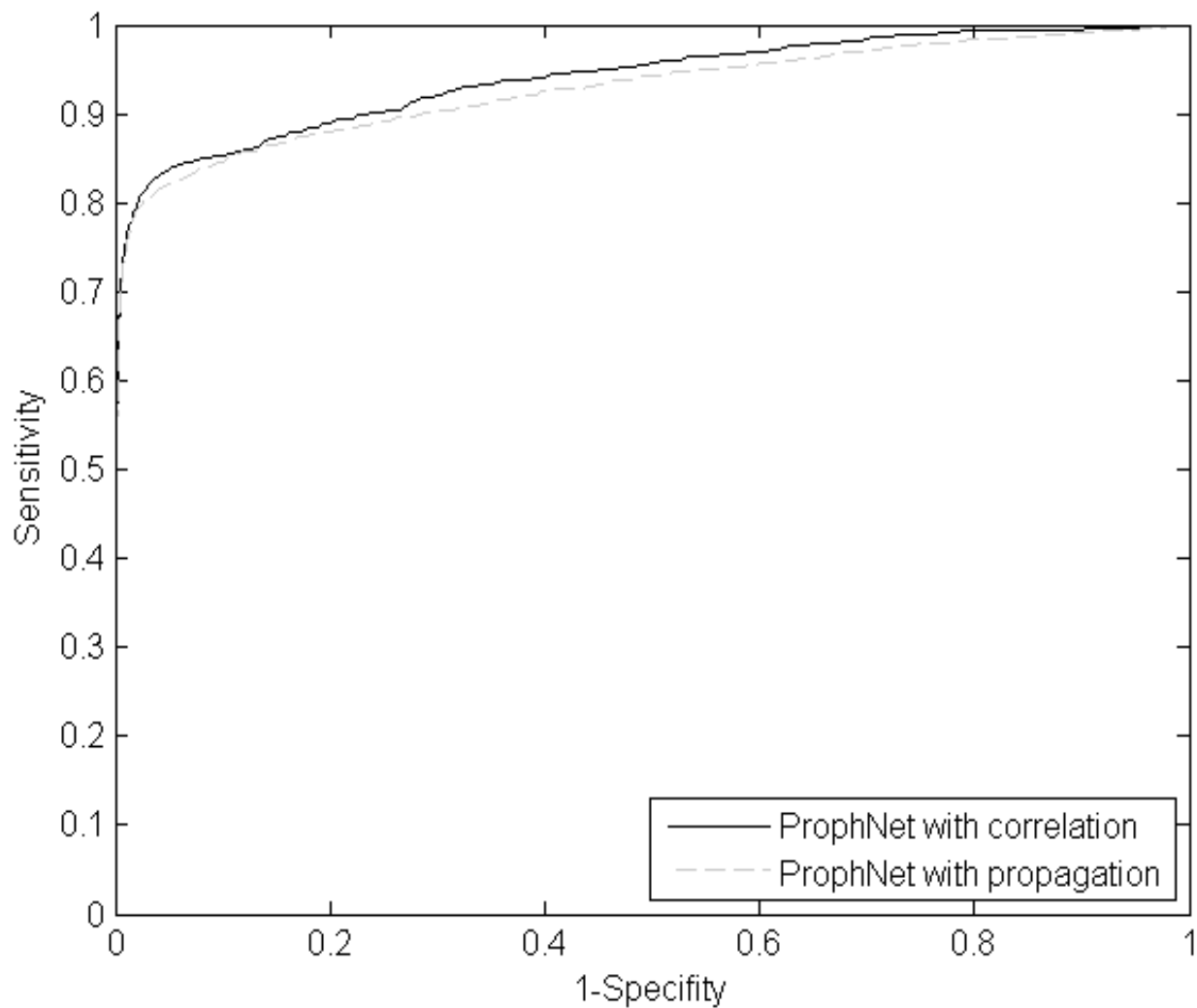


# **PROPAGATION AND CORRELATION ANALYSIS RESULTS** **(GENE-DISEASE LEAVE-ONE-OUT VALIDATION)**



<b>Method</b>	<b>Correlation</b>	<b>Propagation</b>
<b>AUC</b>	0.9393	0.9277
<b>Mean ranking</b>	309	368
<b>Standard deviation</b>	811	944