

**Supplementary Table 3. Ranked percentage of SDE transcripts (mRNAs, miRNAs, and lncRNAs) and known MI-related transcripts contained in dysregulated LmiRM-CTs obtained using our method.**

	<b>Top 5%</b>	<b>Top10%</b>	<b>Top1 5%</b>	<b>Top 20%</b>	<b>Top 30%</b>	<b>Top 40%</b>	<b>Top 50%</b>	<b>100%</b>
SDE_transcripts	96.77%	97.14%	96.82%	96.34%	94.86%	93.55%	93.50%	92.35%
MI_transcripts	8.06%	7.62%	9.55%	8.38%	8.69%	8.06%	7.32%	6.82%