

**Supplementary Table 1A.** Cynomolgus macaque and human primer/probe sets used to generate custom Taqman-based assay for qRT-PCR.

Gene	Accession No. <sup>a</sup>	Species <sup>b</sup>		Primer/Probe Sequence (5' to 3') <sup>c</sup>
<i>ESR1</i>	DQ469336	Mf	F	TCACATGATCAACTGGGCAAAGA
			R	AGAAGGTGGACCTGATCATGGA
			P	CACAAAGccTGGCACCC
<i>ESR2</i>	HQ702565	Mf	F	GTATCTCTGTGTCAAGGCCATGAT
			R	CATCCTGGGTCGCTGTGA
			P	CCTGCTCAATTCCaaTATGT
<i>PGR-B*</i>	-	Hs	F	GCCAGACCTCGGACACCTT
			R	CAGGGCCGAGGGAAAGAGTAG
			P	CCTGAAGTTCCGGCCATACCTATCTCCCT
<i>TFF1</i>	DQ464113	Mf	F	GTGCTTCCATCCTAATACCATCGA
			R	GCAGATCCATGCAGAAGTGTCTAAA
			P	TCCCTCCAGAA <u>Ag</u> GGAGTGT
<i>CYP19</i>	DQ529980	Mf	F	CCCCAAACCCAATGAATTACTCTTG
			R	GGCCCAAAGCCAAATGGTT
			P	AAAGTACCTATAAGGA <u>A</u> caTTCTT
<i>HSD17B1</i>	DQ529983	Mf	F	CAGGCCTGGGCCTACTG
			R	CGTTCACGTCCAGCACAGA
			P	<u>CCCACCGCCTCCAGC</u>
<i>HSD17B2</i>	DQ529984	Mf	F	AAAGGAAGGCTGGTGAATGTCA
			R	ATGCCAGCTTGCCATTGG
			P	CCCCT <u>cc</u> TCCCAGCTG
<i>STS</i>	DQ529981	Mf	F	GGGATGCTGTTGAGGAAATGGA
			R	CCAATCTCAGCTCATCTAGAAGGTT
			P	ATCTGC <u>cc</u> CACACTCC
<i>SULT1E1</i>	DQ529982	Mf	F	GAGAGAACGGGCCAGGTT
			R	CGAGAGGTGTCCCTGGATCAG
			P	ATGTT <u>cc</u> TGAGCTCTCG

Gene	Accession No. <sup>a</sup>	Species <sup>b</sup>		Primer/Probe Sequence (5' to 3') <sup>c</sup>
<i>CYP1A1</i>	D17575	Mf	F	CCACAGCCACCTCCAAGATC
			R	TGGCCGACATGGAGATTG
			P	CCTACACTGATCATGCTTT
<i>CYP1B1</i>	AB179009	Mf	F	CCCGCGCTGCAGTGG
			R	CACTCGAGCCTGCACATCA
			P	CTCCTCTTCATCAggtATCC
<i>CYP3A4</i>	S53047	Mf	F	CACACCTTGCCCTTATTGGAAAT
			R	AGCCCCACACTTTCCATACTTT
			P	CaaACGTCCAAAAGCC
<i>GAPDH</i>	DQ464111	Mf	F	TGCCCTCAATGACCACTTGTC
			R	ACCCTGTTGCTGTAGCCAAAT
			P	TCGTTGTCATAccAGGAAAT
<i>ACTB</i>	DQ464112	Mf	F	ACCCCAAGGCCAACCG
			R	CCTGGATGGCCACGTACATG
			P	AAGATGACCCAgtaTCATG

<sup>a</sup> Genbank identification code or cited reference. <sup>a</sup>*Hs*, *Homo sapiens*; *Mf*, *Macaca fascicularis* (cynomolgus macaque). <sup>c</sup> Bold letters indicate differences between macaque and human sequences. Italicized lowercase of base pairs indicate exon boundaries. Underlined sequence represents an 8-base deletion in the HSD17B1 pseudogene. F, forward primer; R, reverse primer; P, probe. Asterisk (\*) indicates that the assay was based on that reported in Sakaguchi et al. Gynecol Oncol. 2004 May;93(2):394-9.

**Supplementary Table 1B.** Applied Biosystems (ABI) Taqman gene expression assays used in qRT-PCR

Gene ID	NCBI Accession no.	Species <sup>a</sup>	ABI assay ID
<i>PGR</i> <sup>b</sup>	XM_001095317.1	<i>Mm</i>	Rh02830094_m1
<i>GREB1</i>	XM_001089374.1	<i>Mm</i>	Rh02866842_m1

<sup>a</sup>*Mm*, *Macaca mulatta* (rhesus macaque). <sup>b</sup> PGR was used to generate PGR-A expression level; quantification of PGR-A was done by subtraction of PGR-B (custom assay) from PGR, as previously described in Sakaguchi et al. Gynecol Oncol. 2004 May;93(2):394-9.