<table>
<thead>
<tr>
<th>Gene*</th>
<th>ERK regulator/responsive†</th>
<th>NFκB regulator/responsive†</th>
<th>Fold activation‡</th>
</tr>
</thead>
<tbody>
<tr>
<td>Csf2</td>
<td>+</td>
<td>-</td>
<td>3.3</td>
</tr>
<tr>
<td>Ereg</td>
<td>+</td>
<td>-</td>
<td>2.4</td>
</tr>
<tr>
<td>Dusp6</td>
<td>+</td>
<td>-</td>
<td>2</td>
</tr>
<tr>
<td>Osm</td>
<td>+</td>
<td>-</td>
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<tr>
<td>Fosl1</td>
<td>+</td>
<td>-</td>
<td>1.8</td>
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<tr>
<td>Nfkbia</td>
<td>-</td>
<td>+</td>
<td>1</td>
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<tr>
<td>Relb</td>
<td>-</td>
<td>+</td>
<td>1.1</td>
</tr>
<tr>
<td>Nfkb2</td>
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</tr>
<tr>
<td>Ikbke</td>
<td>-</td>
<td>+</td>
<td>1.1</td>
</tr>
</tbody>
</table>

*Genes upregulated > 1.5 fold over unstimulated  
† Gene known to regulate or respond to ERK or NFκB signaling pathway  
‡ Fold activation of *Dusp5*−/− (normalized to *Dusp5*+/+)