

Supplemental Figure 2. Southern hybridization analysis of putative HDR T0 plants. Genomic DNA of seven *DD*43 and two *DD*20 T0 plants were digested with *Nde*I or *Mfe*I to check the 5' and 3' borders using a *HPT* gene probe. Three *DD*43 plants D5-9, F9-7, and G1-2 contained the predicted HDR-specific 5' border 5136 bp *Nde*I band and 3' border 4375 bp *Mfe*I band (circled). A predicted 1748 bp common *Nde*I band was detected in all *DD*43 samples (arrow). One *DD*20 plant A12-36 contained the predicted HDR-specific 5' border 3927 bp *Nde*I band and 3' border 5278 bp *Mfe*I band (circled). A predicted 2427 bp common *Nde*I band was detected in both *DD*20 samples (arrow). The displayed DIGVII markers are 8576, 7427, 6106, 4899, 3639, and 2799 bp.