

Table S7 GO term enrichment for cellular component in the Australian cline for genes that overlap windows in the top 1% tail of Fst.

GO term	1 % Tail Cellular Component	
	q-value	description
GO:0005886	1.97E-06	plasma membrane
GO:0045211	2.47E-04	postsynaptic membrane
GO:0042600	6.03E-04	chorion
GO:0005887	6.03E-04	integral to plasma membrane
GO:0005578	2.51E-03	proteinaceous extracellular matrix
GO:0043234	2.51E-03	protein complex
GO:0016020	2.51E-03	membrane
GO:0005622	1.45E-02	intracellular
GO:0005634	2.82E-02	nucleus
GO:0016021	3.70E-02	integral to membrane
GO:0005875	3.70E-02	microtubule associated complex