

Appendix A Primer Sequences of Enzyme Genes Related to Grain Starch Synthesis

| Gene name | Accession No. | Primer sequences (5'-3') |
|-----------|---------------|--|
| AGPS2 | AK071826 | F:TCTTTTGTGGCCATTCATCTGG R:TGATTCCAAGCACACTCTCATCGAC |
| AGPL2 | AK071497 | F:TAGATAGGCCTTGGAATCGCACC R:TAGAGTTCCCATTCCAAAACAAACC |
| SSI | AK109458 | F:GGGCCTTCATGGATCAACC R:CCGCTTCAAGCATCCTCATC |
| SSIIa | AK101978 | F:GGCCAAGTACCAATGGTGAA R:GCATGATGCATCTGAAACAAAGC |
| SSIIIa | AK061604 | F:GCCTGCCCTGGACTACATTG R:GCAAACATATGTACACGGTTCTGG |
| GBSSI | X62134 | F:GATGAGATACGGAACGCCCT R:GCCCATGTGGAAACCAGTCT |
| SBEI | AK065121 | F:TGGCCATGGAAGAGTTGGC R:CAGAAGCAACTGCTCCACC |
| SBEIIb | D16201 | F:ATGCTAGAGTTTGACCGC R:AGTGTGATGGATCCTGCC |
| ISA | | F:5GGGTCAATTTGCGCGTCTAC R:CATTCCCCGTCGATTGAAC |
| Actin | AY212324.1 | F:GTGATGGTTGGTATGGGGCA R:CTCAGTCAGCAACACAGGGT |

Note: F: Forward primer; R: Reverse primer

Appendix B Correlation of enzyme activities of starch synthesis and the starch content in superior and inferior spikelets of rice

| | Amylose content | | Amylopectin content | | Total starch content | | Amylose/Amylopectin ratio | |
|--------|-----------------|----------|---------------------|----------|----------------------|----------|---------------------------|----------|
| | Superior | Inferior | Superior | Inferior | Superior | Inferior | Superior | Inferior |
| AGP | 0.124 | 0.990** | 0.528 | 0.977** | 0.433 | 0.980** | -0.091 | -0.914** |
| SSS | 0.251 | 0.977** | 0.709 | 0.970** | 0.624* | 0.973** | -0.030 | -0.919** |
| GBSS | 0.504 | 0.991** | 0.634 | 0.995** | 0.696* | 0.996** | 0.273 | -0.961** |
| SBE | 0.262 | 0.990** | 0.624 | 0.995** | 0.569 | 0.995** | 0.017 | -0.960** |
| DBE | 0.384 | 0.973** | 0.532 | 0.986** | 0.565 | 0.985** | 0.188 | -0.968** |
| AGPS2b | 0.386 | 0.973** | 0.620* | 0.988** | 0.628* | 0.987** | 0.153 | -0.968** |
| AGPL2 | 0.380 | 0.976** | 0.581* | 0.976** | 0.598* | 0.977** | 0.163 | -0.932** |
| SSI | 0.661* | 0.980** | 0.341 | 0.960** | 0.568 | 0.964** | 0.568 | -0.888** |
| SSIIa | 0.270 | 0.839** | 0.477 | 0.828** | 0.470 | 0.831** | 0.088 | -0.777** |
| SSIIIa | 0.223 | 0.961** | 0.420 | 0.968** | 0.406 | 0.968** | 0.061 | -0.936** |
| GBSSI | 0.394 | 0.989** | 0.513 | 0.974** | 0.556 | 0.978** | 0.207 | -0.905** |
| SBEI | 0.542 | 0.978** | 0.609* | 0.989** | 0.697* | 0.989** | 0.324 | -0.967** |
| SBEIIb | 0.290 | 0.970** | 0.468 | 0.970** | 0.473 | 0.971** | 0.114 | -0.922** |
| ISAI | 0.390 | 0.981** | 0.821** | 0.984** | 0.772** | 0.985** | 0.071 | -0.953** |

Note: The symbol of * indicates the correlation significance at the 0.05 probability level; ** indicates the correlation significance at the 0.01 probability level.