

Supplementary Data

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Expression, purification and DNA-binding properties of zinc finger domains of DOF proteins from *Arabidopsis thaliana*

Hakimeh Moghaddas Sani^{1,2}, Maryam Hamzeh-Mivehroud^{2,3}, Ana P. Silva⁴, James L. Walshe⁴, S. Abolghasem Mohammadi⁵, Mahdyieh Rahbar-Shahrouzias², Milad Abbasi², Omid Jamshidi², Jason KK Low⁴, Siavoush Dastmalchi^{2,3,6*}, Joel P. Mackay⁴

¹Faculty of Advanced Medical Sciences, Tabriz University of Medical Sciences, Tabriz, Iran

²Biotechnology Research Center, Tabriz University of Medical Sciences, Tabriz, Iran

³School of Pharmacy, Tabriz University of Medical Sciences, Tabriz, Iran

⁴School of Life and Environmental Sciences, The University of Sydney, NSW 2006, Australia

⁵School of Agriculture, University of Tabriz, Tabriz, Iran

⁶Faculty of Pharmacy, Near East University, POBOX:99138, Nicosia, North Cyprus, Mersin 10, Turkey

	DOF domain
Arabidopsis-AtDOF3.4 (Q39088)	QQEQLPCPRCDSSNTKFCYNNYNSQPRHFCKACRRYWTHGGTLRDVPVGGGTRK----96
Arabidopsis-AtDOF5.8 (Q9FGD6)	QQEQLSCPRCESTNTKFCYNNYNSQPRHFCKSCRRYWTHGGTLRDIIPVGGVSRK----97
Arabidopsis-AtDOF2.1 (Q8LE43)	GGEAQNCPRCESPTNKFCYNNYNSLQPRYFCKSCRRYWTGGTLRNVPVGGGCRK---113
Maize-MNB1A (P38564)	VGEGDPCPRCASRDTKFCYNNYNTSQPRHFCKGCRRYWTGGTLRNVPVGGGTRK---125
Maize-PBF (O24463)	LPEALKCPRCDSSNTKFCYNNYSMSQPRYFCKACRRYWTHGGTLRNVPVGGGCRK---95
Pigeonpea-CcDOF (A0A0K8K646)	PQQQQKCPRCDSMNTKFCYNNYNSLQPRHFCKTCKRYWTHGGTFRNIPVGGGSRK---102
Pigeonpea-CcDOF (A0A0K8K636)	PQQPQKCPRCDSFNTKFCYNNYNSLQPRYFCKTCRRYWTGGTLRNVPVGGGCRK---94
Pigeonpea-CcDOF (A0A0K8K648)	NHQALKCPRCDSLNTKFCYNNYNSLQPRHFCKNCRRYWTGGVLRNVPVGGGCRK---106
Sorghum-SbDOF27 (R4NG43)	EPEGLPCPRCESVNTKFCYNNYNSLQPRYFCKTCRRYWTRGGALRNVPVGGNTRK---104
Sorghum-SbDOF24 (E6PBM6)	LAEALRCPRCNSNTKFCYNNYNTMQPRYFCKGCRRYWTHGGTLRNVPVGGGCRK---163
Sorghum-SbDOF23 (C5WWR6)	PDKILPCPRCNSMDTKFCYNNYNTINQPRHFCKNCQRYWTAGGAMRNVPVGGGRK---107
Sorghum-SbDOF19 (E6PBM6)	LAEALRCPRCNSNTKFCYNNYNTSQPRYFCKGCRRYWTHGGALRNVPVGGGCRK---104
Tobacco-NaDOF2.5 (A0A1J6IT59)	KDQAINCPRCNSNTKFCYNNYNSLTQPRYFCKTCRRYWTEGGTLRNVPVGGGSRK---121
Tobacco-NaDOF5.6 (A0A1J6IT68)	HDQPLKCPRCDSHTKFCYNNYNSLTQPRYFCKSCRRYWTGGTLRNIPVGGGCRK---93
Tobacco-NaDOF1.6 (A0A1J6IGB2)	EPENLPCPRCDSTNTKFCYNNYNSLQPRHFCKSCRRYWTRGGTLRNIPVGGGTRK---87
Tobacco-NaDOF1.7 (A0A1J6K6P9)	EQEILKCPRCDSINTKFCYNNYNSLQPRHFCKNCKRYWTGGTLRNIPVGGASRK---106
Anthurium-AaDOF2.1 (A0A1D1YVB2)	PEEALRCPRCDSTNTKFCYNNYNSLQPRYFCKGCRRYWTGGTLRNVPVGGGCRK---187
Anthurium-AaDOF4.6 (A0A1D1XWV9)	REQSLGCPRCHSTNTKFCYNNYNSLTQPRYFCKTCRRYWTEGGTLRNVPVGGGRKA---147
Barley-HvDOF2 (A5HWF1)	KEKAINCPRCNSNTKFCYNNYNSLQPRYFCKTCRRYWTEGGSLRNVPVGGGSRK---102
Barley-HvDOF10 (A5HWF9)	VEVAPNCPRCDSPNTKFCYNNYNSLQPRYFCKGCRRYWTGGSLRNVPVGGGCRK---164
Barley-HvDOF17 (A5HW93)	AAAPLPCPRCSRSTKFCYNNYNSVQPRHFCKACHRYWTAGGALRNVPVGGGRK---207
Barley-HvDOF1 (A5HWF0)	PEPGLKCPRCESNTKFCYNNYNSLTQPFHFCRTTCRRYWTRGGTLRNVPVGGGCRK---121
Apple-MdDOF (D9ZIX8)	HEQALKCPRCDSTHTKFCYNNYNSLQPRYFCKTCRRYWTGGTLRNIPVGGGCRK---108
Apple-MdDOF (D9ZIX6)	PEQALKCPRCDSTNTKFCYNNYNSLQPRYFCKSCRRYWTGGTLRNVPVGGGCRK---88
Apple-MdDOF (D9ZII6)	KDQALNCPRCNSNTTKFCYNNYNSLQPRYFCKTCKRYWTEGGSLRNVPVGGGSRK---133

Figure S1. Sequence alignment of selected DOF-ZF domains. A) Alignment of the DOF-ZF domains of *A. thaliana* DOF3.4, DOF5.8, and DOF2.1 with a number of DOF domains from other species shows high sequence similarity across the DOF family. The UniProt ID of each protein sequence is given inside the bracket.