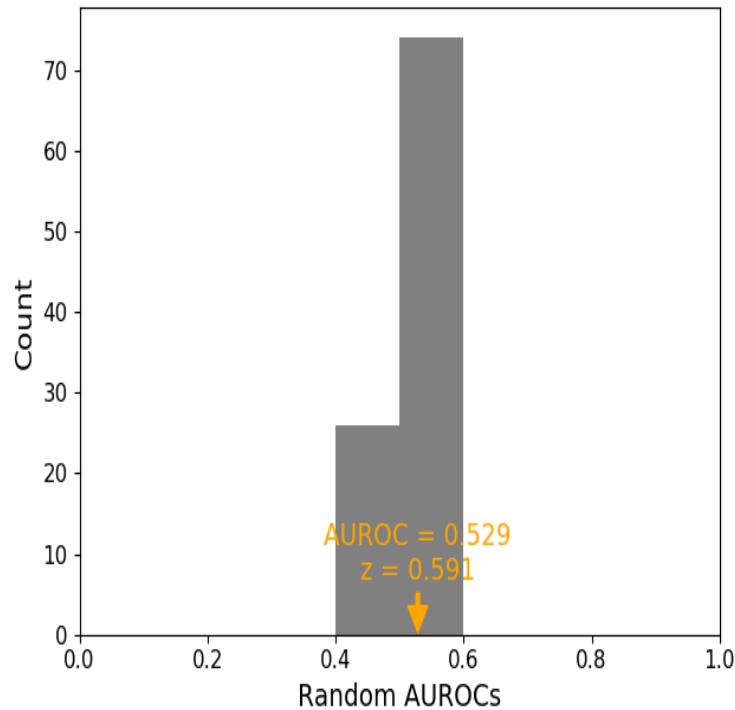
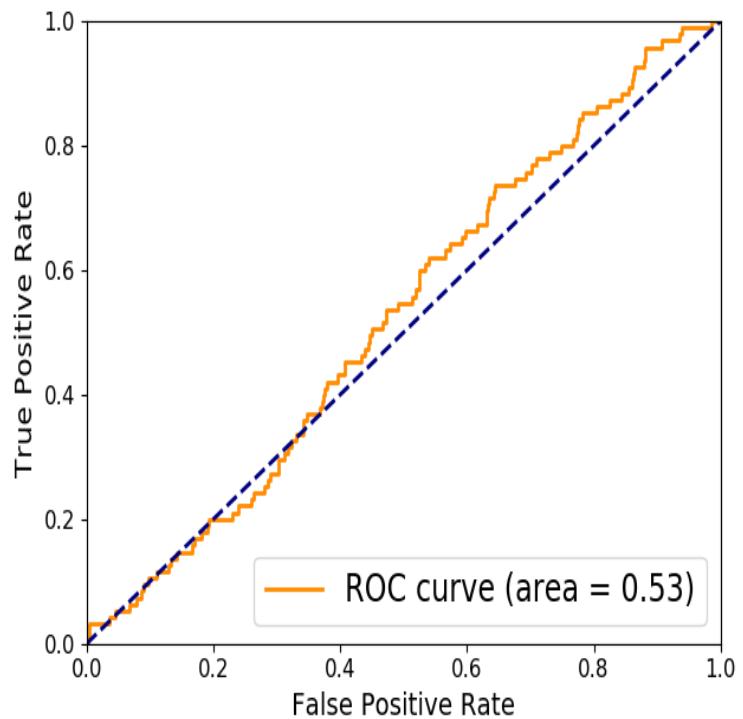


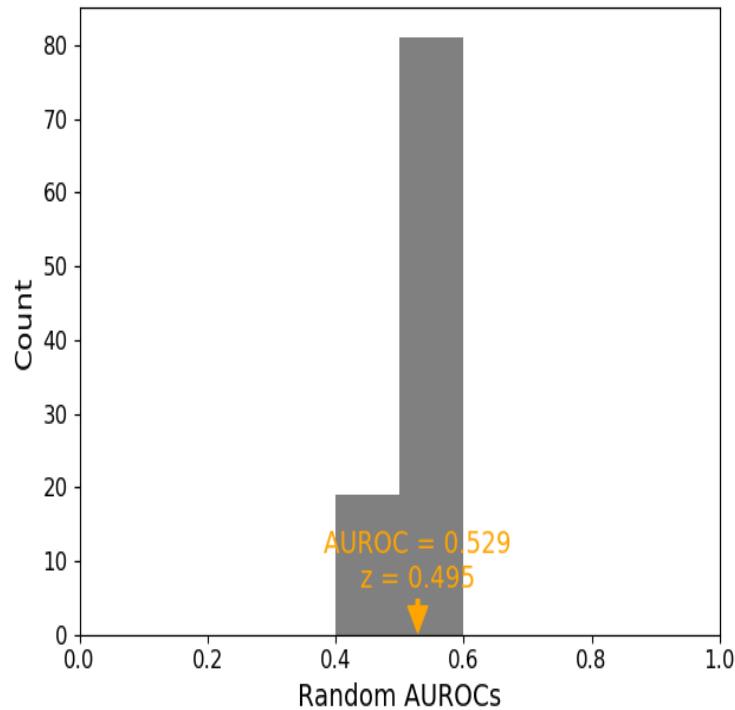
show_21_Zt_degree-matched_from Ismael to GS.png



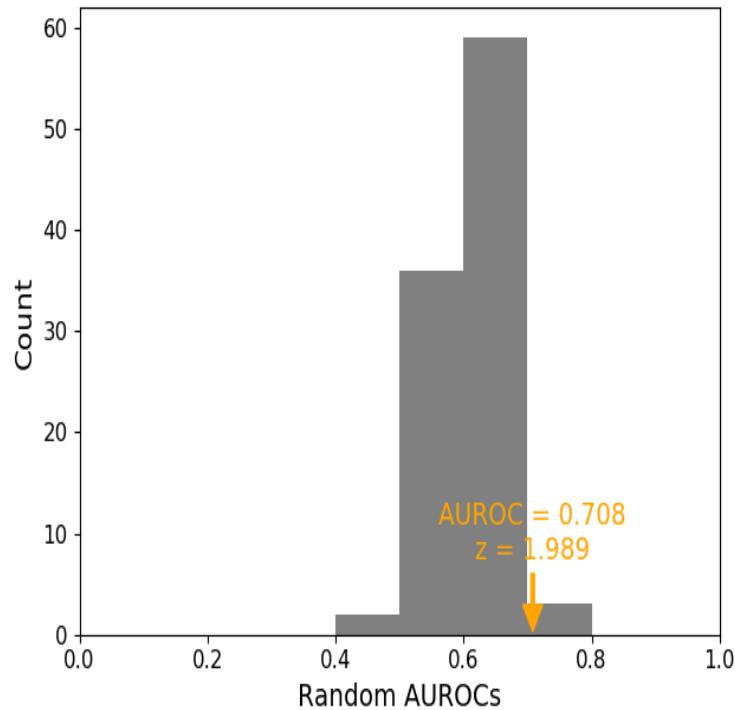
show_2_Diffusion ROC from Ismael to GS STRING10_combined.png



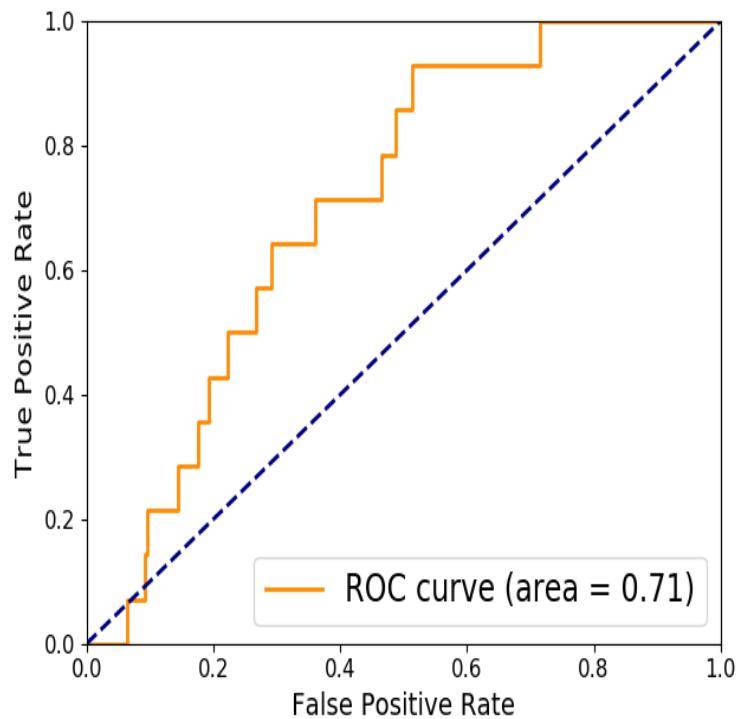
show_22_Zf_degree-matched_from Ismael to GS.png



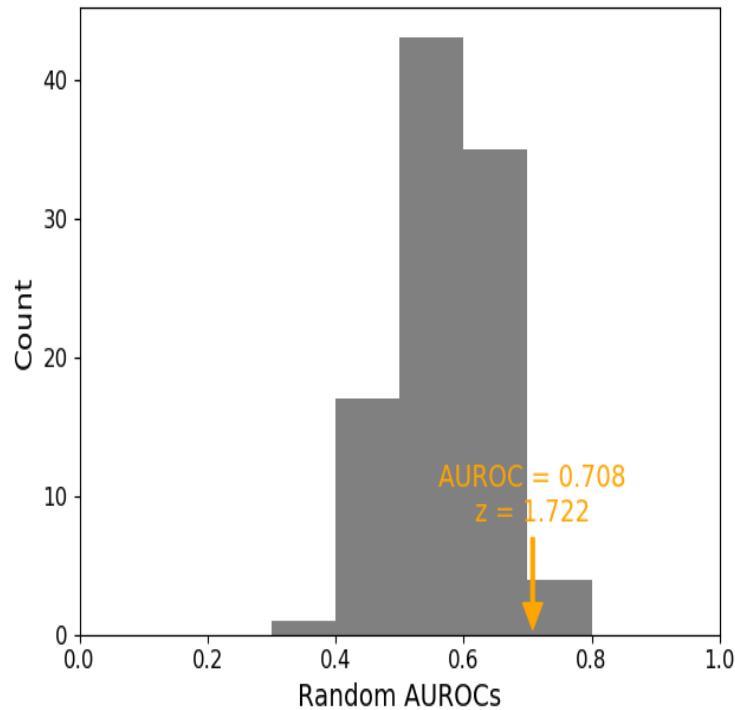
show_11_Zt_degree-matched_from GS to Ismael.png



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	0.708 1.989	2.713	1.722	0.801 *	
Ismael	GS	0.529 0.591	0.885	0.495	0.468	

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	95	100	SSPO;FAM153B;ERV3.1;ZNF735;LY6G5B	N/A	
Ismael	14	14		N/A	

(end of file)