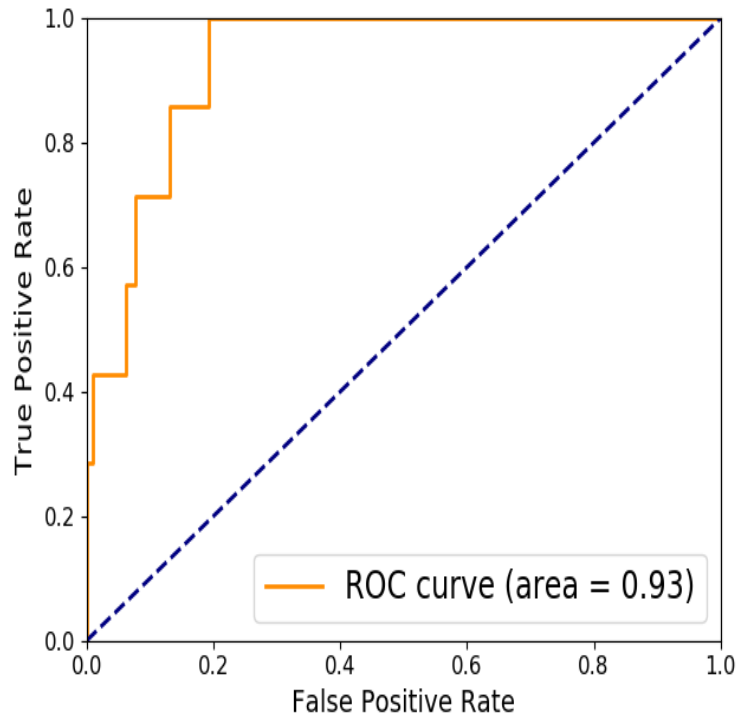
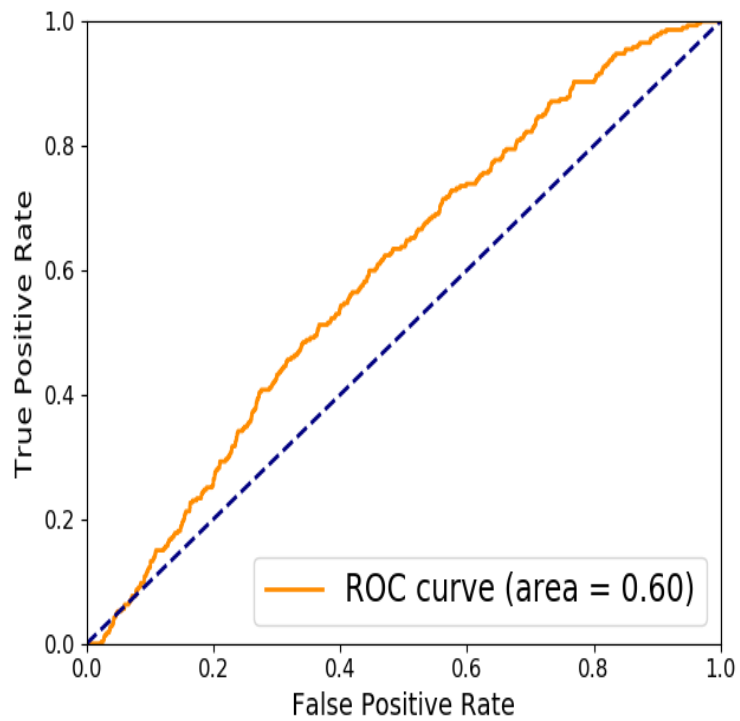


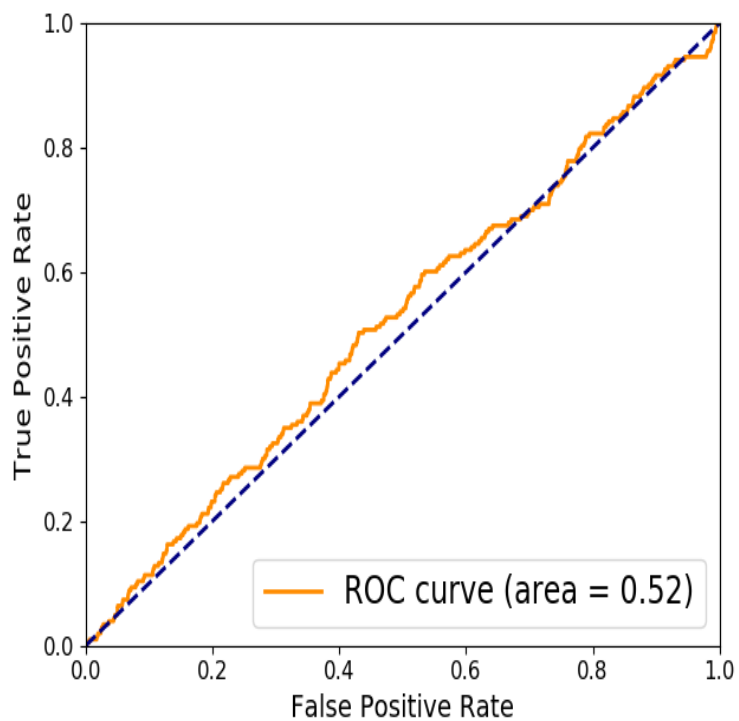
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



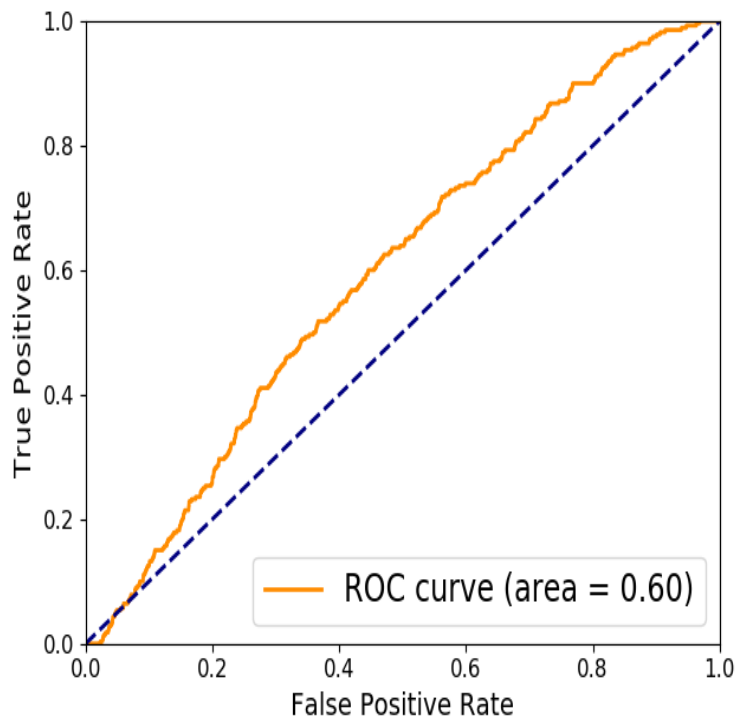
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



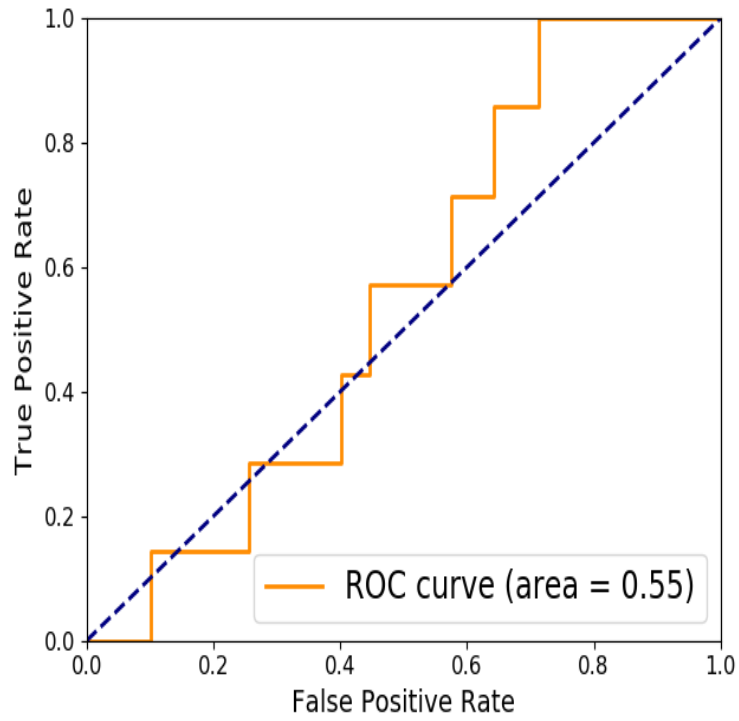
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



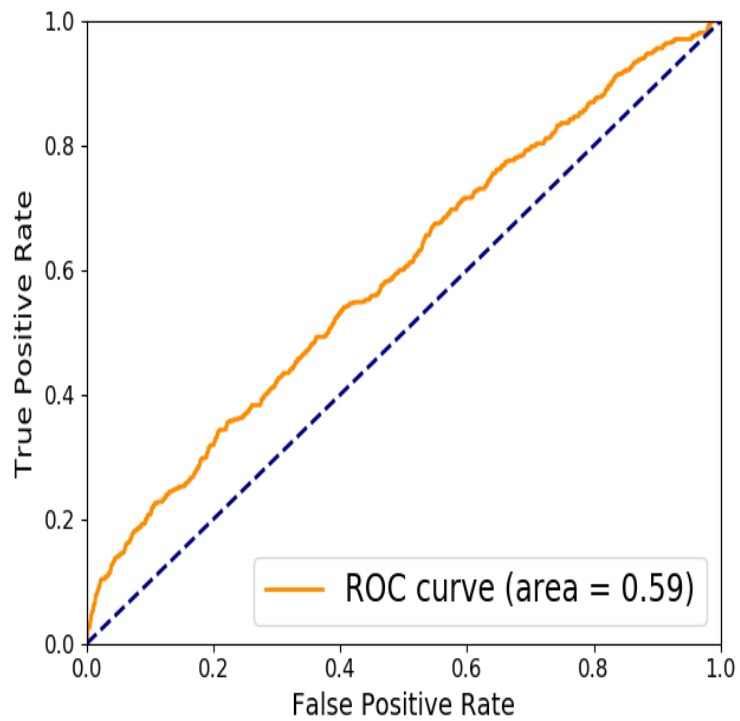
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



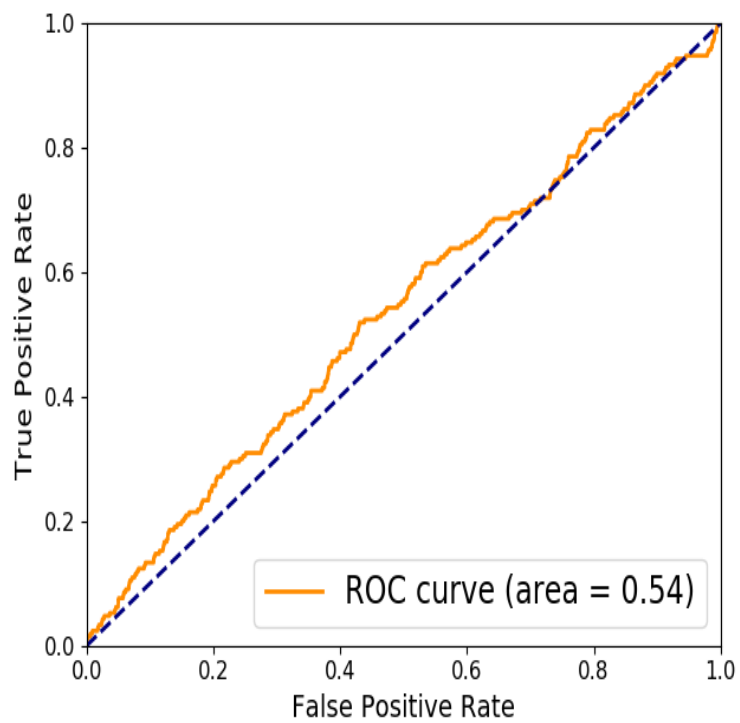
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



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	0.522	1.278	1.011	1.277	0.326
Test_only	GS_only	0.597	2.962	6.155	2.315	2.237
GS_only	Overlap	0.933	5.062	4.473	4.878	4.407
GS_only	Test	0.536	1.890	2.2425401	2.518	1.484
			05217623			
			(invalid)			
Test_only	Overlap	0.552	-0.048	0.490	-0.051	-0.203
Test_only	GS	0.596	2.641	5.637	2.398	1.841416
				71141317		
				05 (inva		
				lid)		
Overlap	Onlies	0.595	5.786	6.5033848	2.708	2.418
			55467717			
			(invalid)			

**	#Mapped	#Total	Not mapped genes
GS	280	295	ABO;PRIM2;C4orf14;PKD N/A
			1L3;PECAM1;MGC34034;C
			6orf105;CDKN2BAS;RPL2
			3AP7;MIR3134;RECQL4;P
			OM121L9P;MIR3169;C4or
			f41;LPAL2
Test_only	203	211	ADGRF3;LENG9;DNAH12;R N/A
			PEL1;AZIN2;CCDC181;HE
			ATR9;FAM177B
Overlap	7	7	FBN2;REST;MIA3;LDLR;S
			MG6;LIPA;HSPB9

(end of file)