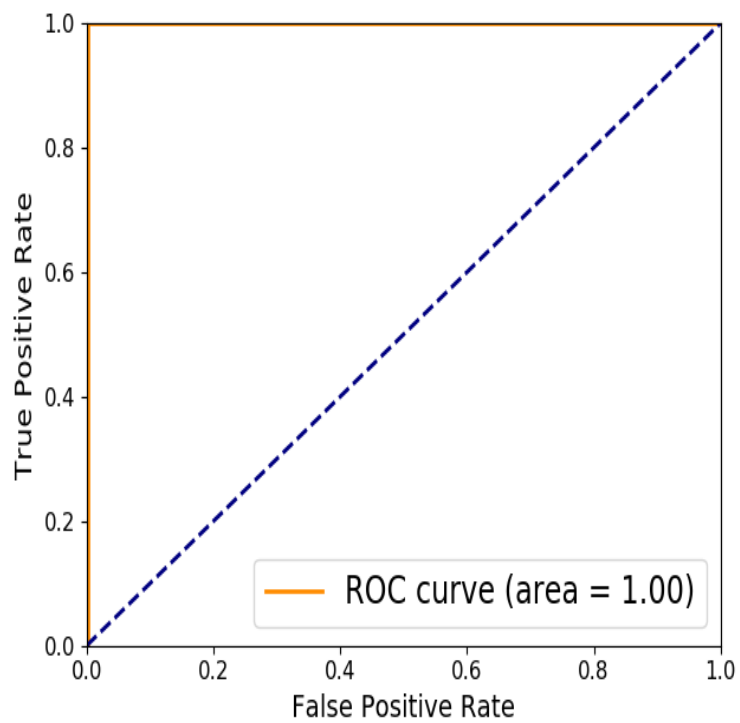
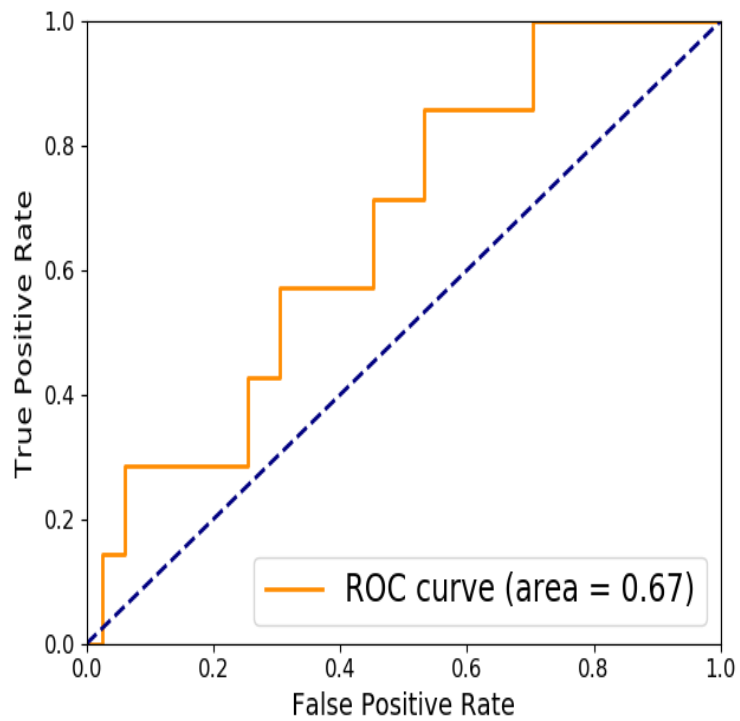


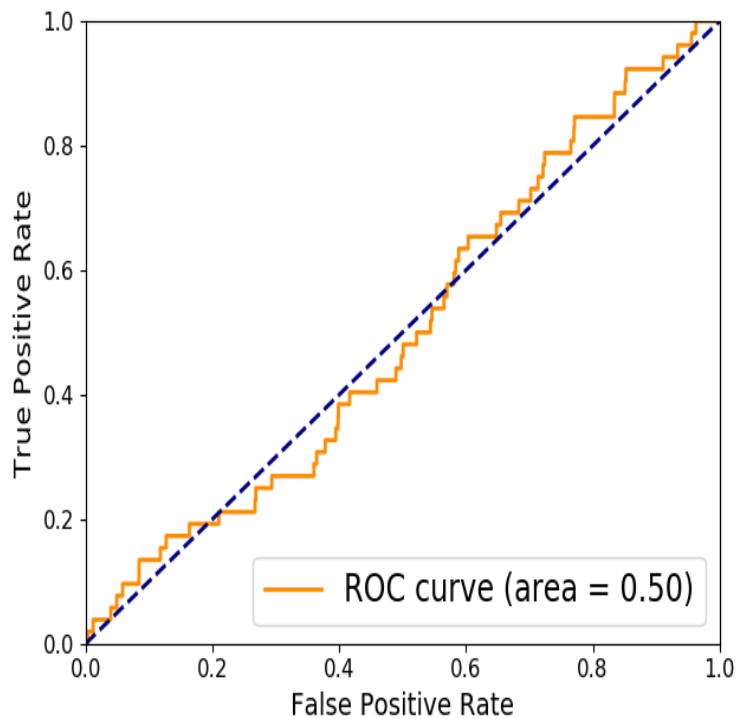
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



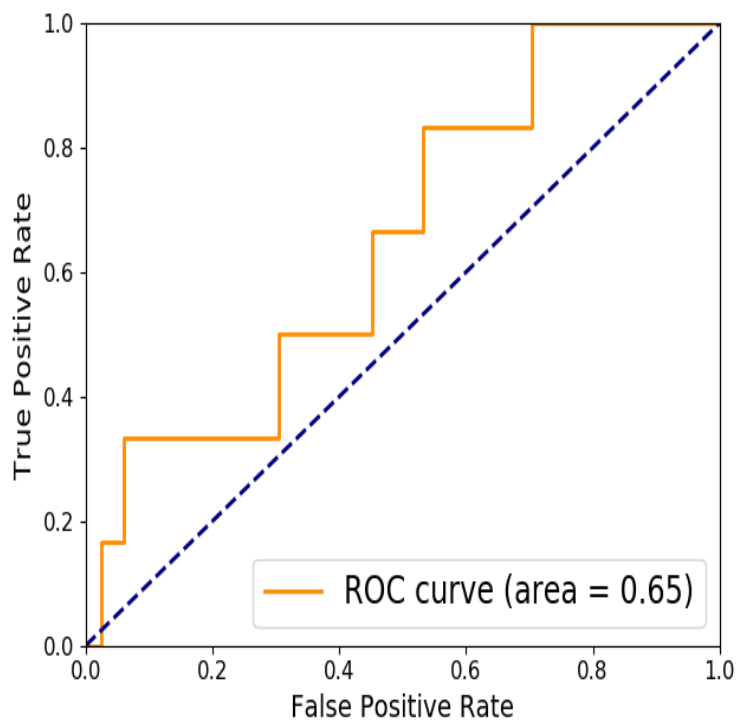
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



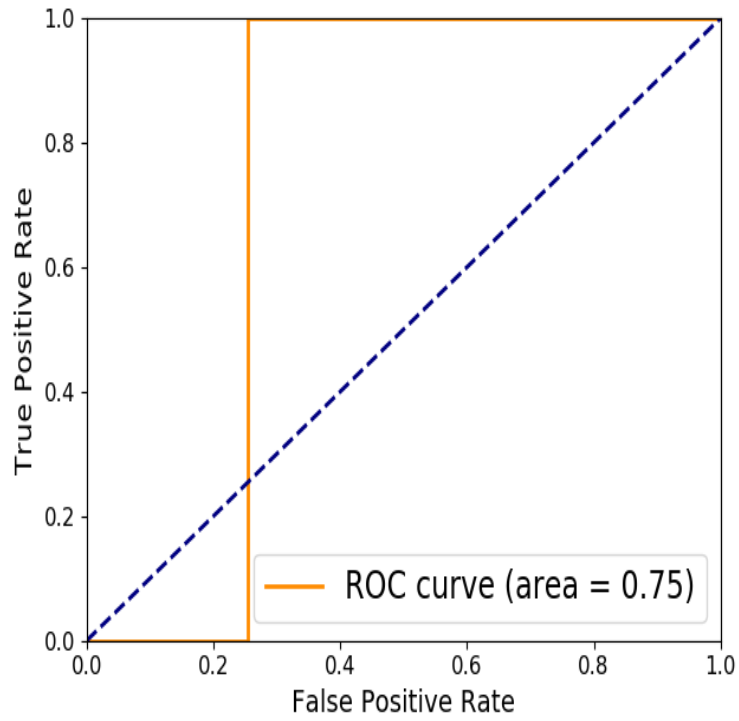
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



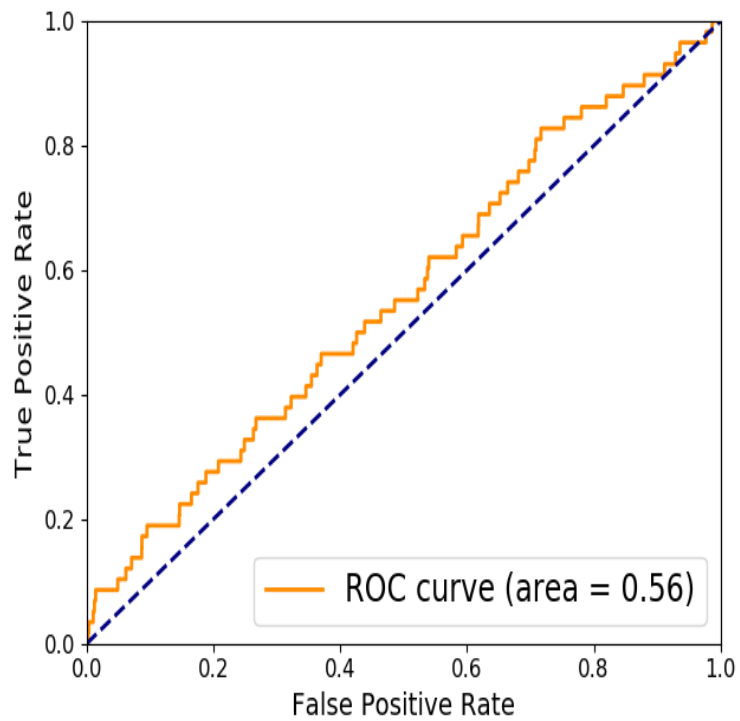
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



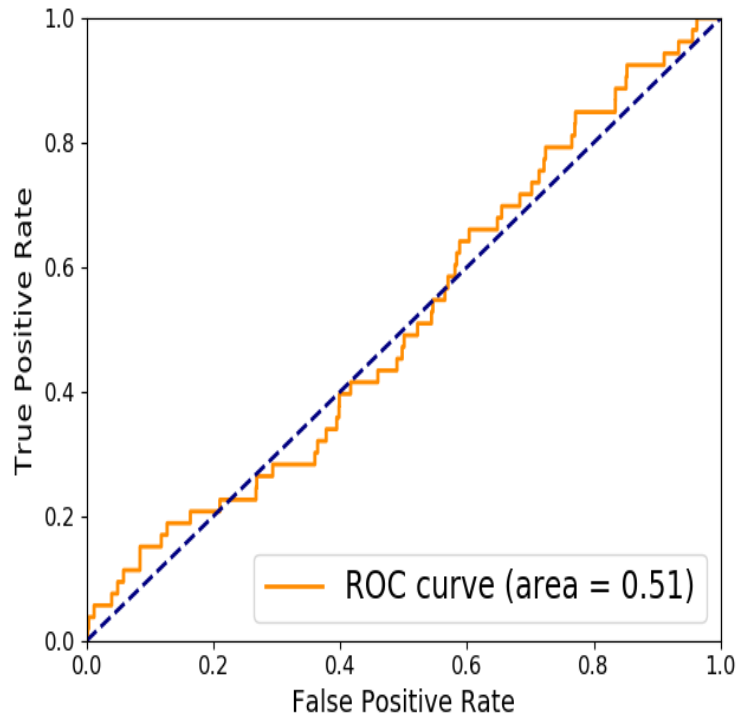
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.503	0.251	0.014	0.814	0.783
Test_only	GS_only	0.654	1.108	1.229	0.639	0.483
GS_only	Overlap	0.998	3.7474606	1.7114571	1.965	1.637808
		07303235	615977714		69636599	
		(invalid)	(invalid)		06 (inva	
					lid)	
GS_only	Test	0.513	0.112	0.452	1.158	1.026
Test_only	Overlap	0.746	-1.220139	0.8057939	-1.243505	-0.78481
		212700807	080705562	347429740	66936716	
		3	(invalid)	8	021 (inv	
		(invalid)		(invalid)	alid)	
Test_only	GS	0.667	0.715	1.450	0.250	0.261
Overlap	Onlies	0.557	1.801	1.296	8.5857097	2.008885
				74504918	87550021	
				(invalid)	4 (inval	
					id)	

**	#Mapped	#Total	Not mapped genes
GS	6	6	N/A
Test_only	52	53	IQCF6
Overlap	1	1	APOE

(end of file)