

Table S17 GO term enrichment for cellular component in the North American cline for genes that overlap windows in the top 2.5% tail of Fst.

2.5% Tail Cellular Component		
GO term	q-value	description
GO:0005634	2.94E-12	nucleus
GO:0005886	8.90E-10	plasma membrane
GO:0005737	1.72E-04	cytoplasm
GO:0045211	8.60E-04	postsynaptic membrane
GO:0016021	8.60E-04	integral to membrane
GO:0005622	8.60E-04	intracellular
GO:0005578	8.60E-04	proteinaceous extracellular matrix
GO:0045179	8.60E-04	apical cortex
GO:0005887	1.11E-03	integral to plasma membrane
GO:0045202	2.75E-03	synapse
GO:0005918	3.83E-03	septate junction
GO:0005875	4.55E-03	microtubule associated complex
GO:0016459	6.83E-03	myosin complex
GO:0005794	1.15E-02	Golgi apparatus
GO:0005938	1.15E-02	cell cortex
GO:0005576	2.29E-02	extracellular region
GO:0005912	2.30E-02	adherens junction
GO:0005700	2.37E-02	polytene chromosome
GO:0016020	4.28E-02	membrane
GO:0008021	4.90E-02	synaptic vesicle