

**Supplementary Table S2.** Mapping statistics of CLC trimmed reads from Illumina and PGM Ion Torrent platforms.

	<i>Oryza sativa</i> spp. <i>japonica</i>		Wild rice		
	var. <i>Nipponbare</i>		<i>(Oryza rufipogon-like)</i>		
	GAll Illumina	PGM Ion Torrent	HiSeq Illumina	HiSeq Illumina subset	PGM Ion Torrent
Mapped reads	525,452	223,605	12,424,965	327,403	169,665
(Forward/Reversed)	(262,567/262,885)	(112,154/111,451)	(6,211,497/6,213,468)	(164,024/163,898)	(85,317/84,348)
% mapped reads	5.42	3.39	3.43	3.46	3.17
Non-specific matches	204,877	64,560	3,601,138	93,502	50,521
(% non-specific matches)	(39.9)	(28.9)	(29.0)	(28.5)	(29.8)
Non-perfect matches	18,385	164,176	1,745,820	47,444	130,855
(% non-perfect matches)	(3.5)	(73.4)	(14.1)	(14.5)	(77.1)
Min coverage	7	96	1,203	75	69
Max coverage	688	534	13,231	360	431
Average coverage	127.7	333.4	8,143.2	218.4	243.6
% GC	39	39	39	39	39
Consensus length	134,551	134,520	134,531	134,529	134,525