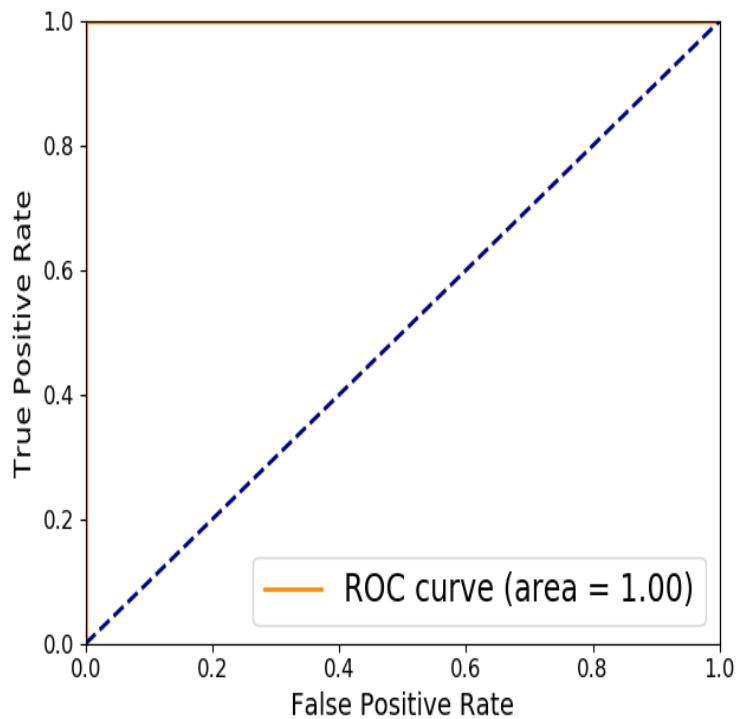
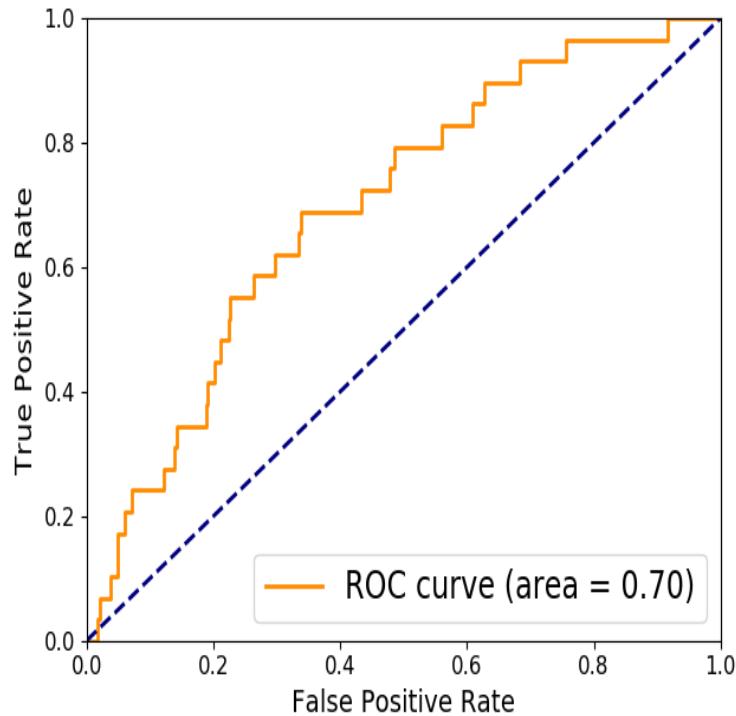


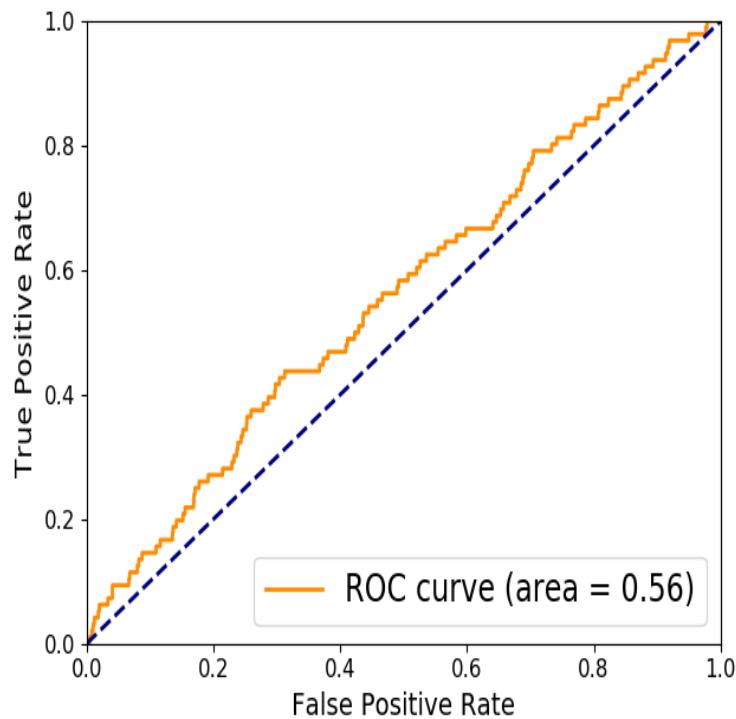
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



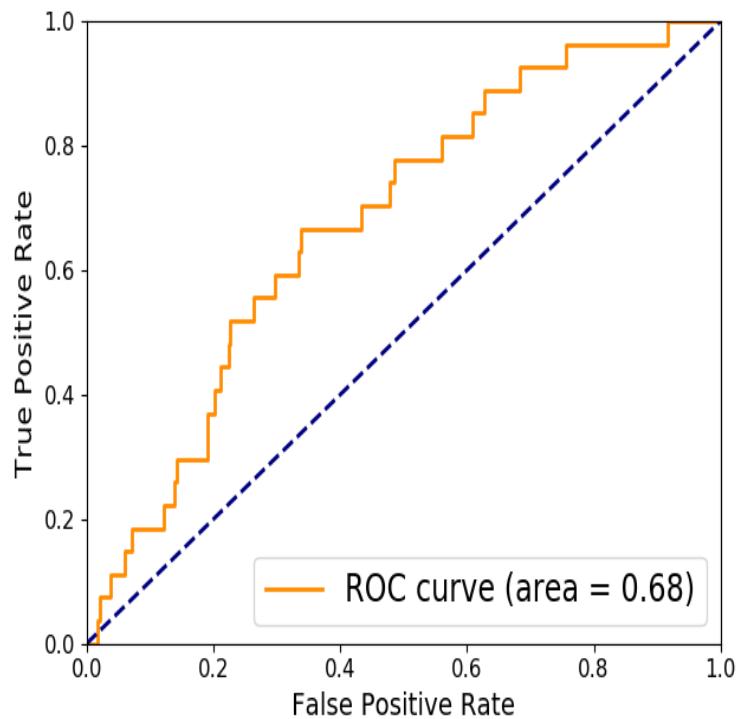
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



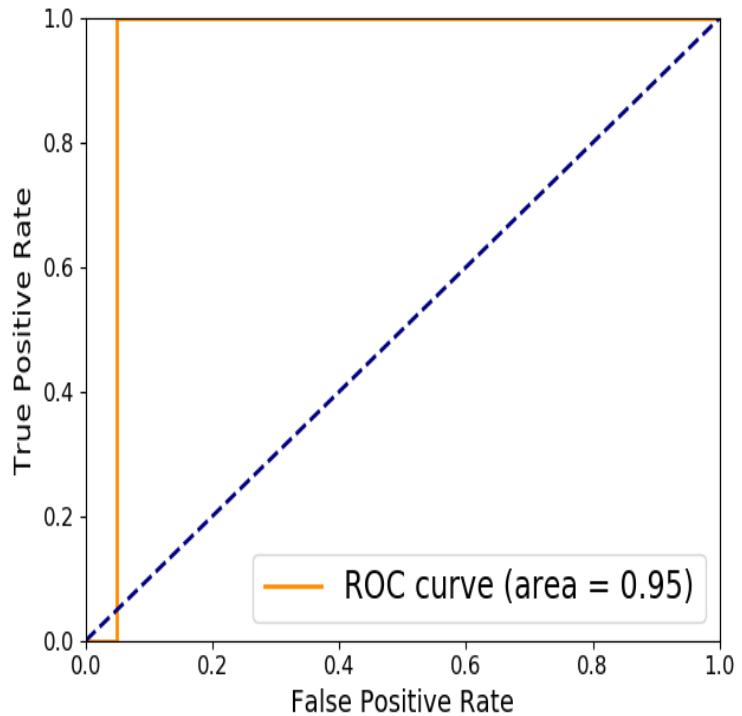
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



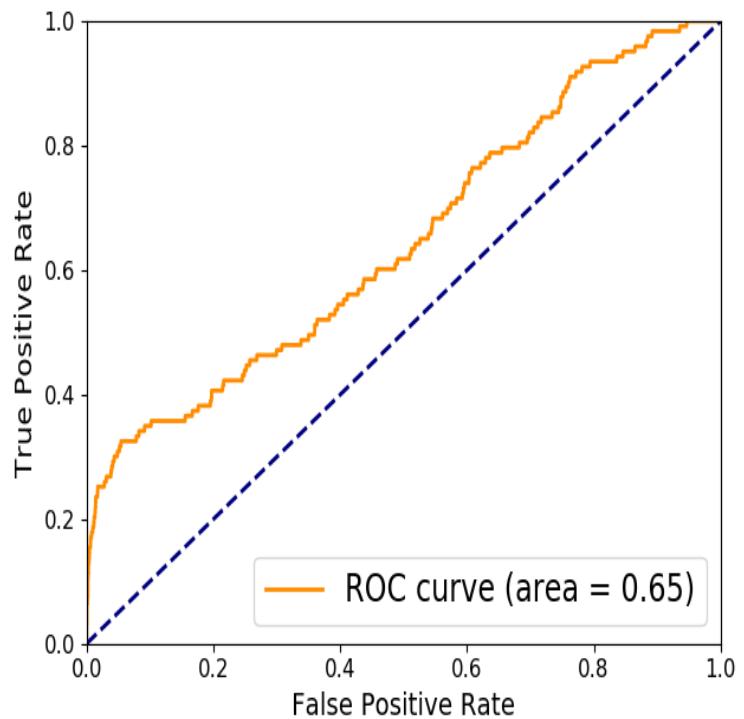
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



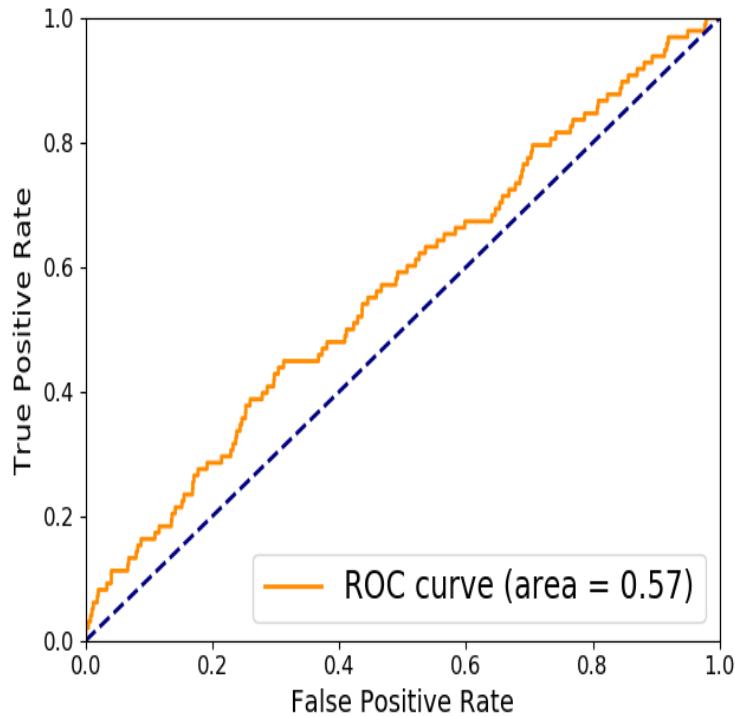
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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_wdegre</i>	<i>Zt</i>	<i>Zf_wdegre</i>	<i>Zf</i>
/	/	e	/	e	/	/
GS_only Test_only 0.561 1.036 2.192 2.430 1.453						
Test_only GS_only 0.680 1.443 3.4599925 1.027 0.844563						
/ / / 073473963 53783341						
/ / / (invalid) 75 (inva						
/ / / / lid)						
GS_only Overlap 1.000 2.248 2.484 2.5979752 2.825285						
/ / / / 01704758 88952983						
/ / / / (invalid) 43 (inva						
/ / / / lid)						
GS_only Test 0.570 1.380 2.315 2.286 1.588						
Test_only Overlap 0.951 2.788 2.213 2.2751013 1.991755						
/ / / / 338509756 88691141						
/ / / / (invalid) 27 (inva						
/ / / / lid)						
Test_only GS 0.699 2.169 3.418 1.0868096 1.500752						
/ / / / 720650664 23046853						
/ / / / (invalid) 77 (inva						
/ / / / lid)						
Overlap Onlies 0.646 4.0117140 6.357 2.9331974 2.699469						
/ / / 59217249 81283344 74310032						
/ / / (invalid) (invalid) 7 (inval						
/ / / / id)						

**	#Mapped	#Total	Not mapped genes		
GS 27 37 RBMS1/ITGB6;DGKB/TMEM195;CENT N/A					
/ / / D2;gene;KCNJ11/ABCC8;TSPAN8/L					
/ / / GR5;OASL/HNF1A;CDC123/CAMK1D;					
/ / / CDKN2A/B					
Test_only 96 98 CXorf22;MROH7 N/A					
Overlap 2 2 SLC30A8;HNF1B					

(end of file)