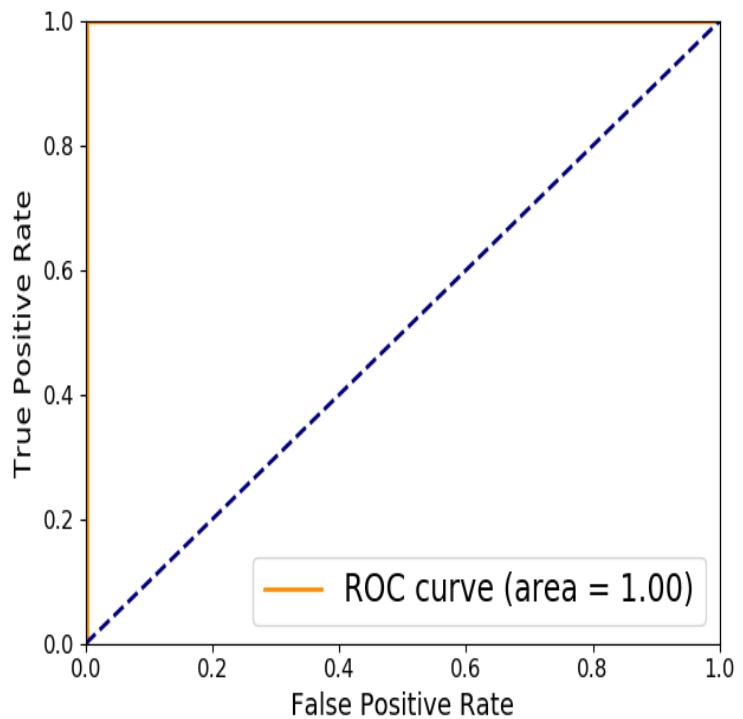
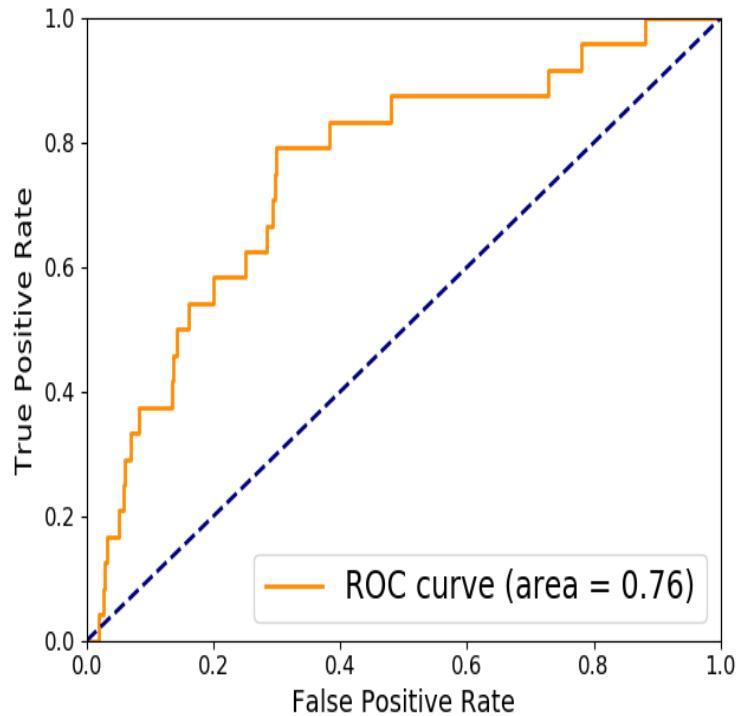


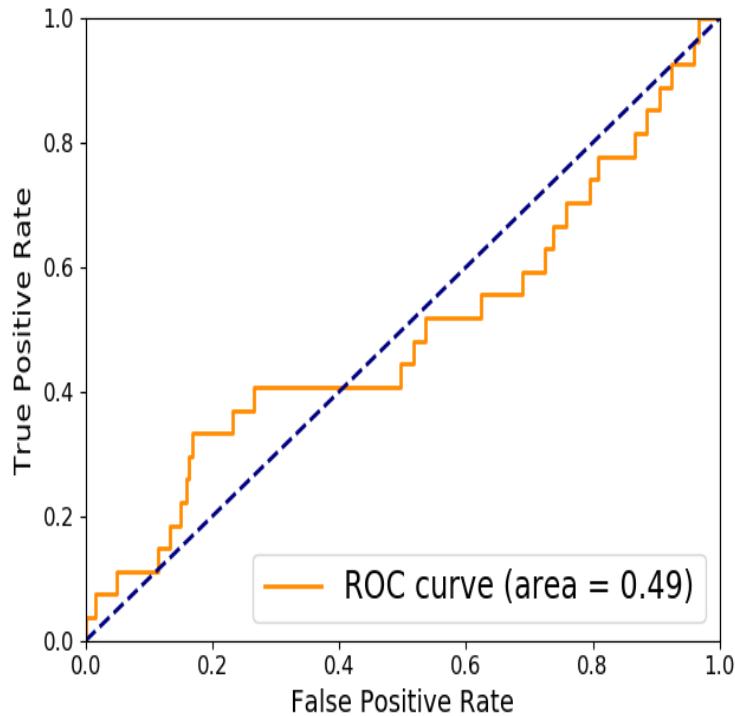
## Diffusion ROC from fGSonly\_tOverlap STRING10\_combined.png



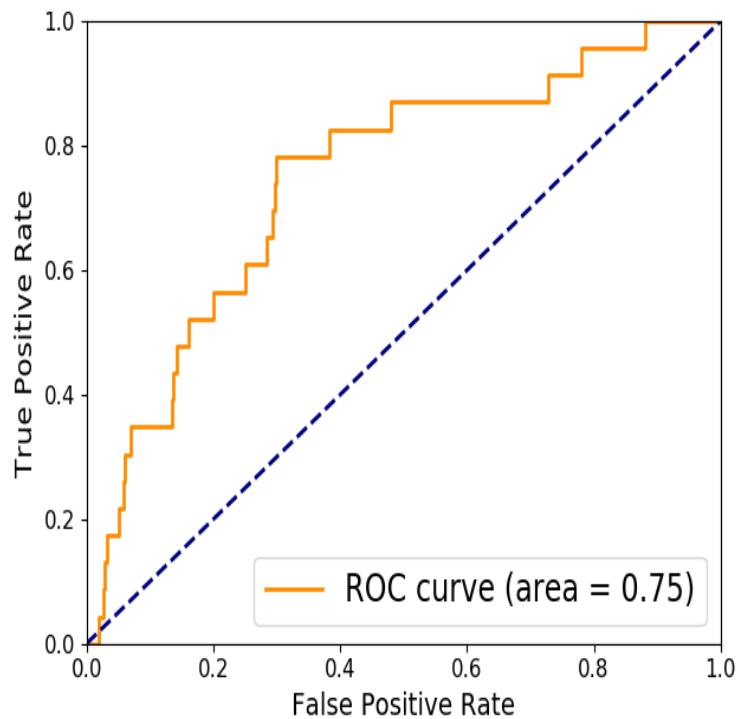
## Diffusion ROC from fTestOnly\_tGS STRING10\_combined.png



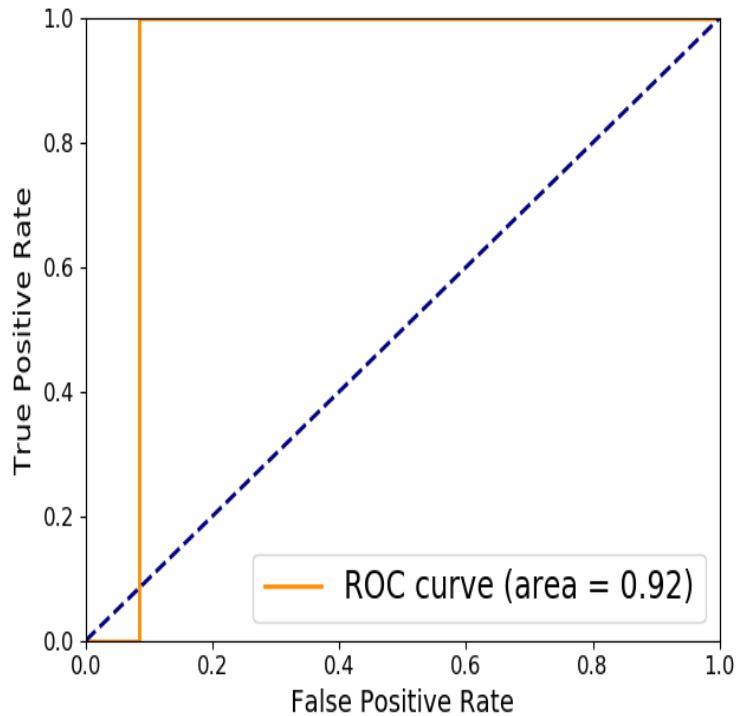
## Diffusion ROC from fGSonly\_tTestOnly STRING10\_combined.png



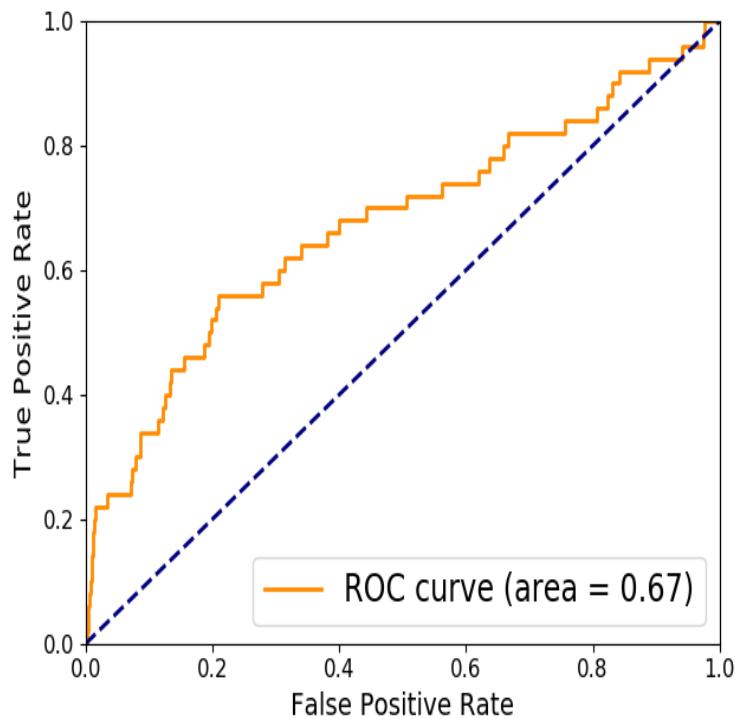
## Diffusion ROC from fTestOnly\_tGSonly STRING10\_combined.png



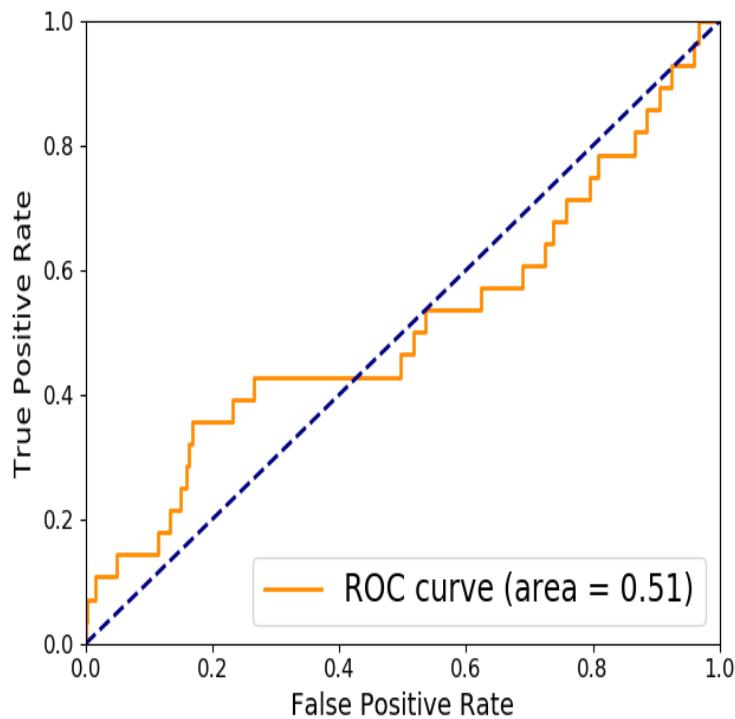
## Diffusion ROC from fTestOnly\_tOverlap STRING10\_combined.png



## 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_wdegre</i>	<i>Zt</i>	<i>Zf_wdegre</i>	<i>Zf</i>
/	/	e	/	e	/	
GS_only	Test_only	0.495   1.453	-0.266	1.405	1.302	
Test_only	GS_only	0.748   3.553	3.662	2.4404393	2.771	
/	/	/	/	893383864	/	
/	/	/	/	/	(invalid)	/
GS_only	Overlap	0.998   4.9956373	1.7088069	1.5500283	1.818199	
/	/	/	20102209	766830656	946019024	34133810
/	/	/	(invalid)	(invalid)	(invalid)	97 (inva
/	/	/	/	/	lid)	
GS_only	Test	0.513   1.907	0.185	1.501	1.411	
Test_only	Overlap	0.916   2.2028445	1.2604971	1.013	0.996	
/	/	/	97302967	789235624	/	/
/	/	/	(invalid)	(invalid)	/	/
Test_only	GS	0.755   3.427	4.065	2.1014566	3.152	
/	/	/	/	10100006	/	
/	/	/	/	(invalid)	/	
Overlap	Onlies	0.675   5.668	4.494	6.0624390	2.248	
/	/	/	/	27790198	/	
/	/	/	/	(invalid)	/	

**	#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)