Supplemental table 1

cloning primers		Forward	Reverse	Source
Lentiviral cloning	ERRα	TGAGCATCCCAGGCTTCTCGTTCAAGAGATGAGAAGCCTGGGATGCTCTTTTTTC	TCGAGAAAAAAGAGCATCCCAGGCTTCTCATCTCTTGAACGAGAAGCCTGGGATGCTCA	Sladek et al, 1997
	HNF4α	TGATCAGCACTCGAAGGTCGTTCAAGAGATGACCTTCGAGTGCTGATCTTTTTTC	TCGAGAAAAAAGATCAGCACTCGAAGGTCATCTCTTGAACGACCTTCGAGTGCTGATCA	
DR1 mutagenesis	-2032 DR1	TGATTATTTCCCATGATCAGTCTCGTTGTA	AAAAACACTTTCTGGGTAGGGAAGATAATT	
	-878 DR1	ATTGCAAAAGCAGAACTACCGATAAGG	TTGTTTTTTCCCTCAGAAGCATGTGATCTT	
	hHNF4α2	tattagaattcAGAATGCGACTCTCCAAAACC	tatactctagaCAGCGGCTTGCTAGATAACTTC	NM_000457
qPCR primers	human *36B4*	GGACATGTTGCTGGCCAATAA	GGGCCCGAGACCAGTGTT	
	human *CYP17A1*	CCGTAAGGGTATCGCCTTCG	GATGTCTATGGACTGTCCGTTG	
	human *CYP11A1*	ACCAGAGACCCATAGGAGTCC	CAACAGGGGCAAAAAGTTCTTG	
	human *CYP21*	CTCCACCTCCCGCCTCTTG	AGTTCAGCACCACCACATCTTG	
	human *CYP11B1*	GGGTGGCCTACAGACAACATC	GGCGACAGCACTTCTGGATT	
	human *3βHSD1*	TGACAGGAGCAGGAGGGTTTC	CAGGCGGTGTGGATGATGAC	
	human *3βHSD2*	GAACGGCCACGAAGAAGAG	TGGGTCTTAACGCACAAGTGT	
	rat *cyclophilin A*	aaggtgaaagaaggcatgagca	agttgtccacagtcggagatgg	
	rat *Pgc-1alpha*	ATGAGAAGCGGGAGTCTGAA	GCGGTCTCTCAGTTCTGTCC	
	rat *Cyp17A1*	TTTCCCAACAAAGGCTTGGAGGTG	AATCAGAATGTCCGTCAGGCTGGA	
	rat *Cyp11A1*	TCCTCAAAGCCAGCATCAAGGAGA	TACGAAGCACCAGGTCGTTCACAA	
Chip primers	ERRalpha promoter	CCATCCGAGTGGAATTTGAGTCCTAAAG	GAACCGTAGACCCAGTAGCCCCACAGAG	Laganiere et al, 2004
ERREs for CYP11A1	-9298	TGCCCTTTCACTACAGCAGA	ATCCCTCCCAGAACACACCT	
	-8650	TCTGACCCTACCTGCCTGAT	GGCTCTTTAAGGCCAGTTCC	
	-1546	CGTGAGACACTGCCTTCCTT	GAGTGGTTGTTGCTCTTCCC	
	-873	GGGTATGGAAGCAGCTCTGA	GACCTTGGCAAATCTGGTTG	
	-264	GCACAGGCAGATATTTCAGGA	AATTTCTGCAGCGTGAGGTT	
	1480	GGGAGGGAGATTTGAAGGTC	GGAGGAATGCTGGGTCTGT	
	15235	GCTCATGCCTGTAATCCCAGTGC	CTG CCT CAG CCT CCT AAG AAG CT	
	19455	TGTGGACAGGTGAGAAGCTG	GACCAGCAAGTCCTCCACTC	
	24042	TTCTCTGGCTGTGCTCTCCT	TTGACATAGCGTGGGACAAA	
	24348	GAGGGCAAAGGACCAGTGT	CATACCCTGACGAGCCTTCT	
	25789	GAGGCAGCAGTGTGAACG	TTTGTCAGTGGCAGGTGCT	
	26180	AGTGACAGAGGCAAGCACCT	CTCTCTCCTCCTGGCAACAG	
	26646	CCAGGAAGAGGACCTAGCG	TACATGGAGTGGCACACAGC	
	28661	GTCTGAATGAAGGGCAGAGG	CTCAGACCCAGGCAAATCAT	
	34696	GGAGACCCTGGGAAGCTTAG	CCCAGAAATCACCCATGGTA	
	38065	CTCACGTCACTGAGAACCCA	GCCCAGTCACCTTCCAAGT	
ERREs for CYP17A1	-16617	CTTCCGCTTTGCTTGTCTTT	GTCGCCATCATTTCAGTTAGC	
	-8555	ATGGGTGGTGAAGAGGATGT	AATCCAGATGATGACAATATTCCTT	
	72	GGTAAGCAGCAAGAGAGCCA	TCCTTGTGCCCTAGAGTTGC	
	1294	GCAGAGGTGGGAGCTAGGTA	CAGCTTTGATCAAGCCAAGG	
	3292	CAGCTGCCAGAGTCTCTCCT	TTCCATTCTGGCTGAAGGTC	
	9334	GAGAAGGAGGTGGAACCCA	ACTGCCAGGTAAGGGCTTCT	
	12177	AGCTGAGTGCTGAAGAGGGA	GAACTTTGACTCCTGCCCAA	
	14189	GCCACCAGAGGCTGATAGAA	CCTTGTTGACAGGAGGGAAA	
	19307	CCACCAGTGTGCCAGAATC	AGAGGTCAGTCTTAAGCAGATGG	
	22402	TCTGTTGTGTGTGTGCCTCA	TGTGGTTCACGTGGAGAGAA	
	25324	GTCACCCATCTGCTCCTGAT	TCCTCCTAACCTTCCAAATGC	
	25500	AATAAGCCTTGGATGGCAGA	CTACTGTGGCTGGAGGTGGT	
DR1 sites for CYP17A1				
	CYP7A promoter	CCAGGTCCGAATGTTAAGTCA	AAGTCCACAGGTATCAGAAGTGG	
	14644	CTTGGTTCCCAGGTCACAGT	AACGATTCCTTCAGCTCCCT	
	-878	TGGAAGGGAAGGGAGAGATA	CTGCTGATGAGCAAAGAAGG	
	-2032	CTTGCTCGTTCTCTCGGTTT	TGTTTCTCTGTTTAATCGGCTTG	
	-3542	GGTACGCAGCCCAGACTTGA	CTC TCA CCT GGA TGG ATG CTC C	
	-6837	CACCTGAGCCTCAAACCAGC	GCA CCG CTG AAC ATA GAC CC	
	-14075	CGATGGTGCGTACCTGTAGTCC	GTT CAA TAT CAG CAG CCT GGG C	