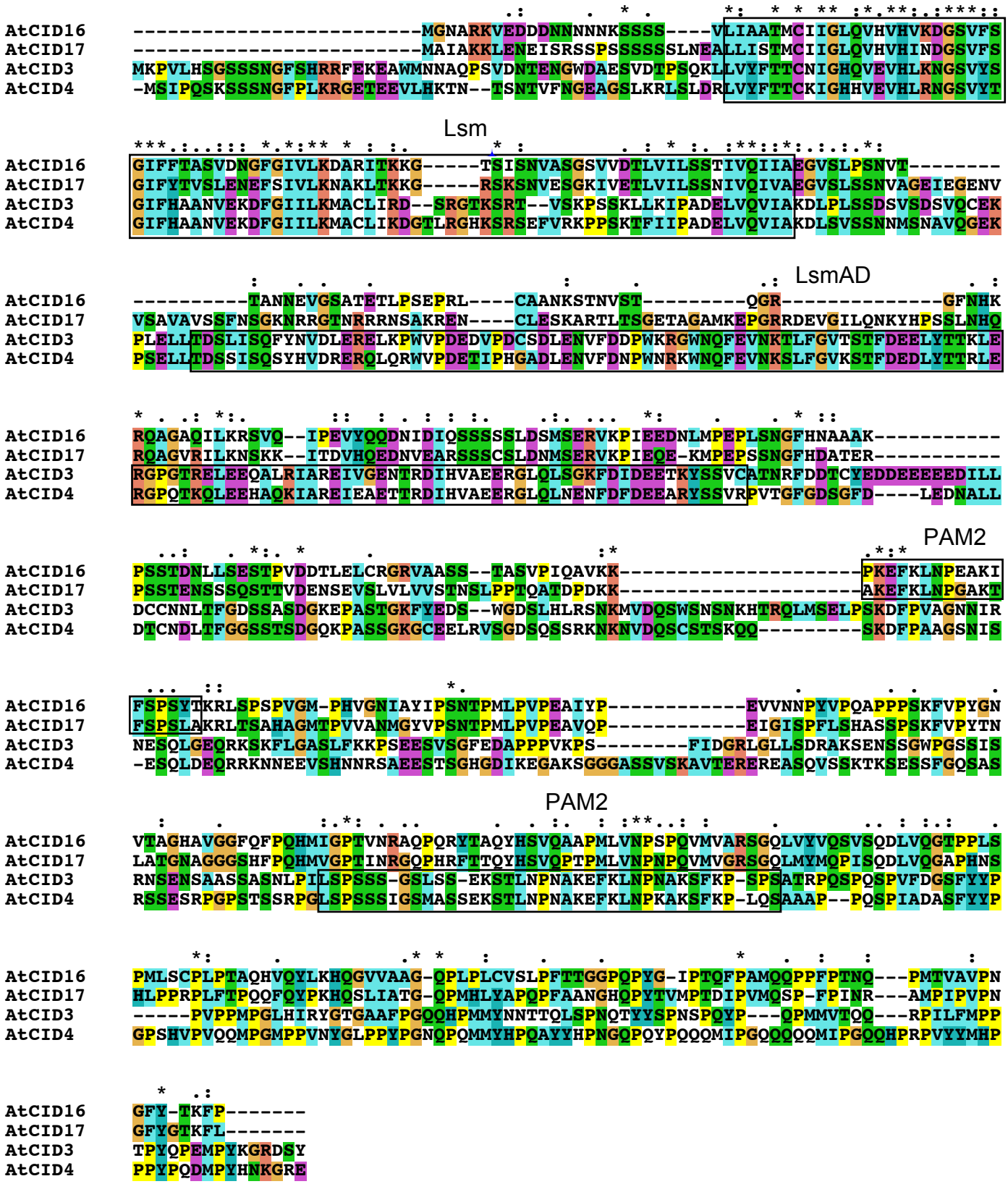

Additional file 2. Protein sequence alignment of the CID3, CID4, CID16 and CID17. The sequence alignments were performed using ClustalX 2.0.12; default colors were used. Regions encompassing sequence Lsm, LsmAD and PAM2 motifs are enclosed by rectangles.