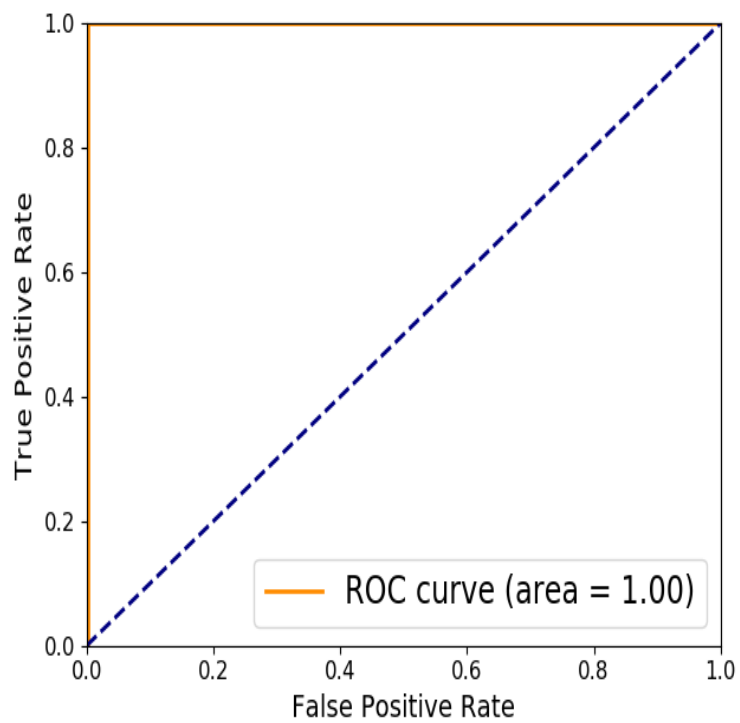
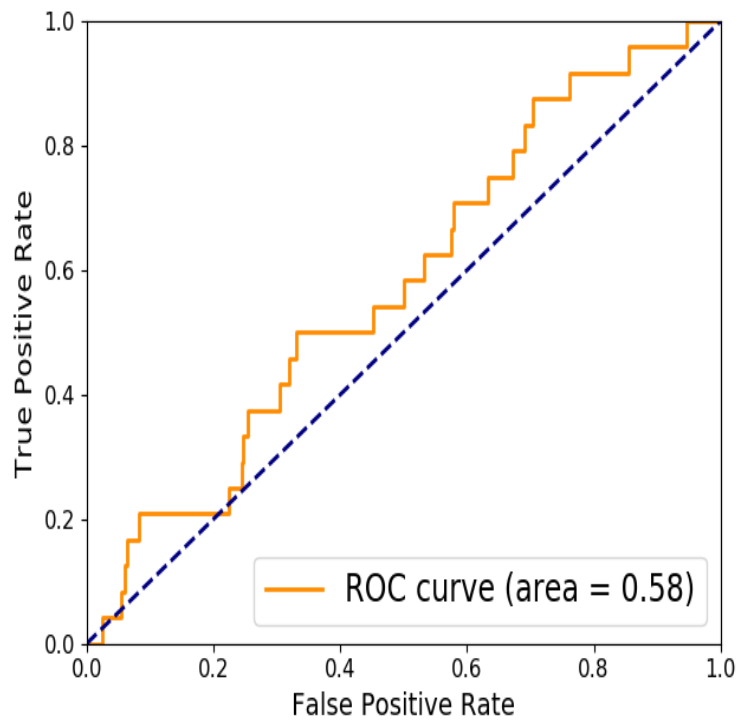


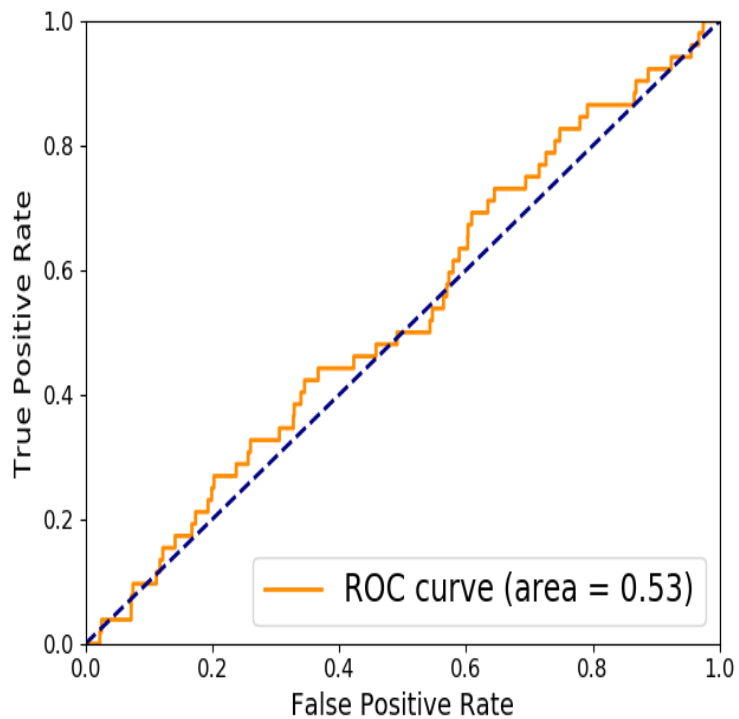
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



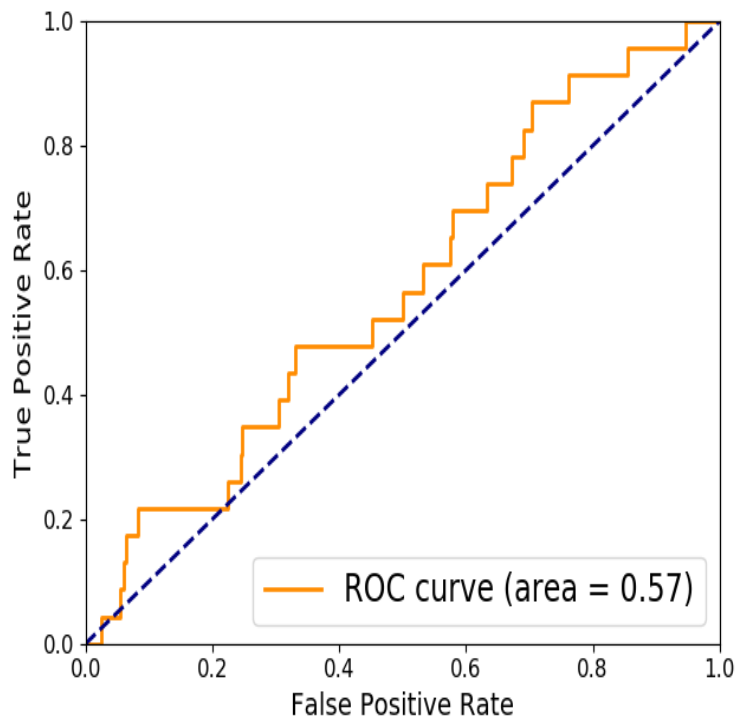
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



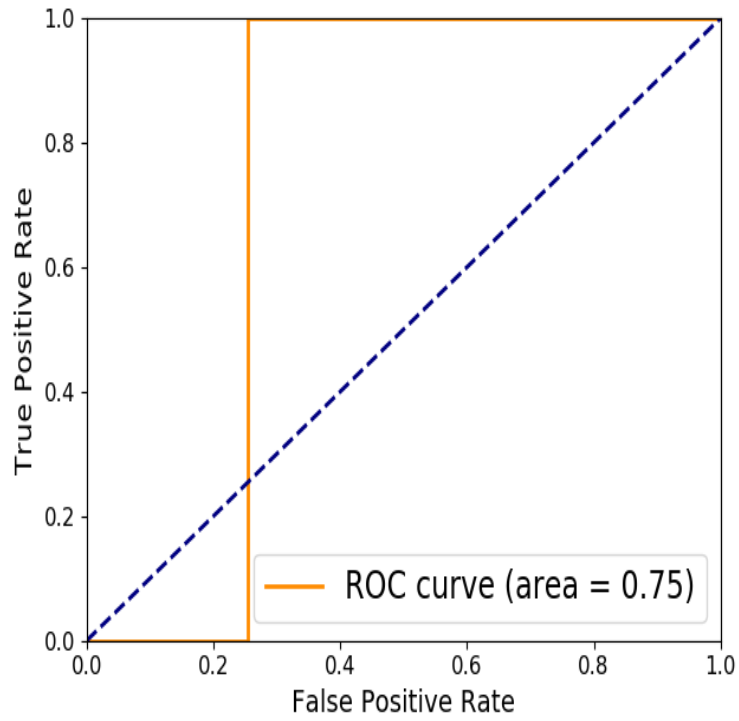
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



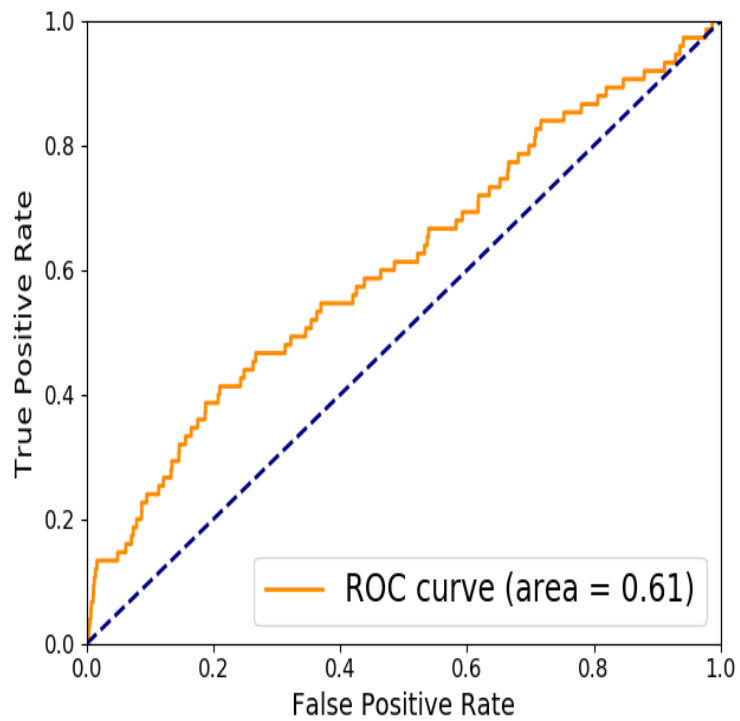
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



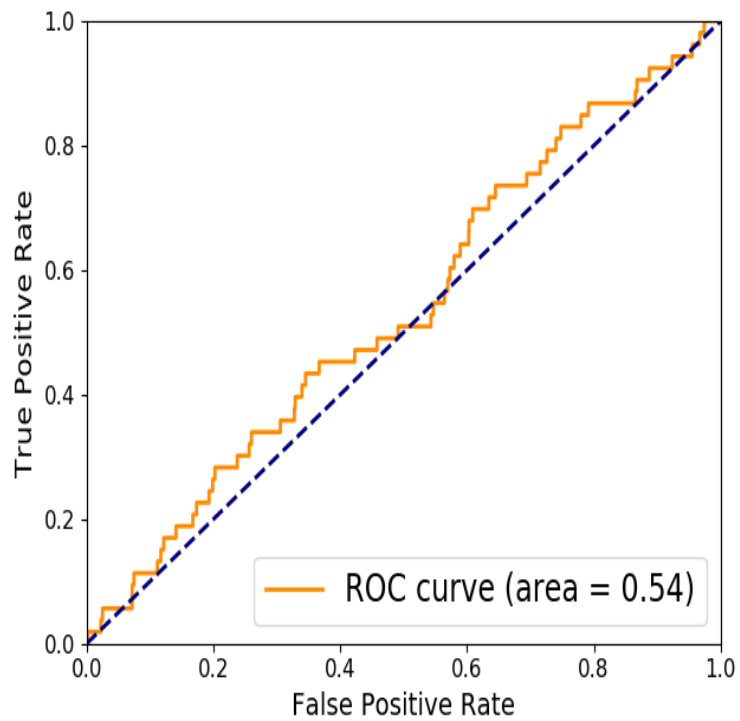
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.528	0.414	0.664	1.062	1.332
Test_only	GS_only	0.572	-0.091	1.080	-0.475	-0.262
GS_only	Overlap	0.998	5.2057860	1.8704939	1.7299362	2.027321
			25460623	519980968	511255523	31773589
			(invalid)	(invalid)	(invalid)	1 (inval
						id)
GS_only	Test	0.537	0.560	0.850	1.294	1.464
Test_only	Overlap	0.746	-1.177672	0.8205637	-1.223582	-0.99869
			010965928	847761259	920693096	71909149
			9	(invalid)	8	187 (inv
			(invalid)		(invalid)	alid)
Test_only	GS	0.579	-0.126	1.972	-0.701	-0.258
Overlap	Onlies	0.609	3.299	3.359	77.686338	1.889
					19682357	
					(invalid)	

**	#Mapped	#Total	Not mapped genes
GS	23	24	HLA -DRB1 N/A
Test_only	52	53	IQCF6 N/A
Overlap	1	1	APOE

(end of file)