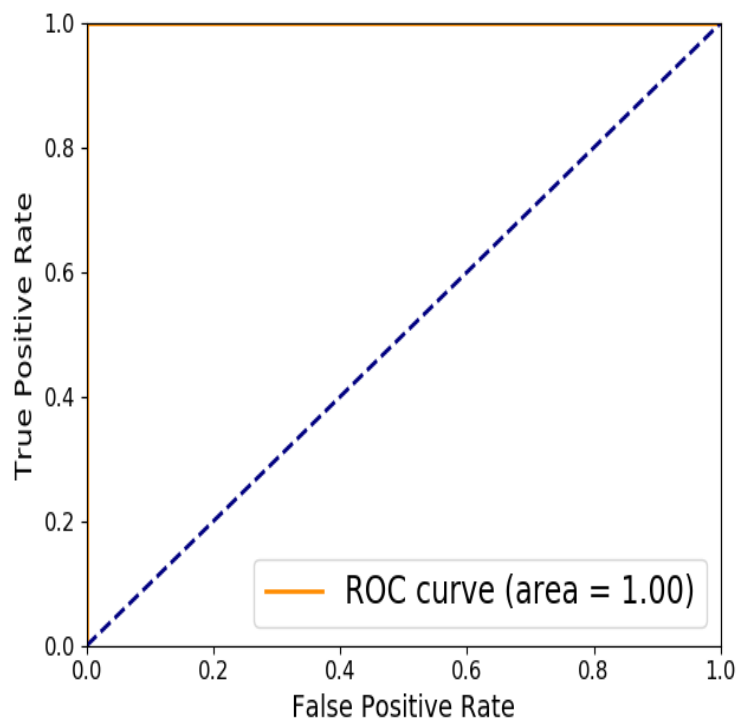
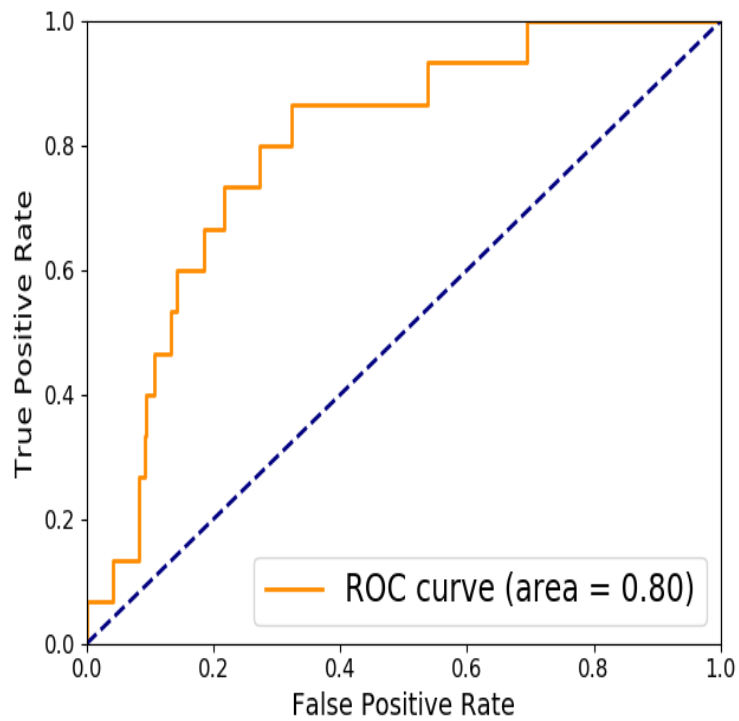


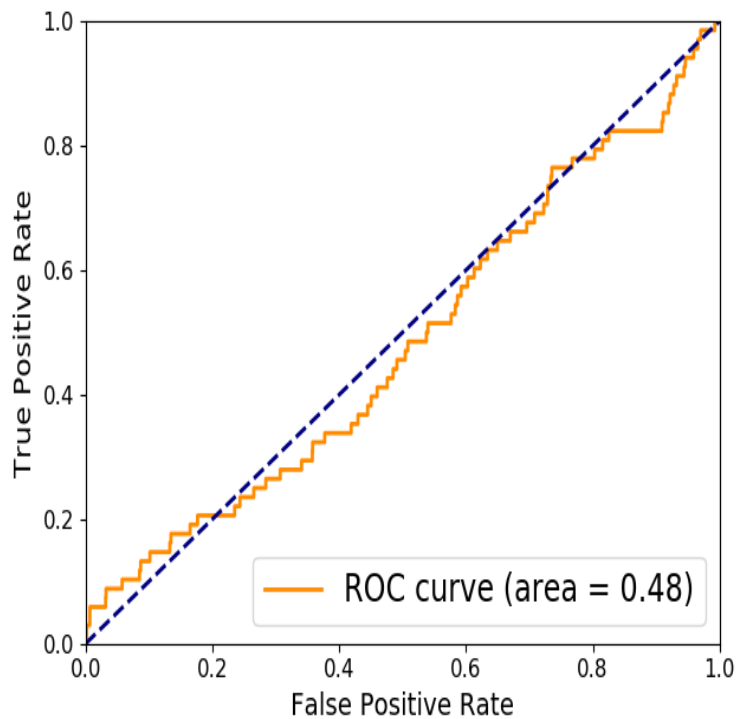
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



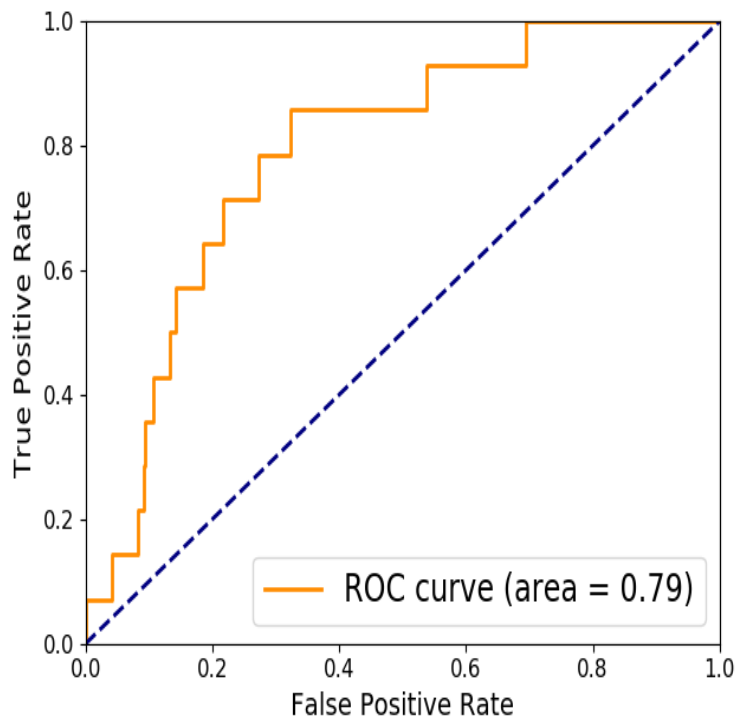
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



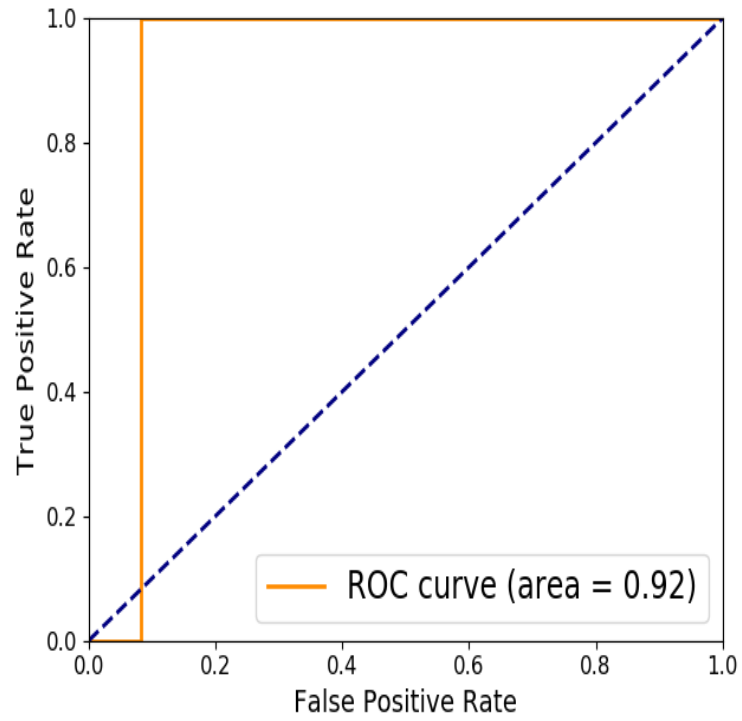
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



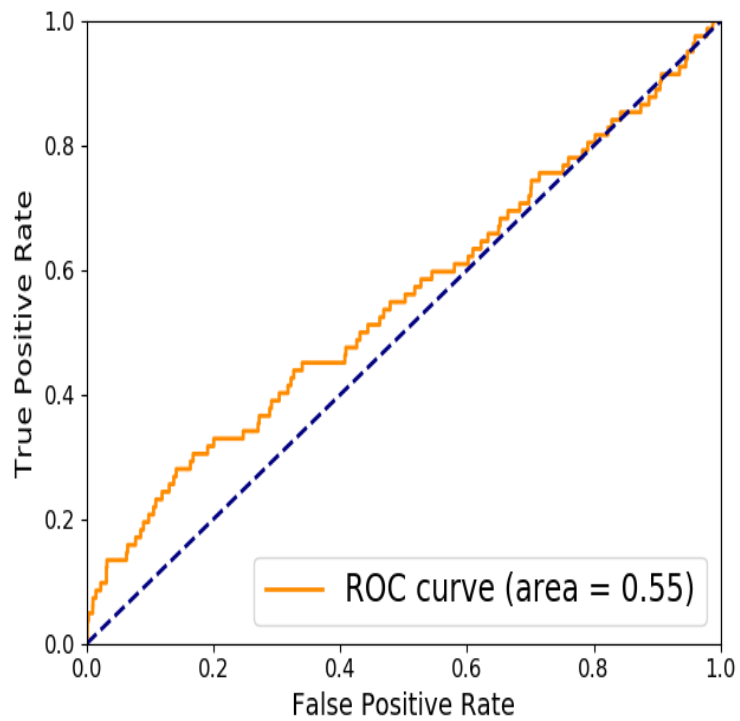
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



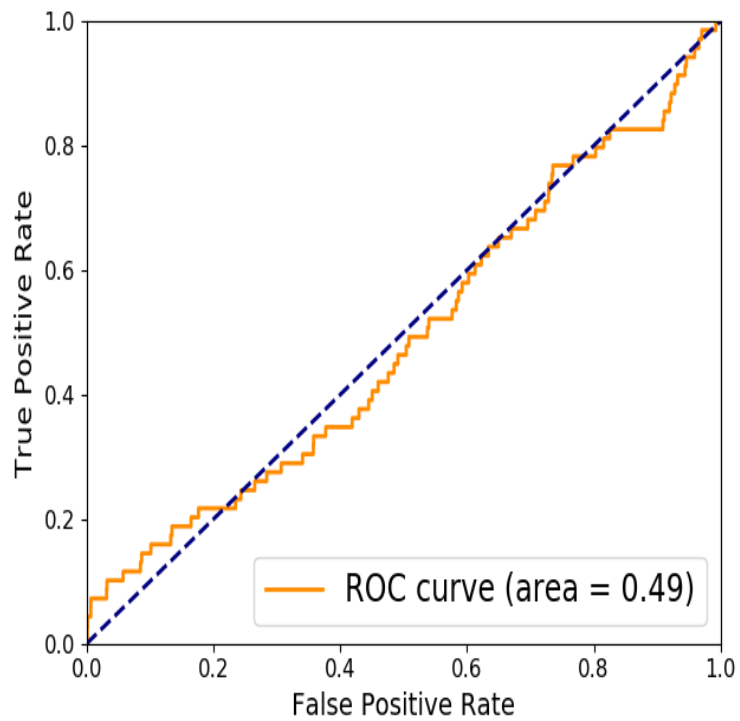
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.482	-0.949	-0.542	-1.072	-0.281
Test_only	GS_only	0.791	2.740	3.489	2.047	2.048
GS_only	Overlap	0.999	1.1068704	1.5361421	2.1339698	2.072068
		411565975	647991789	077306593	22670497	
		(invalid)	(invalid)	(invalid)	15 (inva	
				lid)		
GS_only	Test	0.489	-0.866	-0.399	-1.133110	-0.154
				102742766		
				6		
				(invalid)		
Test_only	Overlap	0.917	1.7369947	1.3076316	0.6994772	0.721098
		373451744	721548675	057612074	12843417	
		(invalid)	(invalid)	(invalid)	41 (inva	
				lid)		
Test_only	GS	0.800	2.4615235	4.256	2.064	1.853
		1588789				
		(invalid)				
Overlap	Onlies	0.552	0.441	1.832	1.4520885	1.133
				573625997		
				(invalid)		

**	#Mapped	#Total	Not mapped genes
GS	14	15	CAMK1V N/A
Test_only	68	71	KMT2C;KRTAP10.9;PRR3 N/A
Overlap	1	1	APOE

(end of file)