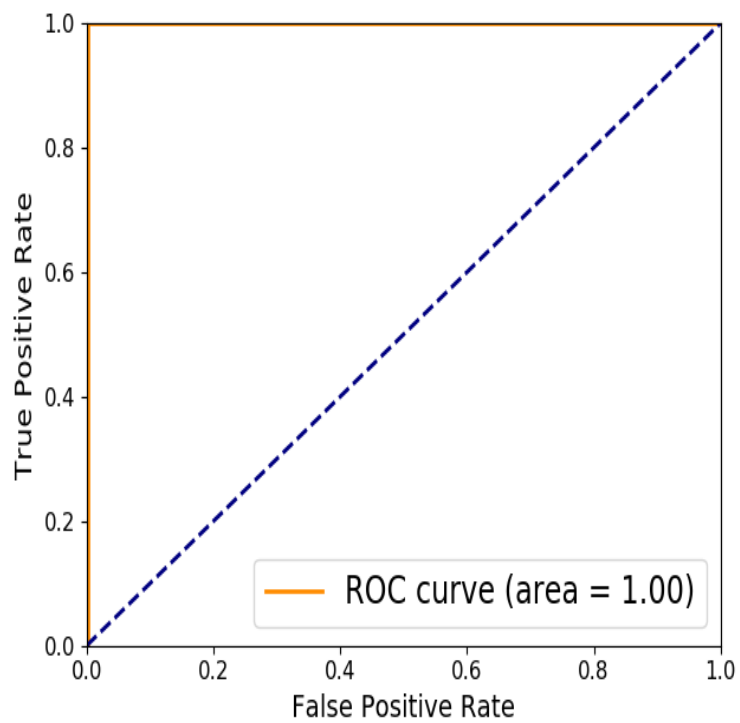
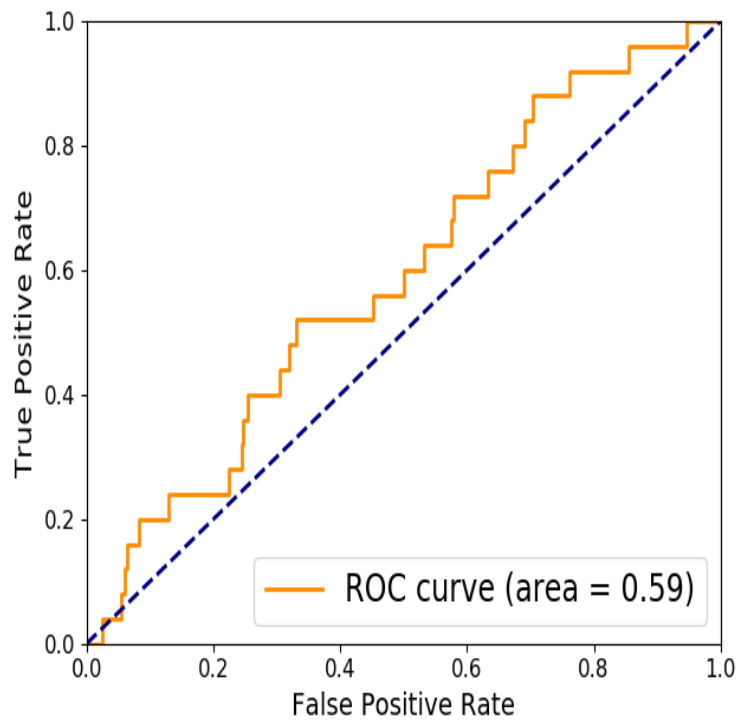


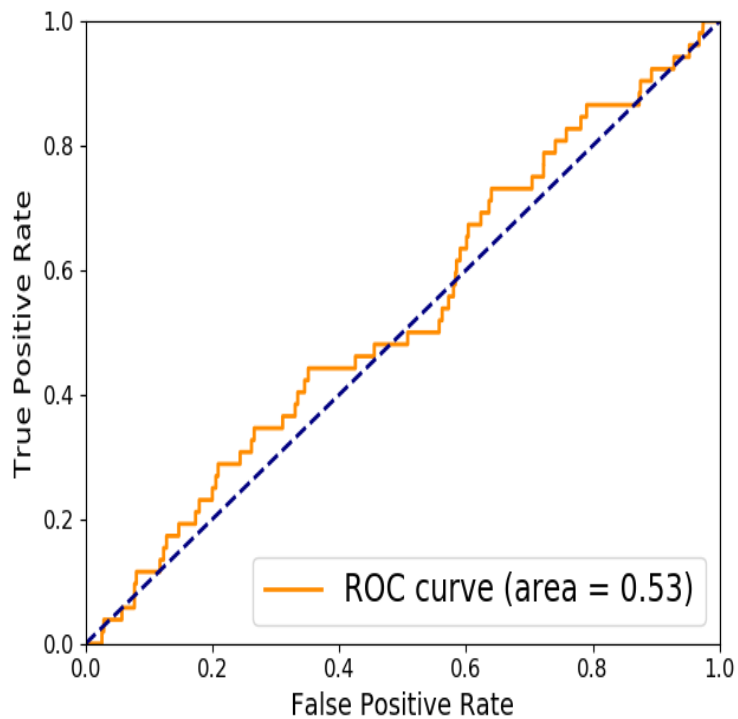
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



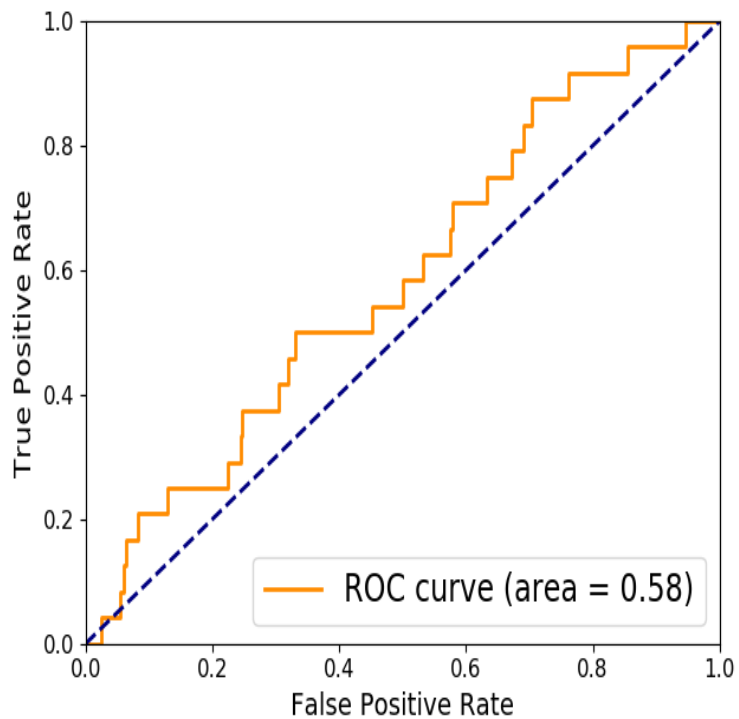
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



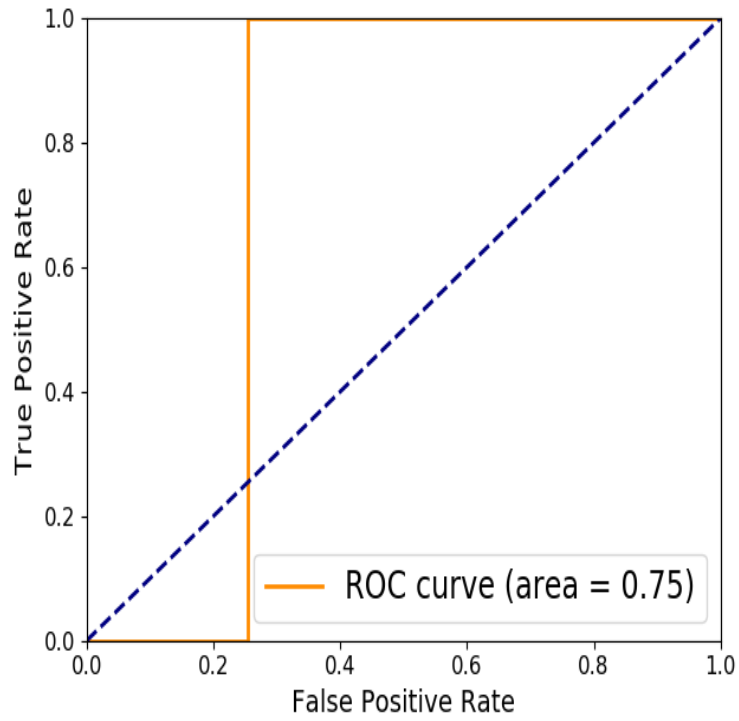
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



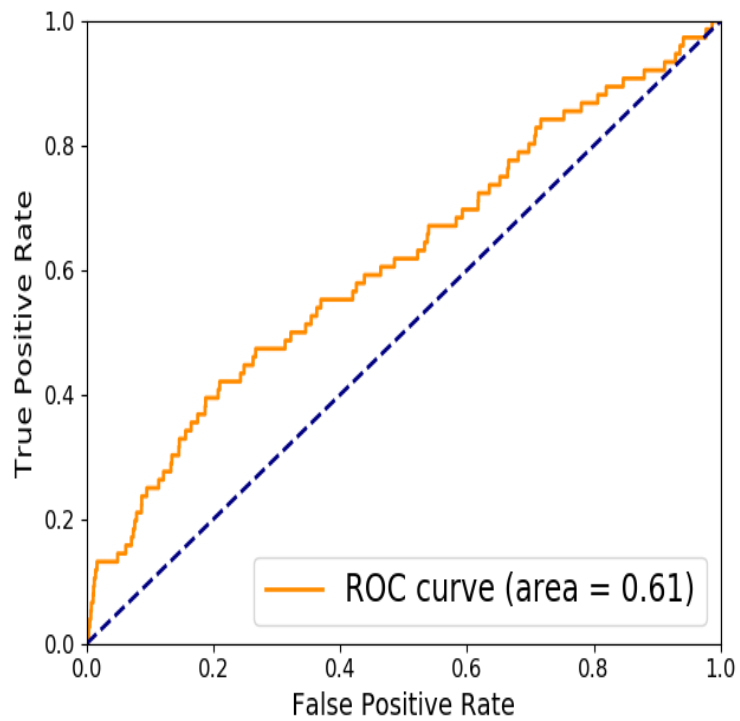
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



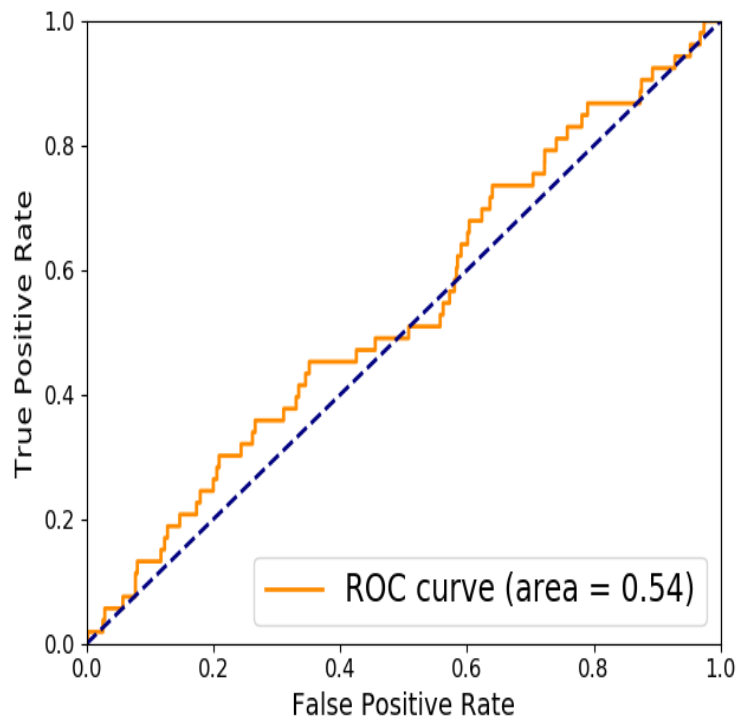
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.529	0.4639862	0.570	1.129	1.106
		413819904				
		(invalid)				
Test_only	GS_only	0.584	0.206	1.690	-0.477	-0.04178
				20447867		
				93 (inva		
				lid)		
GS_only	Overlap	0.998	5.4677458	1.7021189	1.6857453	2.078138
		61967107	602282107	055913894	81411956	
		(invalid)	(invalid)	(invalid)	47 (inva	
				lid)		
GS_only	Test	0.538	0.446	0.909	1.122	1.265
Test_only	Overlap	0.746	-1.224010	1.1077454	-0.830503	-0.82513
		675223229	60842555	875472138	27190069	
		4	(invalid)	3	362 (inv	
		(invalid)		(invalid)	alid)	
Test_only	GS	0.591	0.220	1.759	-0.627291	0.063
				142449923		
				9		
				(invalid)		
Overlap	Onlies	0.613	3.465	3.303	130.09626	2.435
				971797692		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	24	24	N/A
Test_only	52	53	IQCF6
Overlap	1	1	APOE

(end of file)