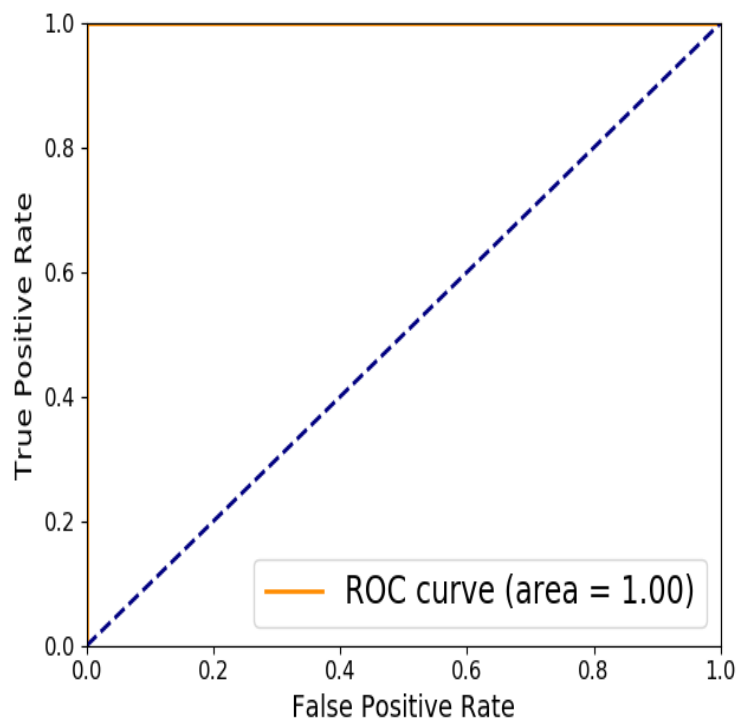
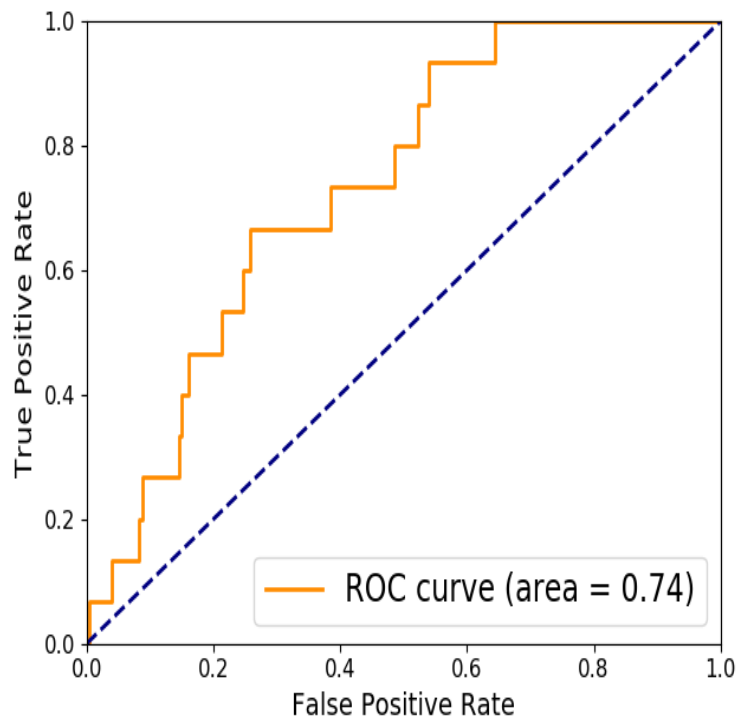


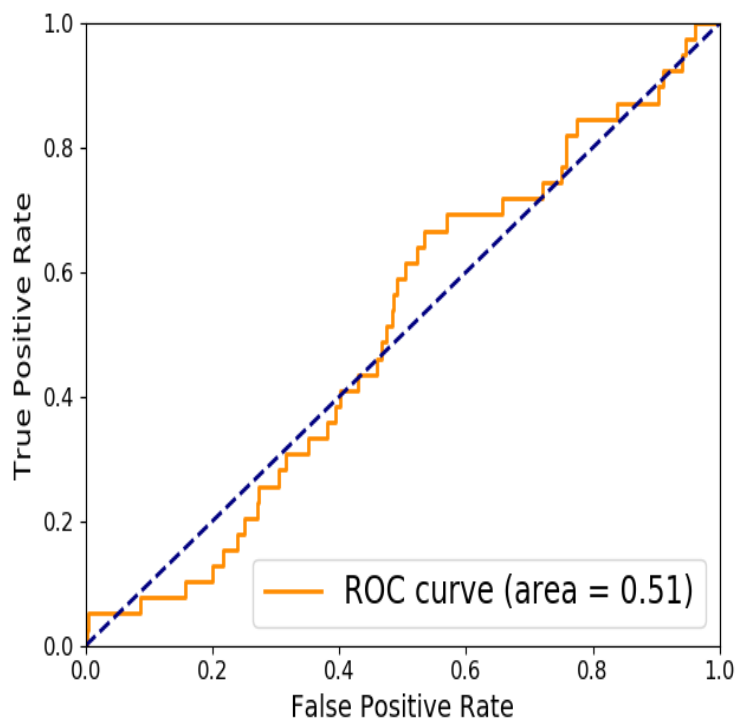
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



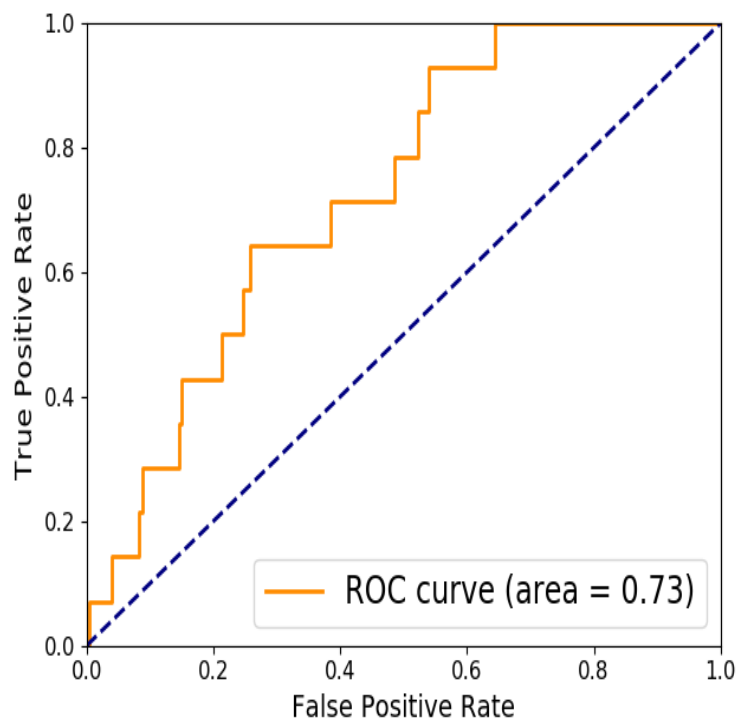
# Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



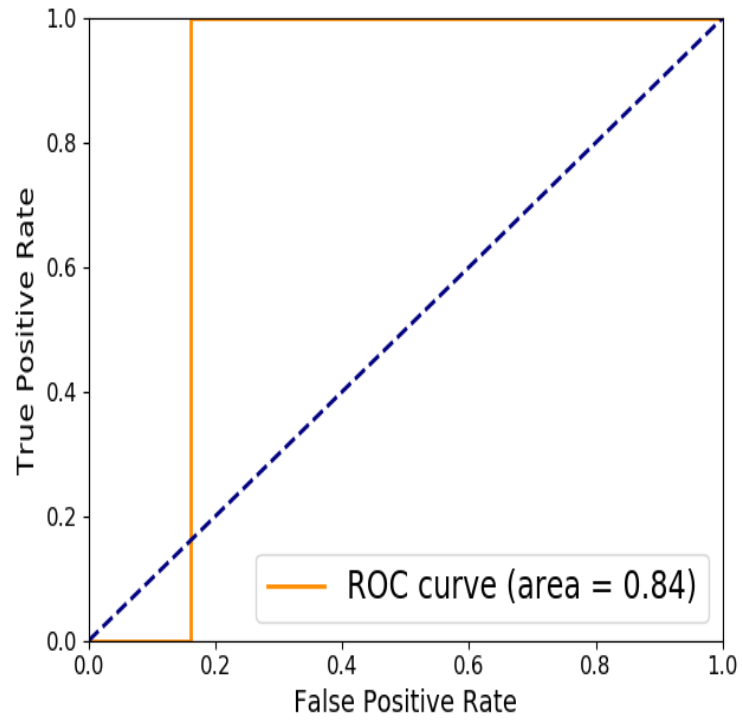
## Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



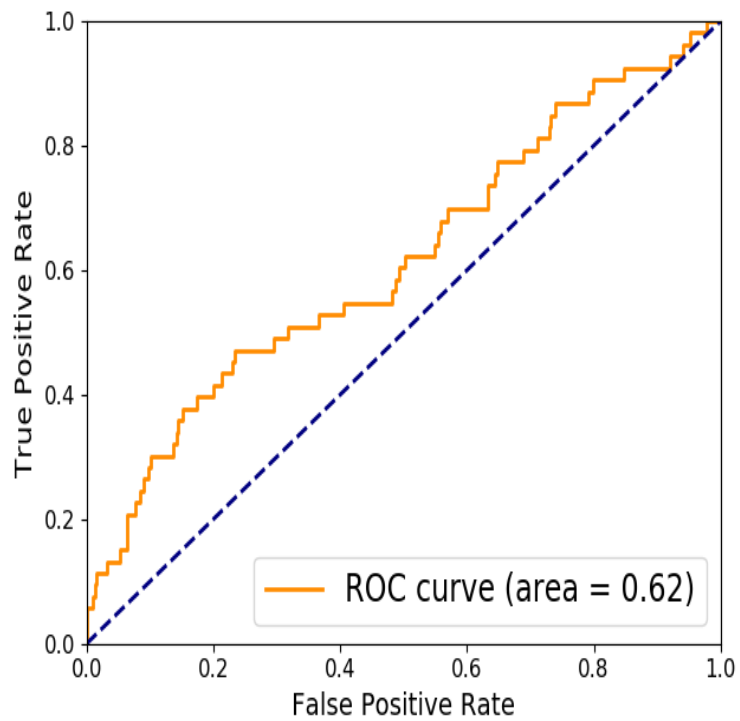
## Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



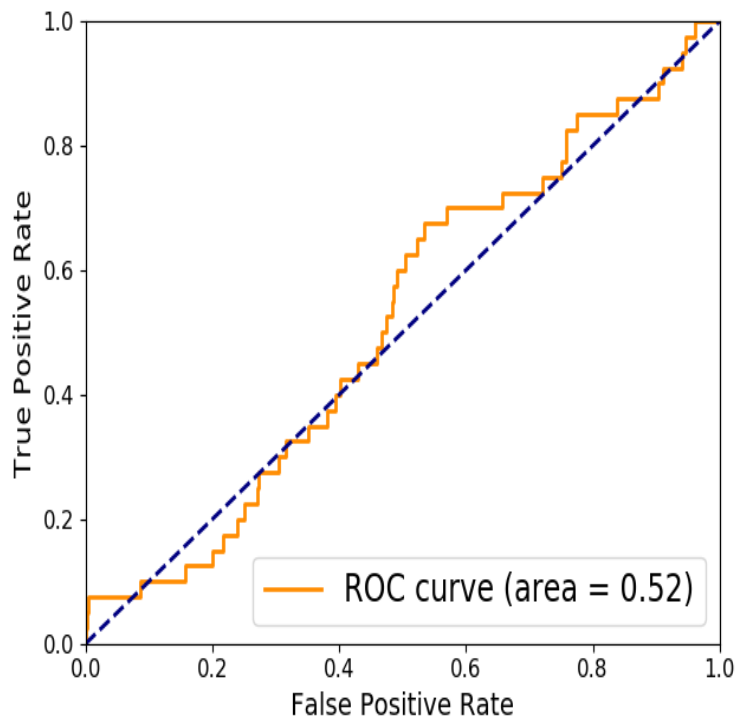
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



# 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	0.508	0.034	0.155	-1.220449	0.179
				751581111		
				5		
				(invalid)		
Test_only	GS_only	0.728	0.252	3.161	0.612	0.718
GS_only	Overlap	0.999	1.4029201	1.7216889	1.8818961	2.026
			658025447	34507038	07700401	
			(invalid)	(invalid)	(invalid)	
GS_only	Test	0.520	0.118	0.413	-0.950065	0.321
				436477809		
				3		
				(invalid)		
Test_only	Overlap	0.839	0.1351006	1.3304017	-0.145613	0.085109
			648395056	583283607	726042715	43263271
			7	(invalid)	(invalid)	924 (inv
			(invalid)		alid)	
Test_only	GS	0.736	0.462	2.8882253	0.809	0.639
				753611002		
				(invalid)		
Overlap	Onlies	0.616	2.017	3.102	4.9725524	1.604
					88725135	
					(invalid)	

**	#Mapped	#Total	Not mapped genes
GS	14	15	CAMK1V   N/A
Test_only	39	39	N/A
Overlap	1	1	APOE

(end of file)