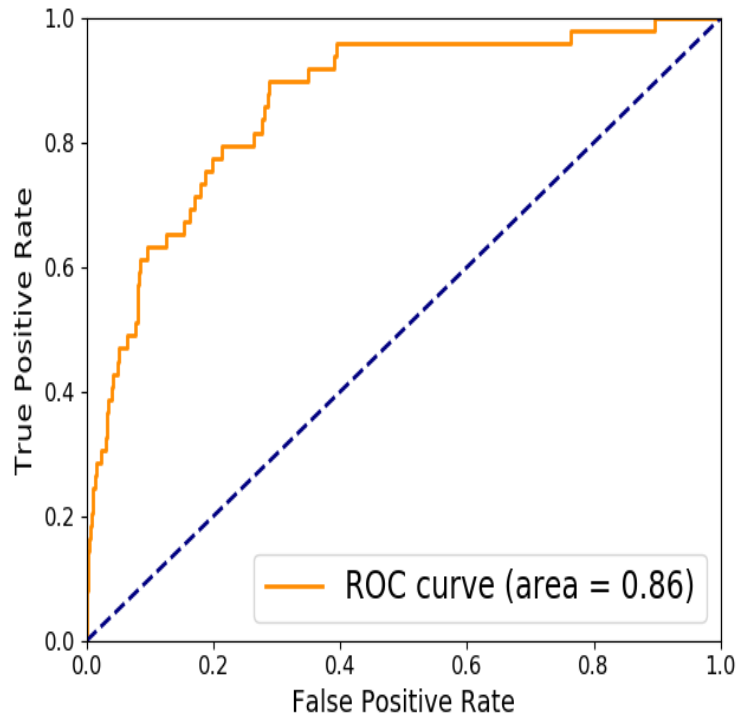
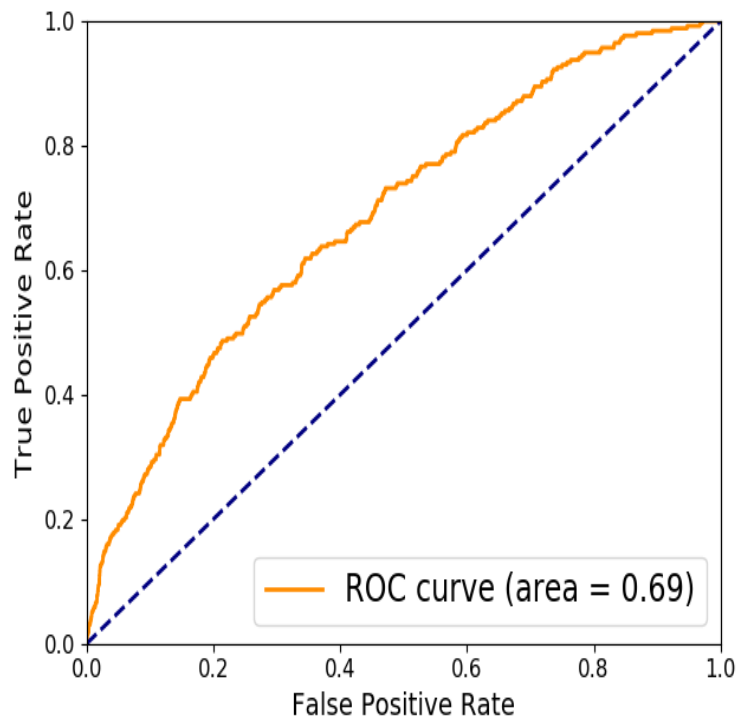


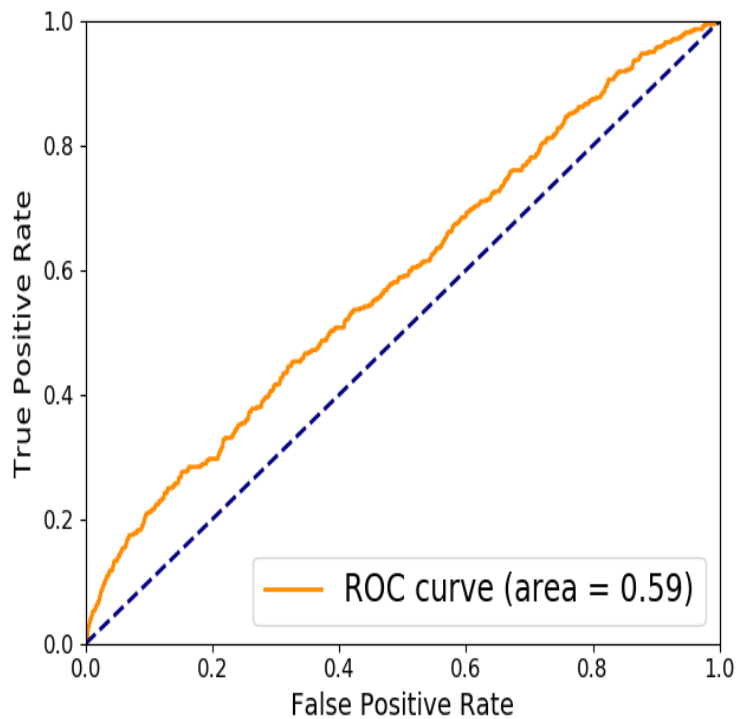
# Diffusion ROC from fGOnly\_tOverlap STRING10\_combined.png



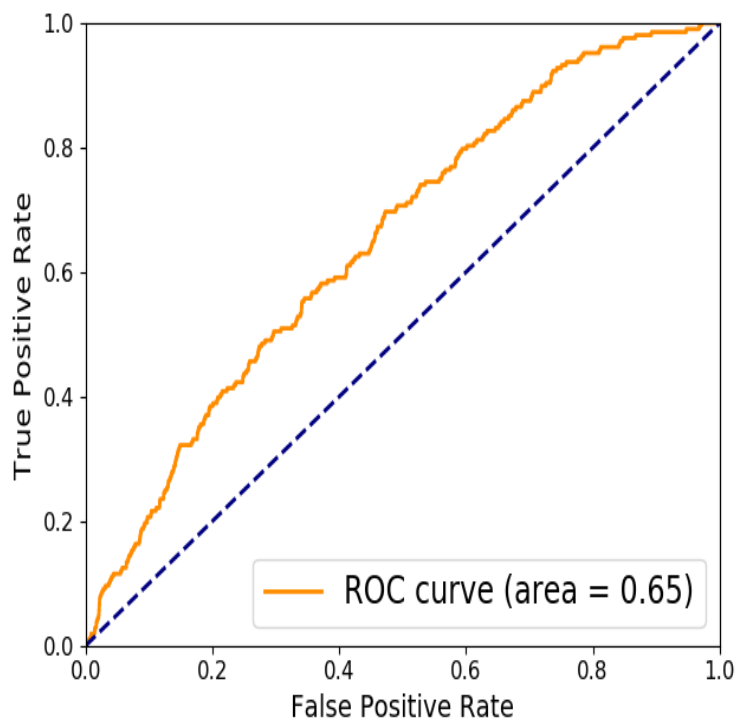
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



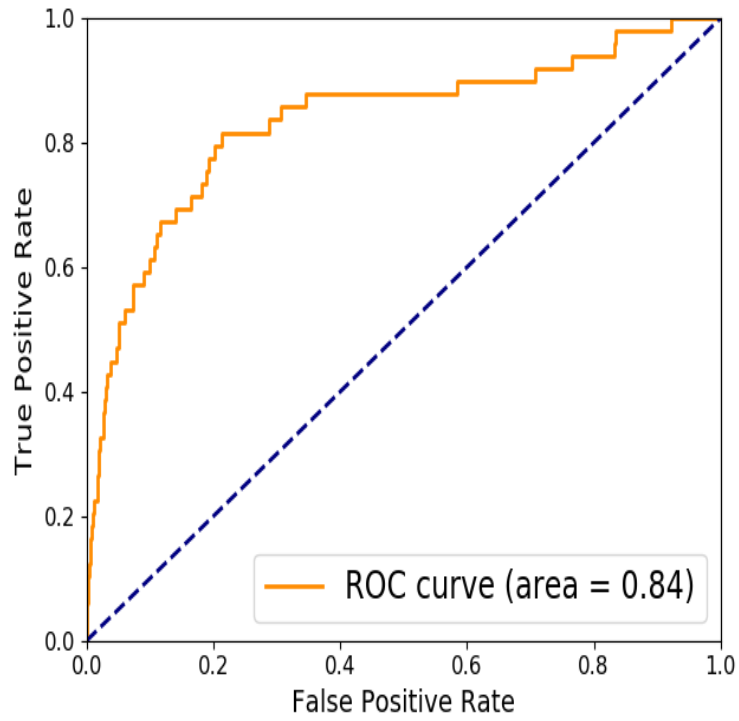
# Diffusion ROC from fGOnly\_tTestOnly STRING10\_combined.png



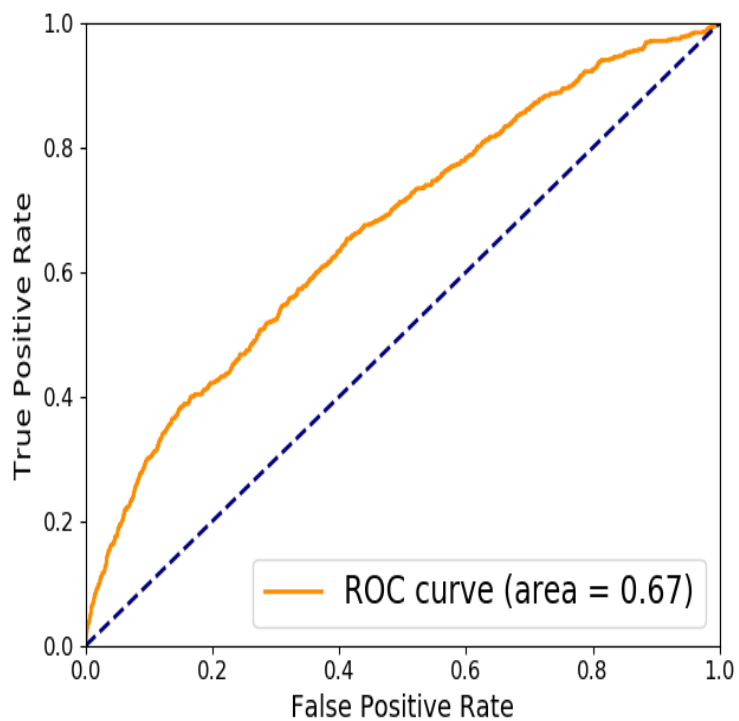
## Diffusion ROC from fTestOnly\_tGOnly STRING10\_combined.png



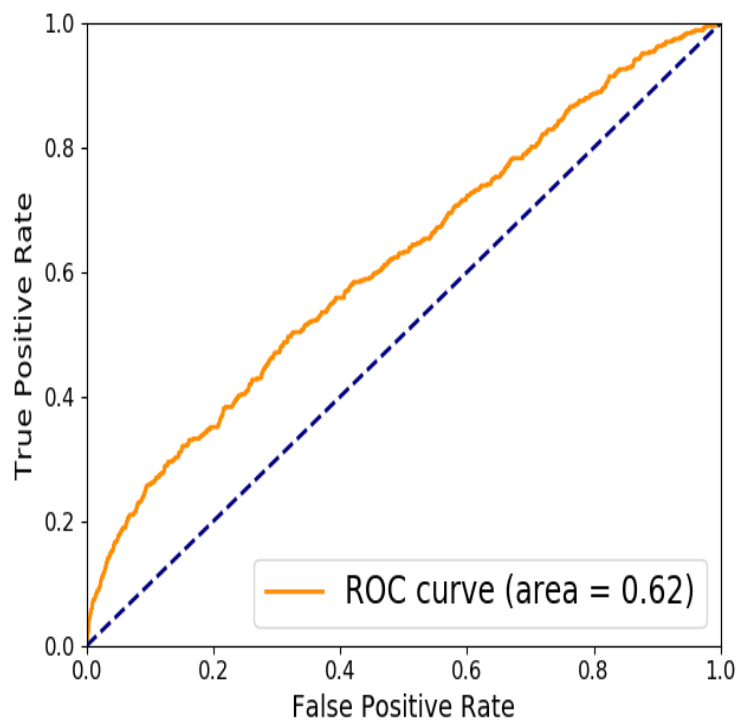
# Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## Diffusion ROC from fOverlap\_tOnlies STRING10\_combined.png



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



Diff_FROM	Diff_TO	AUC	Zt_wdegree	Zt	Zf_wdegree	Zf
GS_only	Test_only	0.585	0.3752828957	6.199	0.051	0.692
		5907153				
		(invalid)				
Test_only	GS_only	0.652	4.664	7.358	3.083	2.912
GS_only	Overlap	0.865	5.389	10.901	3.665	5.093
GS_only	Test	0.618	1.852	8.233	0.875	1.850
Test_only	Overlap	0.835	4.057	8.208	2.6028156562	3.969
				27693		
				(invalid)		
Test_only	GS	0.688	5.264	10.559	3.218	4.210
Overlap	Onlies	0.668	9.388	16.514	5.719	5.945

**	#Mapped	#Total	Not mapped genes
GS	208	229	KDF1;RPA;RFC;MSH5;PMS1
			2L3;AMER1;AK9
Test_only	384	405	KMT2B;Gene;SLF2;PAXBP
			1;ADGRL2;AMER1;BRINP1
			;ACP7;ADGRA3;JCAD;RMD
			N1;CFAP54;KMT2D;DISP3
			;CCDC168;KNSTRN;KMT2C
			;ADGRB3;DNAH12;HLA.A;
			ADGRG4;ADGRV1
Overlap	49	50	AMER1
			CHD8;SIN3A;AXIN1;ZBTB
			20;REG3G;NFE2L2;BRAF;
			CYLD;PPP2R1A;GTF2I;AM
			ER1;PTPRD;RET;PMS1;GN
			A11;SMARCA4;EPAS1;SF3
			B1;GNAS;BAP1;MGA;ERBB
			2;KRAS;RUNX1;PIK3R1;M
			ET;EEF1A1;NOTCH1;SOX9
			;NOTCH2;MED12;RAC1;AT
			M;RASA1;CRYBG3;BRCA2;
			KIT;ARHGAP35;KEAP1;ZN
			F750;TP53;APC;ASXL2;C
			HD4;KDM5C;BRCA1;COL4A
			3;RBBP8;AMOT;MACF1



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*(end of file)*