

Population	Latitude (N), Longitude (E)	Population size	Sample size (morph freq)	Floral Morph			Sample size (pop genetics)	F_{is}	$s(BES)$	$G'st$
				L	S	H				
D01	-2.58°, 51.18°	273	100	0.51	0.49	0	30	0.131	0.180	0.06
D02	-2.63°, 50.98°	~300	96	0.64	0.36	0	30	0.066	0.154	0.07
D03	-2.33°, 50.85°	~500	110	0.44	0.56	0	30	0.071	0.131	0.08
D04	-2.20°, 50.95°	65	65	0.48	0.52	0	24	0.143	0.212	0.08
D05	-2.22°, 51.05°	260	105	0.46	0.54	0	30	0.105	0.318	0.07
D06	-2.49°, 51.23°	~350	101	0.58	0.42	0	30	0.009	0.122	0.09
D07	-2.59°, 51.21°	~250	100	0.58	0.42	0	30	0.149	0.348	0.07
D08	-2.48°, 51.16°	150	94	0.37	0.63	0	29	-0.01	0.222	0.07
D09	-2.49°, 51.05°	~300	100	0.57	0.43	0	29	0.002	0.168	0.07
D10	-2.71°, 51.09°	~500	104	0.52	0.48	0	29	-0.085	0.052	0.08
D*11	-2.42°, 51.13°	58	50	0.34	0	0.66	30	0.065	0.218	0.07
T01	-2.44°, 51.00°	94	94	0.63	0.31	0.06	30	0.117	0.295	0.08
T02	-2.38°, 50.92°	~150	108	0.62	0.32	0.06	24	0.124	0.255	0.07
T03	-2.43°, 51.18°	88	88	0.41	0.38	0.22	30	0.052	0.219	0.07
T04	-2.40°, 51.16°	~200	104	0.36	0.16	0.48	31	0.389	0.529	0.10
T05	-2.25°, 50.95°	20	15	0.33	0.07	0.53	17	0.284	0.473	0.10
T06	-2.56°, 51.14°	79	79	0.41	0.06	0.53	28	0.225	0.422	0.07
T07	-2.30°, 51.08°	103	85	0.29	0.09	0.61	30	0.227	0.394	0.08
T08	-2.58°, 51.09°	161	95	0.36	0.02	0.62	30	0.126	0.314	0.15
T09	-2.55°, 51.05°	87	87	0.33	0.03	0.63	31	0.175	0.318	0.10
T10	-2.42°, 51.10°	~150	82	0.26	0.02	0.72	30	0.251	0.367	0.09
M01	-2.63°, 51.03°	19	19	0	0	1	11	0.272	0.461	0.23
Mean (\pm SD)								0.131 (\pm 0.11)	0.281 (\pm 0.13)	0.088 (\pm 0.03)

F_{is} varies from -1 (heterozygotes excess) to + 1 (homozygotes excess due to inbreeding); values closer to 0 indicate random outcrossing under HWE; population level estimates of selfing rates vary from 0 (maximum outcrossing) to 1 (maximum self-fertilization).