

Additional file 3. Estimates of Evolutionary Divergence between among NAC homologs. Analyses were conducted using the Poisson correction model. All positions containing gaps and missing data were eliminated. There were a total of 241 positions in the final dataset. Evolutionary analyses were conducted in MEGA6 .

	<i>Ec</i> NAC67 <i>Trichy1</i>	<i>E. coracana</i> NAC1	<i>S. italica</i> NAC67	<i>S. bicolor</i> SNAC1	<i>O. rufipogon</i> SNAC1	<i>O. sativa</i> SNAC1	<i>Z. mays</i> NAC67	<i>O. sativa</i> NAC67	<i>O. nivara</i> SNAC1	<i>B. distachyon</i> NAC67	<i>O. sativa</i> NAC20	<i>O. brachyantha</i> NAC67
<i>E. coracana</i> NAC1	0.297											
<i>S. italica</i> NAC67	0.096	0.303										
<i>S. bicolor</i> SNAC1	0.091	0.326	0.047									
<i>O. rufipogon</i> SNAC1	0.124	0.314	0.100	0.082								
<i>O. sativa</i> SNAC1	0.119	0.314	0.096	0.078	0.008							
<i>Z. mays</i> NAC67	0.114	0.320	0.064	0.042	0.100	0.096						
<i>O. sativa</i> NAC67	0.119	0.314	0.096	0.078	0.008	0.000	0.096					
<i>O. nivara</i> SNAC1	0.119	0.314	0.096	0.078	0.008	0.000	0.096	0.000				
<i>B. distachyon</i> NAC67	0.177	0.320	0.207	0.197	0.202	0.197	0.222	0.197	0.197			
<i>O. sativa</i> NAC20	0.162	0.373	0.142	0.128	0.064	0.055	0.147	0.055	0.055	0.243		
<i>O. brachyantha</i> NAC67	0.119	0.320	0.091	0.078	0.021	0.013	0.091	0.013	0.013	0.202	0.064	
<i>T. aestivum</i> NAC2	0.181	0.320	0.197	0.177	0.177	0.172	0.186	0.172	0.172	0.119	0.233	0.177