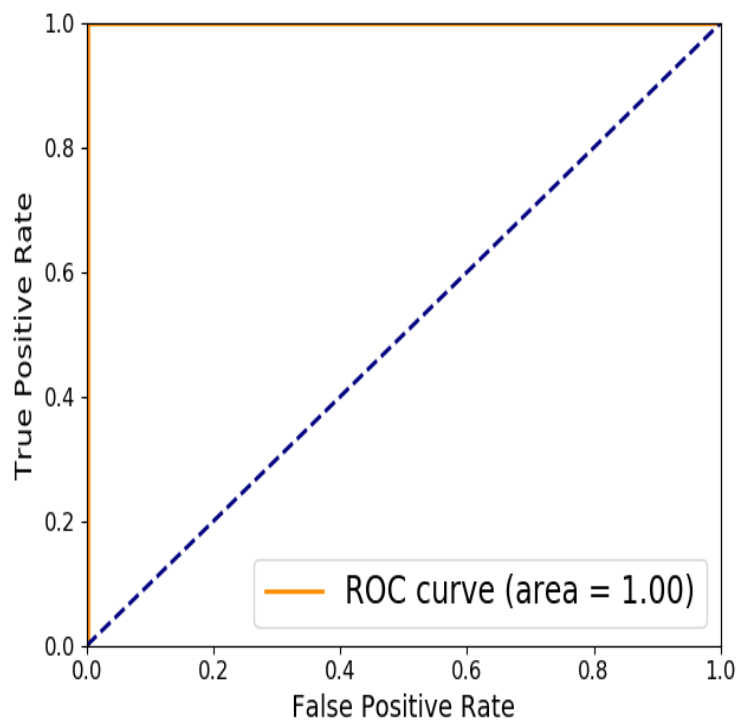
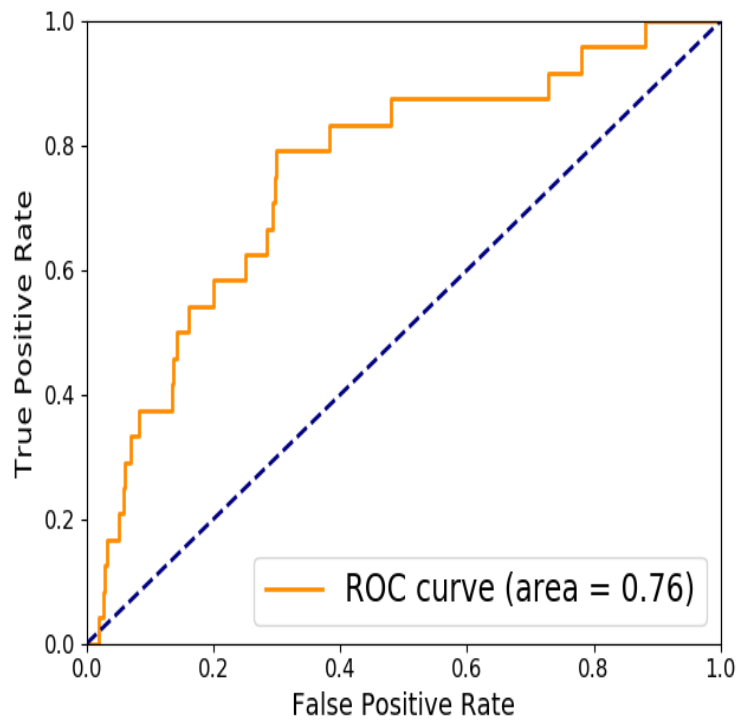


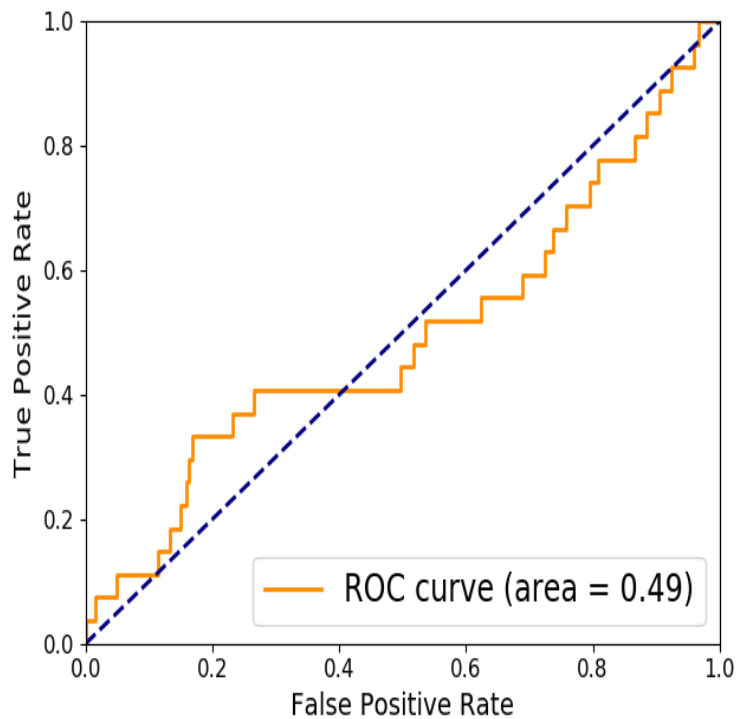
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



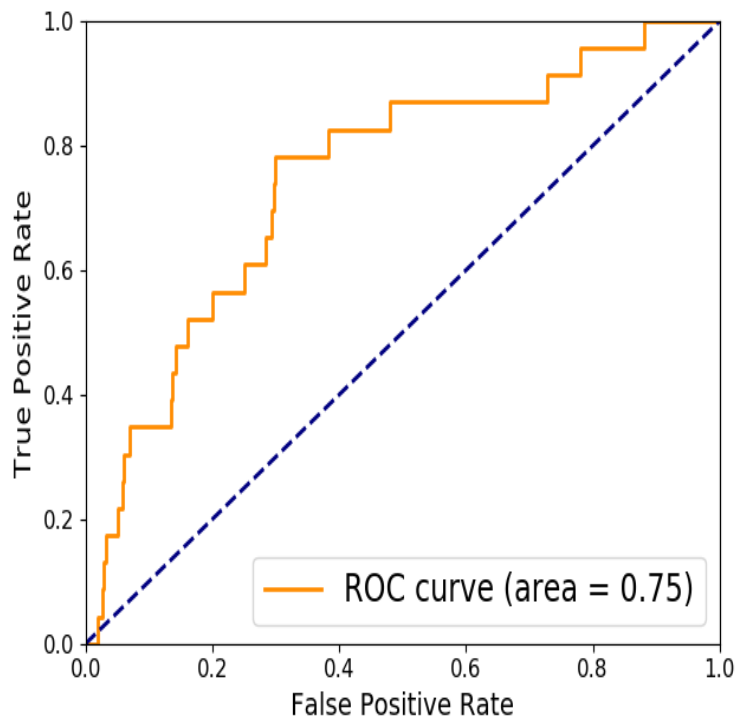
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



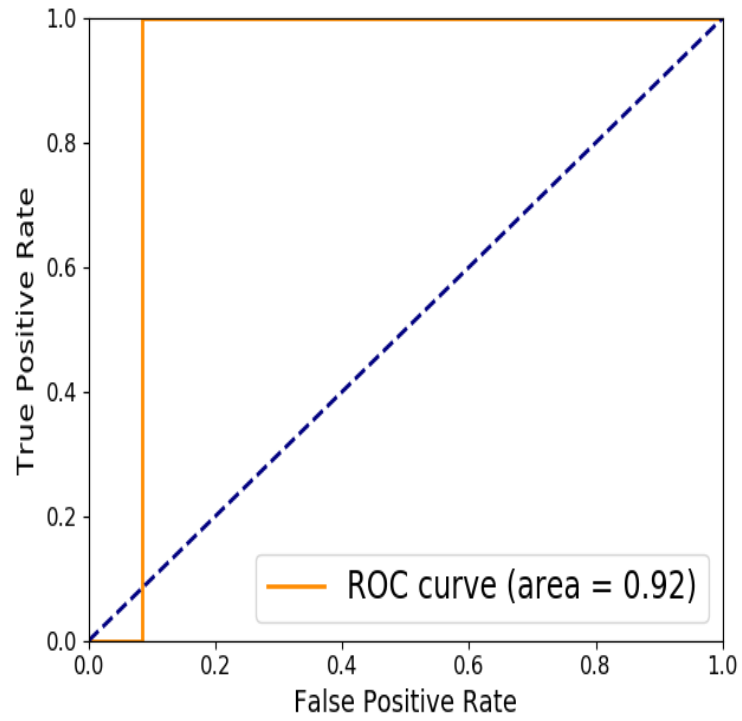
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



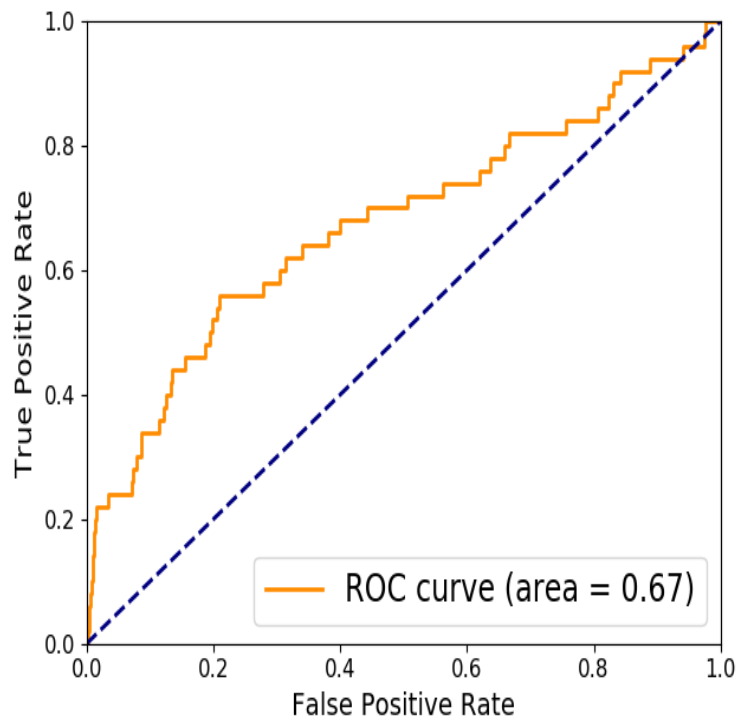
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



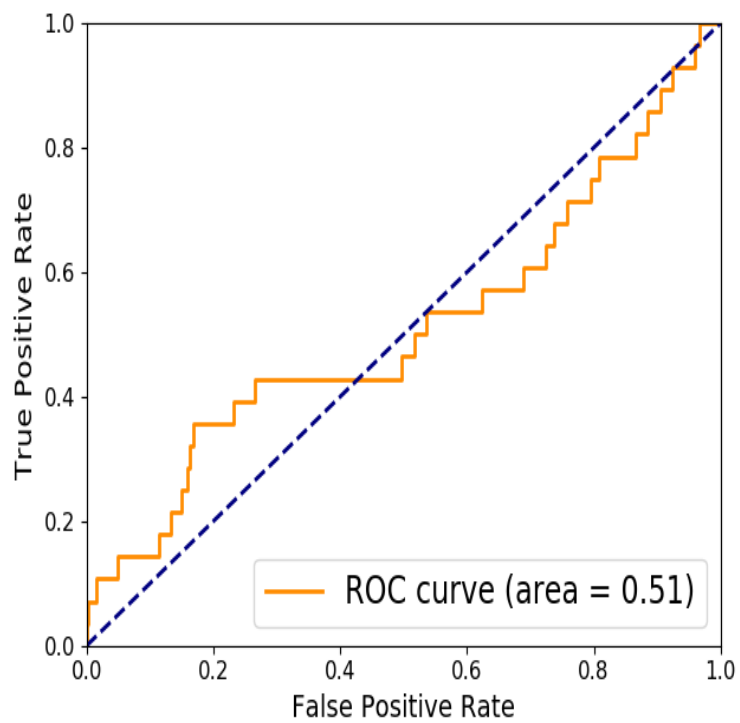
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.495	1.597	-0.221	1.304	1.550
Test_only	GS_only	0.748	3.434	4.493	2.139	2.531
GS_only	Overlap	0.998	5.1386184	1.7785802	1.6659353	1.713829
			32789018	003781561	215137314	06846686
			(invalid)	(invalid)	(invalid)	8 (inval
						id)
GS_only	Test	0.513	1.474	0.201	1.704	1.426448
						60451678
						46 (inva
						lid)
Test_only	Overlap	0.916	2.2135086	1.4368914	1.138	0.960922
			181472476	547179248		63805162
			(invalid)	(invalid)		98 (inva
						lid)
Test_only	GS	0.755	3.837	4.787	2.4386547	2.581660
						06351434
						07375484
						(invalid)
						6 (inval
						id)
Overlap	Onlies	0.675	4.861	4.173	5.9611964	2.437287
						22780455
						10844516
						(invalid)
						4 (inval
						id)

**	#Mapped	#Total	Not mapped genes
GS	23	24	HLA -DRB1
			N/A
Test_only	27	27	
			N/A
Overlap	1	1	APOE

(end of file)