

	<b>Gene</b>	<b>Fold Change (FOLFOX vs. Vehicle)</b>
<b>Cell Cycle Regulation</b>	<i>Cdkn1a (p21<sup>Cip1</sup>)</i>	+ 8.2
	<i>CXCL1</i>	+ 3.3
	<i>GADD45G</i>	+ 2.6
	<i>KLF6</i>	+ 2.3
	<i>STAT3</i>	+ 1.8
	<i>Gas6</i>	+ 1.7
	<i>GADD45B</i>	+ 1.7
	<i>Bax</i>	+ 1.6
	<i>Ccng1</i>	+ 1.5
	<i>Gas7</i>	- 1.5
<b>Oxidative Stress</b>	<i>Mt1</i>	+ 12.4
	<i>Hmox1</i>	+ 2.4
	<i>Gpx3</i>	+ 2.0
	<i>Mgst3</i>	+ 1.7
	<i>Sod3</i>	+ 1.6
	<i>Gstp1</i>	- 2.3
	<i>Gstm6</i>	- 1.8
	<i>Gstm4</i>	- 1.6
<b>Angiogenesis</b>	<i>Angptl3</i>	+ 3.3
	<i>Fgf1</i>	+ 2.0
	<i>Ang</i>	+ 1.5
	<i>Flt1</i>	+ 1.5
<b>Coagulation</b>	<i>Cxcl9</i>	+ 2.1
	<i>Vwf</i>	+ 1.6
<b>Matrix Remodelling</b>	<i>Mmp14</i>	+ 2.0
	<i>Col6a1</i>	+ 1.9
	<i>Col4a1</i>	+ 1.8
	<i>Col4a2</i>	+ 1.5

**Supplementary Table 7. Summary of genes implicated in key enriched pathways**