

TaPDIL4D	31	DEVLALTEST	FEKEVGQDRG	ALM ^E FYAPWC	GHCKKLAPEY	EKLAASFKKA	KSVLIAKVDC	90	
GmPDIS-1	28	DDVVVLSEDN	FEKEVGQDRG	ALM ^E FYAPWC	GHCKKLAPEY	EKLGSFKKA	KSVLIGKVDC	87	
GmPDIS-2	27	DDVVALTEET	FENEVGKDRA	ALM ^E FYAPWC	GHCKRRLAPEY	EQLGASFKKT	KSVLIAKVDC	86	
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TaPDIL4D	91	DEHKSVCSKY	GVSGYPTIOW	FPKGSLEPKK	YEGERTAEAL	TEYVNSEAAT	NVKIAAVPSS	150	
GmPDIS-1	88	DEHKSLCSKY	GVSGYPTIOW	FPKGSLEPKK	YEGERTAESL	VEFVNTEGGT	NVKIATVPSN	147	
GmPDIS-2	87	DEHKSVCGKY	GVSGYPTIOW	FPKGSLEPKK	YEGERTAEAL	AAFVNIEAGT	NVKIASVASS	146	
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TaPDIL4-1	151	VVVLTEETFD	SVVLDETKDV	L ^M ^E FYAPWCG	HOKSLAPIYE	KVASVFKQDE	GVVIANLDAD	210	
GmPDIS-1	148	VVVLTPENFN	EVVLDETKDV	L ^M ^E FYAPWCG	HOKSLAPTYE	KVATAFKLEE	DVVIANLDAD	207	
GmPDIS-2	147	VVVLSPNNFD	EVVFDETKDV	L ^M ^E FYAPWCG	HOKALAPIYE	KVAAAFNLDK	DVVIANVDAD	206	
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TaPDIL4D	211	KYTSLAEKYG	VSGFPTLKFF	PKGNKAGEEY	ESGEELDDFV	KFINEKSGTS	RDSKGQLTSE	270	
GmPDIS-1	208	KYRDLAEKYD	VSGFPTLKFF	PKGNKAGEDY	GGGRDLDDFV	AFINEKSGAS	RDGKGQLTSQ	267	
GmPDIS-2	207	KYKDLAEKYG	VSGYPTLKFF	PKSNKAGENY	DGGEDLDDFV	AFINEKCGTY	RDGKGQLTSK	266	
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TaPDIL4D	271	AGLVASLDAL	VKEFHSAADD	KRKEILSKIE	EEAAKLSGPA	VKHGKIYVNV	AKKILQKGSD	330	
GmPDIS-1	268	AGIVESLDVL	VKEFVAASDE	EKKSVFTRLE	EEVVVKLGSA	SRYGKIYLKA	AKNYREKGSD	327	
GmPDIS-2	267	AGIIASLDL	VKEFVSADSN	EKKAVYSRLE	EEVKKLKGSS	ARHGDLYLKL	AKKGMEKGAD	326	
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TaPDIL4D	331	YTKKETERLH	RLLEKSISPS	KADEFAIKKN	ILSAFSS	367			
GmPDIS-1	328	YAKNEIQRLQ	RILDKSISPA	KADETLKKN	ILSTYAA	364			
GmPDIS-2	327	YAKNEIQRL	RMLEKSISPA	KADEFTLKKN	ILSIFA	362			
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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.