

Table S6. The predicted cis-acting elements in the promoter region of *NtZF-HD* genes

Gene name	CAT-box	ABRE	ERE	TCA-element	CGTCA-motif	MBS	TC-rich repeats	LTR	WUN-motif	ARE
<i>NtZF-HD1</i>	2	2	0	1	1	0	0	0	0	1
<i>NtZF-HD2</i>	0	3	1	0	1	0	0	1	0	2
<i>NtZF-HD3</i>	0	6	0	0	0	2	0	0	1	0
<i>NtZF-HD4</i>	0	7	0	0	1	1	0	2	1	1
<i>NtZF-HD5</i>	0	6	1	4	3	0	0	2	2	3
<i>NtZF-HD6</i>	0	2	2	0	0	0	3	0	0	0
<i>NtZF-HD7</i>	0	0	1	0	0	0	0	0	1	0
<i>NtZF-HD8</i>	0	6	0	0	1	2	0	0	0	1
<i>NtZF-HD9</i>	0	3	5	0	2	0	2	0	1	1
<i>NtZF-HD10</i>	1	2	1	1	0	3	0	0	3	4
<i>NtZF-HD11</i>	1	6	1	1	1	0	1	0	1	2
<i>NtZF-HD12</i>	0	0	0	1	1	0	1	0	2	2
<i>NtZF-HD13</i>	0	7	1	0	1	0	0	0	0	1
<i>NtZF-HD14</i>	0	0	1	0	0	0	0	0	0	0
<i>NtZF-HD15</i>	0	1	1	0	2	0	1	1	0	1
<i>NtZF-HD16</i>	0	3	1	2	1	0	0	1	0	3
<i>NtZF-HD17</i>	0	0	0	1	0	0	2		1	0
<i>NtZF-HD18</i>	0	1	0	1	1	1	0	1	1	0
<i>NtZF-HD19</i>	0	2	0	1	3	1	0	1	0	1
<i>NtZF-HD20</i>	0	10	8	1	2	2	0		0	0
<i>NtZF-HD21</i>	0	1	1	1	1	0	0	1	3	3
<i>NtZF-HD22</i>	0	2	2	1	1	1	1	0	0	1

<i>NtZF-HD23</i>	0	0	1	0	0	0	2	0	1	1
<i>NtZF-HD24</i>	1	4	2	0	0	0	0	1	2	0
<i>NtZF-HD25</i>	0	1	1	0	0	1	0	0	0	3
<i>NtZF-HD26</i>	1	2	0	1	1	1	1	0	1	1
<i>NtZF-HD27</i>	0	2	1	1	0	1	1	1	1	2
<i>NtZF-HD28</i>	0	0	2	0	0	0	1	0	0	3
<i>NtZF-HD29</i>	0	0	0	0	0	0	0	0	0	0
<i>NtZF-HD30</i>	0	2	1	0	2	2	0	2	1	1
<i>NtZF-HD31</i>	1	8	4	1	1	0	0	1	1	0
<i>NtZF-HD32</i>	1	1	0	1	1	0	0	0	0	3
