```
LRR1(66-89)
              TRI
                     GHSGLQG.TLSPDLRNLSE.
LRR2(90-112)
              ER
                     OWNN
                             S
                                . PVP
                                     . S
                                          SGLAS.
LRR3(113-136)
              QV
                     SNNN
                            D
                                . I
                                  PSDVF
                                         QGLTS.
                     DNNP
LRR4 (137-161)
              QS
                            K
                              SWEIPESLRNASA.
LRR5(162-187)
              QN
                  S
                     NSAN
                             S
                                . SL
                                     . GFLGPDEFPG
              SI
LRR6(188-210)
                     AFNN
                            E
                                . ELPMSI
                  Н
                                         AGSQ.
                                     . VI
LRR7(211-233)
              QS
                  W
                     NGQK
                            T
                                         QNMTG.
                                . DIT
              ΚE
                     HSNK
                                     .DFSGLKE.
LRR8(234-256)
                  W
                             S
                                .PL
              ES
                  S
                     RDNS
                             Т
                                . PVPASI
                                         LSLES.
LRR9(257-280)
LRR10(281-301)
              ΚV
                  N
                     TNNH
                                     . VFKSS.
                             Q
                                . PVP
LRR11(363-387)
              TV
                  S
                     EKME
                             T
                                . TISPEFGAIKS
LRR12(388-411)
              QRIII
                     GINNL
                            TG
                               .MIPQELTTLPN.
LRR13(412-432)
             LKTLDVSSNKLF
                                     . GFRSN.
```

## Supplementary information, Figure S3

Sequence alignment of LRRs in TMK1

The boundary of each LRR is shown on its right. The conserved residues are shown with yellow background. The solid pink squares indicate the amino acids specific to plant LRR proteins.