Table 2. Polymorphic loci of exons and adjacent intronic sequences of the carnitine

 acyltransferases genes in Siberian populations

Gene	Polymorphic loci	
	Exons	Adjacent introns
CPTIA	rs80356779 (P479L), rs2228502	rs146937052, rs11228340, rs2278908,
	(F417F), rs2229737 (E321E),	rs545441003, rs22780907,
	rs2229738 (A275T)	rs531240052, rs574929166
CPT1B	<u>rs470117</u> (E531K), <u>rs8142477</u>	rs15195, rs7238, rs17848472,
	(S427C), rs2269383 (G320D),	rs17848473, rs117635543,
	rs17848457 (D268D), <u>rs3213445</u>	rs17848459, rs131758
	(I66V)	
CPT1C	rs763273578 (T740A)	rs1075453, rs115254624, rs892149,
		rs116967749, <u>rs11881828</u> ,
		<u>rs72626236</u> , rs12462023, <u>rs72626237</u> ,
		rs200917939
CPT2	rs2229291 (F352C), rs1799821	
	(V368I), rs1799822 (M647V)	
CRAT	<u>rs17459086</u> (A603P), rs2228304	rs2229419, <u>rs3780691</u>
	(S118S), новый (9: 131866581)	
	(S99F)	

Non-synonymous substitutions are shown in bold. Amino acid substitutions are given in parentheses. The linked polymorphic variants (in lines) are underlined. For the coupled variants at the loci rs470117, rs8142477 and rs3213445 of the *CPT1B* gene haplotypes CCC and TGT were identified; at the loci rs11881828, rs72626236 and rs72626237 of the *CPT1C* gene - haplotypes GAG and AGA; at the loci rs17459086 and rs3780691 of the *CRAT* gene - haplotypes GC and CT.