

Genetic differentiation in weedy (*O. sativa* f. *spontanea*) and wild rice (*O. rufipogon*) revealed by DNA sequence in the QTL *SD7-1/Rc* locus

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**Supporting data 1 (S1).** Photos: Habitats and localities of rice accession collected and stored in laboratory. For wild rice (*O. rufipogon*) at the three top photos, Vientiane province, Laos (Leftmost), Phisanuloke province (middle), Sakon Nakhon province (rightmost). For weedy rice at two bottom photos, Thung Kula Ronghai, Mahasarakham province (left), and Pichit province (right).

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**Table 2.** Four primer pairs used for PCR and DNA sequencing of the *Pi-ta* gene in wild, weedy and cultivated rice (according to Haung et al., 2008).

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Code	Forward primer	Reverse primer	Expected PCR product (bp)
2322F/3370R	5'-GGGCGATCCATGCTGTCAAATC-3'	5'-CCTTTTATCTTGCAAATGCGTCCG-3'	1048
3241F/4615R	5'-GCCTGACATGACGAAGATCC-3'	5'-GCCGACTACTAGCTCCAAATC-3'	1374
4473F/6012R	5'-CCTACAGATCTGTAGCCAGC-3'	5'-GTACCTGTGACAGTGAGGGAGC-3'	1539
5877F/6922R	5'-CCAAGGACTACAACACTTGC-3'	5'-GTTCTTTGATCCAAGTGTAGG-3'	1045

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