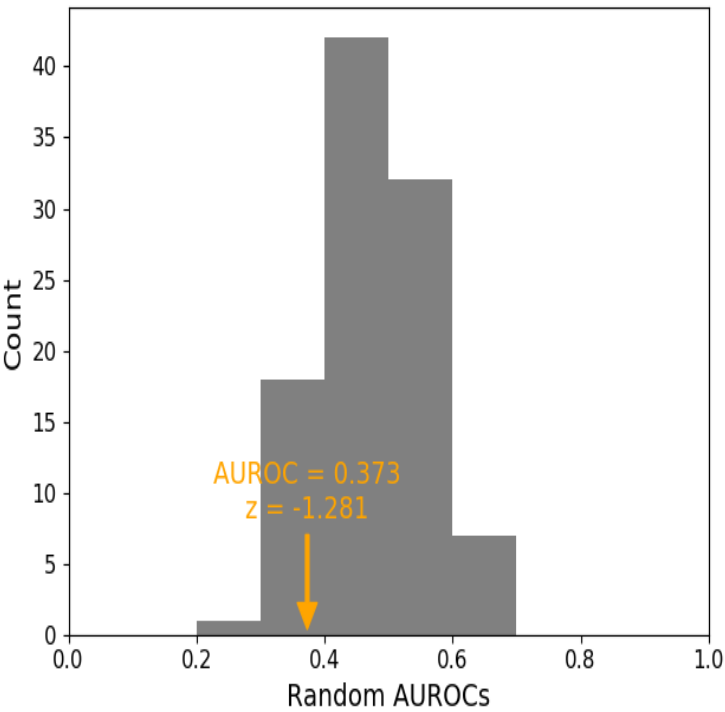
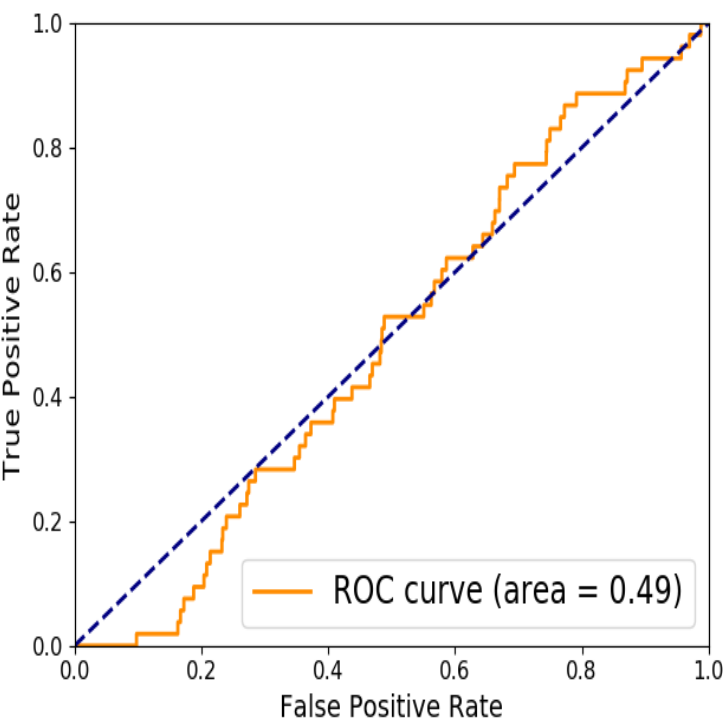


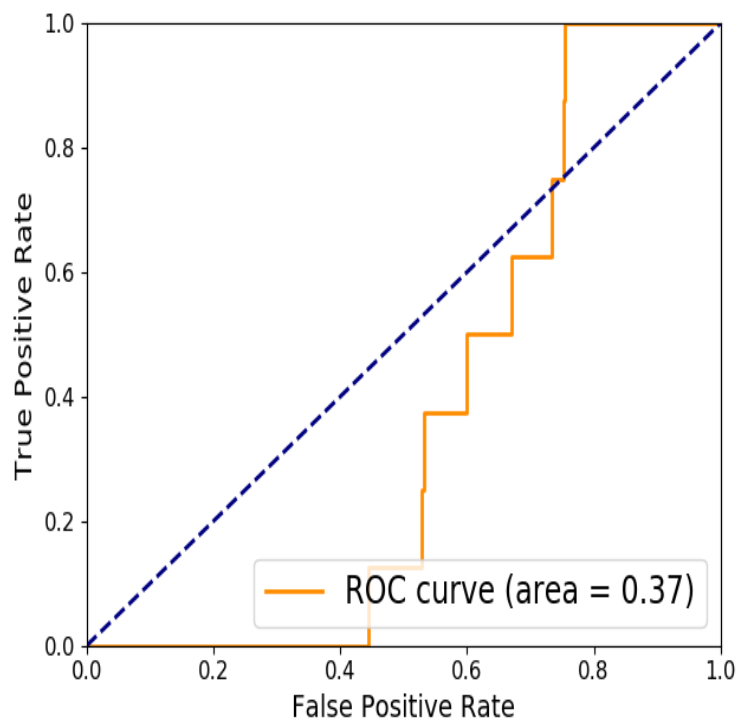
show_11_Zt_degree-matched_from ADSP to Marttinen_protein_decisionTr



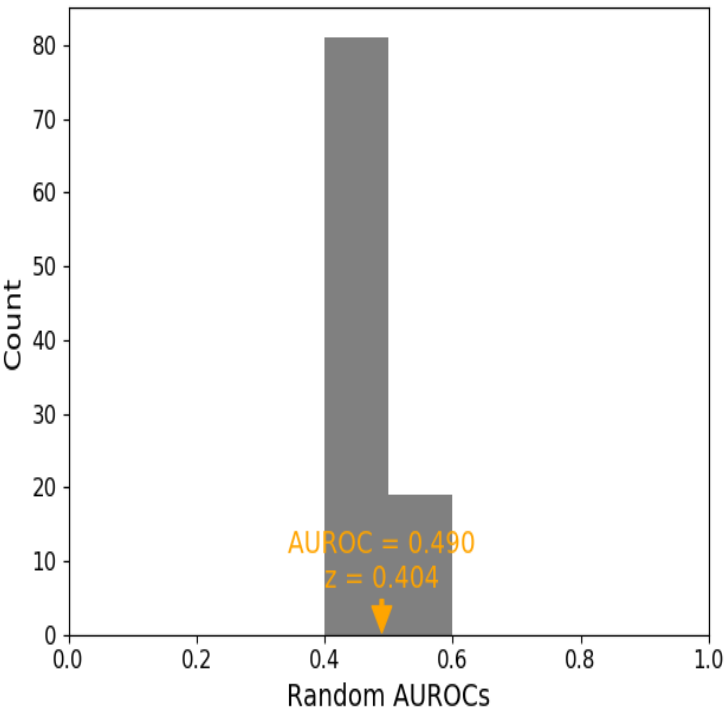
show_2_Diffusion ROC from Marttinen_protein_decisionTree to ADSP STR



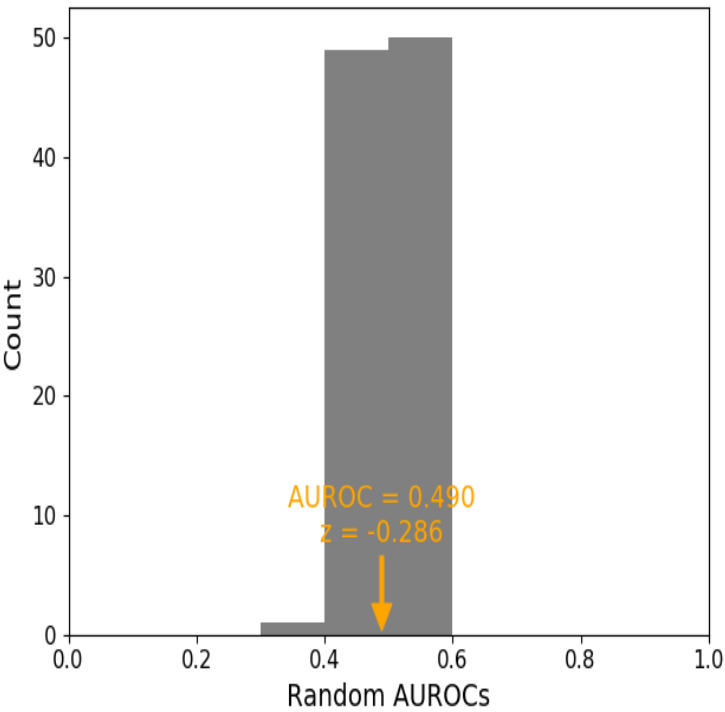
show_1_Diffusion ROC from ADSP to Marttinen_protein_decisionTree STR



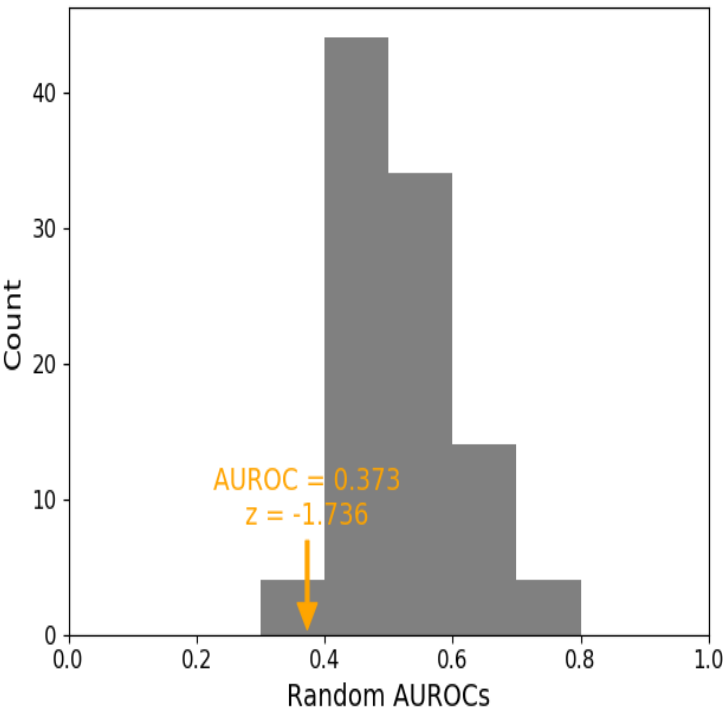
show_22_Zf_degree-matched_from Marttinen_protein_decisionTree to AD



show_21_Zt_degree-matched_from Marttinen_protein_decisionTree to AD



show_12_Zf_degree-matched_from ADSP to Marttinen_protein_decisionTr



Diffusion FROM	Diffusion TO	AUROC	Zt (degree-matched)	Zt (uniform)	Zf (degree-matched)	Zf (uniform)
ADSP	Marttinen	0.373	-1.281	-1.228	-1.736	-1.820
protein						
decisionT						
ree						
Marttinen	ADSP	0.490	-0.286	-0.351	0.404	-0.046 *
protein						
decisionT						
ree						

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
ADSP	53	54	IQCF6
Marttinen_protein_decisionTree	8	8	N/A

(end of file)