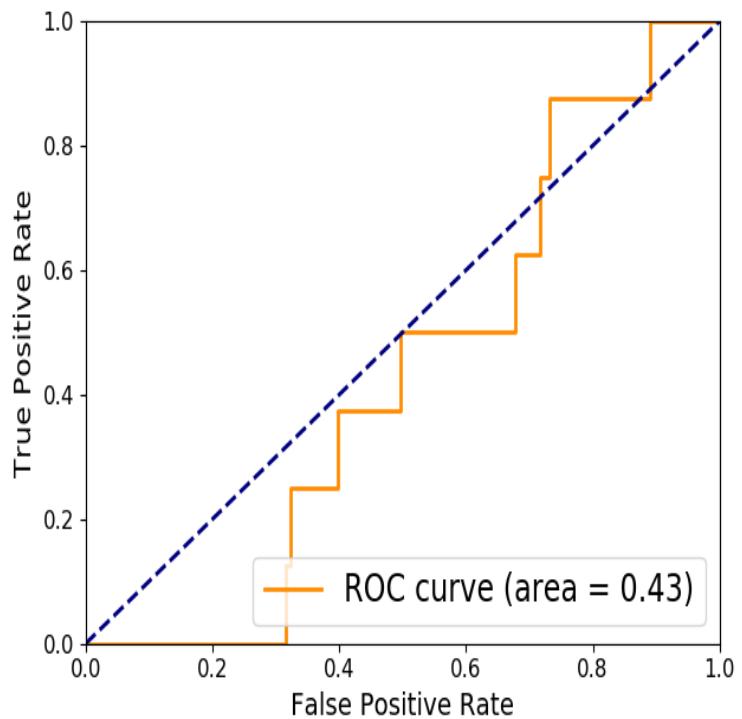
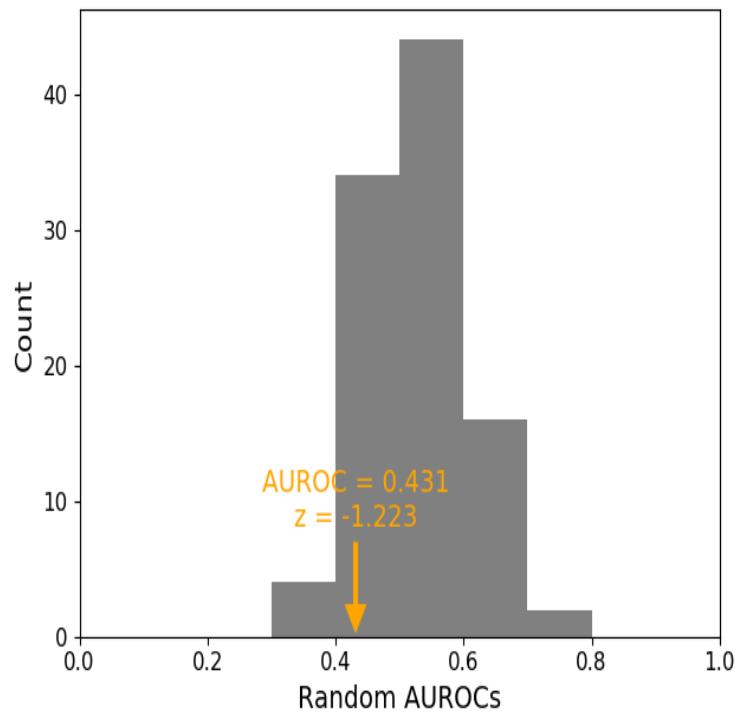


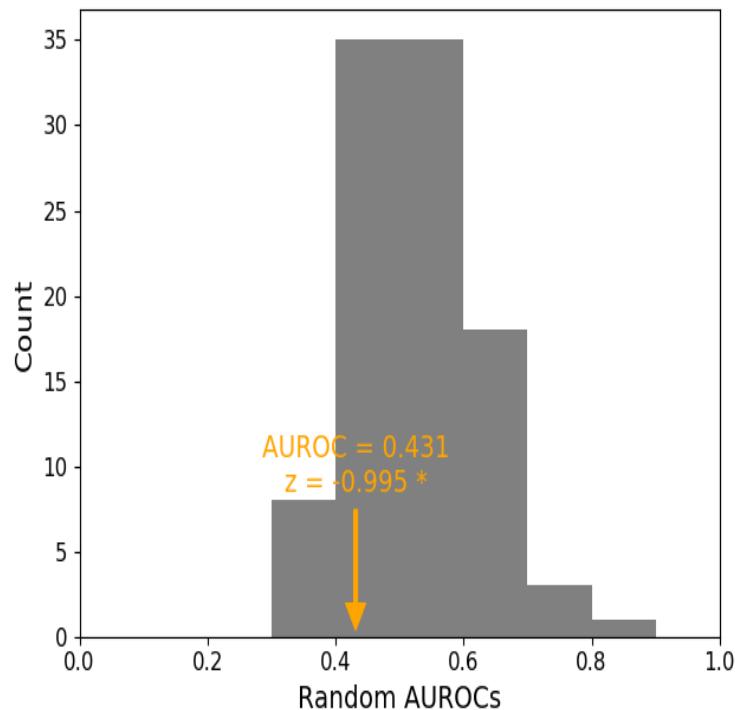
## show\_1\_Diffusion ROC from protein to Marttinen STRING10\_combined.png



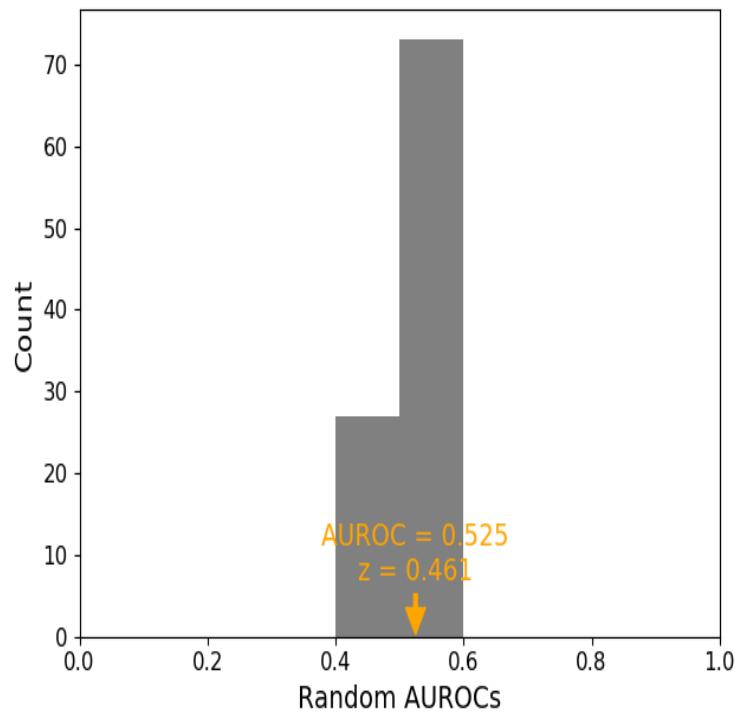
## show\_12\_Zf\_degree-matched\_from protein to Marttinen.png



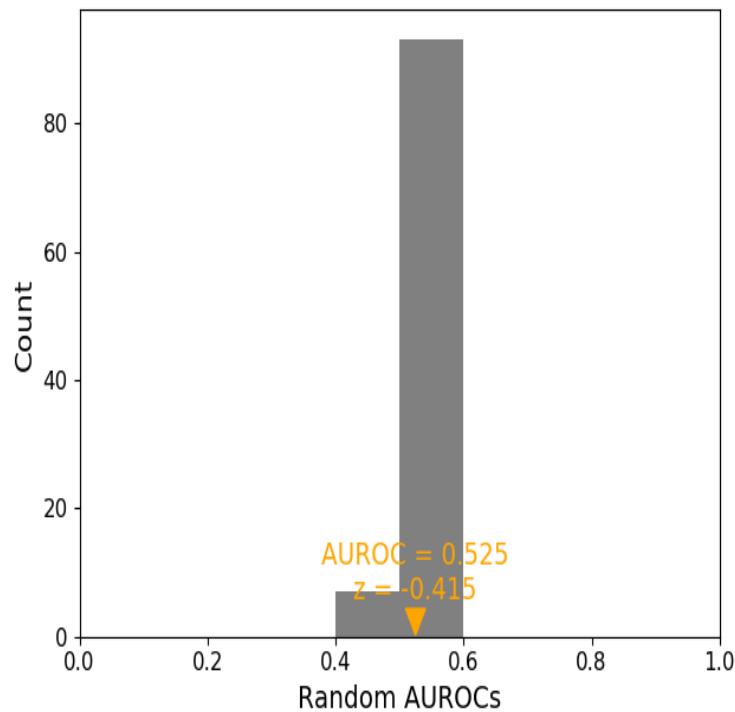
## show\_11\_Zt\_degree-matched\_from protein to Marttinen.png



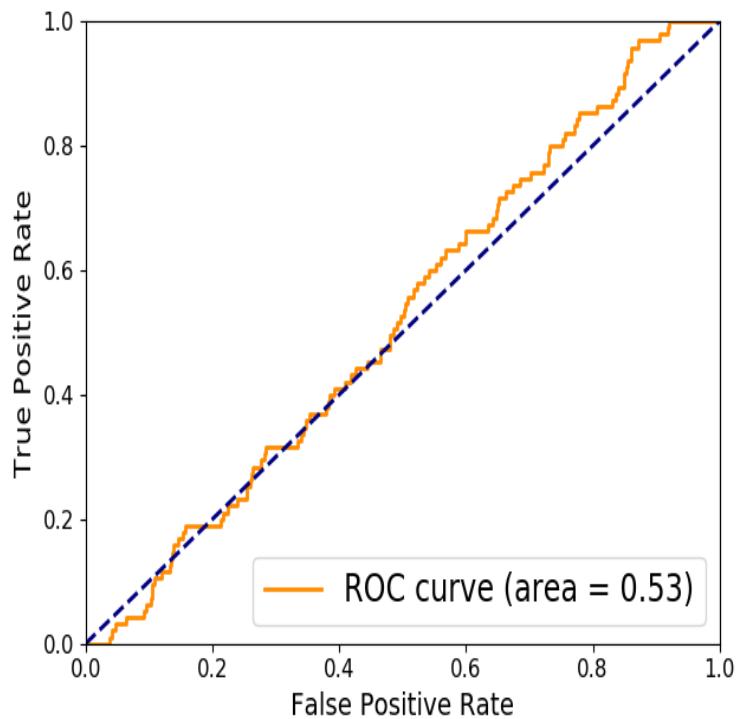
## show\_21\_Zt\_degree-matched\_from Marttinen to protein.png



## show\_22\_Zf\_degree-matched\_from Marttinen to protein.png



## show\_2\_Diffusion ROC from Marttinen to protein STRING10\_combined.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)
<i>protein</i>	<i>Marttinen</i>	0.431	-0.995 *	-0.707	-1.223	-1.276
<i>Marttinen</i>	<i>protein</i>	0.525	0.461	0.876	-0.415	0.133

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

<i>**</i>	<i>#Mapped</i>	<i>#Total</i>	<i>Not mapped genes</i>		
<i>protein</i>	95	100	ERV3.1;LY6G5B;SSPO;FAM153B;ZNF735	N/A	
<i>Marttinen</i>	8	8		N/A	

(end of file)