Figure S6. Alignment of the amino acid sequences of TaPDIL4D, soybean GmPDIS-1, and soybean GmPDIS-2. The active center CGHC motifs (shaded in black), conserve arginine (box with straight lines), and glutamic acid (boxes with dotted lines) are indicated. Asterisks indicate positions that have a single, fully conserved residue. Colon indicates conservation between groups of strongly similar properties. Period indicates conservation between groups of weakly similar properties.