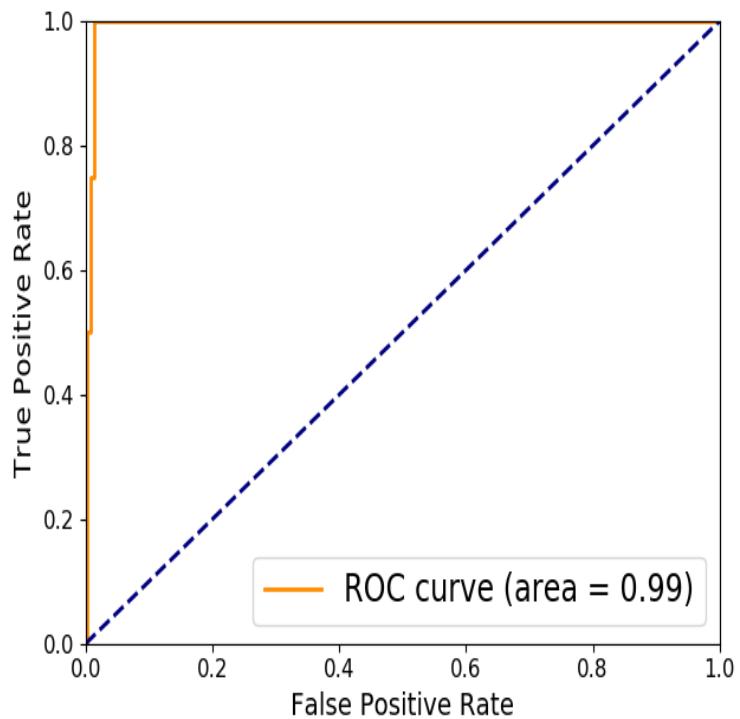
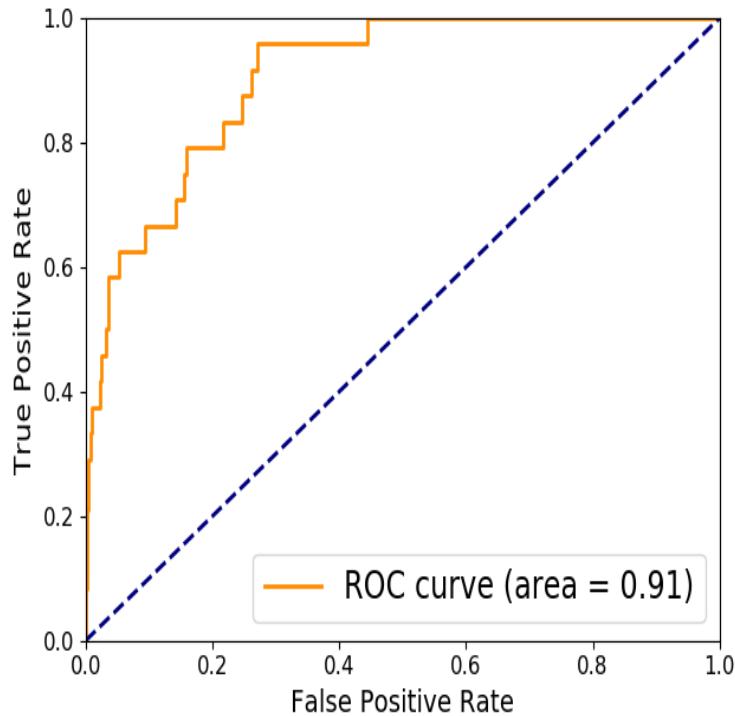


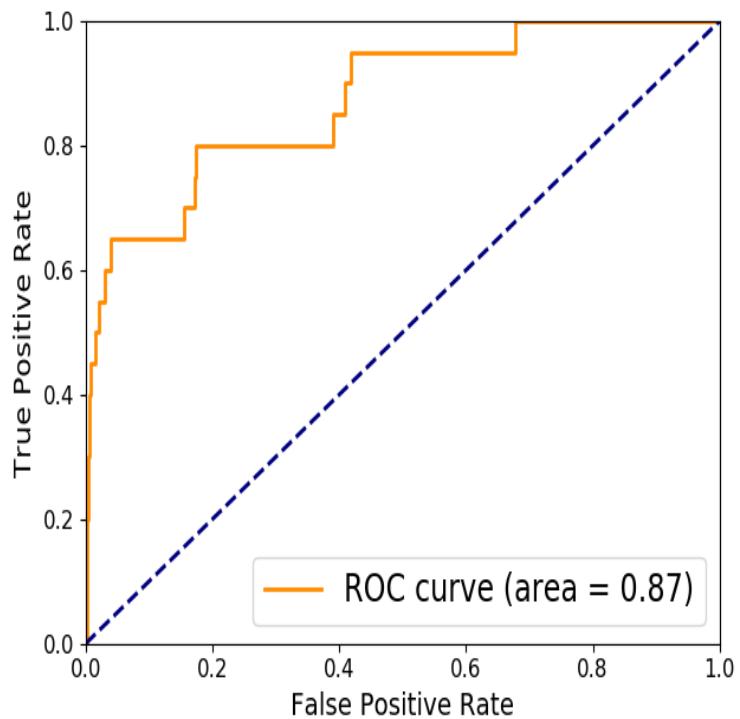
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



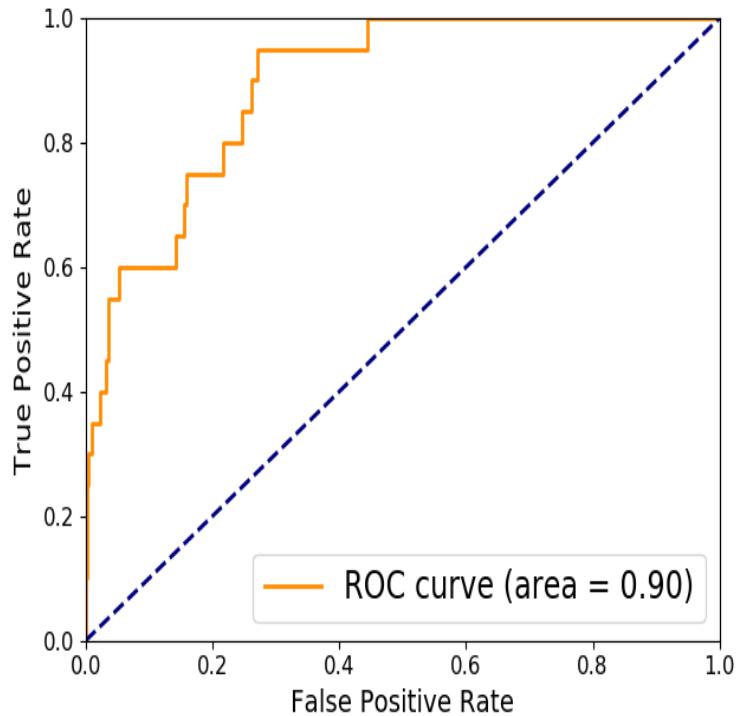
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



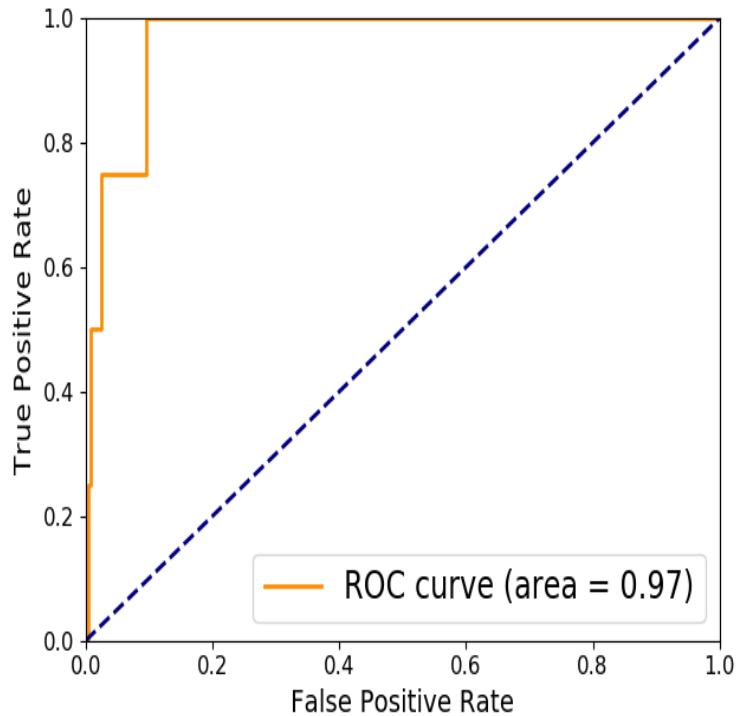
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



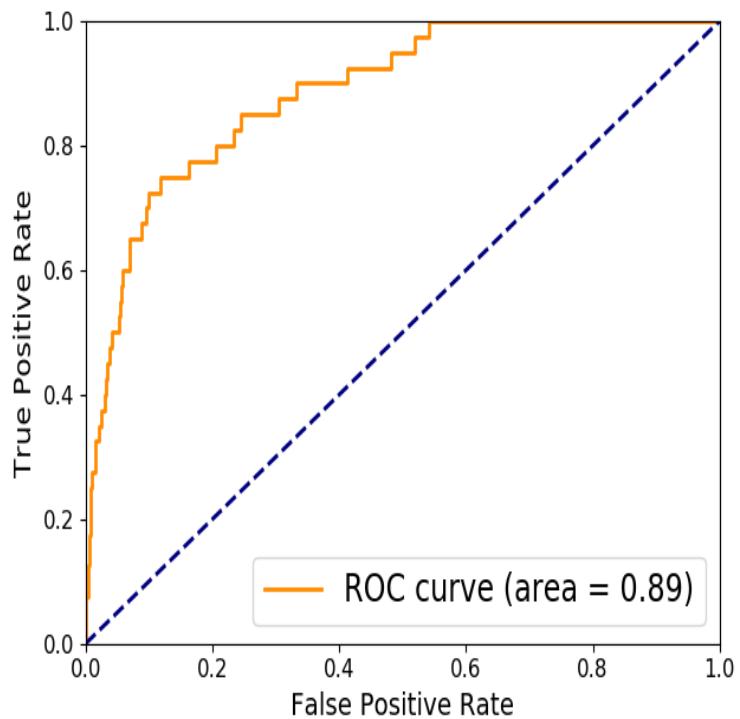
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



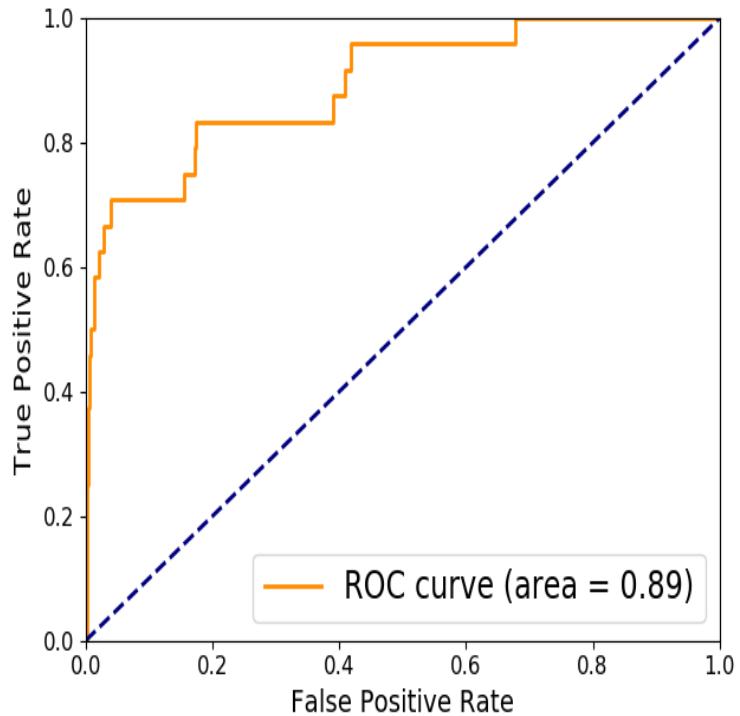
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



Diffusion ROC from fOverlap_tOnlies STRING10_combined.png



Diffusion ROC from fGSonly_tTest STRING10_combined.png



<i>Diff_FROM</i>	<i>Diff_TO</i>	<i>AUC</i>	<i>Zt_wdegree</i>	<i>Zt</i>	<i>Zf_wdegree</i>	<i>Zf</i>
<i>GS_only</i>	<i>Test_only</i>	0.873 5.034	5.768 3.559	2.594		
<i>Test_only</i>	<i>GS_only</i>	0.895 5.842	6.284 4.713	3.783		
<i>GS_only</i>	<i>Overlap</i>	0.993 3.253	3.209 3.234	3.476		
<i>GS_only</i>	<i>Test</i>	0.893 5.708	7.127 3.776	3.592		
<i>Test_only</i>	<i>Overlap</i>	0.967 3.466	3.018 3.456	3.461		
<i>Test_only</i>	<i>GS</i>	0.907 6.812	6.683 4.331	4.149		
<i>Overlap</i>	<i>Onlies</i>	0.887 6.909	8.871 3.221	2.503		

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>	
<i>GS</i>	20 21	<i>DISC2</i>	N/A	
<i>Test_only</i>	20 21	<i>GRK3</i>	N/A	
<i>Overlap</i>	4 4		<i>RELN;CACNA1C;ANK3;MTHFR</i>	

(end of file)