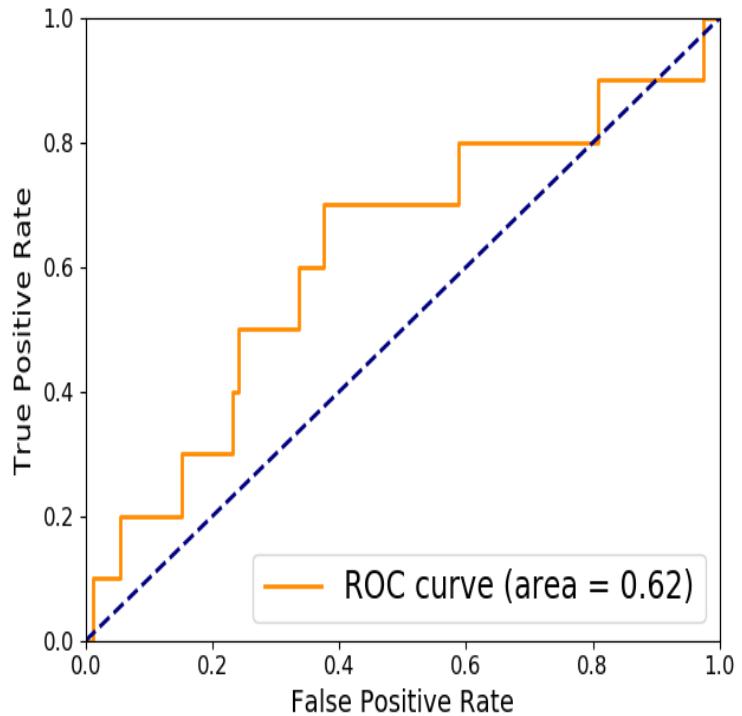
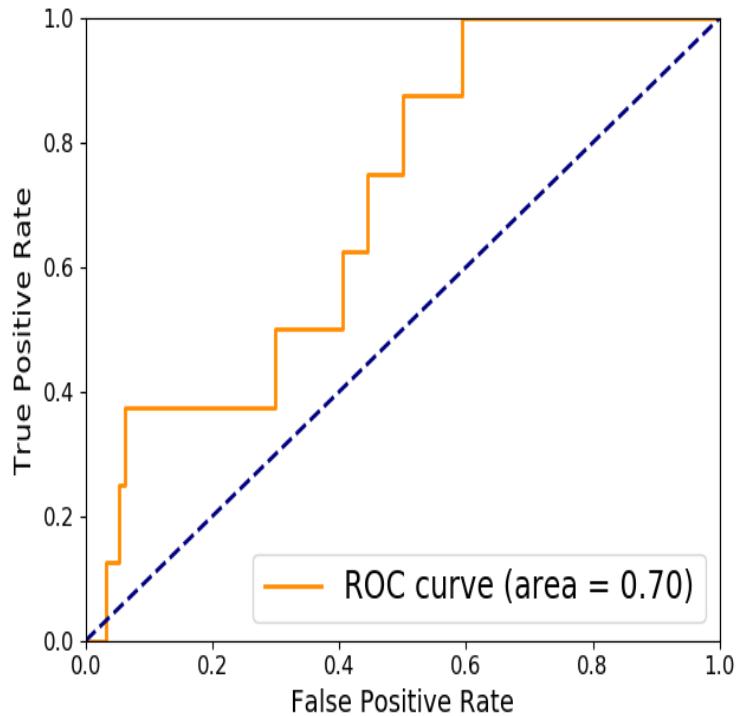


## show\_1\_Diffusion ROC from Gene2axis to Gene1axis STRING10\_combined



## show\_2\_Diffusion ROC from Gene1axis to Gene2axis STRING10\_combined



<i>Diffusion</i>	<i>Diffusion</i>	<i>AUROC</i>	<i>Zt</i>	<i>Zt</i>	<i>Zf</i>	<i>Zf (unif)</i>
<i>FROM</i>	<i>TO</i>		(degree- matched)	(uniform)   matched)	(degree- matched)	orm)
<i>Gene2axis</i>	<i>Gene1axis</i>	0.623	-0.167	1.366	0.368	1.184
<i>only</i>	<i>only</i>					
<i>Gene1axis</i>	<i>Gene2axis</i>	0.701	0.849	1.696	0.457	0.587
<i>only</i>	<i>only</i>					

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>	
<i>Gene2axis Only</i>	8	10	<i>YTHDF3;B3GNTL1P1</i>	N/A
<i>Gene1axis Only</i>	10	10		N/A

(end of file)