

**Supplemental Table 2.** *CYP transcripts detected by microarray analysis in the enterocytes of WT and IE-Cpr-null mice.*

CYPs that were called *present* by the MAS5.0 program in all four enterocyte RNA samples of each mouse strain (IE-Cpr-null or WT) are shown. The results are grouped by *Cyp* subfamily when at least two subfamily members were detected.

<b>Gene Subfamily</b>	<b>Gene Symbol</b>
<i>Cyp1a</i>	<i>Cyp1a1, Cyp1a2<sup>a</sup></i>
<i>Cyp2b</i>	<i>Cyp2b10, Cyp2b9</i>
<i>Cyp2c</i>	<i>Cyp2c29, Cyp2c38, Cyp2c44, Cyp2c55, Cyp2c65, Cyp2c68</i>
<i>Cyp2d</i>	<i>Cyp2d22, Cyp2d26</i>
<i>Cyp2j</i>	<i>Cyp2j6, Cyp2j9</i>
<i>Cyp3a</i>	<i>Cyp3a11, Cyp3a13, Cyp3a25, Cyp3a41a/b, Cyp3a44</i>
<i>Cyp4f</i>	<i>Cyp4f13, Cyp4f14, Cyp4f16</i>
<i>Other</i>	<i>Cyp2s1, Cyp4a10, Cyp4b1, Cyp4v3, Cyp27a1, Cyp51</i>

<sup>a</sup> Detected only in IE-Cpr-null mice