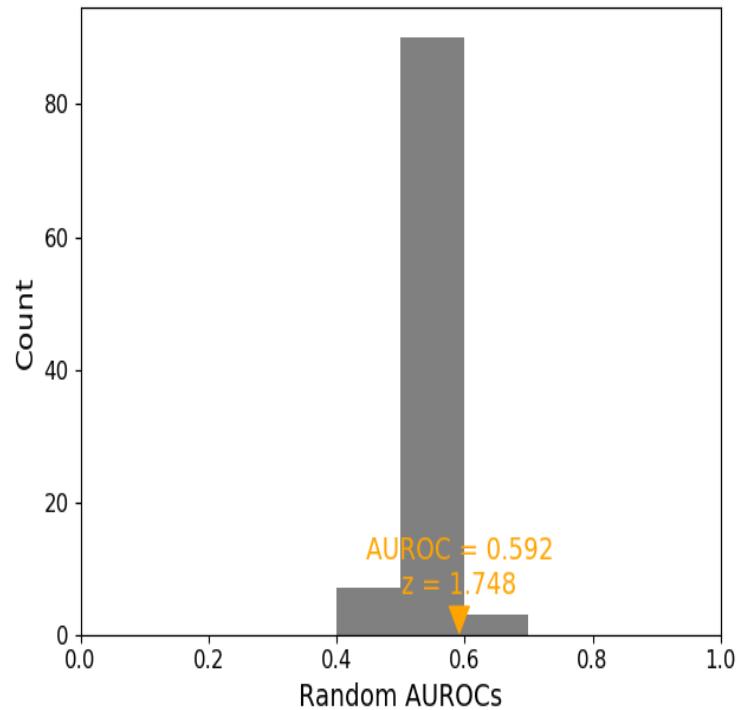
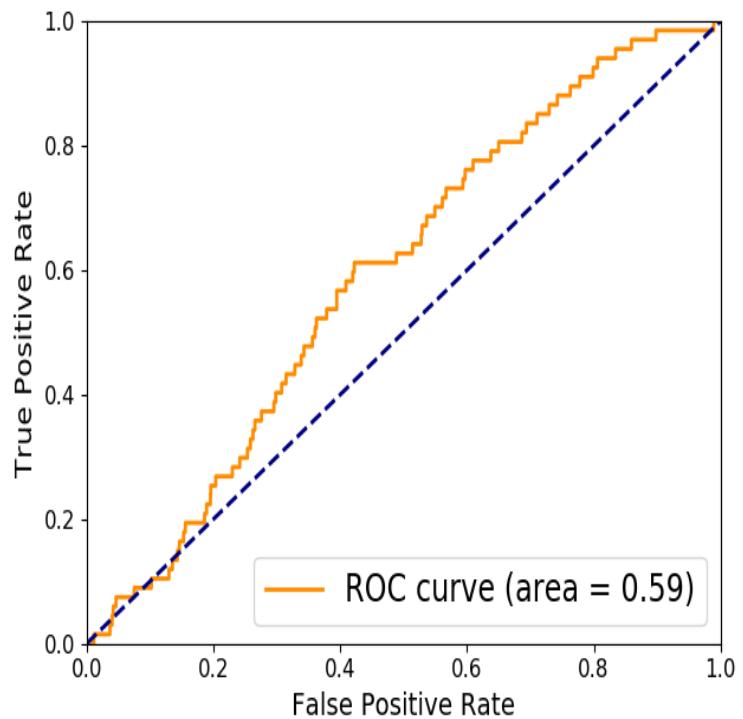


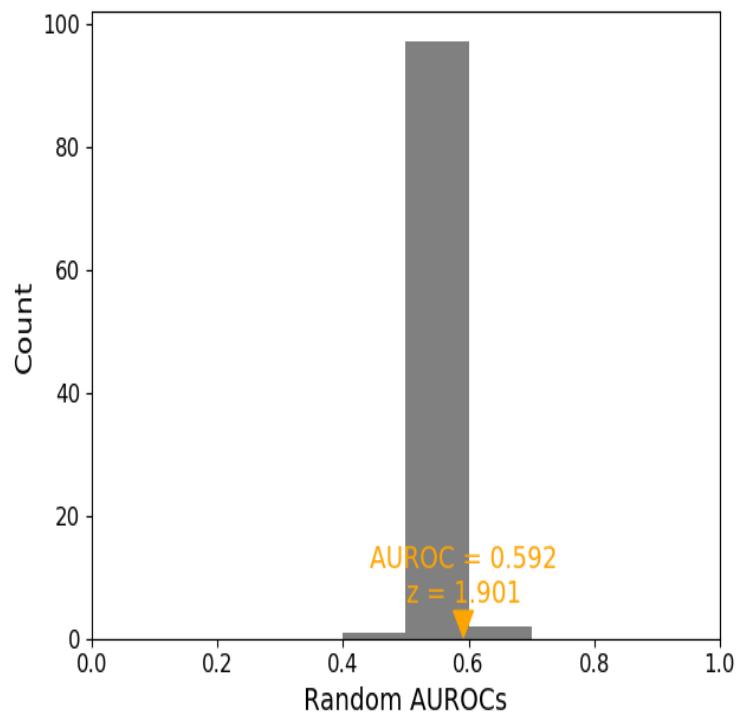
## show\_21\_Zt\_degree-matched\_from Ismael to GS.png



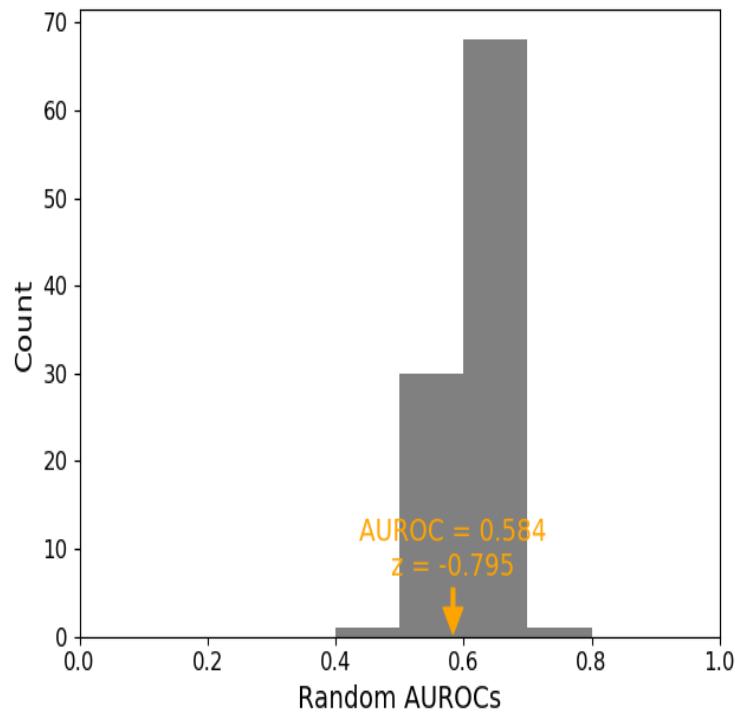
**show\_2\_Diffusion ROC from Ismael to GS STRING10\_combined.png**



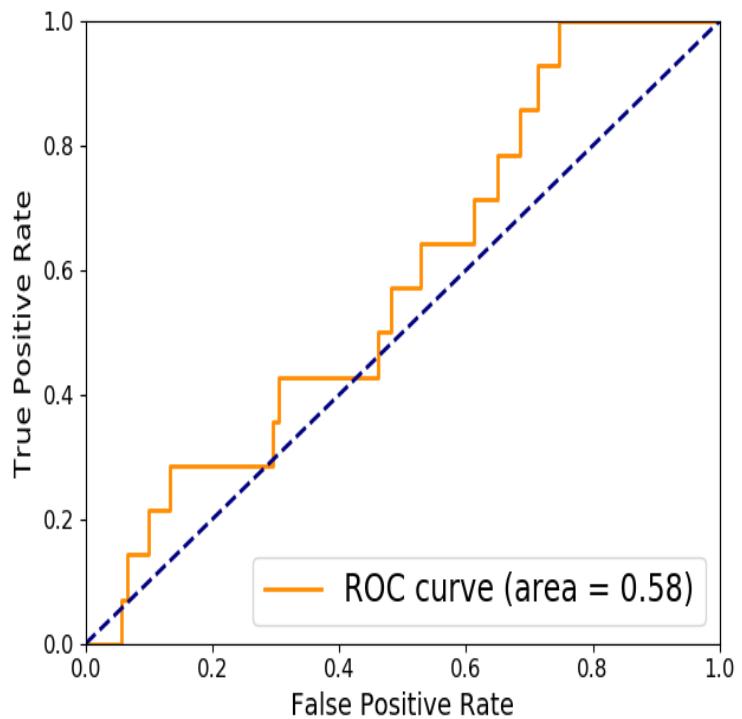
## show\_22\_Zf\_degree-matched\_from Ismael to GS.png



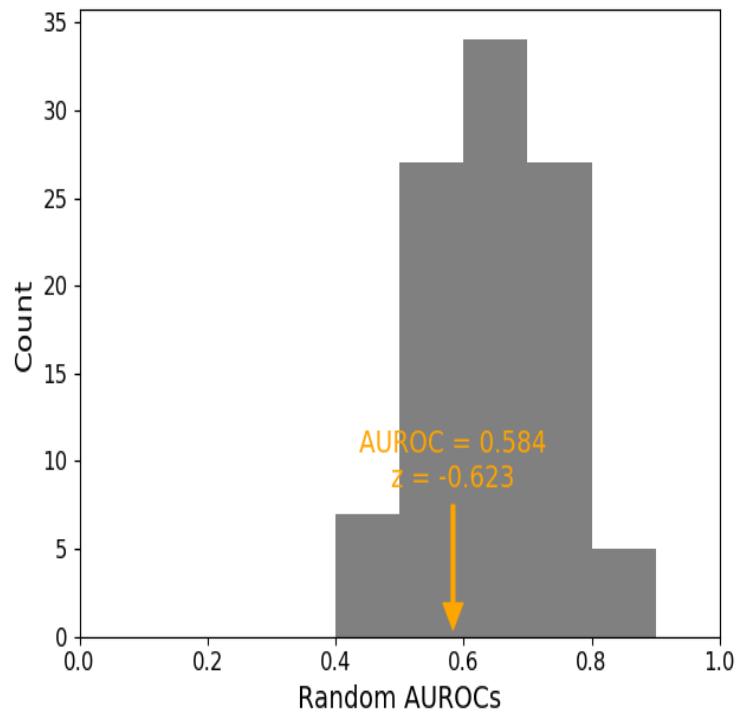
## show\_11\_Zt\_degree-matched\_from GS to Ismael.png



**show\_1\_Diffusion ROC from GS to Ismael STRING10\_combined.png**



## show\_12\_Zf\_degree-matched\_from GS to Ismael.png



<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)
GS	Ismael	0.584	-0.795	1.161	-0.623	-0.336
Ismael	GS	0.592	1.748	2.955	1.901	1.894

Z-scores are computed for the experimental AUROC based on distributions of the random AUROCs

*Zf* : when randomizing source genes (*Diffusion FROM*)

*Zf* : when randomizing recipient genes (*Diffusion TO*)

Random genes are selected either uniformly or degree matched

\* : Distribution of random AUROC is not Gaussian

**	#Mapped	#Total	Not mapped genes	
GS	67	68	SLC35G6	N/A
Ismael	14	14		N/A

(end of file)