0.00110		P6	0.39718	0.00431	0.00011
<i>Osh36</i>	P1	0.28576	0.00200	0.00024	<i>Osh57</i>	P1	0.07766	0.00030	0.00057
	P2	0.06973	0.18836	0.09556		P2	0.42444	0.14991	0.10553
	P3	0.09495	0.03763	0.00303		P3	0.05360	0.00755	0.03679
	P4	0.14037	0.00776	0.02279		P4	0.28908	0.01280	0.03181
	P5	0.41133	0.00202	0.00029		P5	0.15907	0.00212	0.00118
	P6	0.33346	0.00039	0.00109		P6	0.33346	0.00377	0.00697