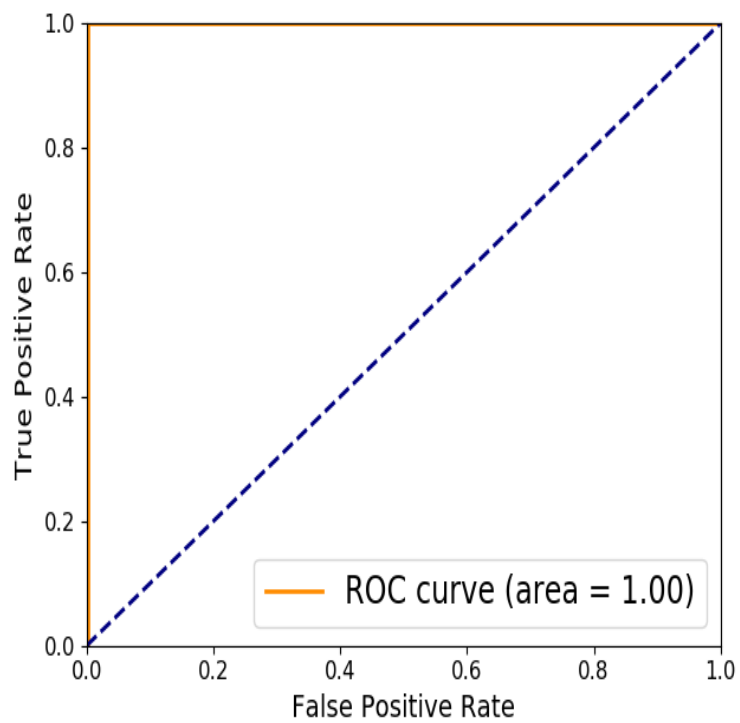
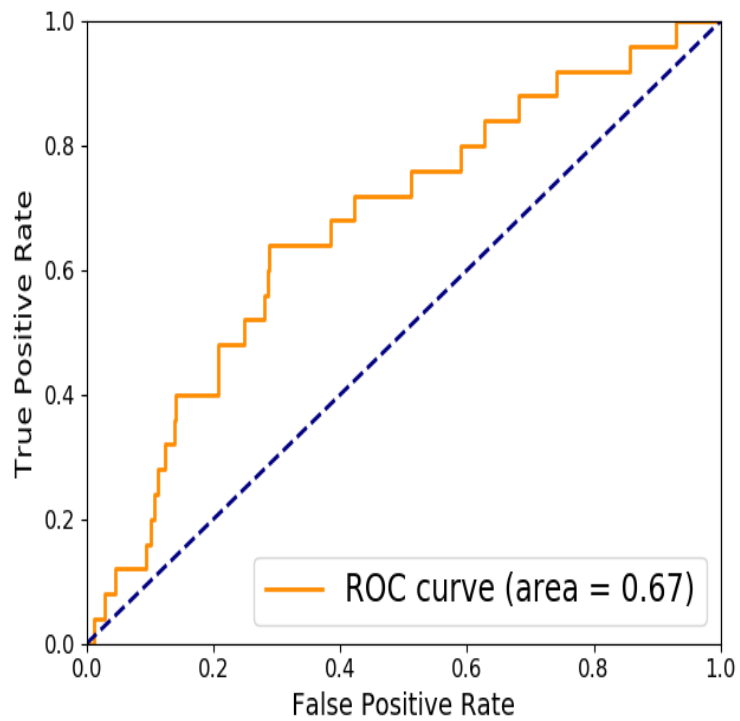


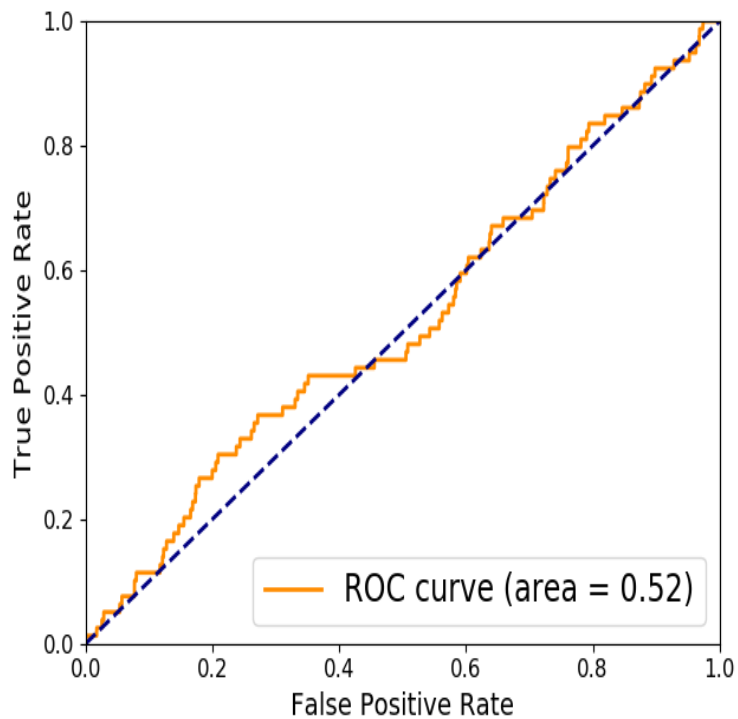
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



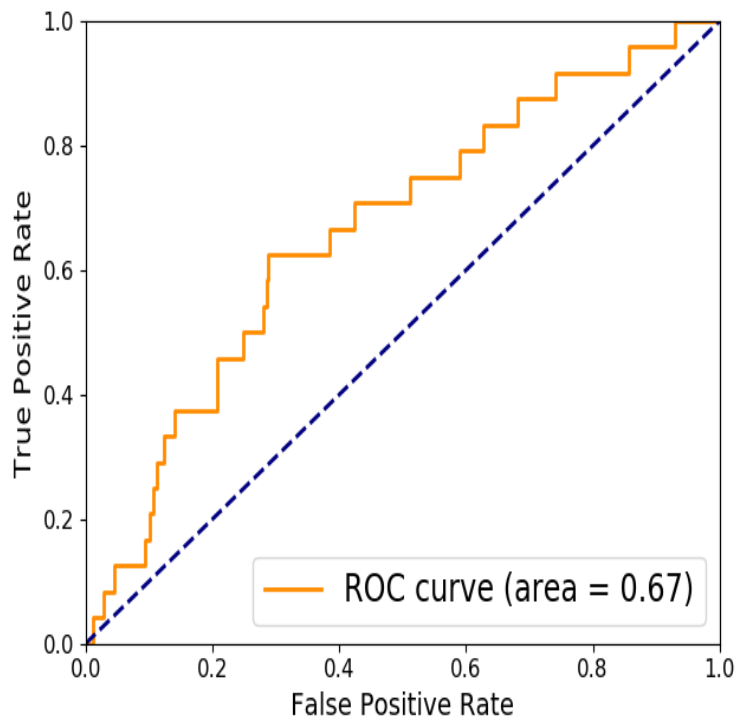
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



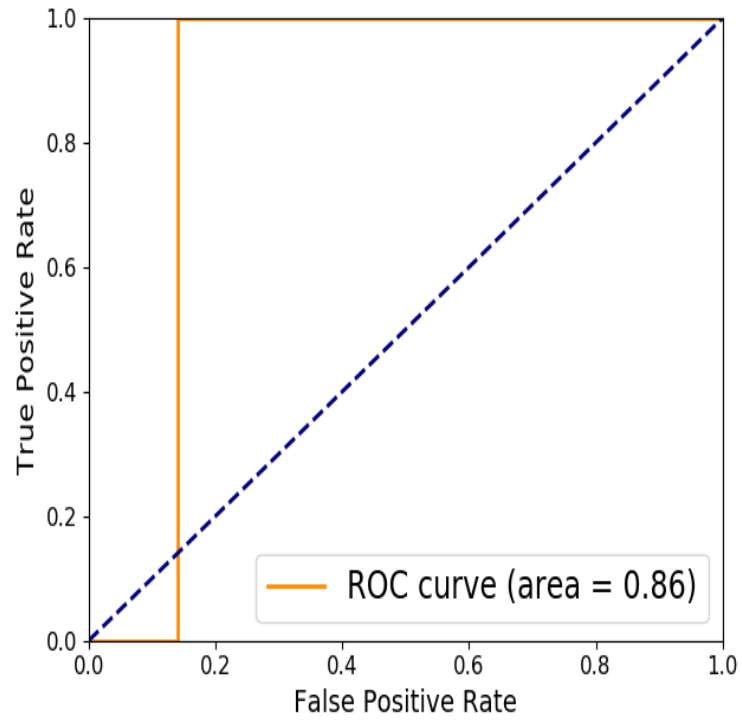
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



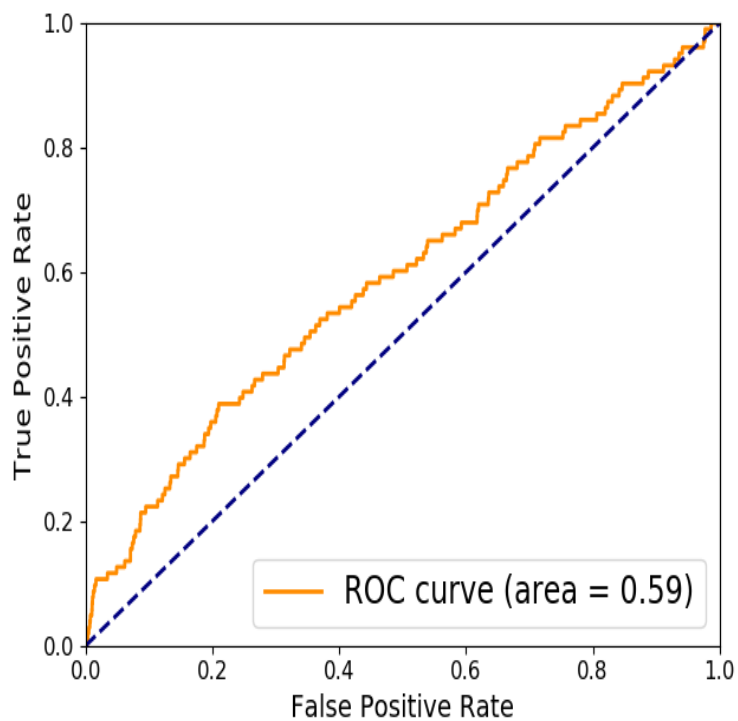
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



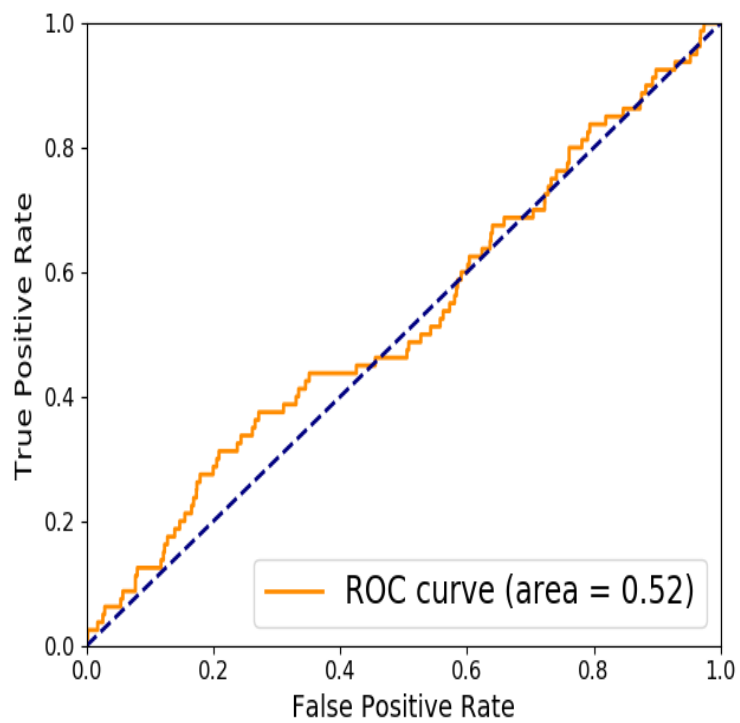
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.519	1.327	0.601	1.8744657	2.053
				334599126		
				(invalid)		
Test_only	GS_only	0.666	1.917	2.924	1.384	1.248736
				06059230		
				12 (inva		
				lid)		
GS_only	Overlap	0.998	5.3502291	1.8091156	1.5297159	1.709839
			18874036	576062786	619401732	96783169
			(invalid)	(invalid)	(invalid)	38 (inva
						lid)
GS_only	Test	0.525	1.359	0.919	2.1178855	2.127
				463535373		
				(invalid)		
Test_only	Overlap	0.860	1.2943939	1.0709606	-0.088232	-0.06589
			03301861	979614295	037523164	95065263
			(invalid)	(invalid)	1	7563 (in
				(invalid)	valid)	
Test_only	GS	0.673	2.373	2.842	1.127	1.405720
				82467683		
				3 (inval		
				id)		
Overlap	Onlies	0.594	3.911	2.987	11.758894	2.672043
				193165894	74419173	
				(invalid)	84 (inva	
					lid)	

**	#Mapped	#Total	Not mapped genes
GS	24	24	N/A
Test_only	79	80	IQCF6
Overlap	1	1	APOE

(end of file)