

Supplemental Table 1: Description of primers used for real-time PCR.

<b>Gene</b>	<b>Strand</b>	<b>Primer sequence (5'- 3')</b>	<b>Source</b>
<i>Adh1</i>	<i>Sense</i>	<i>GCTCTGCCGTCAAAGTCGCCA</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>ACCGACACCTCCGAGGCCAA</i>	<i>PrimerBlast</i>
<i>Cyclophilin</i>	<i>Sense</i>	<i>CAGAACAT ATCCCTGCAT</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>G TTCAGCTCTGGGATGACCTT</i>	<i>PrimerBlast</i>
<i>Cyp2a4</i>	<i>Sense</i>	<i>AGCAGGCTACCT TCGACTGG</i>	(1)
	<i>Antisense</i>	<i>GCTGCTGAAGGCTATGCCAT</i>	
<i>Cyp2b9</i>	<i>Sense</i>	<i>CTGAGACCACAAGCGCCAC</i>	(1)
	<i>Antisense</i>	<i>CTTGAGCATGAGCAGGACTCC</i>	
<i>Cyp2d9</i>	<i>Sense</i>	<i>AGTCTCTGGCTTAATTCCTGAT</i>	(1)
	<i>Antisense</i>	<i>CGCAAGAGTATCGGGAATGC</i>	
<i>Cyp7b1</i>	<i>Sense</i>	<i>TGAGGTTCTGAGGCTGTGCTC</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>TCCTGCACTTCTCGGATGATG</i>	<i>PrimerBlast</i>
<i>Cyp3a16</i>	<i>Sense</i>	<i>AGCACCGCGTGGACTTTATT</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>GGGCTGTGATCTCGATTCAG</i>	<i>PrimerBlast</i>
<i>Cyp3a44</i>	<i>Sense</i>	<i>TTGTGGAGGAAGCCAAAAAGTTT</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>TGAGAAGAGCAAAGGATCAAAAAAGT</i>	<i>PrimerBlast</i>
<i>Cyp3a41</i>	<i>Sense</i>	<i>GTGGAGAAAGCCAAAGGGATT</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>GAAGACCAAAGGATCAAAAAAGTCA</i>	<i>PrimerBlast</i>
<i>Mup</i>	<i>Sense</i>	<i>GACTTTTTCTGGAGCAAATCCATG</i>	(2)
<i>1/2/6/8*</i>	<i>Antisense</i>	<i>GAGCACTCTTCATCTCTTACAG</i>	
<i>Mup1</i>	<i>Sense</i>	<i>CAAAACAGAAAAGGCTGGTGA</i>	<i>Designed by</i>

	<i>Antisense</i>	<i>TTGTGCAAACCTTTCCTTGA</i>	<i>PrimerBlast</i>
<i>Ghr</i>	<i>Sense</i>	<i>CCAACTCGCCTCTACACCG</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>GGGAAAGGACTACACCACCTG</i>	<i>PrimerBlast</i>
<i>Prlr</i>	<i>Sense</i>	<i>CACAGTAAATGCCACGAACG</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>GGCAACCATTTTACCCACAG</i>	<i>PrimerBlast</i>
<i>Igf1</i>	<i>Sense</i>	<i>TGTAAACGACCCGGACCTAC</i>	(3)
	<i>Antisense</i>	<i>CACGAACTGAAGAGCATCCA</i>	
<i>Igfbp3</i>	<i>Sense</i>	<i>ACCCAGAACTTCTCCTCCGA</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>CCGCTTCTGCCTTTGGAAG</i>	<i>PrimerBlast</i>
<i>Glucor</i>	<i>Sense</i>	<i>CGGGACCACCTCCCAA</i>	<i>Designed by</i>
	<i>Antisense</i>	<i>CCCCATAATGGCATCCCGAA</i>	<i>PrimerBlast</i>

\* Mayor urinary protein (MUP) primers will potentially amplify four different *Mup* mRNA (if expressed), i.e., *Mup1*, *Mup2*, *Mup6* and *Mup8*.

#### Reference List

1. Wiwi CA, Gupte M, Waxman DJ . Sexually dimorphic P450 gene expression in liver-specific hepatocyte nuclear factor 4alpha-deficient mice. *Mol Endocrinol* 2004; 18:1975-1987.
2. Holloway MG, Laz EV, Waxman DJ . Codependence of growth hormone-responsive, sexually dimorphic hepatic gene expression on signal transducer and activator of transcription 5b and hepatic nuclear factor 4alpha. *Mol Endocrinol* 2006; 20:647-660.
3. Eleswarapu S, Gu Z, Jiang H . Growth hormone regulation of insulin-like growth factor-I gene expression may be mediated by multiple distal signal transducer and activator of transcription 5 binding sites. *Endocrinology* 2008; 149:2230-2240.

Supplemental Table 2: PCR conditions for different genes.

<b>Gene</b>	<b>Denaturation</b>	<b>Annealing/Extension</b>	
<i>Adhl</i>	95°C for 15 s	60°C for 1 min	
<i>Cyclophilin</i>	95°C for 15 s	60°C for 1 min	
<i>Cyp2a4</i>	95°C for 15 s	60°C for 1 min	
<i>Cyp2b9</i>	95°C for 15 s	60°C for 1 min	
<i>Cyp2d9</i>	95°C for 15 s	60°C for 1 min	
<i>Cyp7b1</i>	95°C for 15 s	60°C for 1 min	
<i>Mup 1/2/6/8*</i>	95°C for 15 s	60°C for 1 min	
<i>Mup1</i>	95°C for 15 s	60°C for 1 min	
<i>Ghr</i>	95°C for 15 s	60°C for 1 min	
<i>Igfbp3</i>	95 C for 15 s	60°C for 1 min	
<i>Glucor</i>	95 C for 15 s	60°C for 1 min	
<i>Igf1</i>	95 C for 15 s	60°C for 1 min	
<b>Gene</b>	<b>Denaturation</b>	<b>Annealing</b>	<b>Extension</b>
<i>Cyp3a16</i>	95°C for 15 s	56°C for 20 sec	72°C for 20sec
<i>Cyp3a44</i>	95°C for 15 s	56°C for 20sec	72°C for 20sec
<i>Cyp3a41</i>	95°C for 15 s	56°C for 20sec	72°C for 20sec
<i>Prlr</i>	95°C for 30 s	61°C for 1 min	72°C for 30sec

\* Mayor urinary protein (MUP) primers will potentially amplify four different *Mup* mRNA (if expressed), i.e., *Mup1*, *Mup2*, *Mup6* and *Mup8*.

**Supplemental Figure 1: Expression of liver *Igfbp3* and *Glucor* expression in female and male *Drd2*<sup>-/-</sup>, neuroDrd2KO and lacDrd2KO mice.** Percentage of target mRNA levels normalized to cyclophilin mRNA levels, in relation to control females are represented in the Y axis. N= 5 in *Drd2*<sup>-/-</sup>, 8-10 in neuroDrd2KO and 6-8 lacDrd2KO mice. Two way analysis of variance for the effects of sex and genotype was performed. *P* of interaction was not significant in any case. \* *P* < 0.05 vs sex-matched *Drd2*<sup>+/+</sup> mice.

