

Table S1. The list of primer sequences of 6 starch synthesis gene used in interval mapping and marker regression analysis

| Locus | Chr | Forward primer ^a | Reverse primer | size ^b |
|-----------------|-----|--------------------------------|---------------------------|----------------------|
| <i>PUL</i> | 4 | CCACCATTAAAGCATCATCAAC | AGTTGTTATATTTTAGGATGGATGG | -20 |
| <i>SBE4</i> | 4 | CCATCACCTCAAATACATCACTC | AGACTGGAATGCCCCCTTAGG | -13 |
| <i>Wx</i> | 6 | TGTTGTTTCATCAGGAAGAACATCTCCAAG | TTAATTTCCAGCCCAACACC | dCAPS(<i>Sty1</i>) |
| <i>SBE1</i> | 6 | TGCTACATAACACGCATACAAAGT | AGACAAAAGCGAAAGGTAATGAG | -331 |
| <i>ISA1</i> | 8 | ATAGATGCTAATGTGATGTGGC | TGGTATAGGCACAACCGTAGA | -10 |
| <i>SSII-1_3</i> | 10 | ATCTTTAGACGATTAGCG | AAGTCACAAGTAGAAGGG | -6 |

^a primer sequences from Yan et al. (2011)

^b the amplicon size of TNG 78 differed from TCS 17 which + and – indicate the amplicon size of TNG 78 is larger than which of TCS 17.

Table S2. The list of primer sequences of genes used in detection gene expression levels in immature grains by real-time PCR

| Gene | Accession | Chr. | Forward | Reverse | Size |
|---------------------------|---------------------|------|---------------------------|---------------------------|------|
| <i>EFla-Q^a</i> | <i>Os03g0178000</i> | 3 | CGTTATTATCATATGCTCGCTGTGT | CAGCCGAAGGGCAATAATCA | 101 |
| <i>SBE4^b</i> | <i>Os04g0409200</i> | 4 | GCCAATGCCAGGAAGATGA | GCGCAACATAGGATGGGTTT | 128 |
| <i>PUL^c</i> | <i>Os04g0164900</i> | 4 | AATCTCCGATGGCTGTTCCTT | TTGAGTTCACACCCGATGCTAA | 90 |
| <i>ISA2^b</i> | <i>Os05g0393700</i> | 5 | TAGAGGTCCTCTTGGAGG | AATCAGCTTCTGAGTCACCG | 170 |
| <i>GBSSI^c</i> | <i>Os06g0133000</i> | 6 | GATGCGTTTCAGCCTTCTTTG | AGTATGGGTTGTTGTTGAGGTTTAG | 74 |
| <i>SSII-3^b</i> | <i>Os06g0229800</i> | 6 | GCTTCCGGTTTGTGTGTTC | CTTAATACTCCCTCAACTCCACCAT | 54 |
| <i>CPE</i> | <i>Os07g0604300</i> | 7 | AAACCAGGAAGCTGTGTAGA | CTGGACTAGAGGTGTCAAGC | 173 |
| <i>AGP</i> | <i>Os07g0604800</i> | 7 | TGCGATGTACTTTGGTCTTA | AGTGATCGGTAATCACCTTG | 95 |
| <i>MADS18</i> | <i>Os07g0605200</i> | 7 | ATCCCGACAACAAATAACAG | AGACGGTTGAGCTTCTGACT | 66 |
| <i>SSII-1^c</i> | <i>Os10g0437600</i> | 10 | GATGAAGTTCAGACGATGATGA | TGACTTAGTTGTGCCTCATAATCCT | 113 |

^{a, b, c} indicate the primer sequences adopted Jain et al. (2006), Ohdan et al. (2005), and Zeng et al. (2007), respectively.

Table S3. Descriptive statistics for eight grain quality-related physiochemical properties of rice harvested from the two crop seasons

| Year and trait | Parents | | RIL population | |
|----------------|---------|--------|------------------------------|-----------|
| | TNG 78 | TCS 17 | Mean±SD | Range |
| 2010-II | | | | |
| PLS | 43 | 43 | 47.46±11.22 | 19-72 |
| PKV (cp) | 2844 | 2453 | 2369.50±396.18 | 1355-3595 |
| HPV (cp) | 1591 | 2182 | 1742.01±363.12 | 1028-2816 |
| BDV (cp) | 1253 | 272 | 654.49±453.46 | 5-1712 |
| CPV (cp) | 2911 | 4001 | 3313.19±683.81 | 2173-5115 |
| SBV (cp) | 67 | 1548 | 916.69±752.88 | -566-2194 |
| PeT (min) | 5.67 | 6.3 | 6.04±0.27 | 5.5-6.8 |
| PaT (°C) | 86.45 | 87.65 | 88.49±2.16 | 74-92.9 |
| 2011-I | | | | |
| PLS | 55 | ND | 46.73±10.6 ^{a**} | 21-69 |
| PKV (cp) | 3066 | 2954 | 2579.10±458.14 | 1149-3523 |
| HPV (cp) | 1620 | 2556 | 1848.12±517.42 ^{**} | 605-3060 |
| BDV (cp) | 1447 | 398 | 730.98±428.08 ^{**} | 12-1766 |
| CPV (cp) | 2830 | 4486 | 3556.75±963.8 ^{**} | 1371-5488 |
| SBV (cp) | -237 | 1532 | 985.18±813.09 [*] | -628-2610 |
| PeT (min) | 6.03 | 6.17 | 5.95±0.3 ^{**} | 5.2-6.7 |
| PaT (°C) | 79.6 | 86.43 | 86.57±2.65 ^{**} | 75.6-91.3 |

^a paired difference between the two crop seasons of 8 traits were detected by t-test which * and ** are indicated at significant levels of 0.05 and 0.01 with 2-tails, respectively.

Fig. S1.



