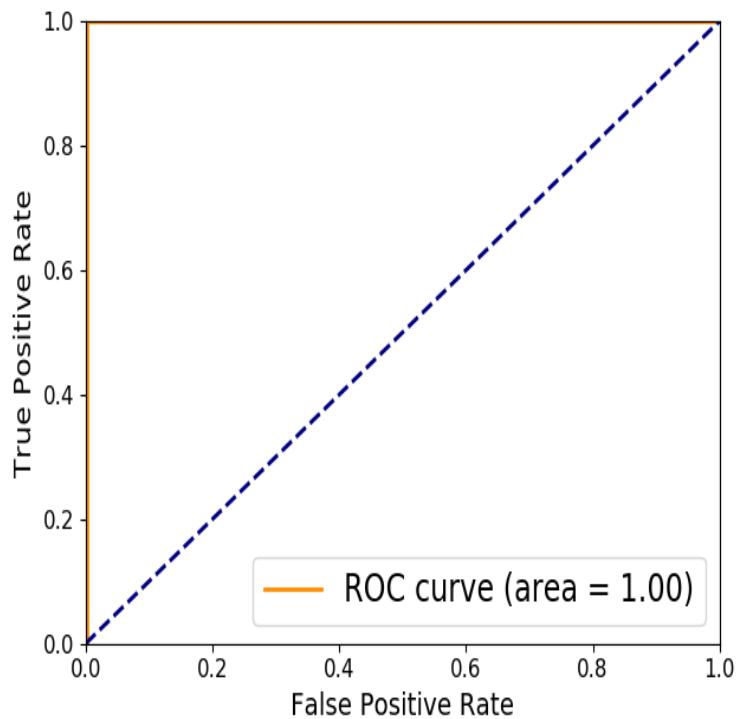
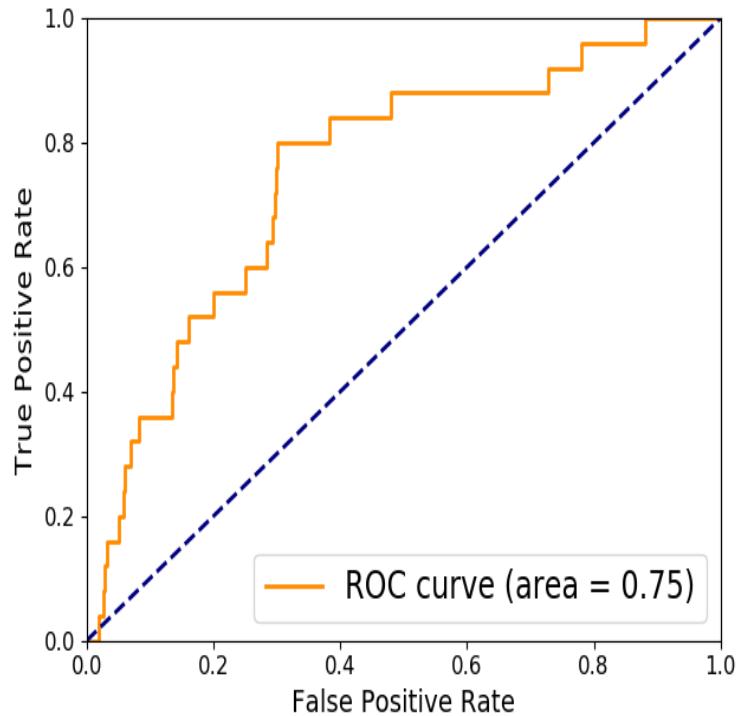


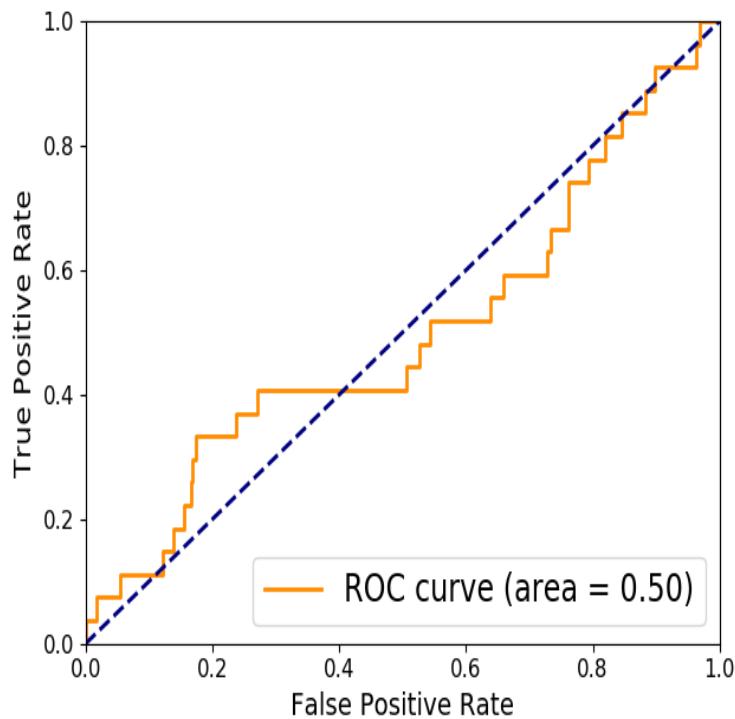
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



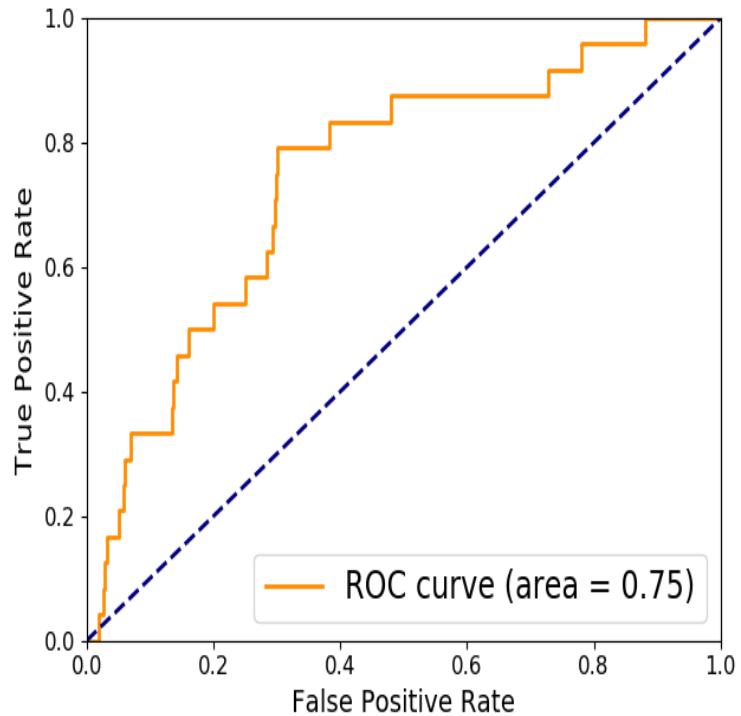
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



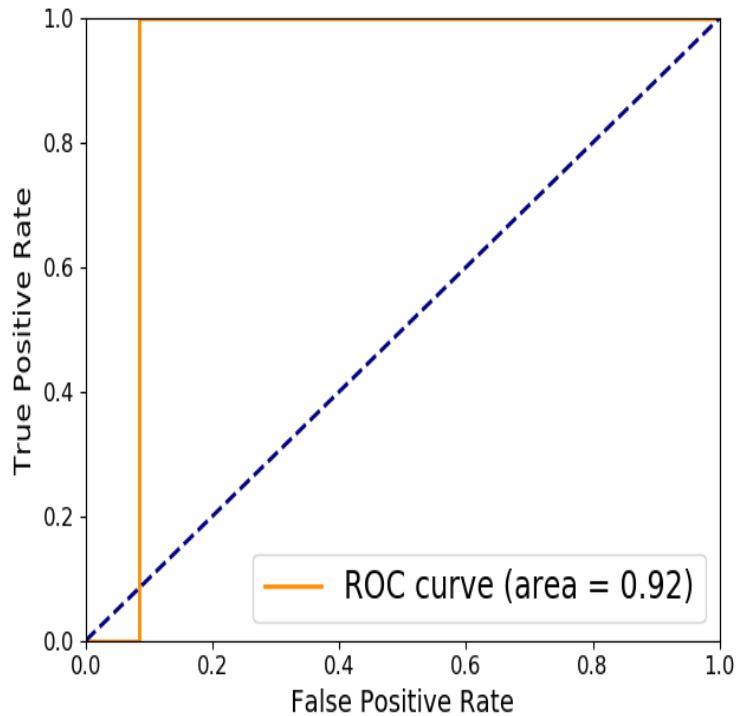
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



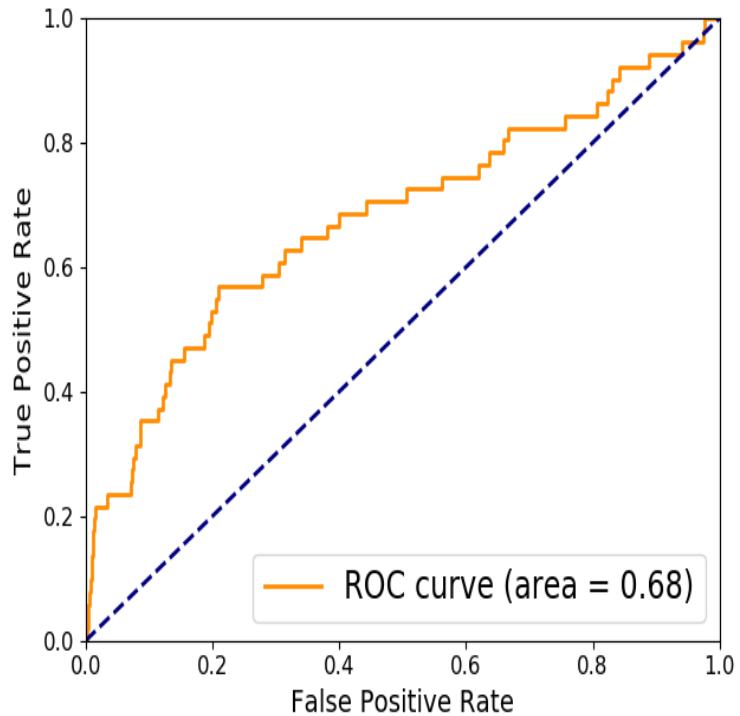
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



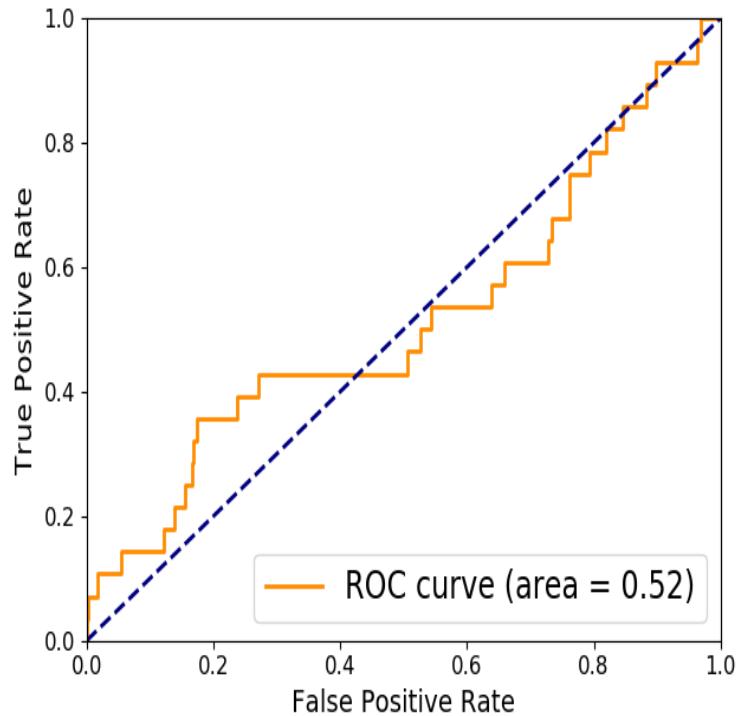
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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>
/ GS_only	/ Test_only	0.499	1.407	-0.141	1.320	1.420484
/ / / / / /				78573786		
/ / / / / /				66 (<i>inva</i>		
/ / / / / /				<i>lid</i>)		
/ Test_only	GS_only	0.746	3.483	3.657	2.594	2.648333
/ / / / / /				28543955		
/ / / / / /				5 (<i>invalid</i>		
/ / / / / /				<i>id</i>)		
/ GS_only	Overlap	0.998	5.3366543	1.6986292	1.960	1.829
/ / / /	41956066	69135178	/	/	/	
/ / / /	(<i>invalid</i>)	(<i>invalid</i>)	/	/	/	
/ GS_only	Test	0.517	1.481	0.296	1.545	1.814
/ / / /						
/ Test_only	Overlap	0.916	2.2185754	1.6658158	1.267	0.931151
/ / / /	877141384	978428739	/		55476200	
/ / / /	(<i>invalid</i>)	(<i>invalid</i>)	/	31 (<i>inva</i>		
/ / / /				<i>lid</i>)		
/ Test_only	GS	0.753	3.725	4.145	2.7684120	2.481
/ / / /			876262006			
/ / / /			(<i>invalid</i>)			
/ Overlap	Onlies	0.680	5.456	4.540	5.7566902	2.584
/ / / /			00701707			
/ / / /			(<i>invalid</i>)			

**	#Mapped	#Total	Not mapped genes	/
GS	24	24	/	N/A
Test_only	27	27	/	N/A
Overlap	1	1	/	APOE

(end of file)