

## **Column Assignments for fugu-based stickleback SAPs**

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

1. ref - stickleback chromosome
2. rPos - position on stickleback chromosome
3. trns - ENSEMBL transcript name
4. pep - ENSEMBL peptide name
5. AA1 - one amino acid
6. loc - location in the peptide sequence
7. AA2 - variant amino acid