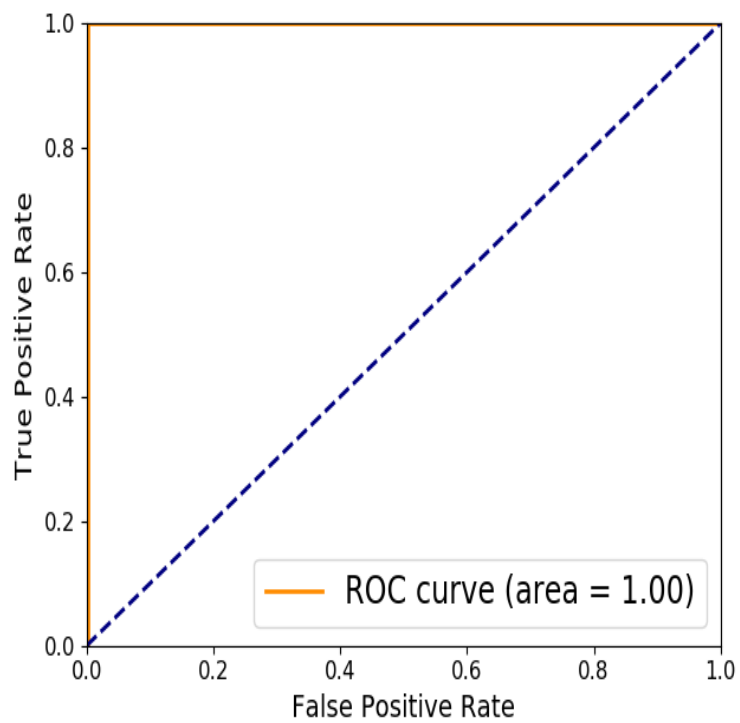
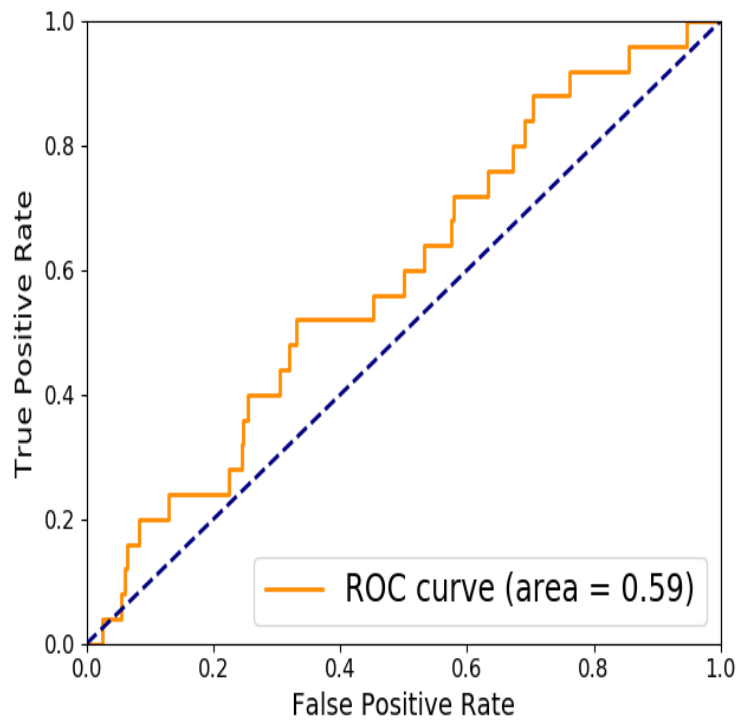


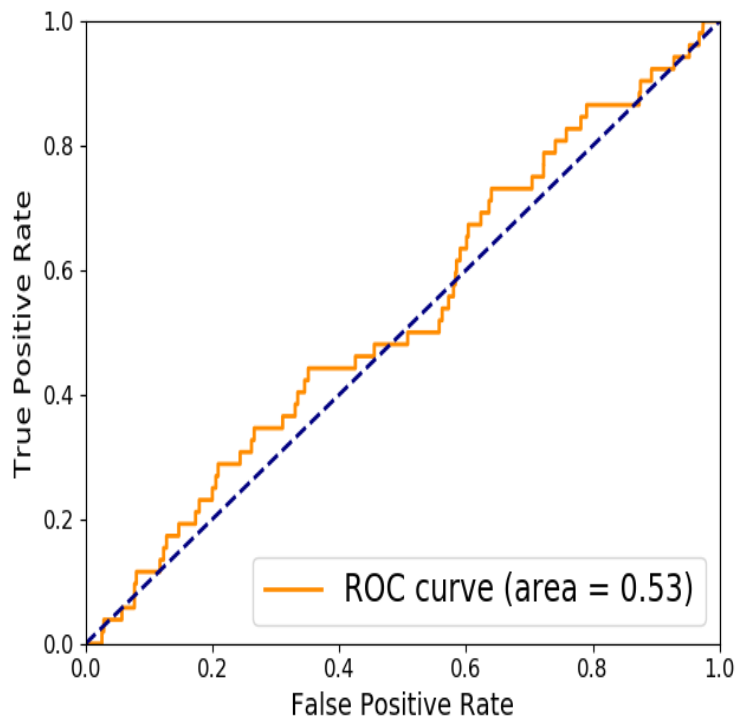
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



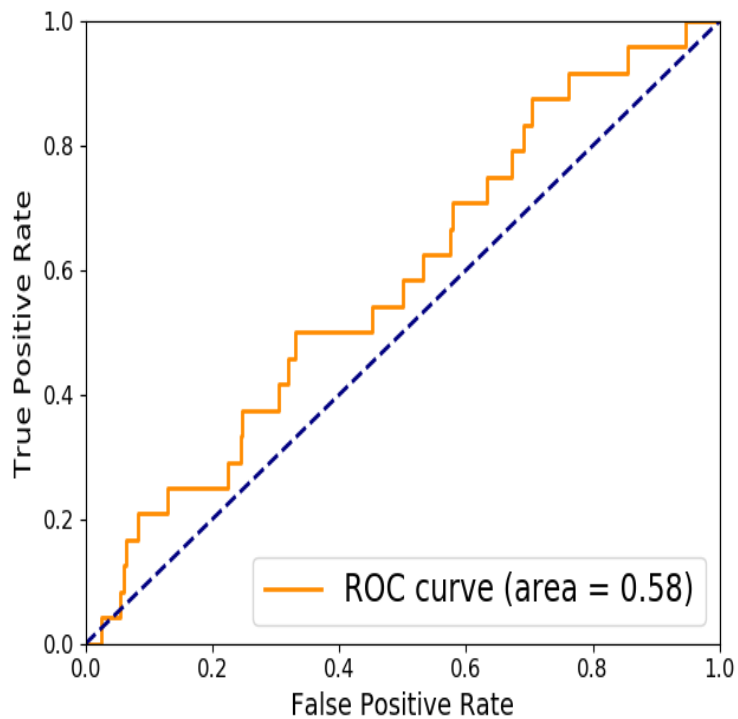
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



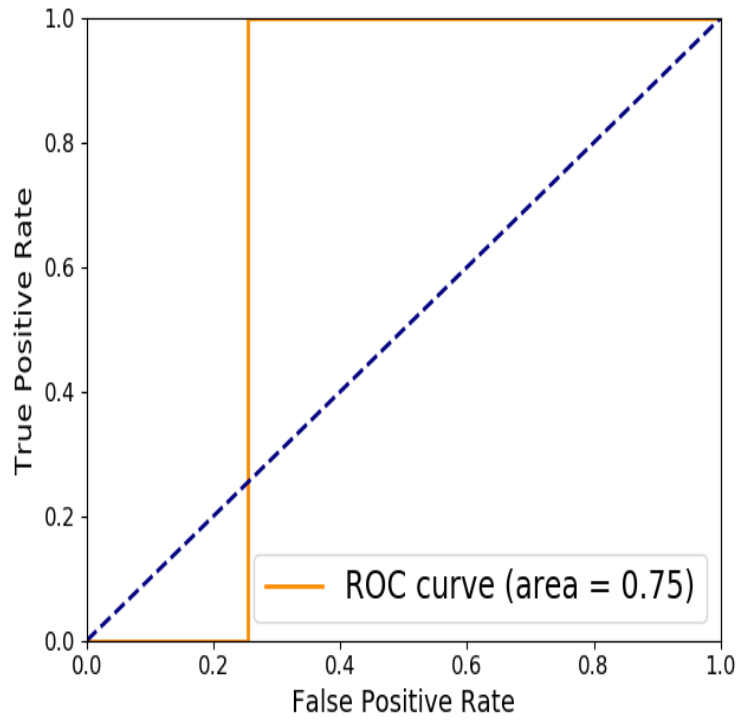
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



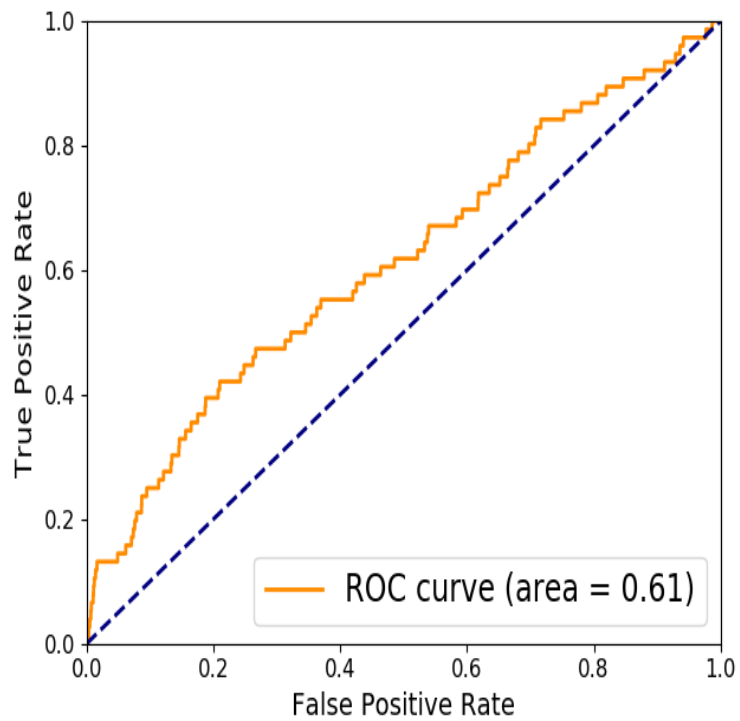
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



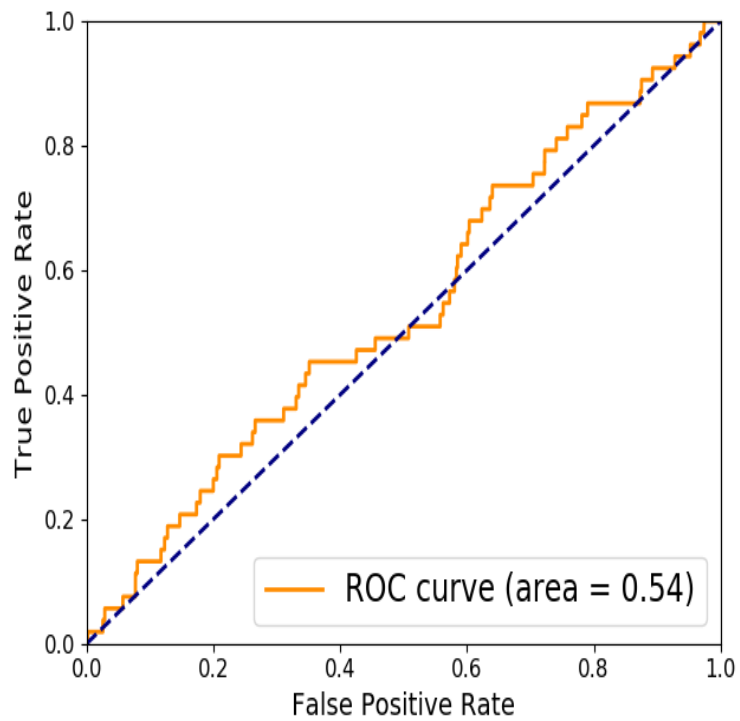
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.244	0.623	0.922	1.315597
				80406913		
				72 (inva		
				lid)		
Test_only	GS_only	0.584	0.290	1.434	-0.412	-0.08074
				71069021		
				411 (inv		
				alid)		
GS_only	Overlap	0.998	5.2684568	1.7878689	1.5337404	1.774985
			38925757	79349857	577055618	89719422
			(invalid)	(invalid)	(invalid)	67 (inva
						lid)
GS_only	Test	0.538	0.6001197	1.069	1.166	1.452
			572185477			
			(invalid)			
Test_only	Overlap	0.746	-1.261932	0.9772910	-0.959288	-0.84325
			379951785	743076567	515027625	66194986
			(invalid)	(invalid)	4	86 (inva
				(invalid)		lid)
Test_only	GS	0.591	0.198	1.611	-0.480030	-0.08913
				476668409	42593510	
				26	5639 (in	
				(invalid)	valid)	
Overlap	Onlies	0.613	3.933	3.088	132.26434	2.219
				998063436		
				(invalid)		

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

(end of file)