



Fig.S1. Identification of *oslpr5-1* and *oslpr5-2*. (a) The gene structure of *OsLPR5*. Black boxes, exon; Grey boxes, UTR; Line, intron. (b) The mutation sites of two independent lines and the comparative nucleotide sequences of the WT, and *OsLPR5* mutants. The red letters in the nucleotide sequences indicate PAM. (PAM: protospacer adjacent motif) (c) The peak maps of sequences in *OsLPR5* mutants.