

Table 1: Estimates of the DFE obtained from troughs in diversity around functional elements. Standard errors are shown in square brackets

BGS	Gene Conversion	Protein-Coding Exons		Conserved Non-Coding Elements	
		γ_a	p_a	γ_a	p_a
+	+	[]	[]	[]	[]
-	+	[]	[]	[]	[]
+	-	8,350 [1,680]	1.29×10^{-5} [4.23×10^{-6}]	228 [12.8]	2.27×10^{-3} [2.40×10^{-4}]
-	-	20,200 [1,460]	8.61×10^{-6} [9.52×10^{-7}]	504 [18.2]	1.27×10^{-3} [7.12×10^{-5}]