Table 3 Allele frequencies across reporting units of the two late-father alleles most common amongst late-fall Chinook: nucleotide G at is the total number of gene copies available for estimating the allele Otsh_v2.0.

	Chr34:954,054		Chr34:1,151,868	
Reporting Unit	freq.	N	freq.	N
CV-Late-Fall	0.750	516	0.733	592
CV-Fall	0.054	910	0.030	908
CV-Spring	0.023	396	0.003	396
CV-Winter	0.005	222	0.005	222
Cent-Cal-Coast	0.000	92	0.004	278
Klamath-Trinity	0.000	172	0.000	358
SO-NCal-Coast	0.000	192	0.000	304