Table S4. Summary of SNP-based clonal identity validation for *Fix10* plants.

Plants	Number of SNPs	Number of heterozygous SNPs	Resolution of SNPs	Total SNP analyzed (16A vs. C927)	Heterozygous SNP identical to CY927	Heterozygous SNP shared with CY927(%)
16A	503,034	144,291	742.0	−	−	−
C927	2,723,730	446,568	137.0	−	−	−
CY927	2,823,993	2,615,241	132.2	−	−	−
CY927 F2 #1	2,367,588	1,610,583	157.6	−	−	−
CY927 F2 #2	2,479,421	1,905,622	150.5	−	−	−
Clonal *Fix10* T_1 #2	2,580,465	2,352,864	144.6	2,131,467	1,913,387	89.8
Clonal *Fix10* T_1 #6	2,525,930	2,294,281	147.8	2,131,467	1,874,784	88.0
Clonal *Fix10* T_1 #7	2,640,509	2,418,139	141.4	2,131,467	1,959,908	92.0
Clonal *Fix10* T_2 #2	2,814,808	2,598,063	132.6	2,131,467	2,078,411	97.5
Clonal *Fix10* T_2 #6	2,674,430	2,448,766	139.6	2,131,467	1,974,892	92.7
Clonal *Fix10* T_2 #7	2,756,744	2,538,003	135.4	2,131,467	2,028,741	95.2