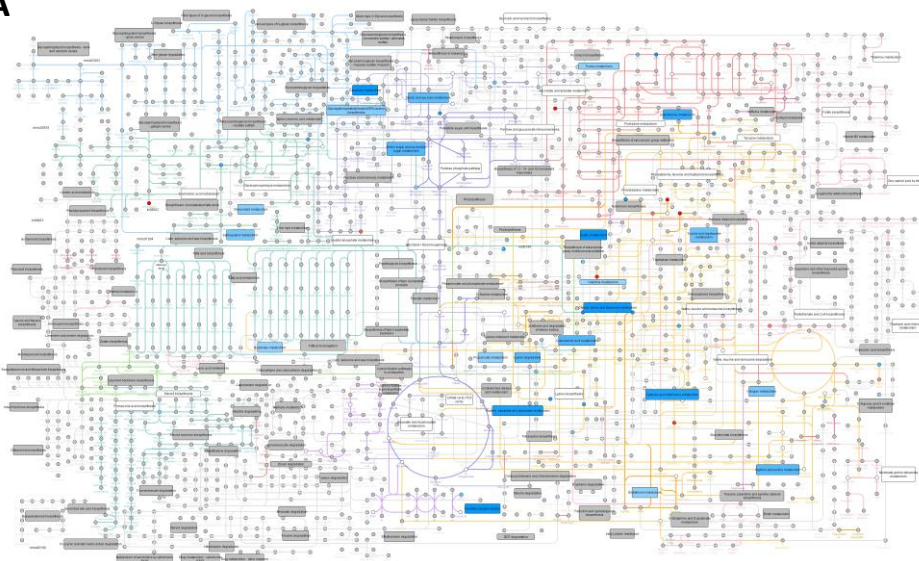
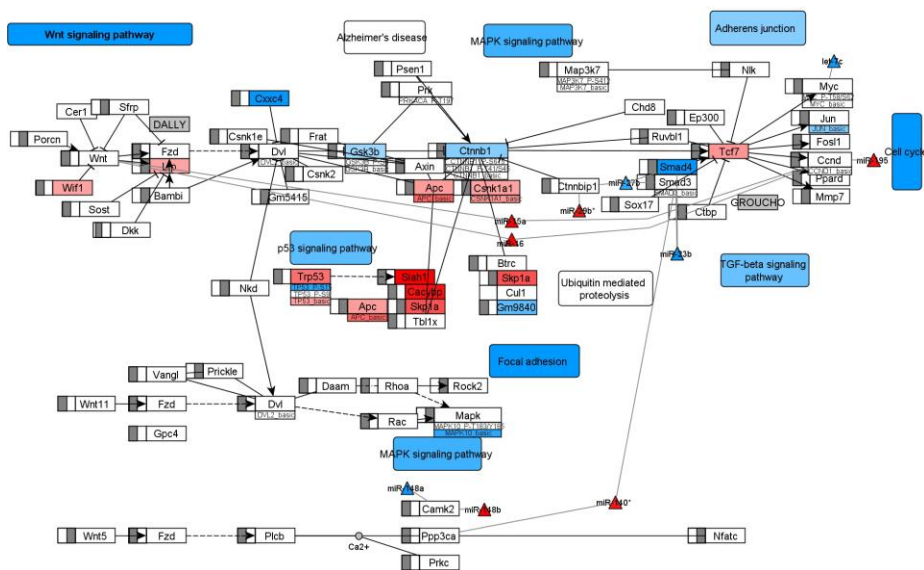


A



B



C

Data types		
DNAm	<div> <div><div></div><div></div><div></div></div> <div>Skp1a</div> </div> <div> <div><div></div><div></div><div></div></div> <div>Cul1</div> </div> <div> <div><div></div><div></div><div></div></div> <div>Gm9840</div> </div>	hypermethylated no difference hypomethylated
mRNA	<div> <div></div> <div>Skp1a</div> </div> <div> <div></div> <div>Cul1</div> </div> <div> <div></div> <div>Gm9840</div> </div> <div> <div></div> <div>DALLY</div> </div>	upregulated no difference downregulated no data
miRNA		same colors as mRNA
Protein modification	<div> <div></div> <div>Trp53</div> </div> <div> <div></div> <div>TP53_P-S117</div> </div> <div> <div></div> <div>TP53_P-S9</div> </div> <div> <div></div> <div>TP53_basic</div> </div>	same colors as mRNA
Metabolite	Taurine ● Pyruvate ○ L-Alanine ● Acetate ●	same colors as mRNA
Pathway	<div> <div></div> <div>Galactose metabolism</div> </div> <div> <div></div> <div>Histidine metabolism</div> </div> <div> <div></div> <div>Pyrimidine metabolism</div> </div> <div> <div></div> <div>Pyruvate metabolism</div> </div>	high significance small significance non significant contains no significant feature
Color keys		
\log_2 fold change (e.g. Metabolite) <div> <div></div> <div>-1.5</div> <div>0</div> <div>1.5</div> </div>		
p-value (e.g. Pathway) <div> <div></div> <div>0.0005</div> <div>0.05</div> </div>		