

Table II – Differential IPA genes in M1+CRC vs M1. Alphabetical lists of genes showing upregulated and downregulated IPA that are common to the two macrophage datasets correspondent to the co-cultures with RKO and HCT-15 cell lines.

Upregulated IPA	Downregulated IPA
<i>CUX1</i>	<i>ACYP2</i>
<i>EIF2A</i>	<i>ATP6V0A1</i>
<i>EIF4G3</i>	<i>C15orf38</i>
<i>ETNK1</i>	<i>CHPF</i>
<i>GRK3</i>	<i>COX17</i>
<i>JDP2</i>	<i>FAM91A1</i>
<i>MAP3K8</i>	<i>MAP3K20</i>
<i>MESD</i>	<i>NOL4L</i>
<i>NRF1</i>	<i>PTGS1</i>
<i>PLEKHA6</i>	<i>RAB6A</i>
<i>SIPA1L</i>	<i>RERE</i>
<i>SMG6</i>	<i>SLC9B2</i>
<i>TIA1</i>	<i>SOGA1</i>
<i>THOC2</i>	<i>TBC1D5</i>
<i>TMED7</i>	<i>WDR33</i>
<i>WDR20</i>	<i>AP3S2</i>
<i>TICAM2</i>	