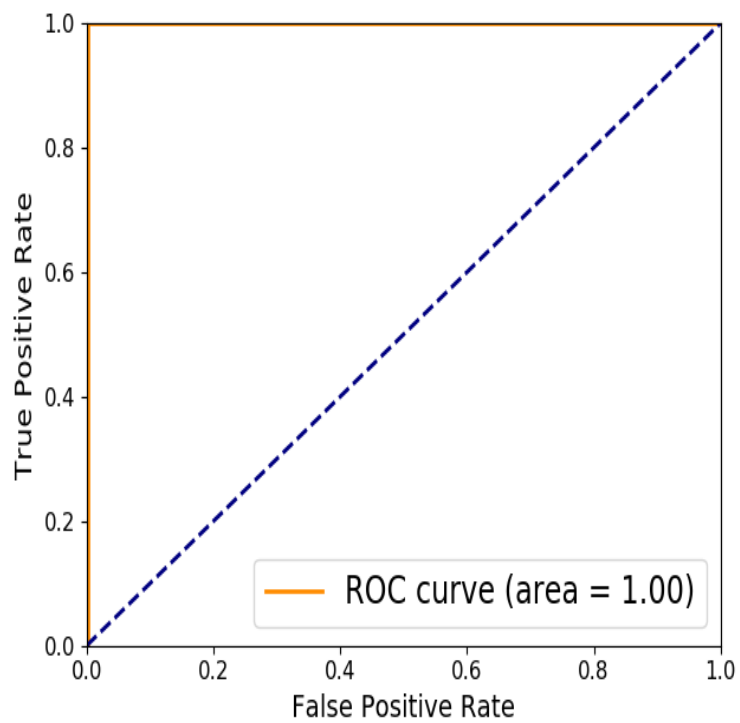
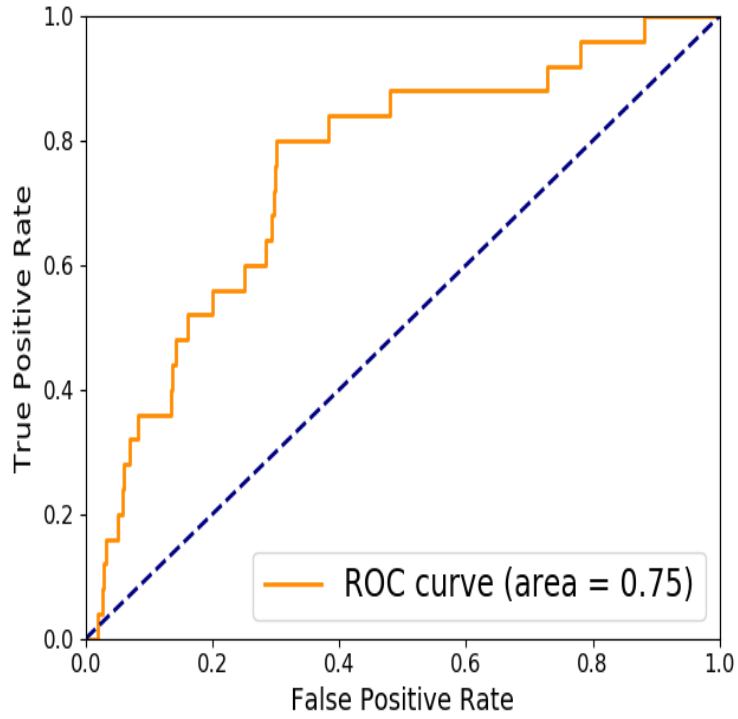


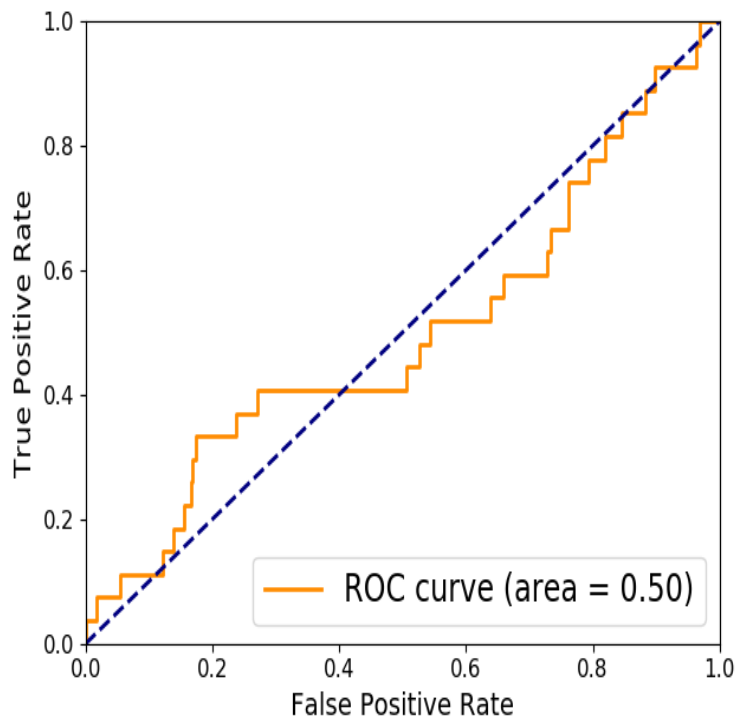
## Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



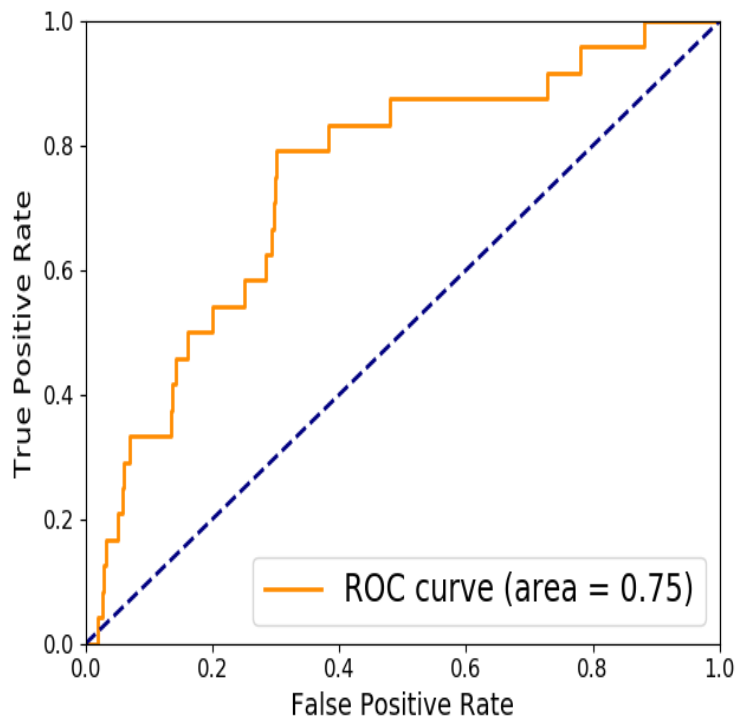
# Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



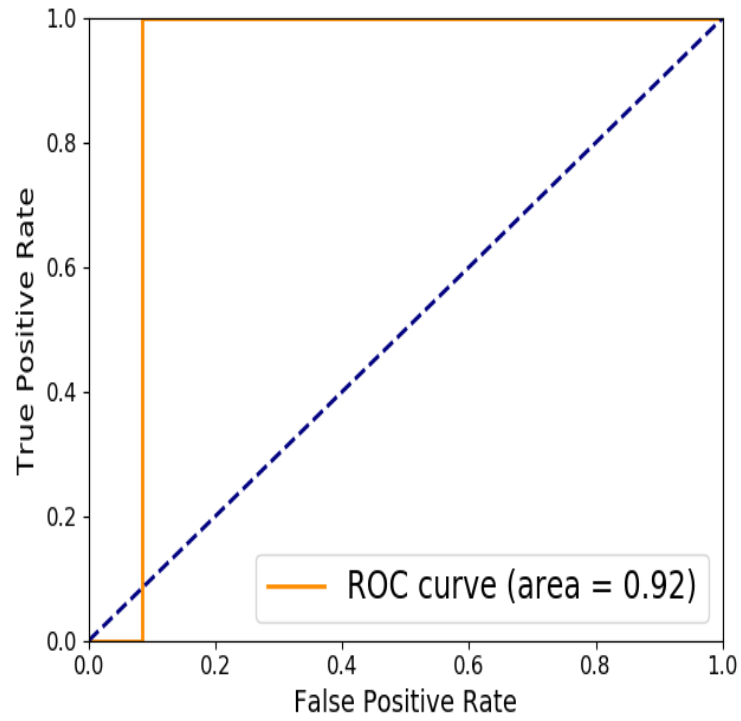
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



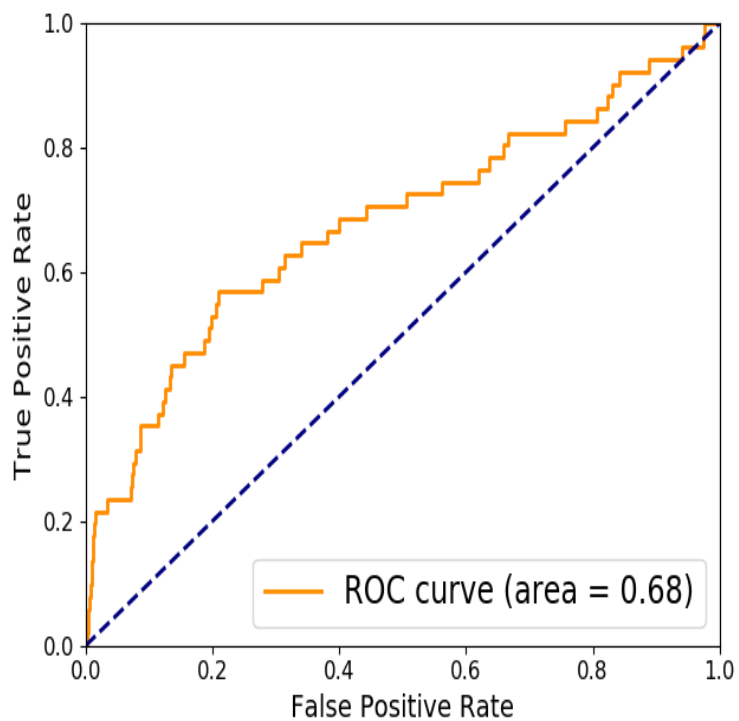
# Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



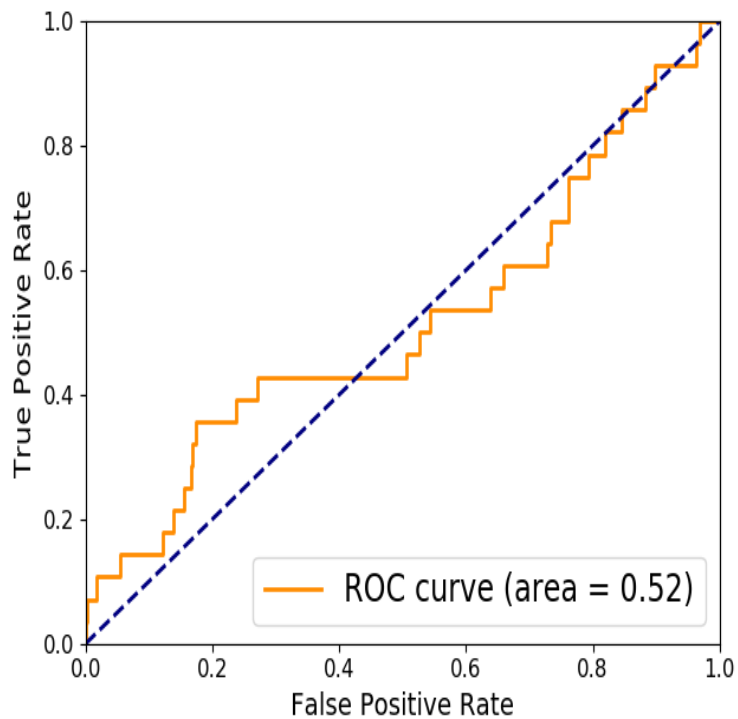
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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	0.499	1.664	-0.095	1.549	1.564
Test_only	GS_only	0.746	3.884	4.0838276	2.422	2.487356
			26825669		23715427	
			(invalid)		9 (inval	
					id)	
GS_only	Overlap	0.998	5.3506295	2.0748890	1.7398185	1.867945
			42852921	35455198	59770431	04126353
			(invalid)	(invalid)	(invalid)	32 (inva
						lid)
GS_only	Test	0.517	1.884	0.298	1.351	1.501
Test_only	Overlap	0.916	2.2187528	1.4711324	1.0323726	0.980023
			741729095	547935951	534832784	89283045
			(invalid)	(invalid)	(invalid)	61 (inva
						lid)
Test_only	GS	0.753	3.510	4.393	2.5761951	2.881277
					13375177	41448007
					(invalid)	7 (inval
						id)
Overlap	Onlies	0.680	6.113	4.678	5.8115012	2.391
					12419781	
					(invalid)	

**	#Mapped	#Total	Not mapped genes
GS	24	24	N/A
Test_only	27	27	N/A
Overlap	1	1	APOE

(end of file)