

## Supplementary Table S7

Table S7 QTLs that overlapped with cloned panicle-related genes

| The QTLs identified in this study |     |               |           | Genes related to ear development that previous studies have cloned |     |                 |                 |                                     |                          |
|-----------------------------------|-----|---------------|-----------|--|-----|-----------------|-----------------|-------------------------------------|--------------------------|
| Loc                               | Chr | Start         | End       | Gene-ID  | Chr | start           | end             | Function                            | reference                |
|                                   |     |               |           | GRMZM2G011357  | 1   | 239,608,<br>944 | 239,612,26<br>7 | TF                                  | Colasanti et al.         |
|                                   |     |               |           | GRMZM2G081554  | 1   | 241,218,<br>603 | 241,226,85<br>4 | Kaurene synth.<br>(GA synthesis)    | Bensen et al.,<br>1995   |
|                                   |     |               |           | Zm00001d032961   | 1   | 244,858,<br>795 | 244,867,41<br>7 | ent-copy<br>diphosphate<br>synthase | Dellaporta et al<br>1994 |
| Sub-pop3_19DH                     | 1   | 13961614<br>9 | 252952675 |  |     |                 |                 |                                     | Bensen et al.<br>1995    |
|                                   |     |               |           | Zm00001d031068   | 1   | 175,807,<br>851 | 175,809,43<br>2 | serine-threonine<br>protein kinase  | McSteen et al.<br>2001   |
|                                   |     |               |           | Zm00001d032922   | 1   | 243,201,<br>405 | 243,204,86<br>5 | indeterminate<br>domain protein     | McSteen et al.<br>2007   |
|                                   |     |               |           |  |     |                 |                 |                                     | Skirpan et al.<br>2008   |
|                                   |     |               |           |  |     |                 |                 |                                     | Skirpan et al.<br>2009   |
|                                   |     |               |           |  |     |                 |                 |                                     | Colasanti et al.<br>1998 |

|               |   |                |                 |                |   |                 |                 |                                    |   |
|---------------|---|----------------|-----------------|----------------|---|-----------------|-----------------|------------------------------------|---|
|               |   |                |                 | Zm00001d031459 | 1 | 190,964,<br>811 | 190,978,36<br>9 | RNA<br>polymerase IV               | Erhard et al.<br>2009                                     |
|               |   |                |                 | Zm00001d031451 | 1 | 190,382,<br>866 | 190,386,58<br>9 | SBP-box<br>transcription<br>factor | Chuck et al.<br>2014                                      |
|               |   |                |                 | GRMZM2G104925  | 4 | 133,662,<br>510 | 133,664,99<br>8 | LRR<br>receptor-like<br>protein    | Taguchi-Shiob<br>ara et al., 2001                         |
| Sub-pop3_21JH | 4 | 75,627,56<br>9 | 151,339,53<br>6 | Zm00001d051012 | 4 | 136,765,<br>871 | 136,767,71<br>2 | LRR protein<br>kinase              | Taguchi-Shiob<br>ara et al<br>2001                        |
|               |   |                |                 | GRMZM2G104925  | 4 | 133,662,<br>510 | 133,664,99<br>8 | LRR<br>receptor-like<br>protein    | Taguchi-Shiob<br>ara et al., 2001                         |
| Sub-pop4_22YS | 4 | 46,403,89<br>5 | 172,368,60<br>9 | Zm00001d051012 | 4 | 136,765,<br>871 | 136,767,71<br>2 | LRR protein<br>kinase              | Taguchi-Shiob<br>ara et al 2001<br>Bommert et al<br>2013b |
| Sub-pop4_21YS | 5 | 76615734       | 181154647       | Zm00001d016075 | 5 | 142,503,<br>737 | 142,509,12<br>9 | Negatively<br>regulating KRN       | An YX et al<br>2022                                       |