


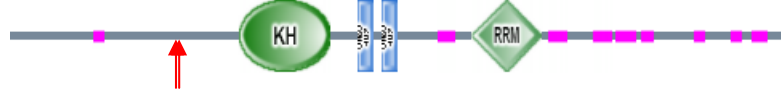


Supplemental Table 2. Phosphorylation of Arabidopsis RNA metabolism-related proteins

<p>At1g04080</p>	<p>Similar to yeast Prp39 splicing factor</p>	
<p>At4g17720</p>	<p>RRM-containing protein</p>	
<p>At5g04430</p>	<p>NOVA-like RNA-binding protein</p>	
<p>At5g51300</p>	<p>SF1 splicing factor-related</p>	

Phosphorylation sites are indicated by red arrows. Pink regions indicate areas of low compositional complexity. HAT, half-a-tetratricopeptide repeat; RRM, RNA recognition motif; KH, K homology RNA-binding domain; ZnF_C2HC, zinc finger. Protein structures are generated using the SMART database (<http://smart.embl-heidelberg.de/>).