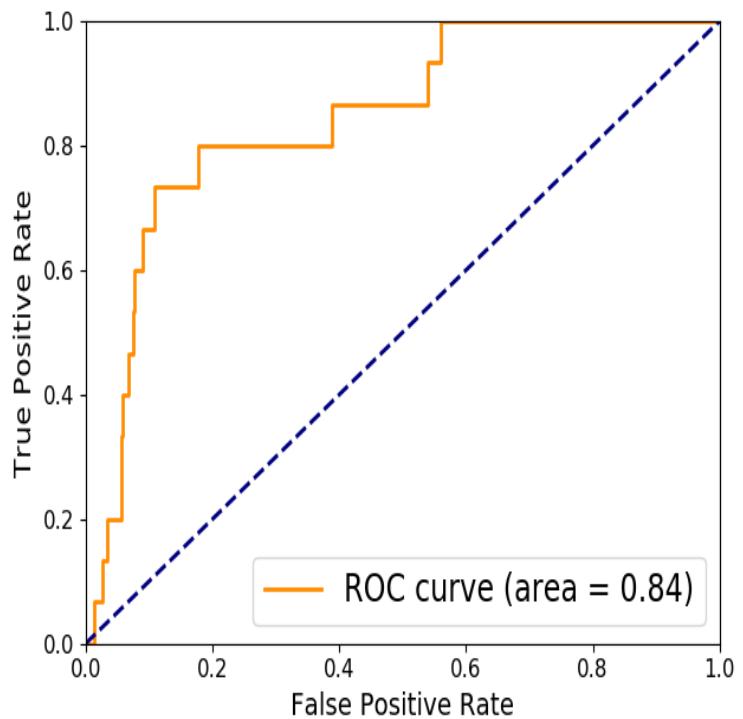
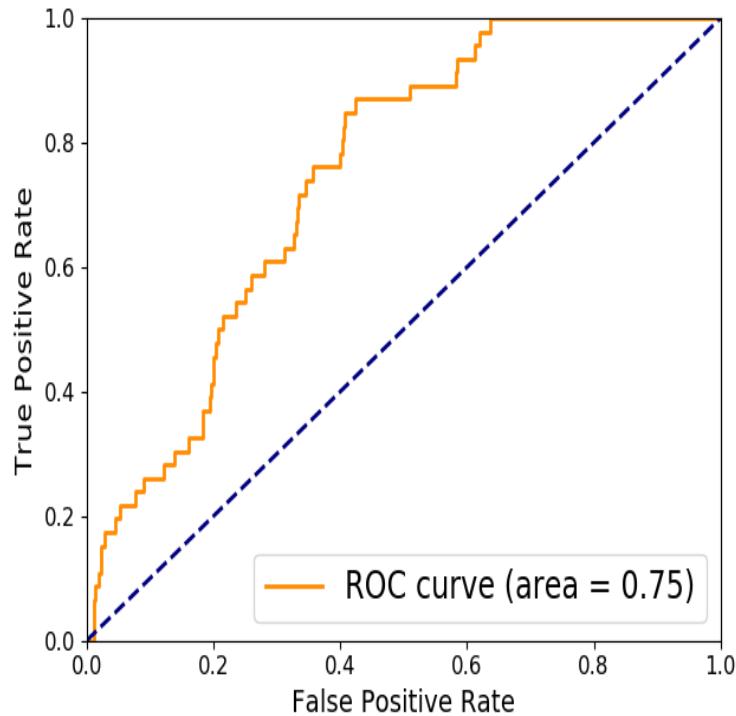


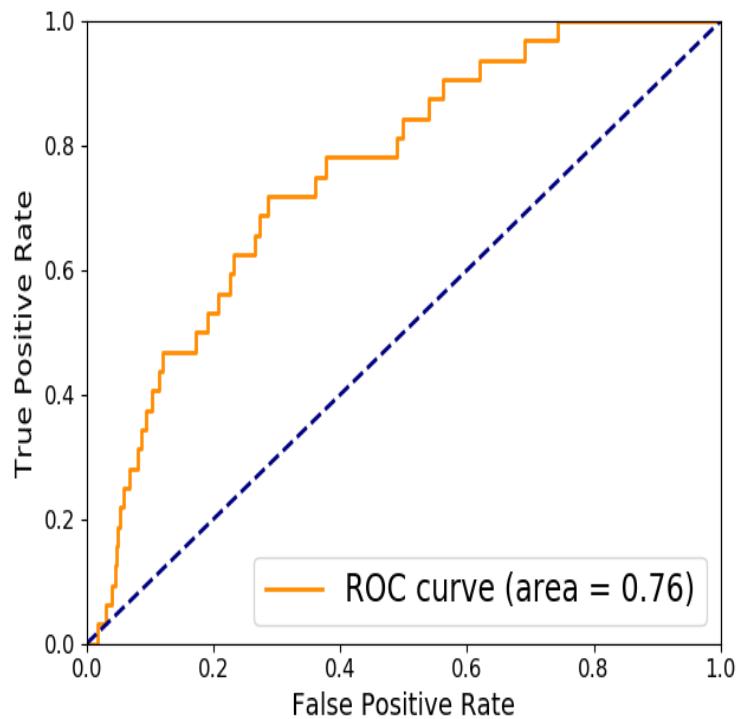
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



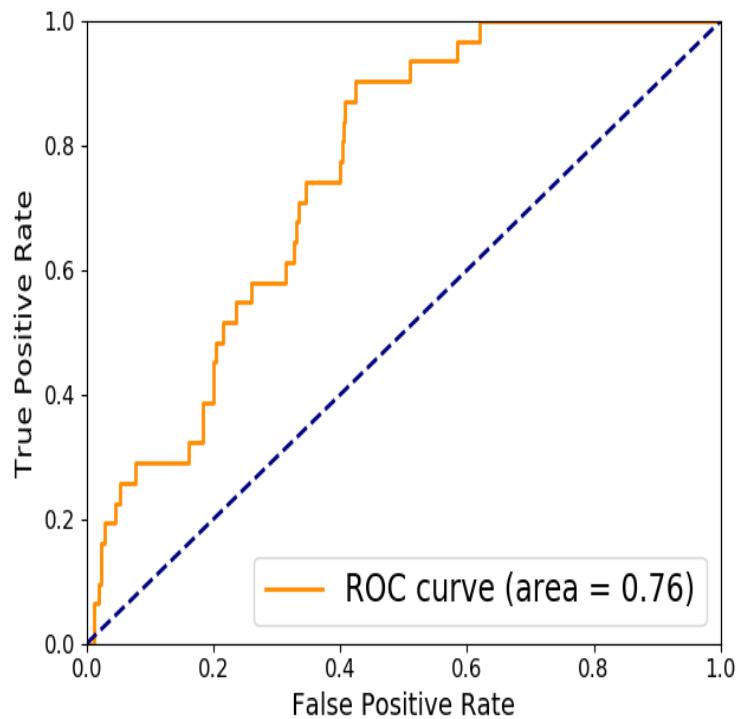
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



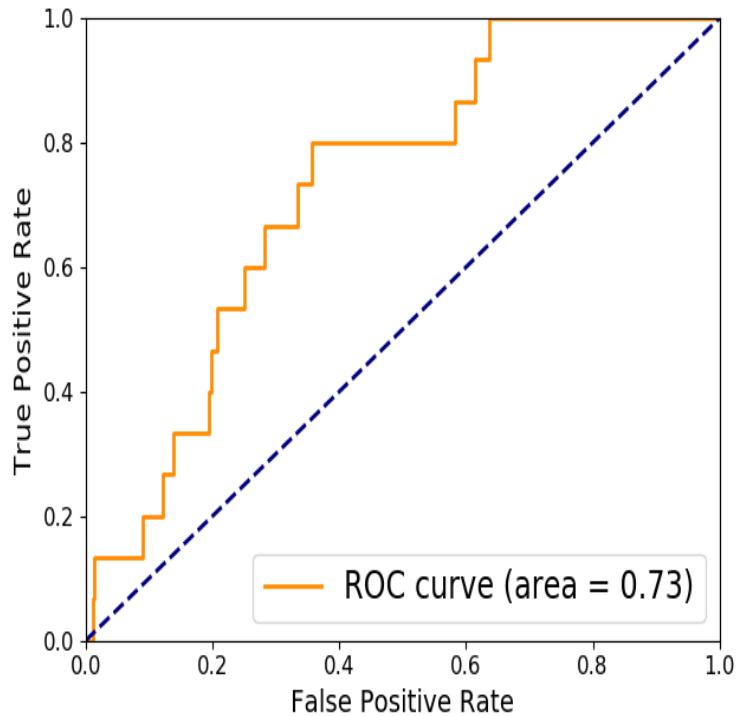
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



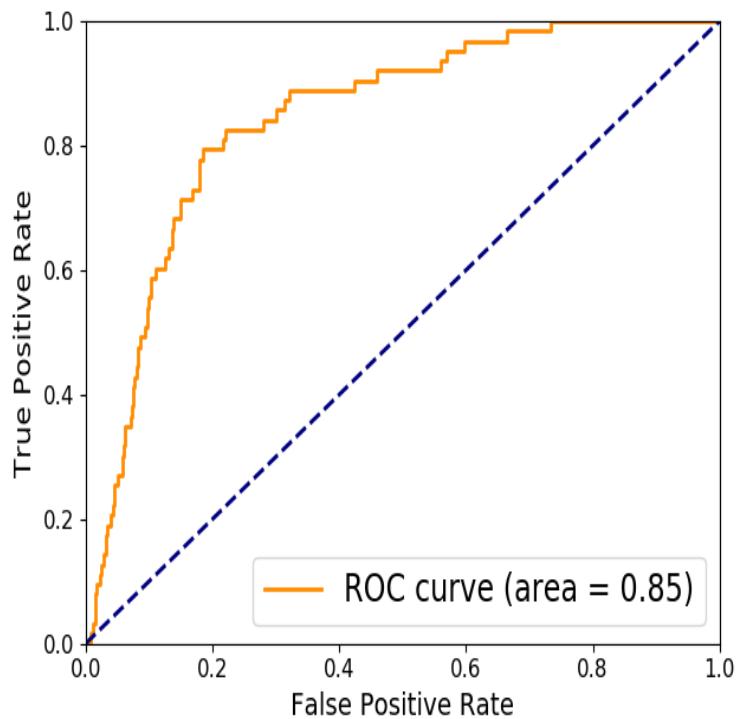
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



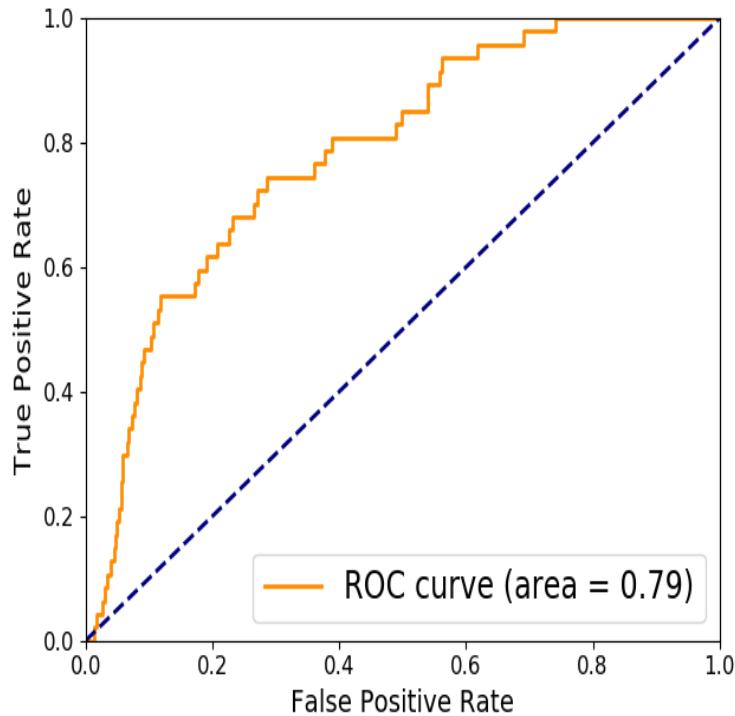
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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>	0.758	4.017	5.593	3.175	2.147	
<i>Test_only</i>	<i>GS_only</i>	0.757	3.082	4.860	0.74818368	0.962	
				64401258			
				(invalid)			
<i>GS_only</i>	<i>Overlap</i>	0.845	3.803	4.505	2.52817868	1.985	
				0356985			
				(invalid)			
<i>GS_only</i>	<i>Test</i>	0.786	5.897	7.941	2.93197282	2.982	
				9688678			
				(invalid)			
<i>Test_only</i>	<i>Overlap</i>	0.731	2.795	3.446	0.113	0.92731722	
				74894577			
				(invalid)			
<i>Test_only</i>	<i>GS</i>	0.749	3.955	6.298	0.588	1.283	
<i>Overlap</i>	<i>Onlies</i>	0.846	7.326	8.773	3.059	2.730	

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>		
<i>GS</i>	31	32	MTND2P28;MT-RNR2;MT-	N/A	
			RNR1;MTATP6P1		
<i>Test_only</i>	32	32	MT-RNR1;MT-	N/A	
			RNR2;MTATP6P1		
<i>Overlap</i>	15	18	MT-RNR1;MT-	MTATP6P1;MT-ND5;MT-	
			RNR2;MTATP6P1	CO3;MT-CO1;MT-ND1;FTL	
				;MT-ND6;MT-RNR2;MT-	
				ATP8;MT-ND3;ALDOB;MT-	
				RNR1;MT-ND4L;MT-CYB	
				;MT-ND2;MT-ND4;MT-	
				ATP6;MT-CO2	

(end of file)