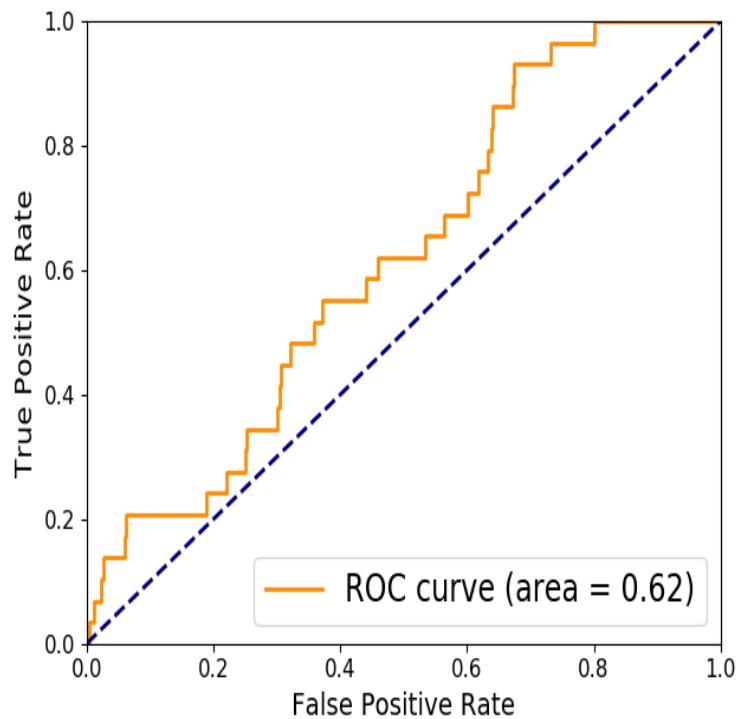
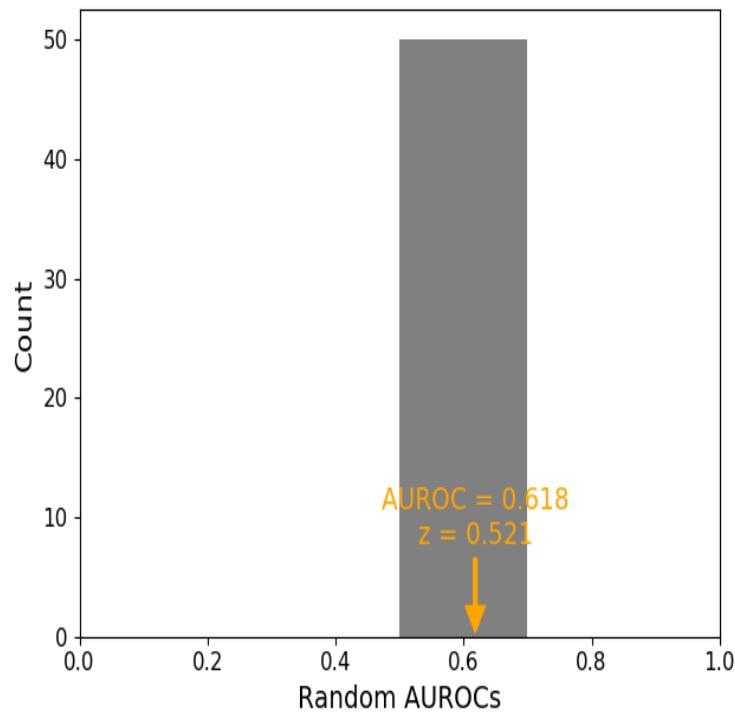


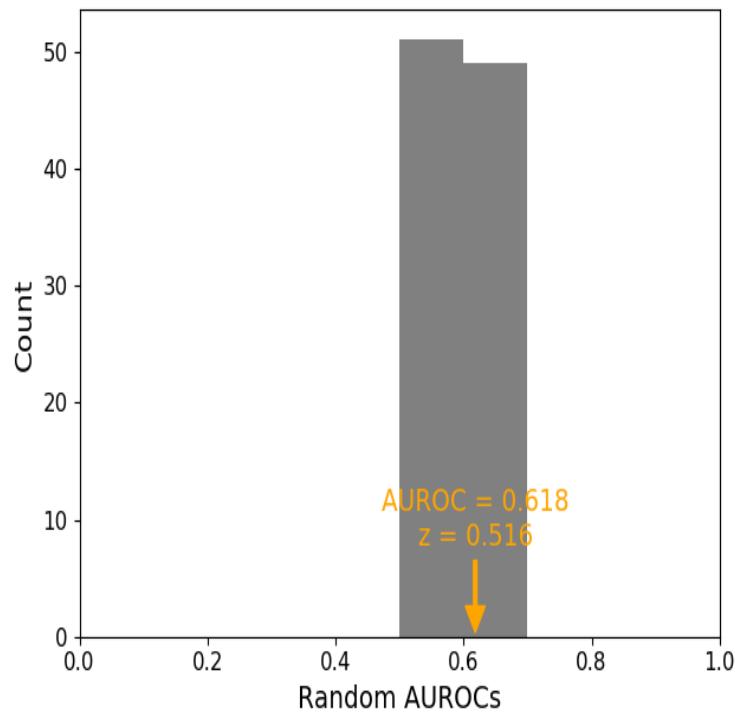
## show\_1\_Diffusion ROC from protein to Marttinen STRING10\_combined.png



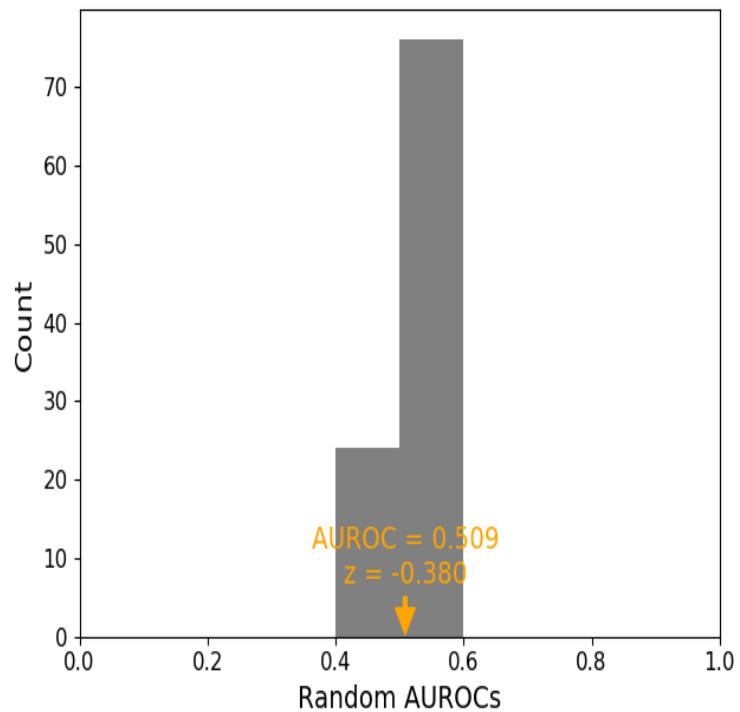
## show\_12\_Zf\_degree-matched\_from protein to Marttinen.png



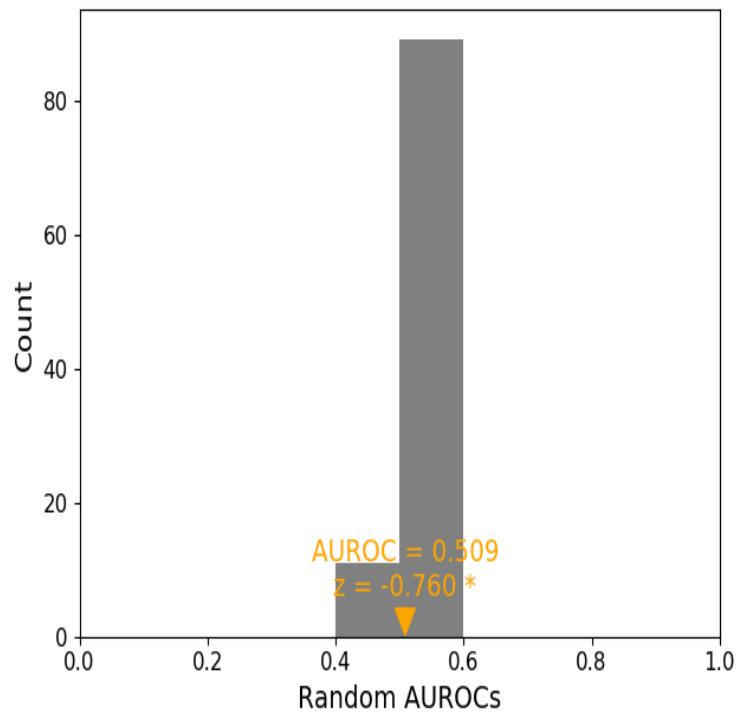
## show\_11\_Zt\_degree-matched\_from protein to Marttinen.png



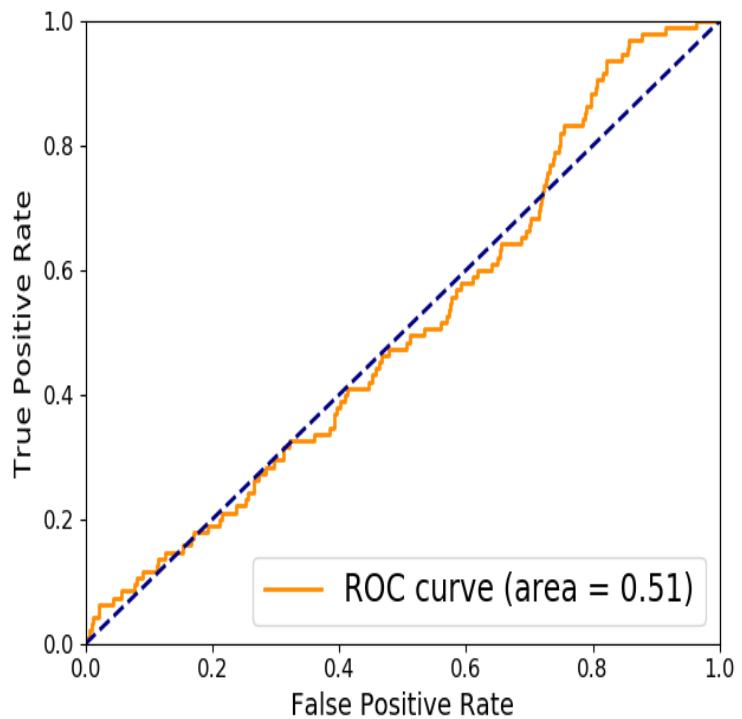
## show\_21\_Zt\_degree-matched\_from Marttinen to protein.png



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



## show\_2\_Diffusion ROC from Marttinен 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>	0.618	0.516	2.384	0.521	0.027
<i>Marttinen</i>	<i>protein</i>	0.509	-0.380	0.293	-0.760 *	-0.397

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	LY6G5B;ZNF735;ERV3.1;FAM153B;SSPO	N/A	
<i>Marttinen</i>	29	30	C9orf172	N/A	

(end of file)