

**Table S16 GO term enrichment for cellular component in the North American cline for genes that overlap windows in the top 1% tail of Fst.**

1 % Tail Cellular Component		
GO term	q-value	description
GO:0005634	6.87E-09	nucleus
GO:0005622	4.74E-03	intracellular
GO:0005886	4.74E-03	plasma membrane
GO:0016021	2.47E-02	integral to membrane
GO:0005875	2.47E-02	microtubule associated complex
GO:0005938	2.47E-02	cell cortex
GO:0005887	2.47E-02	integral to plasma membrane
GO:0016020	2.93E-02	membrane
GO:0005737	4.09E-02	cytoplasm
GO:0005794	4.95E-02	Golgi apparatus