

Table S2: List of putative loci under incomplete selective sweeps in *D. melanogaster* Rwanda population inferred from nS_L test.

Chromosome	Cluster start – end	Minimum standardized nS_L	Site of minimum nS_L	Derived allele frequency
2L	1147263-1152901	-3.77	1147263	8/22
	1545621-1548645	-3.68	1545621	14/22
	5812475-5816415	-3.15	5815486	8/22
	6596769-6603746	-3.99	6603091	9/22
	9433919-9610983	-3.33	9433919	17/22
	12372465-12377872	-3.46	12375980	11/21
	12718277-12844111	-3.10	12834789	8/21
	16798951-16894905	-3.35	16894905	12/21
	17234502-17245049	-3.45	17235226	9/21
	17602170-17619087	-3.72	17603916	16/21
	19727261-19768977	-3.44	19730058	8/21
2R	3786646-3886834	-2.42	3886479	8/21
	5254576-5283840	-2.54	5266445	8/21
	5548485-5552974	-2.42	5548485	9/21
	7119351-7133870	-2.47	7126724	12/22
	7816110-8669818	-3.29	8133130	17/22
	12721950-12740626	-2.83	12727369	8/22
	13826522-13834107	-2.46	13826676	8/22
	18089346-18541764	-2.80	18089697	8/22
3L	2997574-2998073	-3.82	2998073	8/22
	3136553-3149179	-4.49	3144835	11/22
	6087588-6129005	-4.51	6129005	16/21
	6538594-6567487	-4.18	6547881	8/21
	11654477-11679598	-3.59	11654544	9/21
	11826677-11839249	-3.88	11833069	12/18

	13427088-13432591	-3.44	13430725	11/22
	16076838-16137484	-4.20	16094714	12/22
	19210376-19236655	-3.80	19210732	8/22
3R	10230101-10904118	-3.14	10333644	14/22
	12062855-12066090	-3.37	12062855	12/22
	13330699-13334656	-3.14	13330699	15/22
	13906691-13911447	-3.41	13908193	10/22
	15432404-15445674	-3.46	15443091	12/22
	16561264-16575140	-3.22	16569519	13/21
	19838638-19884976	-3.17	19839035	12/22
	20872678-20878483	-3.49	20876949	12/22
X	2814198-2838349	-5.45	2833040	9/22
	10210768-10225115	-4.21	10219139	8/22
	14151360-14164063	-4.50	14157513	16/21
	14969055-14996063	-4.16	14975977	13/21
	16948533-17158766	-4.25	17124249	8/19