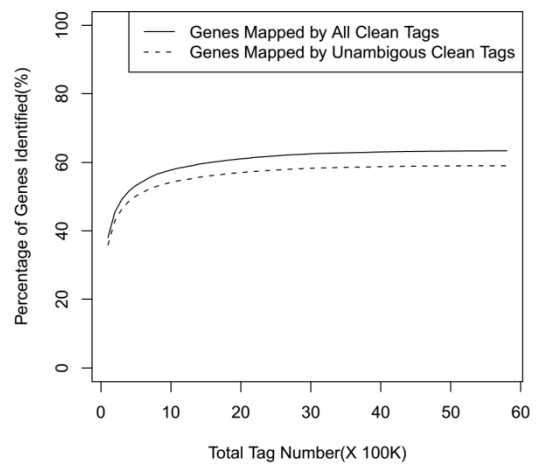
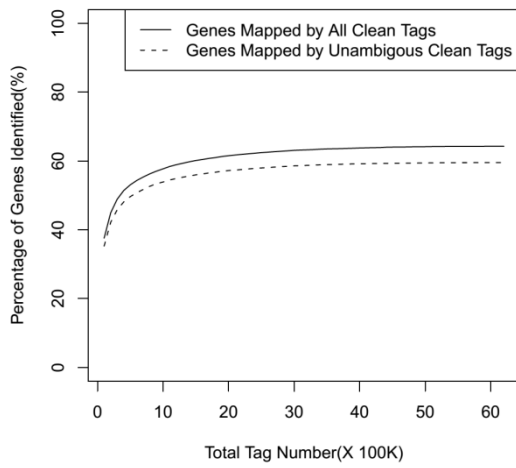


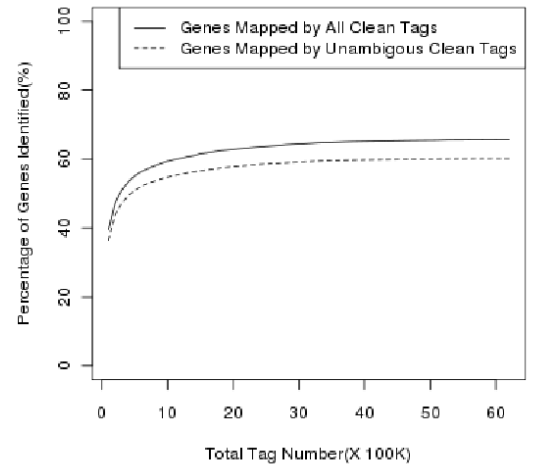
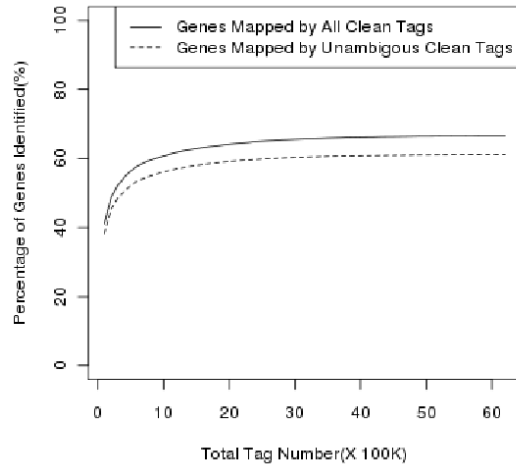
Table S1. Sequencing saturation analysis

Tissue	Control	Drought stress
Root - 1 d - replication 1		
Root - 1 d - replication 2		
Root - 4 d - replication 1		

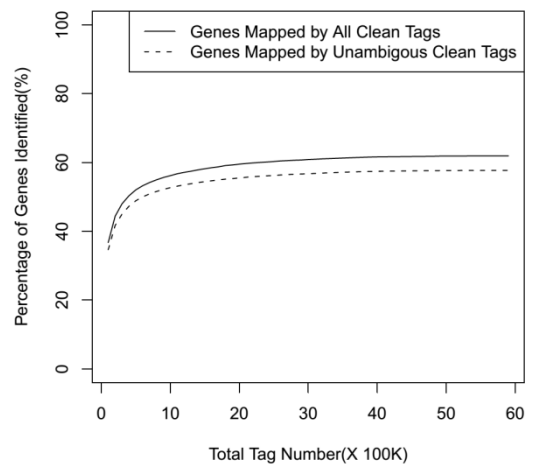
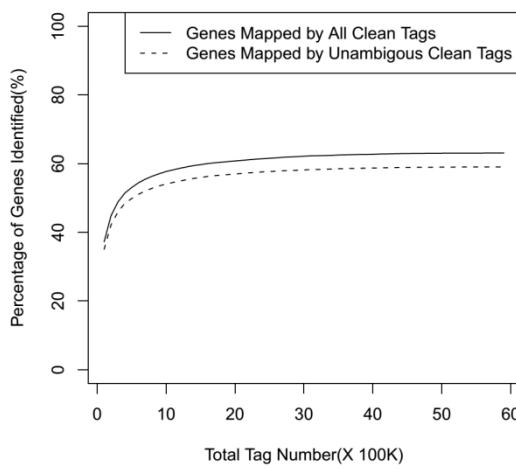
Root - 4 d -  
replication 2



Root - 7 d -  
replication 1



Root - 7 d -  
replication 2



<p>Leaf - 1 d - replication 1</p>		
<p>Leaf - 1 d - replication 2</p>		
<p>Leaf - 4 d - replication 1</p>		
<p>Leaf - 4 d - replication 2</p>		

	<p>Percentage of Genes Identified(%)</p> <p>Total Tag Number(X 100K)</p> <p>— Genes Mapped by All Clean Tags - - - Genes Mapped by Unambiguous Clean Tags</p>	<p>Percentage of Genes Identified(%)</p> <p>Total Tag Number(X 100K)</p> <p>— Genes Mapped by All Clean Tags - - - Genes Mapped by Unambiguous Clean Tags</p>
<p>Leaf - 7 d - replication 1</p>	<p>Percentage of Genes Identified(%)</p> <p>Total Tag Number(X 100K)</p> <p>— Genes Mapped by All Clean Tags - - - Genes Mapped by Unambiguous Clean Tags</p>	<p>Percentage of Genes Identified(%)</p> <p>Total Tag Number(X 100K)</p> <p>— Genes Mapped by All Clean Tags - - - Genes Mapped by Unambiguous Clean Tags</p>
<p>Leaf - 7 d - replication 2</p>	<p>Percentage of Genes Identified(%)</p> <p>Total Tag Number(X 100K)</p> <p>— Genes Mapped by All Clean Tags - - - Genes Mapped by Unambiguous Clean Tags</p>	<p>Percentage of Genes Identified(%)</p> <p>Total Tag Number(X 100K)</p> <p>— Genes Mapped by All Clean Tags - - - Genes Mapped by Unambiguous Clean Tags</p>