Identification and Characterization of a Fatty Acidand Retinoid-Binding Protein Gene (*Ar-far-1*) from the Chrysanthemum Foliar Nematode, *Aphelenchoides ritzemabosi*

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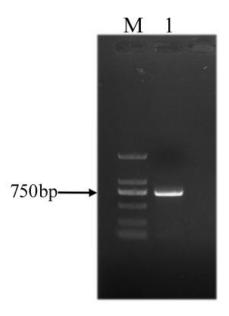


Figure S1. The cDNA sequence amplification results of fatty acid- and retinoid-binding gene (*Ar-far-* 1) of *Aphelenchoides ritzemabosi*.

Translation of DNAMAN1{RF+1}(31-574) Universal code Total amino acid number: 180, MW=20369 Max ORF starts at AA pos 1(may be DNA pos 31) for 180 AA(540 bases), MW=20369																			
1 1	AACO	CCA	AGT	TTG	AGC	AAC	СТС	CAAC	AAA	ACC	 	CTT L					TCT S		
61 11	GCG(A										 	TTG. L							
121 31	CAAC Q											TTC F							
181 51	GACA D										 	GCC. A							
241 71	GCTA A											TTT. F							
301 91	ACT(T											GAT D							
361 111	ATCA I										 	'GAA E							
421 131	ATT(I		GAA E					GTC V		GAC D	 	CAAA K		TTG L	GGC G	GAA E			AAA K
481 151	CAGI Q										 	GTT. V					AAA K		Q Q
541 171	ACT(T							CAAA K			 TAA	ACG	GTC	GTC	GAA	AAC	GGA	TAG	CTT

Figure S2. The amino acid sequence of fatty acid- and retinoid-binding gene (*Ar-far-1*) of *Aphelenchoides ritzemabosi*.

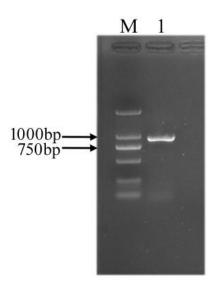


Figure S3. The DNA sequence amplification results of fatty acid- and retinoid-binding gene (*Ar-far-* 1) of *Aphelenchoides ritzemabosi*.

<pre>SEQ DNAMAN1: 947 bp; Composition 322 A; 178 C; 169 G; 278 T; 0 OTHER Percentage: 34.0% A; 18.8% C; 17.8% G; 29.4% T; 0.0%OTHER Molecular Weight (kDa): ssDNA: 292.34 dsDNA: 583.70 ORIGIN</pre>												
	1	AACCCAAGTT	TGAGCAACCT	CAACAAAACC	ATGAGTCTTC	GCACATTCGT	TTCTTTGTTC					
	61	GCGCTCTTCT	GTCTTAGCTA	TGCAGCTACT	CTTCCGTTGA	GCATTCATCA	AGTGCCCGAA					
	121	CAACTTAAAG	GTAAAAATCG	TGATTACCAT	GTTTTTGATT	GTTTTAAATT	TTAGAAGTTG					
	181	TGCCCGAGGA	AGTGAAAAAA	TTCTATGCTG	ACCTCACTGA	AGAAGACAAG	ACCATCTTAA					
	241	AAGAGGTTGC	TGCTAACCAC	GCCAGCTACG	AAAACGAAGA	TCAAGCTATG	GAAGCTCTCA					
	301	AGGCAAAGAG	CGAAAAACTG	TTTAACAAAG	CTACCGAACT	TCGTACTCTC	TTGAAAACCA					
	361	AAATTGACTC	GTTGAAACCC	GATGCGAAAG	CATTTGTTGA	AGGTGTAAGT	TTAAGTCAAT					
	421	TTTGCACGTA	CATAATTTTA	AGAATATTAG	CTAGCGTGCA	AAGCCAATTC	AAACTATTAA					
	481	TTTTCAGATC	ATCAACAAGG	TTCGCGCCTT	GAAACCTAAG	GGTGAAGAGA	AGCCTGACTT					
	541	AAAGAAGGTA	TTCAATTTTG	CATTAAACTT	ATTTTATTAA	AAGAAACTTT	TTTAGATTCG					
	601	TGAAGTTGCT	AACGAAGTCA	TCGACTCCTA	CAAAGCTTTG	GGCGAAGAAT	CTAAACAGAA					
	661	CTTGCAAGAA	ACGTTCCCAC	AAATCACGAA	CGTTATTAAG	AGTTAGTTTG	CATTCTTATA					
	721	AAGCTTTCAA	ATTTTCGATT	TTTTAGACGA	GAAATTCCAA	ACTCTCGCTC	AAGGTTTGAT					
	781	CAAACAAGAA	AATTAAACGG	TCGTCGAAAA	CGGATAGCTT	CGACTAATTA	TGTCAACAAT					
	841	GACCATAGTG	ATTTTTGTAA	TAATGTGTGT	TAAAATTTCG	GATCCTTATA	GTTGTGCTCG					
	901	TTTGTTCGAA	TAATGTCAGA	TCGAAAAAAC	ACGAAATAAA	CTTCATA						

Figure S4. The DNA sequence of fatty acid- and retinoid-binding gene (*Ar-far-1*) of *Aphelenchoides ritzemabosi*. The shaded parts were the locations of the four introns. The arrows and box were the positions of the primers of the southern-blot probe and the restriction enzymes Dra I, respectively.

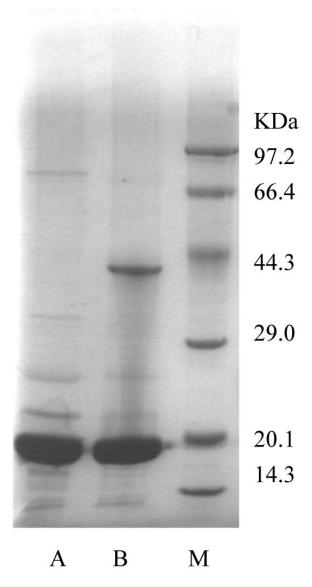


Figure S5. Sodium dodecyl sulphate polyacrylamide gel (SDS-PAGE) of *Escherichia coli* cells expressing fatty acid- and retinoid-binding protein (*Ar*-FAR-1) and its purified protein. M: Protein Marker, A: reduced protein, B: non-reduced protein.



Figure S6. The map of the recombinant vector (pET-32a-Ar-far-1). .