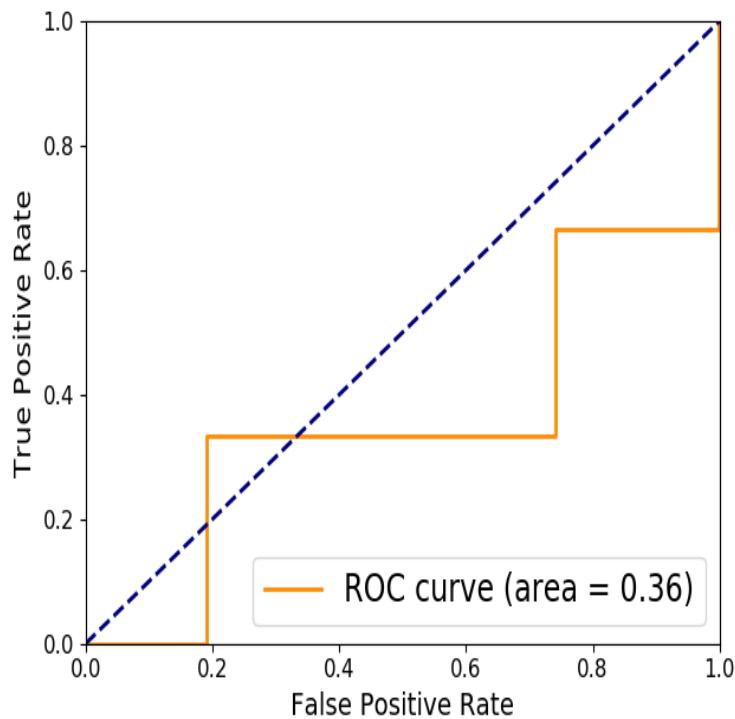
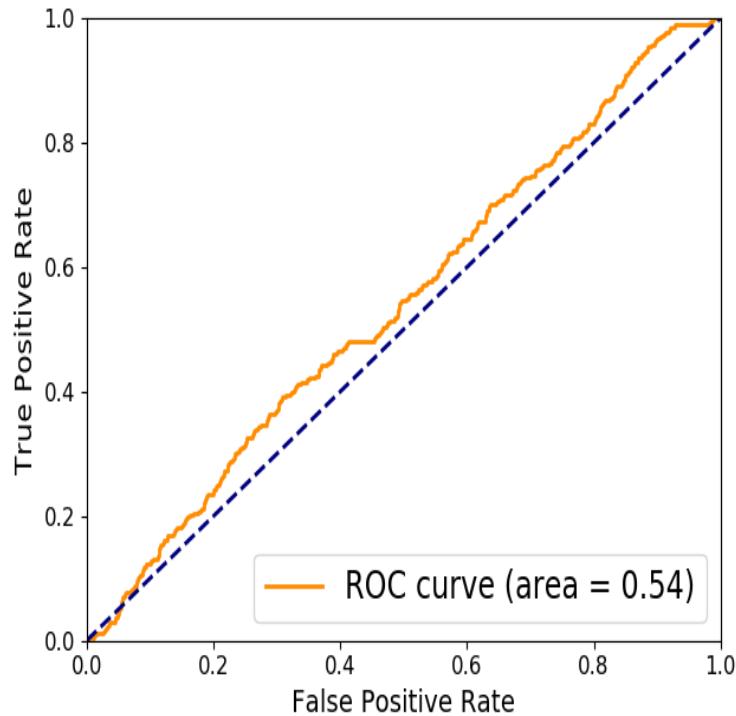


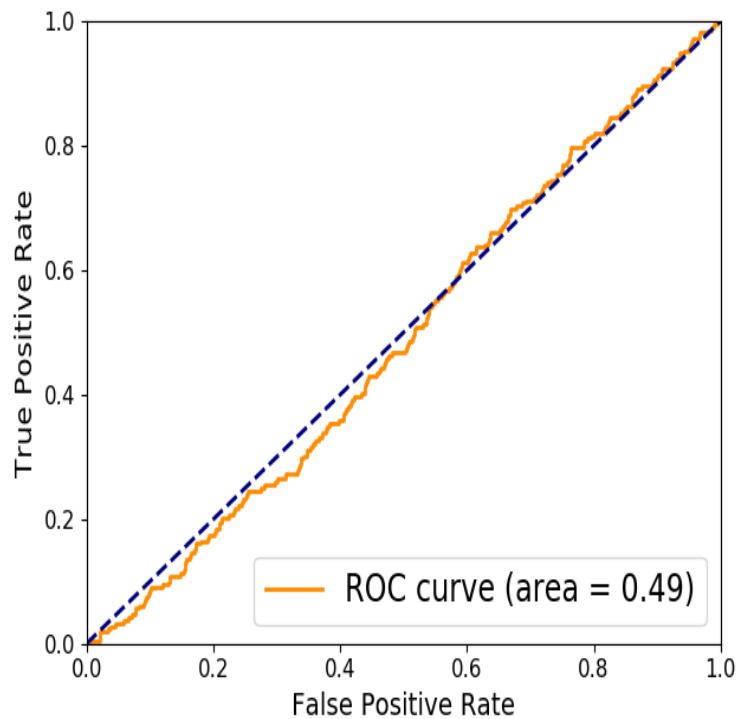
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



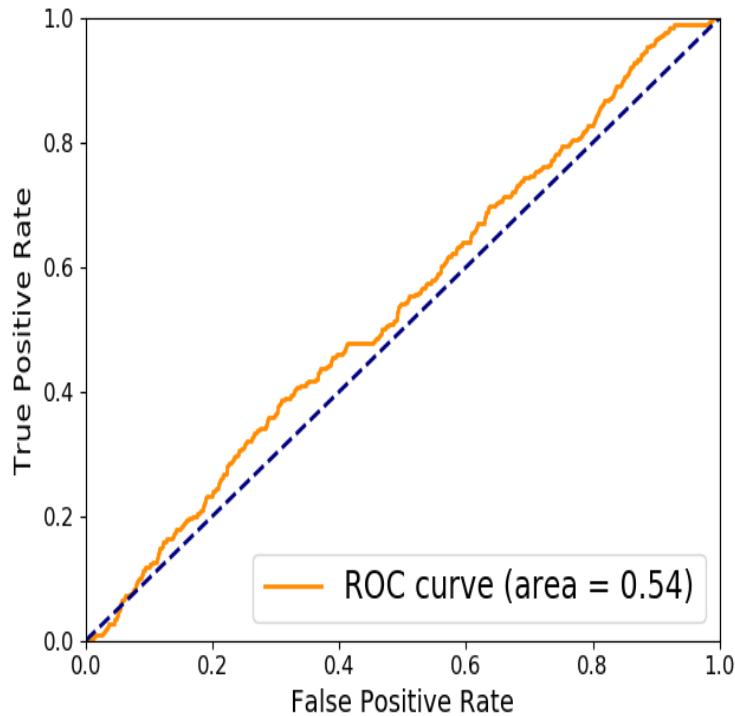
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



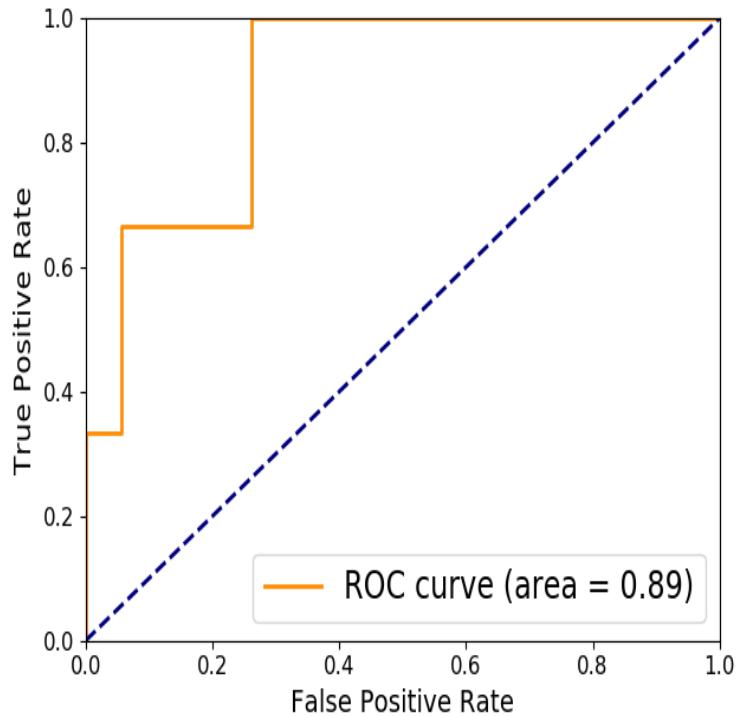
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



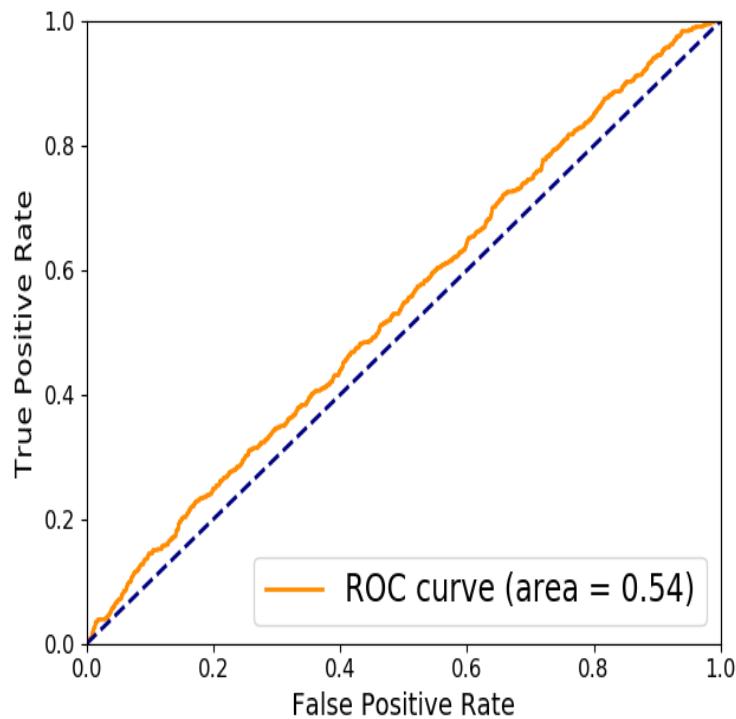
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



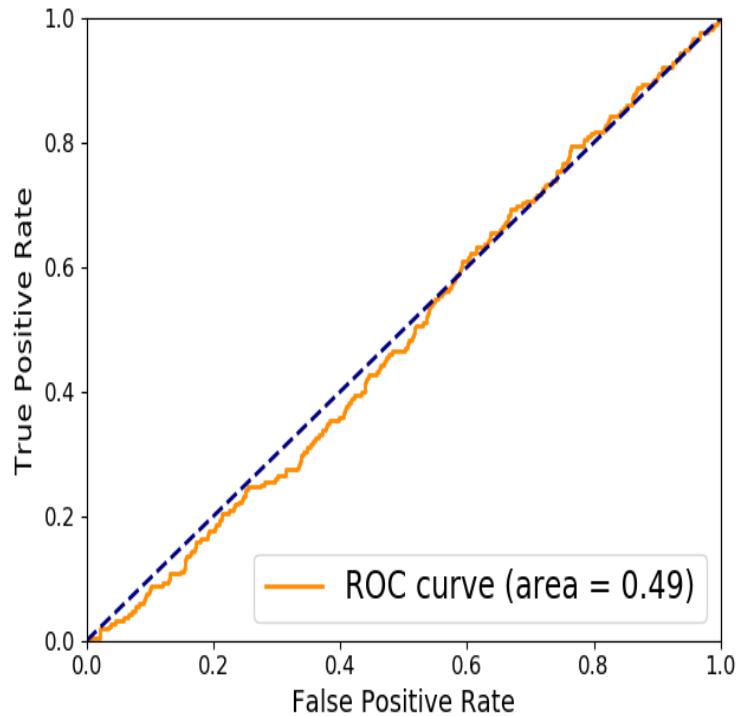
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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>	0.491	-0.251	-0.356	-0.141	0.163	
<i>Test_only</i>	<i>GS_only</i>	0.541	0.382	2.442	0.318	0.180	
<i>GS_only</i>	<i>Overlap</i>	0.357	-0.435	-0.777	0.73057587	0.5513255	
				34262847	33570364		
					(invalid)	(invalid)	
<i>GS_only</i>	<i>Test</i>	0.489	-0.224	-0.588	-0.019	0.139	
<i>Test_only</i>	<i>Overlap</i>	0.894	2.495	2.668	4.321	4.8808156	
				44043601			
					(invalid)		
<i>Test_only</i>	<i>GS</i>	0.544	0.72734856	3.005	0.778	0.601	
		93103832					
			(invalid)				
<i>Overlap</i>	<i>Onlies</i>	0.543	1.820	3.596	0.676	1.658	

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>		
<i>GS</i>	354	379	CFAP46;PCNX2;INTS13;MAP4 N/A		
			K3-DT;DUS2;ATP5PB;C3orf8		
			0;PLPP1;ZNF37BP;C17orf10		
			2;RPSAP69;SEPT7P2;CEP83;		
			PYCR3;ADGRA1-AS1;ERBIN;S		
			MIM10L2A;GMDS-DT;MIGA1;F		
			AM153B;NPNT;CIPC;RAB29;		
			FRMD6-AS1;ATP2B1-AS1		
<i>Test_only</i>	212	217	LENG9;C5orf45;OPLAH;MROH N/A		
			7;LOC100653515		
<i>Overlap</i>	3	3	AEBP1;ARMCX4;SASH1		

(end of file)