Table S4 Polymorphism analysis of rs3951819 (A/G SNP) in BHLHA9

	Polymorphism analysis					Statistical analysis ( <i>P</i> -value)			
	Patients				Carriers	SHFM	Carriers vs.		
	Total	SHFM	SHFLD	GWC	_	VS.	Patients		
	(n=27)	(n=15)	(n=10)	(n=2)	(n=17)	SHFLD	Total	SHFM	SHFLD
<allele frequency*=""></allele>									
A	16	7	7	2	11	0.23	0.49	0.25	0.56
G	11	8	3	0	6				
<genotype frequency=""></genotype>									
AA/A	6	4	1	1	4				•••
AA/G	2	2	0	0	3				
AG/A	5	2	3	0	1				
AG/G	5	4	1	0	3				•••
AAA/A	4	0	3	1	4				
AAA/G	1	0	1	0	0				
AAG/A	0	0	0	0	2				
AAG/G	2	1	1	0	0				
AGA/A	0	0	0	0	0			•••	•••
AGA/G	0	0	0	0	0	•••		•••	•••
AGG/A	1	1	0	0	0				
AGG/G	1	1	0	0	0				

<sup>\*</sup> On the duplications/triplications negative normal chromosome 17.

The 95% confidence intervals for odds ratios include 1.0 in all the comparisons (not shown).

P-values have not been caluculated for GWC, because of the small patient number.

The allele frequency in the Japanese population is 60% for A and 40% for G