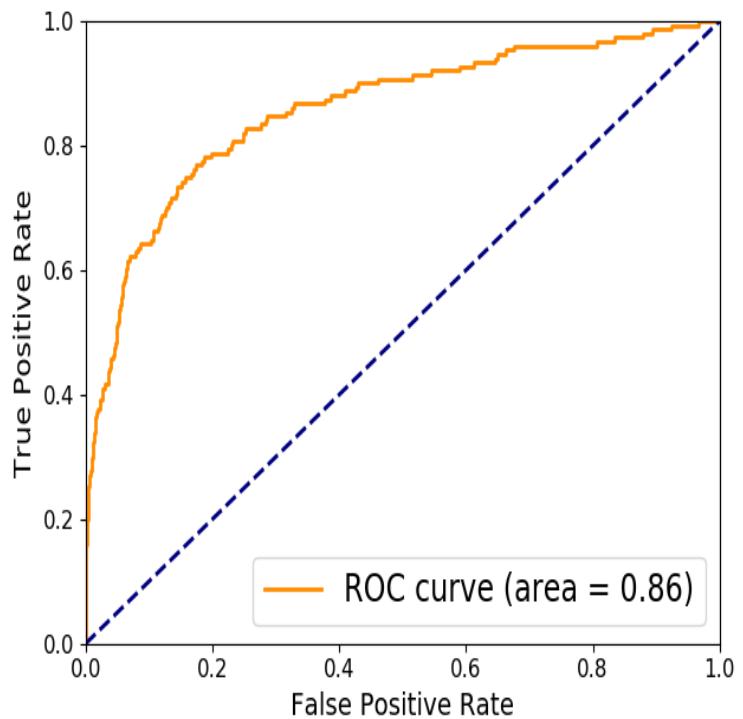
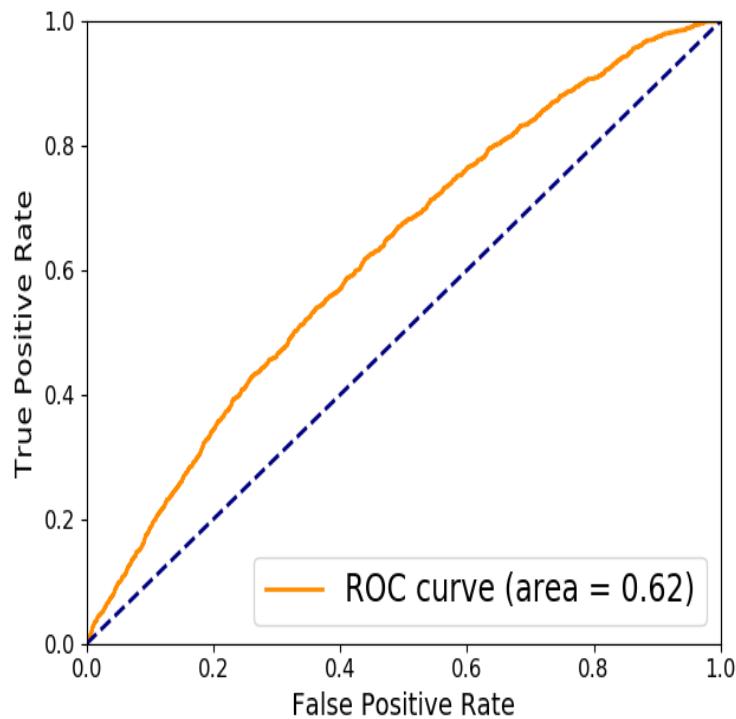


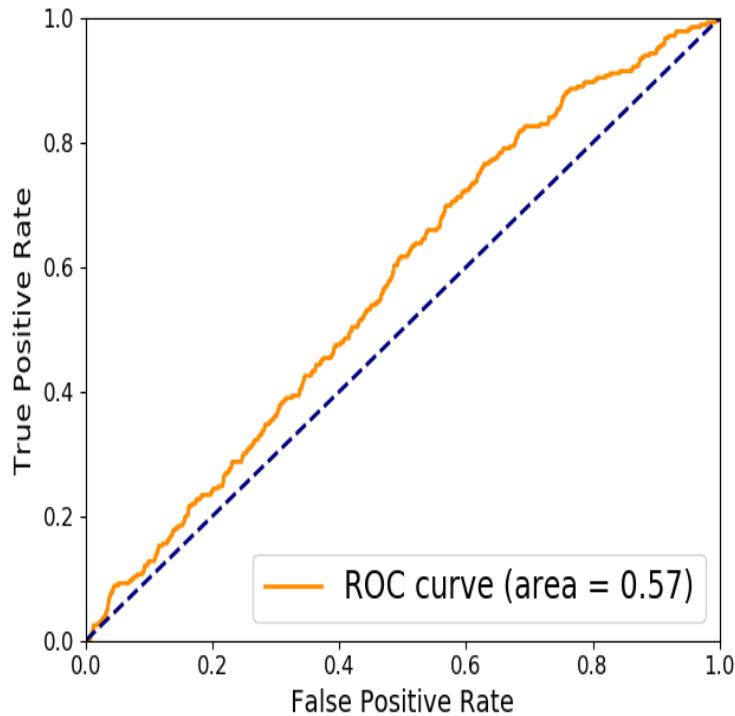
Diffusion ROC from fGSonly_tOverlap STRING10_combined.png



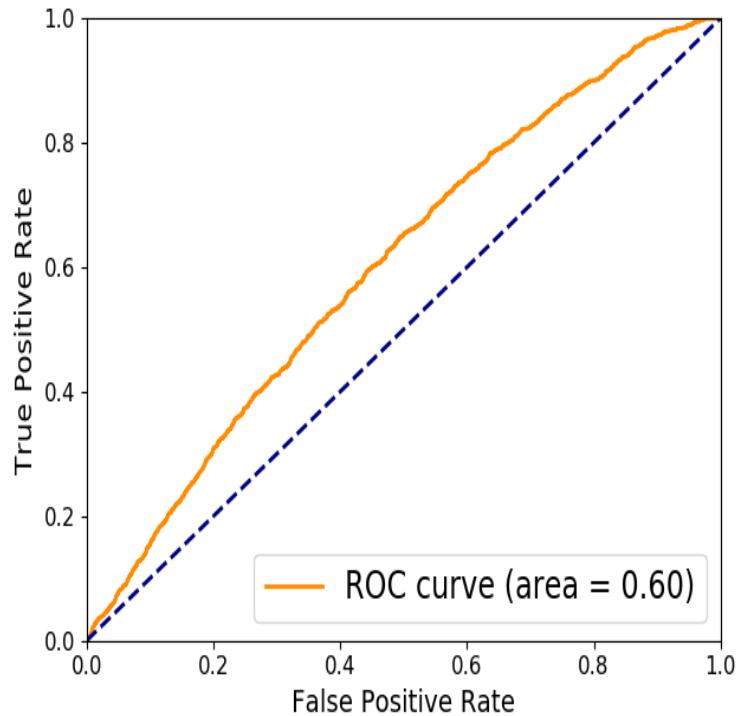
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



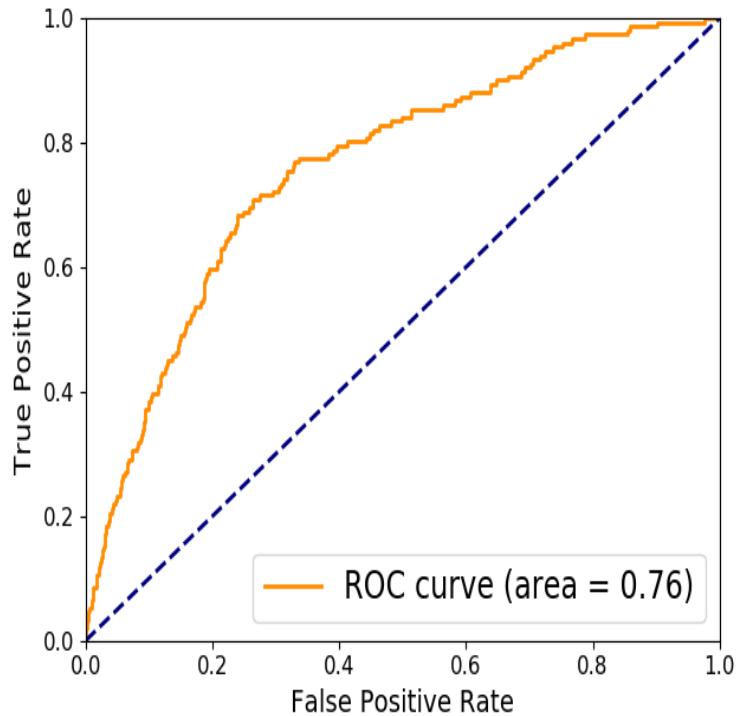
Diffusion ROC from fGSonly_tTestOnly STRING10_combined.png



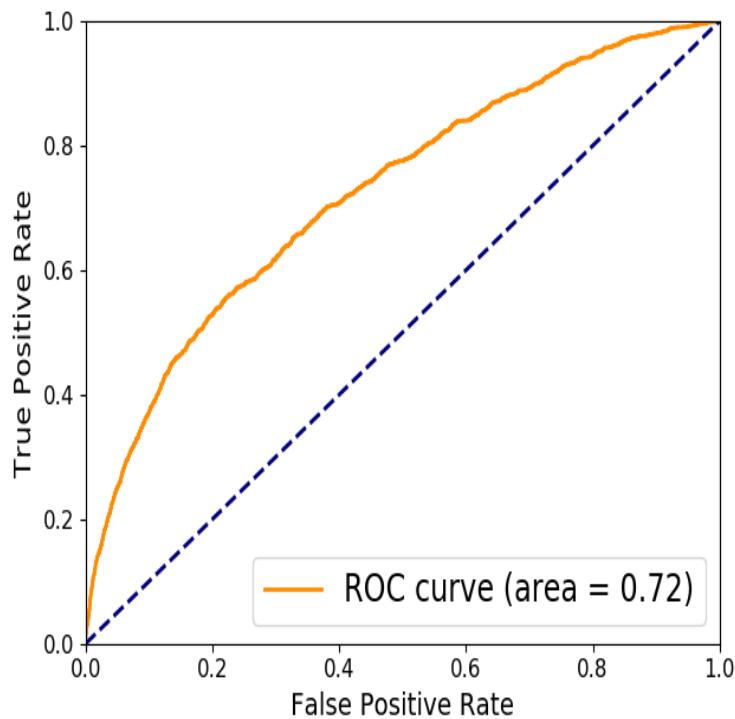
Diffusion ROC from fTestOnly_tGSonly STRING10_combined.png



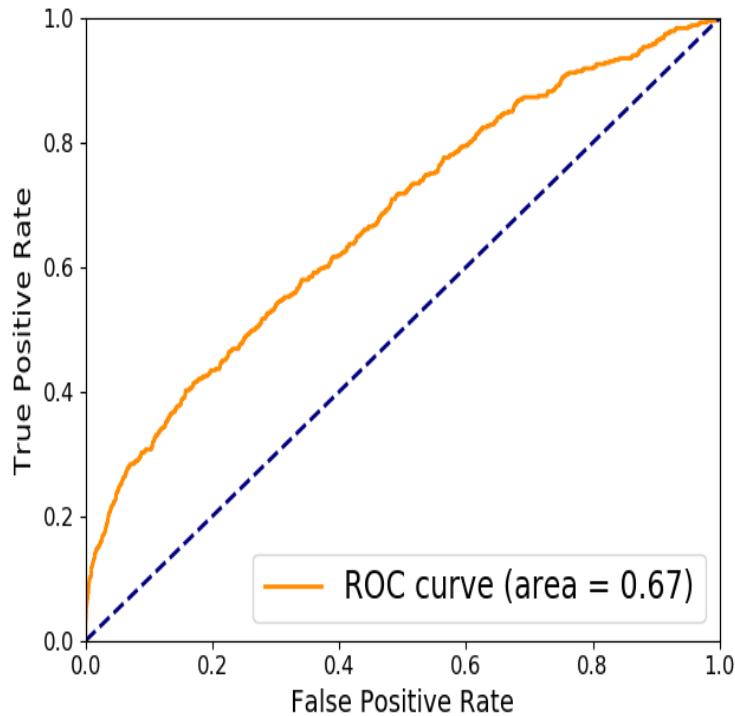
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



Diffusion ROC from fOverlap_tOnlies STRING10_combined.png



Diffusion ROC from fGSonly_tTest STRING10_combined.png



<i>Diff_FROM</i>	<i>Diff_TO</i>	<i>AUC</i>	<i>Zt_wdegree</i>	<i>Zt_e</i>	<i>Zf_wdegre</i>	<i>Zf</i>
<i>GS_only</i>	<i>Test_only</i>	0.572	1.720	4.771	3.089	2.187
<i>Test_only</i>	<i>GS_only</i>	0.604	-0.091	14.585	0.379	1.628
<i>GS_only</i>	<i>Overlap</i>	0.859	12.282	15.836	13.464	9.484
<i>GS_only</i>	<i>Test</i>	0.675	6.05168165	12.594	8.724	6.228
		5875711				
		(invalid)				
<i>Test_only</i>	<i>Overlap</i>	0.764	5.832	13.0482383	3.4528125	3.679
		68235378		639169978		
		(invalid)	(invalid)			
<i>Test_only</i>	<i>GS</i>	0.624	1.217	15.7749793	1.224	1.900
		29625605				
		(invalid)				
<i>Overlap</i>	<i>Onlies</i>	0.724	17.838	33.784	15.949905	12.273
				59808249		
				(invalid)		

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>		
<i>GS</i>	1219	1264	;SFRS2;MYCL;IGK;TRB;R N/A		
			ECQL4;IGL;MSH5;DNAH12		
			;C2orf44;HMGN2P46;PMS		
			2P1;AK9;9-Sep;MIR142;		
			RFC;ZNF278;KMT2D;ZNF1		
			98;NUTM2B;IGH;TRA;RPA		
			;NUTM1;MICU3;MALAT1;T		
			RD;NBPF1;RUNDC2A;11-M		
			ar;BAGE2;PMS2L3;6-Sep		
			;TFPT;KNSTRN;C18orf56		
			;5-Sep;ACKR3;MTCP1;DU		
			X4L1;14-Sep;FAM47C;C2		
			orf43;SSX1;C3orf70;KM		
			T2A;AMER1;KMT2B;KDF1;		
			12-Sep;KMT2C		
<i>Test_only</i>	282	297	ADGRL2;CFAP54;ADGRV1; N/A		
			PAXBP1;ANK3;ADGRB3;J		
			CAD;DNAH12;ADGRA3;CCD		
			C168;KNSTRN;RMDN1;BRI		

			NP1;ACP7;ADGRG4;AMER1	
			;KMT2B;KMT2D;KMT2C;SL	
			F2;DISP3	
+	+	+	+	+
Overlap	151	157	KNSTRN;AMER1;KMT2B;DN ZBTB20;PIK3CA;MSH2;RA	
			AH12;KMT2D;KMT2C SA1;FGFR3;AXIN1;ARID1	
			A;CHD2;SPOP;HLA-A;DNA	
			H12;EPHA2;MGA;COL4A3;	
			NBEA;OR5L2;VHL;SMARCA	
			1;AJUBA;EPAS1;DYRK1A;	
			FAT4;CNBD1;JAK1;TP53;	
			CYLD;SPTAN1;ATM;BAP1;	
			MYC;NUP98;CRYBG3;GNAQ	
			;EPHA7;RAC1;SMC3;PMS1	
			;CDH10;FAT3;GNAS;PPP6	
			C;MAPK1;CHD4;RBM10;SI	
			N3A;PPP2R1A;PTEN;MUC1	
			6;TRAF3;ANO1;EPHA5;CU	
			L1;CDKN2A;LRP1B;EP300	
			;TRIM33;FBXW7;COL11A1	
			;CHD3;EPHA3;FANCM;KMT	
			2B;ARHGAP35;MUC17;KMT	
			2C;RYR2;PREX2;HUWE1;K	
			DM5C;CREBBP;MAP3K1;AS	
			XL2;REG3G;ZMYM3;ATRX;	
			LCA5L;TGFBR2;JAK2;RBB	
			P8;PCM1;RNF213;SPTA1;	
			PDZRN4;ERBB4;DDX10;TT	
			N;CNTRL;AMOT;ZRANB3;M	
			ED12;SOX9;NF1;ERCC2;X	
			IRP2;GNA11;BRAF;SACS;	
			MET;NRAS;KMT2D;NFE2L2	
			;POLE;SMAD4;CASP8;CSM	
			D3;CHD8;MAP2K1;NSD1;K	
			EAP1;PIK3R1;CTNNB1;RB	
			1;ARID2;PRDM9;APC;TRR	
			AP;MUC5B;PBRM1;RET;HR	
			AS;APOB;STK11;EEF1A1;	
			RYR3;NAV3;NOTCH2;PTPR	
			D;BRCA2;VPS13C;NOTCH1	
			;FLG;SMARCA4;KDM6A;KN	
			STRN;MN1;ZNF750;RUNX1	
			;ERBB2;PTPN14;ALK;TBC	
			1D4;CUL3;CSMD1;MACF1;	
			DMD;MAP2K4;GTF2I;KIT;	
			ARHGAP5;SF3B1;FAT1;AM	
			ER1;OR2T33;CIC;BRCA1;	
			MYH9;KRAS	
+	+	+	+	+

(end of file)