

## Column Assignments for fugu-based stickleback SNPs

This is a tab-separated file with the following columns:

1. schr stickleback chromosome
2. spos stickleback position
3. fchr fugu chromosome
4. fpos fugu position
5. A reference nucleotide
6. B variant nucleotide

For each of 21 individuals there are four columns.

- 7-10. fresh1 -- BEPA
- 11-14. fresh2 -- BIGL
- 15-18. fresh3 -- FTC
- 19-22. fresh4 -- HUTU
- 23-26. fresh5 -- MATA
- 27-30. fresh6 -- MUDL
- 31-34. fresh7 -- PAXB
- 35-38. fresh8 -- NOST
- 39-42. fresh9 -- SCX
- 43-46. fresh10 -- SHEL
- 47-50. fresh11 -- TYNE\_8
- 51-54. salt1 -- TYNE\_1
- 55-58. salt2 -- NEU
- 59-62. salt3 -- GORT
- 63-66. salt4 -- ANTL
- 67-70. salt5 -- GJOG
- 71-74. salt6 -- BGR
- 75-78. salt7 -- RABS
- 79-82. salt8 -- SALR
- 83-86. salt9 -- BDGB
- 87-90. salt10 -- JAMA