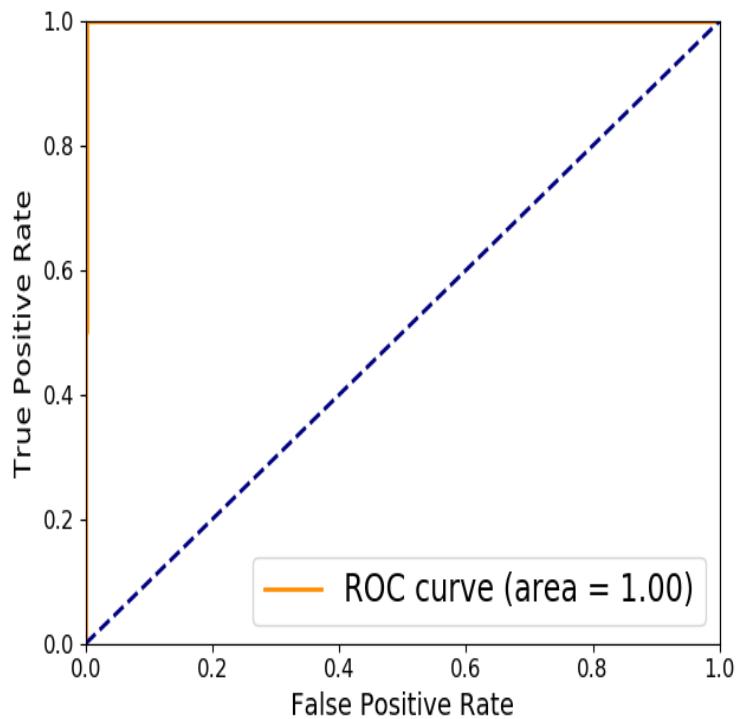
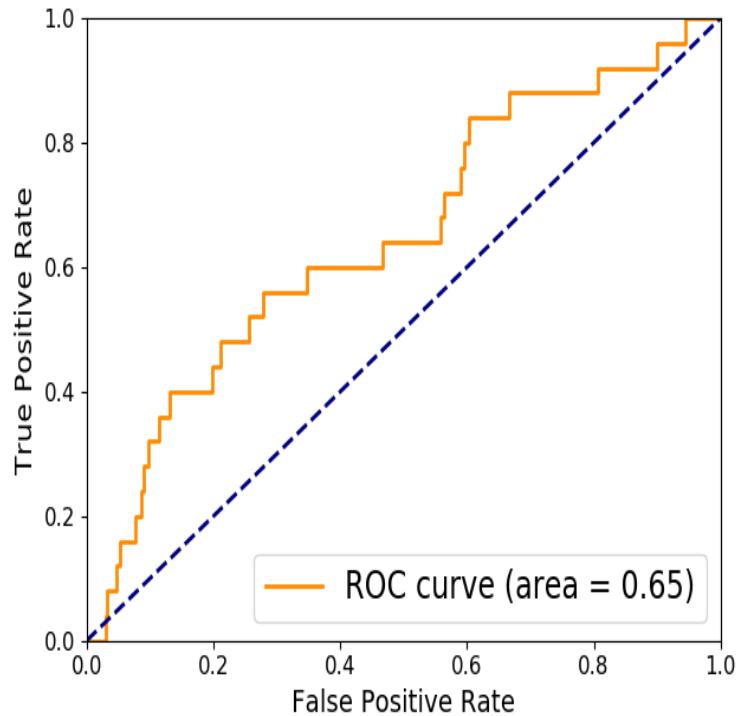


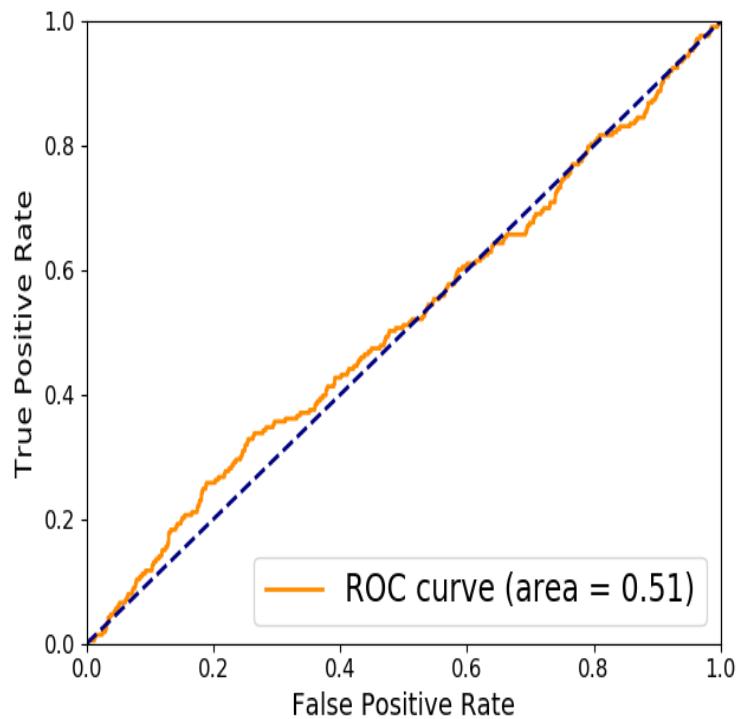
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



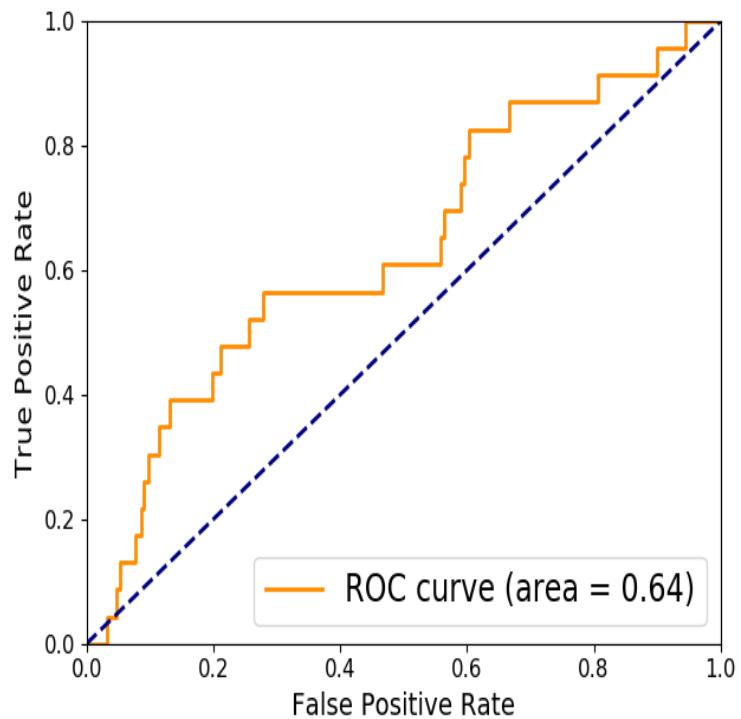
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



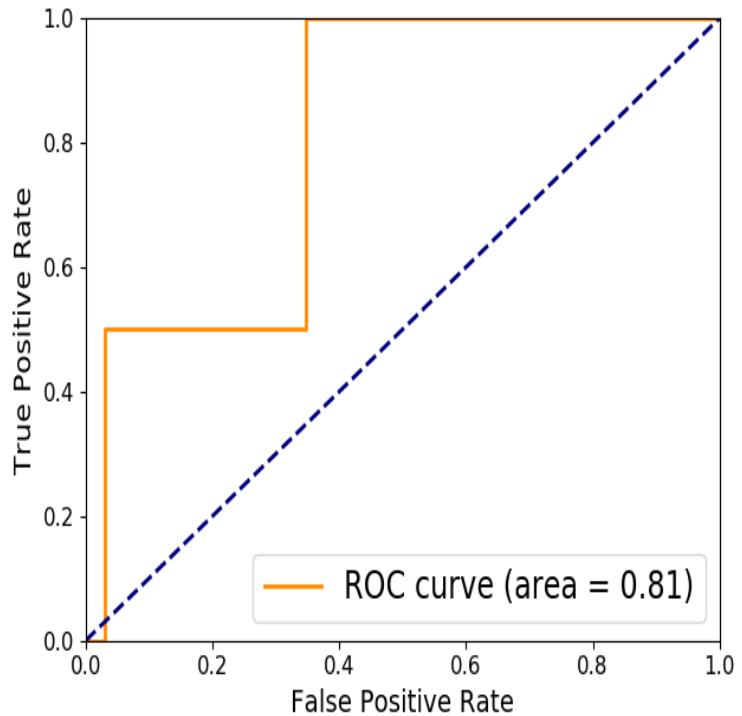
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



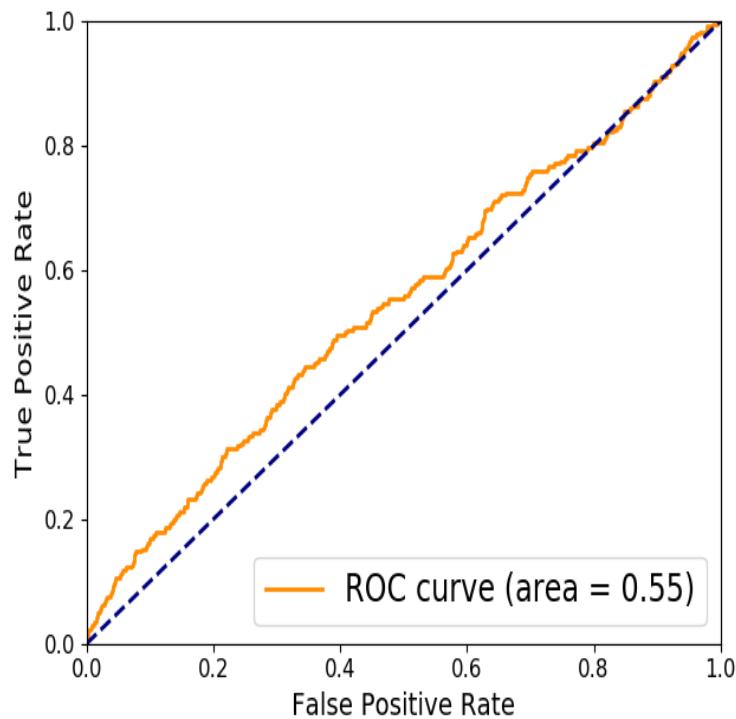
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



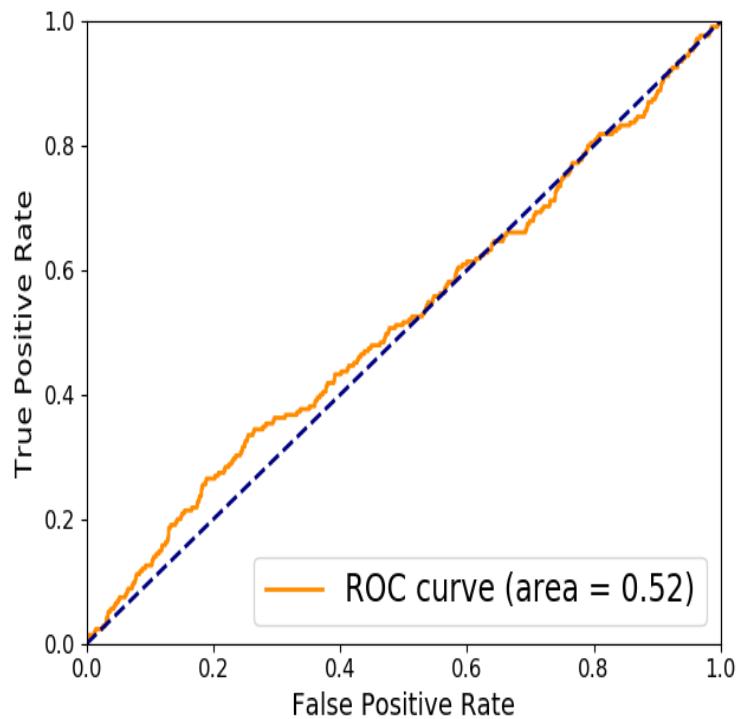
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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>	0.513	1.659	0.642	2.492	1.92281352	
/	/	/	/	/	68545605	/	
/	/	/	/	/	(invalid)	/	
<i>Test_only</i>	<i>GS_only</i>	0.636	1.917	2.251	1.706	1.052	
/	/	/	/	/	6283505	/	
/	/	/	(invalid)	/	/	(invalid)	/
<i>GS_only</i>	<i>Test</i>	0.517	1.674	0.971	2.910	2.002	
/	/	/	/	/			
<i>Test_only</i>	<i>Overlap</i>	0.811	0.80736794	1.527	0.44173518	0.278	
/	/	/	03236397	/	65223335	/	
/	/	/	(invalid)	/	(invalid)	/	
<i>Test_only</i>	<i>GS</i>	0.650	2.081	2.725	1.531	1.026	
/	/	/					
<i>Overlap</i>	<i>Onlies</i>	0.549	3.335	2.336	5.763	2.062	

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>		
<i>GS</i>	23	23		N/A	
/	/	/			
<i>Test_only</i>	213	218	OPLAH;MROH7;LENG9;LOC100653515;C5o	N/A	
/	/	/	rf45	/	
<i>Overlap</i>	2	2		APOE;CR1	

(end of file)