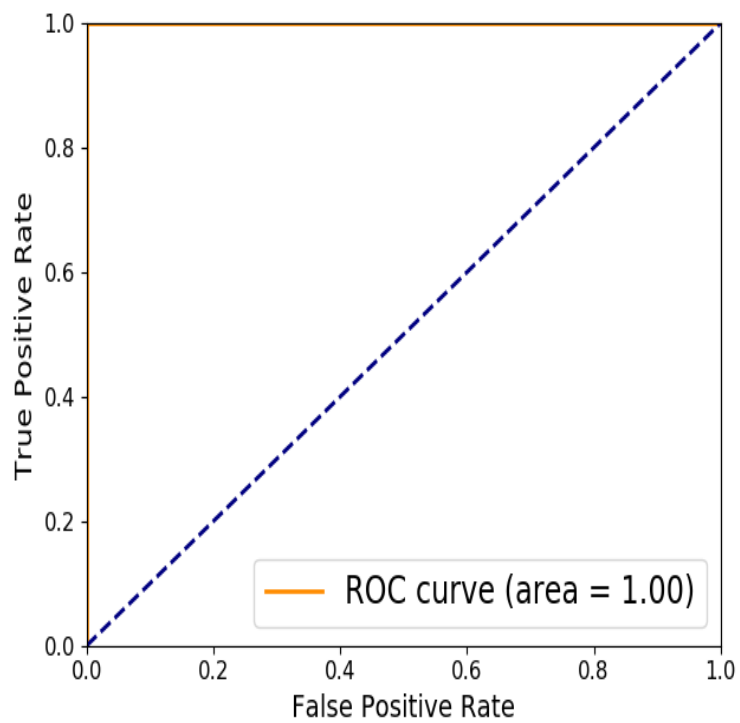
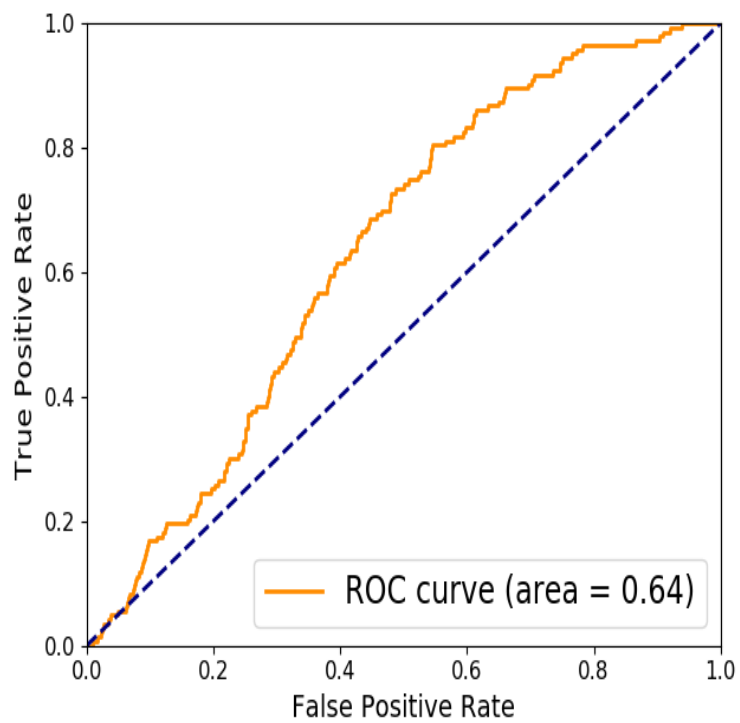


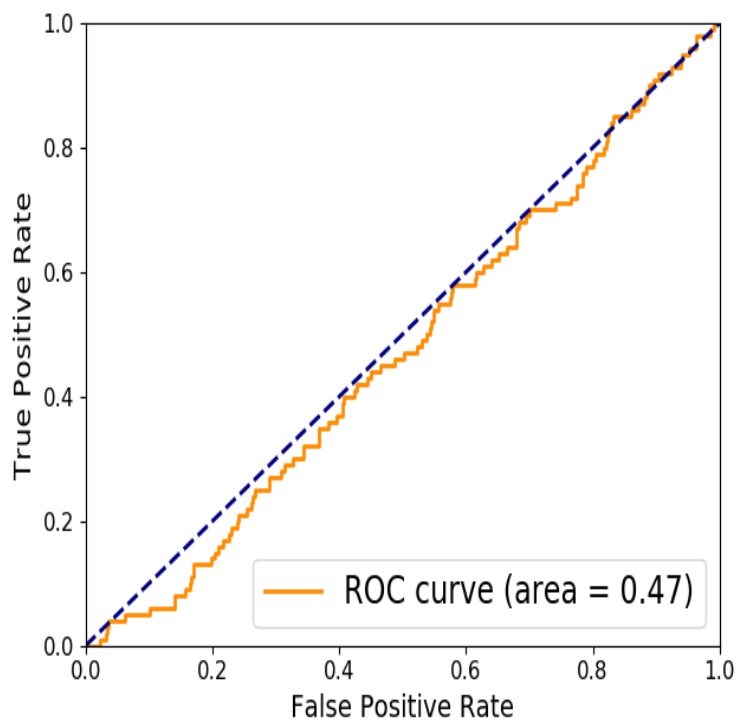
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



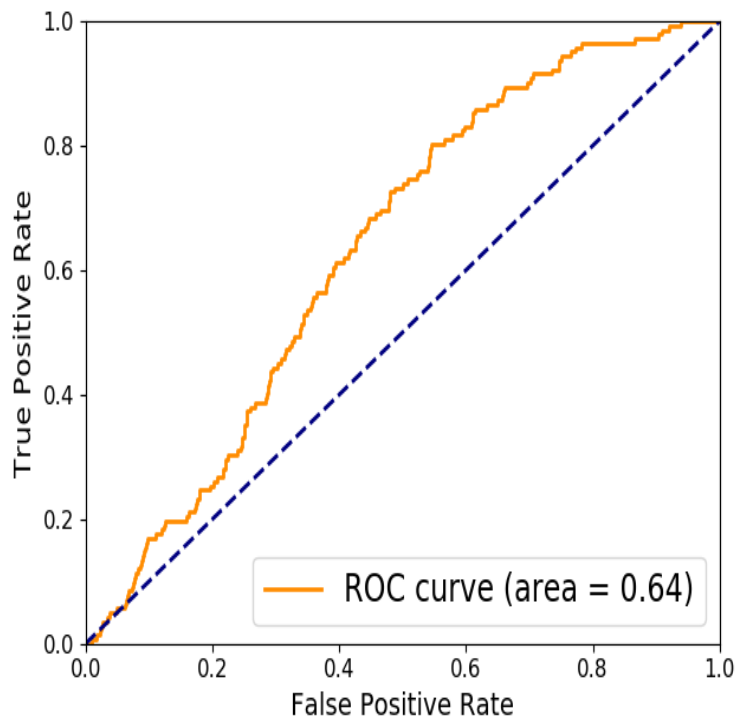
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



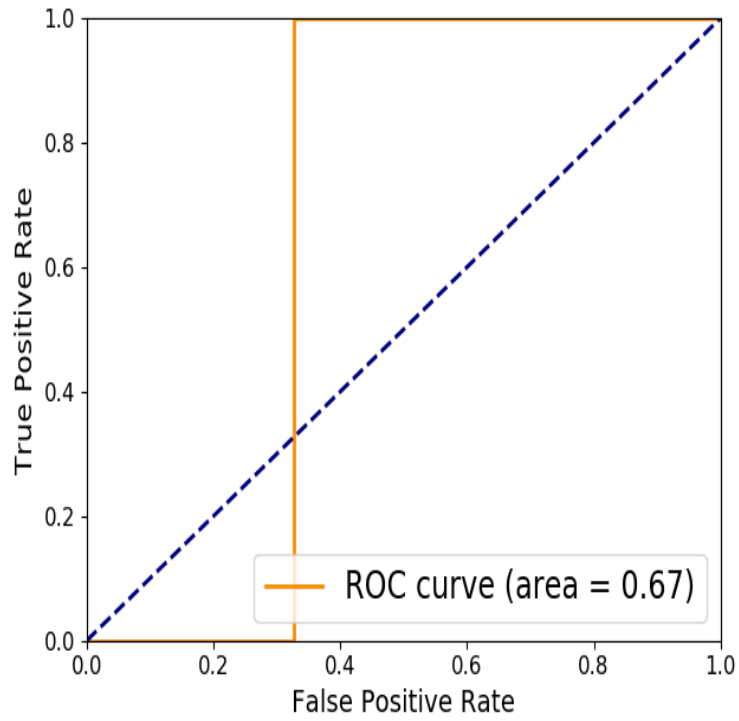
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



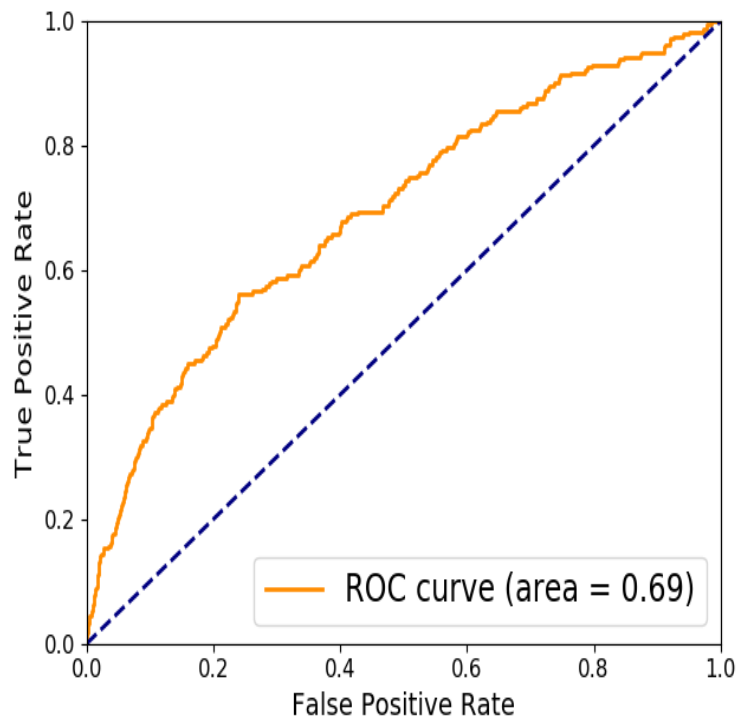
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



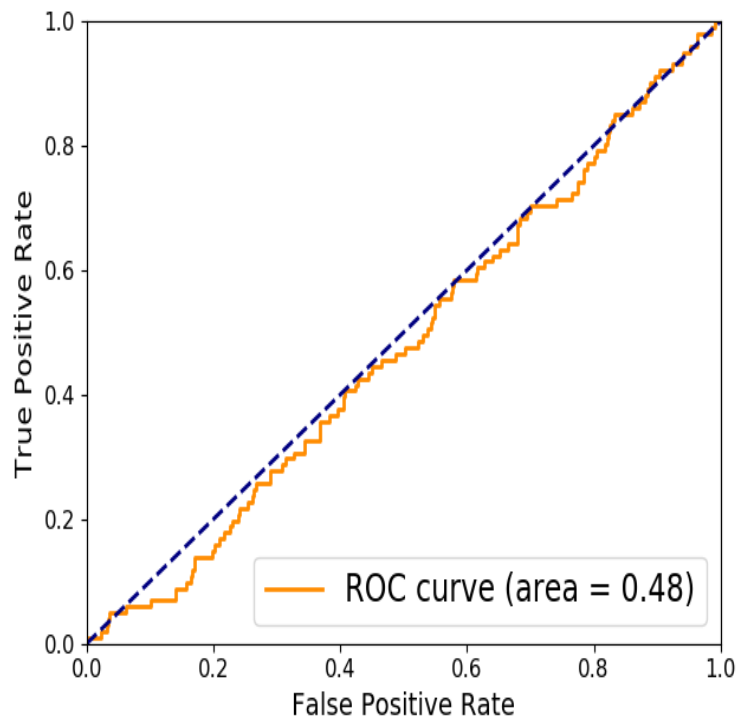
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## Diffusion ROC from fOverlap\_tOnlies STRING10\_combined.png



## Diffusion ROC from fGOnly\_tTest STRING10\_combined.png



Diff_FROM	Diff_TO	AUC	Zt_wdegre	Zt	Zf_wdegre	Zf
		e		e		
GS_only	Test_only	0.475	-1.340	-1.042	-0.789	-0.889
Test_only	GS_only	0.637	-0.355	5.634	-1.382949	-1.034
				723964488		
				(invalid)		
GS_only	Overlap	1.000	4.1782993	1.6949882	1.4037202	1.356510
			01158174	387814224	007882865	49268003
			(invalid)	(invalid)	(invalid)	09 (inva
					lid)	
GS_only	Test	0.480	-0.872	-0.564	-0.638	-0.770
Test_only	Overlap	0.674	-5.623683	0.5940855	-2.551742	-1.85022
			959162842	409755229	486317427	86241943
			(invalid)	(invalid)	(invalid)	577 (inv
					alid)	
Test_only	GS	0.637	-0.433	5.249	-1.096	-0.911
Overlap	Onlies	0.693	8.016	11.061	6.1233678	2.657
				65637137		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	142	157	LOC105378102;EEF1AKMT3;ND2;TERC;LINC01   N/A
			237;LOC100631378;LINC02227;AFG1L;PPP1R
			26-AS1;ND5;LOC105375494;ADA2;RNR2;ATP8
			;LOC107987209
Test_only	100	102	AMER2;ERVFRD.1   N/A
Overlap	1	1	APOE

(end of file)