

Table S1: List of putative loci under incomplete selective sweeps in *D. melanogaster* Rwanda population inferred from *iHS* test.

Chromosome	Cluster start - end	Minimum standardized <i>iHS</i>	Site of minimum <i>iHS</i>	Derived allele frequency
2L	1547008 - 1557247	-3.75	1547204	15/22
	4824296 - 4860577	-4.03	4824431	10/22
	5810884 - 5825009	-3.34	5815486	8/22
	6020532 - 6055868	-3.34	6055868	9/22
	9509499 - 9831699	-5.90	9582539	10/20
	11022970 - 11036917	-3.82	11036917	9/21
	11866168 - 11892750	-3.52	11892750	9/21
	12804885 - 12840609	-3.76	12840609	10/21
	16019930 - 16020004	-3.17	16019930	9/21
	17230328 - 17297935	-3.84	17297624	12/21
	17602339 - 17603635	-3.32	17603635	13/21
2R	5649989 - 5718051	-3.33	5708056	11/20
	7114975 - 7137571	-3.87	7127281	11/22
	7828412 - 8658752	-5.19	7911386	12/22
	10135366 - 10665005	-3.86	10665005	9/22
	11001899 - 11016632	-3.28	11006196	9/22
	12723745 - 12768255	-4.38	12734718	10/21
	13832525 - 13835190	-3.49	13832748	11/22
3L	3103656 - 3145740	-4.84	3142414	9/22
	4472504 - 4490155	-4.35	4490062	8/22
	5960208 - 5974572	-4.04	5966477	8/22
	6072249 - 6129005	-5.13	6109760	10/21
	6537946 - 6595815	-4.26	6548571	12/21
	8126905 - 8182746	-4.18	8134344	10/22
	14430470 - 14434825	-4.12	14431036	11/18
	16070044 - 16095749	-4.48	16095749	12/22
	19210723 - 19236655	-4.14	19210732	8/22
3R	8567616 - 8567660	-3.70	8567660	9/19

	9033505 - 9157259	-3.63	9125222	8/19
	10333644 - 10389581	-3.46	10386821	10/22
	12063858 - 12066090	-3.12	12063858	14/22
	13905933 - 13911447	-3.84	13910288	12/22
	15427616 - 15503321	-3.42	15442743	13/22
	16254275 - 16259857	-3.40	16254275	11/22
	18973474 - 18973529	-3.49	18973529	13/22
X	737336 - 1229075	-4.50	1081402	8/22
	2814828 - 2836819	-5.68	2832651	9/22
	17230521 - 17234450	-4.88	17230521	8/21
	18636049 - 18639500	-5.23	18636156	8/20
	19077625 - 19104409	-5.80	19093150	8/20