

Supplementary Table S6. Microsatellite and SNP analysis of SALTAS broodstock families in which the sex-determining gene housed on Ssa03 (AS-11).

Markers are arranged in order corresponding to their placement on the female European Atlantic salmon linkage map. [www.asalbase.org](http://www.asalbase.org)

Marker	2009_32 <sup>1</sup>	2009_37	Total	LOD	Theta
<b>Ssa02<sup>2</sup></b>					
Ssa1077BSFU	28/35 <sup>3</sup> (2.9) <sup>4</sup>				
Ssa0055BSFU					
Ssa0183BSFU					
Ssa202DU		13/17 (1.1)			
Ssa0233BSFU					
Ssa1265BSFU					
Ssa0182ECIG					
<b>Ssa06</b>					
OMM5037	16/35 (0)				
Ssa0074ECIG		10/17 (0.1)			
Ssa0376BSFU					
Ssa0789BSFU					
Ssa0043BSFU					
Ssa0103aECIG	12/35 (0)				
Ssa0142BSFU					
Ssa0350BSFU					
Ssa1299BSFU		8/17 (0)			
Ssa0752BSFU					
<b>Ssa03</b>					
Ssa0649BSFU	28/33 (3.8)		28/33	3.8	0.15 Ssa0649BSFU
Ssa0017BSFU	30/34 (4.9)	15/16 (3.2)	45/50	8	0.1 Ssa0017BSFU
Ssa0368BSFU	31/35 (5.1)		31/35	5.1	0.11 Ssa0368BSFU
Ssa0670BSFU		13/17 (1.1)	13/17	1.1	0.23 Ssa0670BSFU
OMM1267	24/26 (4.8)	14/15 (2.9)	38/41	7.7	0.07 OMM1267
Ssa0516BSFU	30/34 (4.9)	15/16 (3.2)	45/50	8	0.1 Ssa0516BSFU
OMM5019	29/32 (5.3)		29/32	5.3	0.09 <u>OMM5019</u>
SsaD144	30/34 (4.9)				

<sup>1</sup>Family (2009 year class where family ID's are given as

XX, values below). <sup>2</sup> Chromosome number shown in

Ssa0X. <sup>3</sup>(Number of offspring for which genotype and phenotype agree) / (number of offspring for which

genotype and phenotype disagree). <sup>4</sup>LOD score corresponding to the ratio of correct genotype and phenotype matches:incorrect genotype and pheoptype