

Table S4. P-values per field and overall for distortion towards *L. serriola* and *L. sativa* for the BC_{1S1} generatio. Based on a Monte Carlo algorithm, 50,000x drawing N plants from the expected distribution per locus over all fields and n per field. Employed p-value for significance (1/# informative loci): 1/220 = 0.0047. Red is distortion towards *L. serriola*, blue is distortion towards *L. sativa*. Only significant loci joining all fields in one analyses are reported in Fig. 2, discussion of single field distortion is considered beyond our scope. For two loci below indicated in yellow significant distortion for all fields could be based on one field only (other fields: p >0.05). For all others: at least two fields are significant or providing a similar trend (p<0.05) suggestion a general field exposure based selection of genotypes. m.d. = missing data. See method section for full methodological details

Number (#)	Marker name	Linkage group	cM	All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21		All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21
				<i>L. serriola</i>	<i>L. serriola</i>					<i>L. serriola</i>	<i>L. serriola</i>			
1	E35/M59-F-127-SER	1	4.1	m.d.	m.d.	m.d.	m.d.	m.d.		m.d.	m.d.	m.d.	m.d.	m.d.
2	E45/M48-F-070-SAT	1	16.1	m.d.	m.d.	m.d.	m.d.	m.d.		m.d.	m.d.	m.d.	m.d.	m.d.
3	C09C-CTT-013-SAT	1	16.3	0.46064	0.06816	0.67464	0.59182			0.45398	0.8632	0.22124	0.25118	
4	C09C-ACG-008-SER	1	18	0.99902	0.98868	0.98394	0.5871			0.00032	0.00244	0.00484	0.16042	
5	NBS-33-sat	1	28	0.9999	0.48034	1	0.87422			0.99988	0.9955	0.99234	0.75228	
6	E45/M48-F-295-SAT	1	28.7	0.43614	0.03542	0.81572	0.55754			0.94522	0.20822	0.96656	0.83144	
7	NBS-13-ser	1	29.5	1	1	1	0.98912			0.02372	0.01754	0.53312	0.01088	
8	E44/M48-F-297-SAT	1	33.5	0.63706	0.70766	0.02576	0.97474			0.53108	0.05556	0.9124	0.33214	
9	E44/M48-F-109-SAT	1	40.9	0.2226	0.14378	0.42934	0.34736			0.84908	0.35944	0.61782	0.9026	
10	C09C-ATC-011-SAT	1	71	1	0.99998	1	1			0.00002	0.00002	0	0.00002	
11	E44/M48-F-114-SER	1	78.8	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
12	C09C-ATA-005-SER	1	86.3	1	0.70664	0.99934	0.99928			0.00002	0.09074	0.00008	0.00008	
13	E45/M48-F-297-SAT	1	97.3	1	0.93998	1	1			0.26708	0.30998	0.12784	0.42026	
14	E45/M48-F-235-SER	2	1.9	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
15	NBS-23-ser	2	13.2	0.99992	0.99758	0.99994	m.d.			0.85566	0.07918	0.93354	0.90404	
16	C09C-ACT-012-SAT	2	17	0.8437	0.97364	0.16532	0.67074			0.10974	0.01028	0.7365	0.1897	
17	E35/M59-F-273-SAT	2	25	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
18	NBS-30-ser	2	32.4	1	0.99996	1	0.99818			0.17958	0.60732	0.27478	0.02246	
19	E44/M48-F-147-SAT	2	39.6	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
20	C09C-ACT-004-SER	2	61.4	1	0.99906	1	0.68924			0.24194	0.56412	0.16706	0.14244	
21	C09C-ATC-010-SAT	2	62.4	0.99982	0.89164	0.99952	0.82836			0.03718	0.10836	0.00876	0.66086	
22	C09C-CTT-008-SER	2	65.3	0.99948	0.99224	0.95202	0.94198			0.14112	0.34958	0.22164	0.05802	
23	C09C-ACG-014-SAT	2	67.4	0.56112	0.7845	0.55402	0.13372			0.43888	0.2155	0.31864	0.73266	
24	C09C-GTT-216-SER	2	81.3	1	1	1	0.76372			0.006	0.07828	0.013	0.08274	
25	C09C-CTG-319-SER	2	90.5	0.99986	0.99896	0.98858	0.77448			0	0.0003	0.00344	0.07866	
26	E44/M49-F-212-SAT	3	7.9	0.02908	0.00446	0.66524	0.01062			0.95832	0.61402	0.6906	0.96184	
27	NBS-01-ser	3	8.8	1	0.96814	1	0.32526			0.38512	0.07932	0.30602	0.8714	
28	E44/M49-F-082-SAT	3	11.6	0.04106	0.00396	0.58804	0.0641			0.71928	0.44082	0.24362	0.96694	
29	C09C-AAT-305-SER	3	13.1	0.99988	0.9994	0.9982	0.14742			0.00082	0.01006	0.00042	0.55532	
30	C09C-CTG-328-SER	3	18.8	0.55596	0.35508	0.84418	0.0824			0.32224	0.41016	0.07358	0.6995	
31	E45/M48-F-207-SER	3	21.5	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
32	C09C-CTA-600-SAT	3	24.2	0.10164	0.10966	0.14634	0.33908			0.85636	0.80486	0.76956	0.49202	
33	NBS-34-sat	3	26.5	0.0025	0.00058	0.59798	0.00384			1	1	1	1	
34	E35/M59-F-281-SER	3	27.2	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
35	C09C-CTA-325-SAT	3	29.7	0.00156	0.10352	0.01628	0.00556			0.9973	0.81428	0.96584	0.98132	

Number (#)	Marker name	Linkage group	cM	All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21		All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21
				<i>L. serriola</i>	<i>L. serriola</i>					<i>L. sativa</i>	<i>L. sativa</i>			
Distortion towards preference for:														
36	E44/M49-F-479-SAT	3	37.1	0.98658	0.96128	0.69354	0.88218			0.71176	0.28838	0.75398	0.49944	
37	E44/M49-F-362-SAT	3	44.9	0.48544	0.64502	0.18334	0.5465			0.21352	0.01158	0.47886	0.68162	
38	E45/M48-F-287-SER	3	46.6	0.00054	0.04884	0.00164	0.07982			0.82642	0.3211	0.9647	0.46028	
39	NBS-18-sat	3	47.1	0.5654	0.46782	0.81516	0.08628			1	0.97504	1	0.99924	
40	CO9C-ACG-011-SER	3	50.8	0.46262	0.65326	0.28698	0.2586			0.95624	0.88866	0.89464	0.56848	
41	NBS-20-ser	3	52	1	0.99768	1	0.72288			0.99338	0.7921	0.99634	0.65488	
42	1A36-F-195-SAT	3	52.2	0.00642	0.06088	0.0605	0.02552			0.99698	0.7499	0.99972	0.57016	
43	1A31-F-107-SAT	3	52.4	0.30568	0.35338	0.54292	0.0809			0.72062	0.51002	0.73042	0.3459	
44	E35/M59-F-299-SAT	3	52.5	0.04702	0.5245	0.04218	0.03672			0.98612	0.68322	0.92368	0.89474	
45	CO9C-CTA-142-SAT	3	53.1	0.00002	0.00002	0.00014	0.00002			1	1	0.99974	1	
46	CO9C-GTT-241-SER	3	58.8	0.73082	0.5212	0.50466	0.63488			0.26918	0.3029	0.49534	0.19434	
47	1A31-F-122-SER	3	72.5	0.85074	0.086	0.99096	0.63696			0.60164	0.68984	0.517	0.20712	
48	NBS-32-ser	3	72.5	0.99996	0.98748	0.99998	0.52746			0.99186	0.26444	0.9991	0.85528	
49	1A31-F-124-SER	3	73.1	0.08202	0.00216	0.55744	0.47958			0.6624	0.53786	0.5591	0.41098	
50	NBS-35-ser	3	83	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
51	CO9C-GTG-070-SAT	3	87.8	0.00008	0.00084	0.06712	0.02008			0.99976	0.99712	0.88504	0.9473	
52	CO9C-ATA-006-SER	3	89	1	0.99412	1	0.72726			0.26486	0.74736	0.12396	0.12776	
53	NBS-09-sat	3	106.6	0.0061	0.01378	0.12116	0.03262			1	0.99964	1	0.91468	
54	CO9C-CTG-380-SER	3	112.4	0.42304	0	0.84436	0.58026			0.44592	0.96792	0.074	0.20286	
55	E35/M59-F-193-SAT	3	117.2	0.039	0.02062	0.02582	0.7679			0.63468	0.7399	0.41156	0.27846	
56	E35/M59-F-187-SER	3	117.2	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
57	1A31-F-347-SER	3	125.1	0.01698	0.00018	0.147	0.7513			0.77144	0.97672	0.74398	0.02274	
58	CO9C-CTA-285-SAT	3	129.6	0.0001	0.00054	0.00272	0.26918			0.99976	0.99824	0.99312	0.56282	
59	CO9C-TCC-181-SER	3	130.2	0.96934	0.85424	0.44506	0.98704			0.0143	0.05482	0.3539	0.00282	
60	CO9C-TCC-202-SAT	4	17.5	1	0.99998	0.92972	0.99698			0.00002	0.00002	0.03562	0.0007	
61	CO9C-GTT-148-SAT	4	25	1	0.99992	0.9961	0.99682			0.00002	0.00008	0.02252	0.00064	
62	E35/M59-F-220-SAT	4	36.9	0.97268	0.95498	0.29876	0.99272			0.66914	0.1279	0.53478	0.93116	
63	E35/M59-F-276-SAT	4	49.1	1	0.99998	0.98722	0.99818			0.04602	0.04852	0.03088	0.59904	
64	CO9C-TCC-281-SAT	4	52.1	0.88808	0.82266	0.65562	0.58932			0.11192	0.0986	0.23866	0.41068	
65	CO9C-ATT-009-SAT	4	58.8	0.99984	0.95816	0.99364	0.91758			0.00512	0.08622	0.06316	0.03336	
66	CO9C-ATT-010-SAT	4	60.8	0.9985	0.84082	0.96684	0.96884			0.02756	0.264	0.18572	0.01066	
67	CO9C-GTG-310-SAT	4	63.4	0.95844	0.84226	0.68174	0.86588			0.02584	0.083	0.2161	0.0604	
68	E45/M48-F-071-SER	4	75.8	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
69	CO9C-CTT-012-SAT	4	83.4	0.93548	0.33172	0.8924	0.95322			0.04042	0.51924	0.06092	0.01746	
70	CO9C-ACT-005-SAT	4	94.1	0.00004	0.00026	0.065	0.01924			0.99986	0.99876	0.8843	0.94866	
71	CO9C-CTA-102-SAT	4	100.5	0.00002	0.00002	0.00002	0			1	1	1	0.99996	
72	CO9C-CTC-001-SAT	4	102.9	0.00002	0.00002	0.00032	0.00006			1	1	0.99912	0.99994	
73	CO9C-CTC-020-SAT	4	104.7	0.00002	0.00002	0.00048	0.00004			1	1	0.99984	0.99986	
74	CO9C-CTC-021-SER	4	106.7	0.49074	0.4233	0.85344	0.05336			1	0.99988	0.9888	0.94664	
75	CO9C-TCC-167-SER	4	109.2	0.818	0.99468	0.59044	0.01896			0.77918	0.16136	0.69604	0.92428	
76	1A31-F-259-SAT	4	112.2	0.00002	0.00002	0.00004	0.00066			1	0.99736	0.99436	0.99568	

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				<i>L. serriola</i>	<i>L. serriola</i>				<i>L. serriola</i>	<i>L. serriola</i>		
Distortion towards preference for:												
77	CO9C-CTC-010-SER	4	153.3	0.97346	0.9634	0.9469	0.14946		0.01246	0.01104	0.02	0.55084
78	1A36-F-250-SAT	4	157.8	0.2076	0.06494	0.88782	0.02774		0.95178	0.89784	0.39406	0.94184
79	CO9C-GTG-193-SAT	4	160.5	0.3974	0.2752	0.5303	0.25044		0.51808	0.58382	0.34712	0.58688
80	NBS-24-ser	5	0	1	0.99996	1	0.55056		0.1206	0.0043	0.5591	0.44944
81	CO9C-CTC-017-SAT	5	3.3	0.90514	0.99842	0.79468	0.00532		0.19086	0.0004	0.44194	0.97494
82	CO9C-ATC-006-SER	5	8.3	1	1	1	0.44444		0.00482	0.00218	0.05436	0.27
83	E44/M48-F-223-SER	5	13.5	1	0.99996	1	0.93604		0.0008	0.00338	0.02132	0.10896
84	CO9C-CTC-004-SER	5	18.8	0.99964	0.99996	0.81114	0.5126		0.0036	0.00004	0.18886	0.4874
85	CO9C-ACG-005-SER	5	22.8	0.71038	0.96762	0.35268	0.08866		0.19086	0.0095	0.4437	0.6807
86	CO9C-AAT-162-SER	5	28.4	1	1	1	0.59234		0.00922	0.09072	0.0092	0.19564
87	1A31-F-129-SER	5	35.9	0.9	0.88142	0.82362	0.3826		0.96534	0.17652	0.98768	0.8981
88	1A31-F-224-SAT	5	43.7	0.5228	0.36272	0.84554	0.10304		0.73524	0.07338	0.79414	0.8971
89	CO9C-ATT-014-SER	5	46	0.32866	0.60896	0.34814	0.08848		0.67134	0.39104	0.44798	0.6802
90	CO9C-CTG-96-SAT	5	66	0.05606	0.57246	0.2155	0.00172		0.91378	0.2845	0.6757	0.99114
91	CO9C-ATC-005-SAT	5	66.6	0.14422	0.43704	0.17016	0.17218		0.89886	0.4114	0.7337	0.92102
92	CO9C-ATC-002-SER	5	70.8	0.9933	0.86936	0.99878	0.15624		0.71854	0.43674	0.3864	0.84376
93	CO9C-ACG-013-SER	5	72.6	0.99984	0.9984	0.99986	m.d.		0.4331	0.1337	0.20744	0.93284
94	CO9C-AAT-200-SAT	5	75.6	1	1	0.6797	0.9997		0.00002	0.00002	0.4551	0.00006
95	CO9C-ACT-009-SAT	5	78.5	0.02374	0.0228	0.52494	0.01956		0.99192	0.9772	0.60262	0.9474
96	CO9C-AAT-172-SAT	5	84.6	0.00284	0.02418	0.08748	0.0419		0.9994	0.9971	0.8509	0.90106
97	1A36-F-160-SER	5	86.7	0.95928	0.54902	0.96404	0.77062		0.01564	0.18266	0.00166	0.54762
98	E44/M49-F-316-SER	5	107.9	0.00022	0.01674	0.0001	0.39114		0.99294	0.86142	0.97866	0.59438
99	E45/M48-F-127-SER	5	109.6	1	0.99704	1	0.92674		0.86888	0.57734	0.88512	0.4019
100	E35/M59-F-129-SER	5	110.3	0.4418	0.55568	0.07006	0.8095		0.24208	0.23978	0.5678	0.06282
101	CO9C-GTG-111-SAT	5	111	0.01064	0.31108	0.00144	0.1983		0.9967	0.68892	0.99958	0.6572
102	CO9C-GTT-213-SER	5	111.5	0.42016	0.31038	0.3843	0.57738		0.57984	0.48708	0.6157	0.22402
103	CO9C-CTG-178-SAT	5	112.8	0.00002	0.00366	0.00044	0.05768		0.99994	0.98872	0.9985	0.87288
104	E35/M59-F-270-SAT	5	117.3	0.60318	0.49298	0.14952	0.89604		0.9309	0.87468	0.87242	0.23726
105	NBS-10-sat	5	120.1	0.21646	0.03336	0.53586	0.38876		0.99996	0.98706	1	0.61124
106	E35/M59-F-178-SER	5	126.2	0.931	0.70822	0.70132	0.84444		0.96574	0.8667	0.96606	0.23822
107	E35/M59-F-182-SAT	5	126.5	0.8217	0.72788	0.34304	0.85012		0.99632	0.995	0.97298	0.26632
108	NBS-22-ser	5	129	0.67044	0.1108	0.96832	0.29596		1	0.99994	1	0.70404
109	CO9C-AAT-431-SAT	5	131.5	0.15282	0.68562	0.07296	0.09382		0.78166	0.16136	0.86308	0.7937
110	1A36-F-156-SER	5	133.2	0.97552	0.88578	0.6945	0.95224		0.97878	0.6789	0.99036	0.4165
111	NBS26-sat	5	134.3	m.d.	m.d.	m.d.	m.d.		m.d.	m.d.	m.d.	m.d.
112	CO9C-AAT-340-SER	5	140.2	0.3022	0.2283	0.35602	0.4455		0.79836	0.56322	0.80832	0.31442
113	E35/M59-F-198-SAT	5	163.9	0.00476	0.1491	0.02138	0.04774		0.99762	0.89192	0.9952	0.49506
114	NBS-07-ser	6	7.1	1	1	1	0.04824		0.69954	0.3524	0.45882	0.79762
115	1A36-F-304-SAT	6	7.8	0.62974	0.04938	0.95406	0.49928		0.84464	0.90474	0.13564	0.94596
116	CO9C-ATT-013-SAT	6	11.7	0.00002	0.00002	0.03318	0.0058		1	1	0.93418	0.98
117	CO9C-ACG-004-SER	6	40	0.12476	0.1528	0.74814	m.d.		0.80792	0.6963	0.14962	0.9903

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				<i>L. serriola</i>	<i>L. serriola</i>				<i>L. serriola</i>	<i>L. serriola</i>			
Distortion towards preference for:													
118	CO9C-AAG-005-SER	6	43.4	0.8413	0.501	0.97846	0.0612		0.09978	0.29156	0.00816	0.7576	
119	E35/M59-F-217-SAT	6	45.1	0.9923	0.68446	0.99078	0.82508		0.03626	0.20386	0.00302	0.6818	
120	CO9C-ATT-005-SAT	6	45.7	0.0189	0.02426	0.38858	0.03334		0.9811	0.97574	0.4867	0.91694	
121	CO9C-ACG-012-SAT	6	53.7	0.9999	0.61792	1	0.83964		0.0001	0.38208	0.00002	0.07552	
122	CO9C-AAG-008-SAT	6	55.6	0.99914	0.41394	1	0.49154		0.0005	0.42962	0.00002	0.32818	
123	CO9C-CTT-006-SAT	6	56.6	0.99924	0.62726	0.99946	0.9181		0.00034	0.23426	0.00018	0.03276	
124	E35/M59-F-103-SAT	6	71.6	0.988	0.79918	0.99716	0.328		0.4286	0.4027	0.08942	0.82004	
125	CO9C-CTC-008-SAT	6	82.8	0.99766	0.99494	0.99438	0.12872		0.0044	0.00176	0.0134	0.72584	
126	CO9C-CTA-253-SER	6	82.9	0.96296	0.96748	0.93446	0.08882		0.0187	0.01032	0.02538	0.686	
127	CO9C-GTT-155-SER	7	3.8	0.9643	0.06278	0.99106	0.70166		0.0357	0.75584	0.00894	0.11856	
128	CO9C-CTA-406-SER	7	4.3	0.7421	0.09954	0.9707	m.d.		0.15438	0.6621	0.00972	0.80918	
129	CO9C-ATA-004-SAT	7	6.3	0.00002	0.00014	0.02306	0		1	0.99986	0.95204	0.99996	
130	E44/M49-F-248-SAT	7	14.2	m.d.	m.d.	m.d.	m.d.		m.d.	m.d.	m.d.	m.d.	
131	1A36-F-171-SAT	7	15.3	0.8561	0.28274	0.97878	0.25866		0.41738	0.47226	0.35336	0.471	
132	CO9C-ACG-002-SER	7	15.6	0.29588	0.35022	0.3295	0.08458		0.57624	0.41462	0.47002	0.6966	
133	E45/M48-F-142-SAT	7	15.6	0.68254	0.05154	0.94152	0.47424		0.91718	0.43652	0.86074	0.92594	
134	CO9C-ACT-010-SER	7	15.9	0.97602	0.53782	0.98004	0.44704		0.01086	0.23842	0.0062	0.2723	
135	CO9C-AAT-098-SAT	7	25.6	0.74194	0.72568	0.74324	0.1717		0.41138	0.27432	0.36534	0.68902	
136	CO9C-AAG-007-SAT	7	30.9	0.52292	0.60998	0.86474	0.00766		0.6396	0.39002	0.21722	0.97584	
137	CO9C-ACT-013-SER	7	40.4	0.98566	0.9422	0.96394	0.11586		0.0311	0.01894	0.0884	0.62178	
138	NBS-04-ser	7	42.1	0.99344	0.93978	0.99816	0.07578		0.91986	0.34076	0.95198	0.8287	
139	NBS-16-ser	7	43.5	0.95812	0.9425	0.9878	0.0144		0.89908	0.11117	0.95324	0.9544	
140	CO9C-CTG-112-SER	7	50.6	0.18772	0.5843	0.0535	0.29992		0.70318	0.22138	0.83878	0.42334	
141	CO9C-ACT-006-SER	7	60.5	1	1	1	0.8375		0.00032	0.00248	0.00446	0.1625	
142	1A31-F-170-SER	7	61.6	0.00544	0.08878	0.04344	0.06426		0.60568	0.12666	0.86194	0.37524	
143	E35/M59-F-101-SER	7	63.2	0.0131	0.05044	0.4732	0.00238		0.99862	0.6583	0.98798	0.9644	
144	CO9C-CTT-002-SER	7	66.8	0.6691	0.82884	0.4002	0.34906		0.3309	0.06904	0.39532	0.65094	
145	CO9C-ACT-001-SER	7	68.6	0.37412	0.69046	0.22732	0.22588		0.50282	0.159	0.59344	0.5209	
146	1A36-F-223-SAT	7	90.7	0.83172	0.81784	0.68374	0.34316		0.99988	0.90258	0.99564	0.94286	
147	CO9C-ATA-009-SER	7	97	0.0637	0.47866	0.02014	0.12568		0.9363	0.28854	0.89916	0.87432	
148	CO9C-GTT-270-SAT	8	0	0.32924	0.32186	0.02094	0.97878		0.67076	0.5279	0.97906	0.00698	
149	CO9C-CTC-014-SER	8	0.5	1	1	1	0.96616		0	0.02796	0.00038	0.00664	
150	NBS-11-ser	8	5.1	1	0.8644	1	0.98838		0.0235	0.29608	0.08032	0.01162	
151	E45/M48-F-067-SER	8	9.5	m.d.	m.d.	m.d.	m.d.		m.d.	m.d.	m.d.	m.d.	
152	E44/M49-F-262-SAT	8	18.4	0.99976	0.79082	0.99882	0.95072		0.11794	0.20256	0.0642	0.4045	
153	NBS-08-ser	8	25.6	1	1	1	0.9906		0.00002	0.00066	0.0261	0.0094	
154	CO9C-CTC-005-SAT	8	31.5	0.97624	0.6148	0.92752	0.92616		0.03842	0.24736	0.07248	0.07384	
155	CO9C-ATC-009-SER	8	57.4	1	1	1	0.99358		0.00002	0.0066	0.00002	0.00642	
156	E35/M59-F-153-SAT	8	57.9	0.37722	0.3492	0.86878	0.01424		0.20048	0.69004	0.03988	0.2408	
157	1A36-F-127-SAT	8	60.8	0.32348	0.75226	0.46998	0.0216		0.00118	0.02866	0.00986	0.03582	
158	CO9C-CTG-370-SAT	8	64.2	0.1419	0.34332	0.17922	0.23724		0.80482	0.513	0.7266	0.60588	

Number (#)	Marker name	Linkage group	cM	All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21	All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21
				<i>L. serriola</i>	<i>L. serriola</i>				<i>L. serriola</i>	<i>L. serriola</i>			
Distortion towards preference for:													
159	1A31-F-146-SER	8	72.1	0.00004	0.00298	0.00388	0.03318			0.89142	0.71242	0.43868	0.84448
160	C09C-ATT-004-SER	8	79.3	1	0.9987	1	0.77708			0.00566	0.1257	0.00704	0.05832
161	E44/M48-F-141-SAT	8	80.6	0.39274	0.65808	0.71504	0.01282			0.99962	0.9153	0.96768	0.83008
162	C09C-CTC-012-SAT	8	90.2	0.99526	0.9055	0.88166	0.9703			0.00258	0.04564	0.0651	0.00962
163	1A36-F-323-SAT	8	128.2	0.99946	0.94302	0.97956	0.96502			0.05392	0.03854	0.26866	0.10664
164	C09C-AAT-248-SAT	8	133.6	0.5894	0.99798	0.38352	0.00182			0.33214	0.00046	0.491	0.99174
165	E44/M48-F-192-SAT	8	143.3	0.3227	0.95474	0.14074	0.01394			1	0.98102	0.99582	0.94572
166	NBS-02-ser	9	0	1	0.99156	1	0.98596			0.00074	0.03946	0.009	0.0686
167	C09C-GTG-151-SER	9	20.3	0.89978	0.90998	0.8023	0.30824			0.06266	0.03738	0.1075	0.46128
168	C09C-AAG-003-SAT	9	24.4	1	0.9966	1	0.15824			0.00002	0.00086	0.00002	0.68262
169	C09C-ACT-015-SAT	9	28.1	0.93808	0.92866	0.98334	0.0231			0.06192	0.03228	0.01666	0.92218
170	C09C-ATC-008-SAT	9	36.5	0.95372	0.50748	0.98064	0.6197			0.07116	0.49252	0.0085	0.3803
171	1A36-F-189-SAT	9	46.4	0.85978	0.14538	0.97524	0.669			0.7482	0.76178	0.30672	0.58408
172	C09C-CTC-002-SAT	9	49.5	0.99582	0.56102	0.99428	0.93624			0.01478	0.62184	0.00572	0.02304
173	C09C-ATC-001-SAT	9	53.1	0.9277	0.1241	0.90656	0.982			0.11022	0.75438	0.04884	0.05088
174	E44/M49-F-175-SER	9	60.7	0.93704	0.17588	0.98024	0.87794			0.98462	0.8662	0.85726	0.71882
175	C09C-CTA-098-SER	9	81.4	0.03102	0.354	0.05076	0.03284			0.9875	0.4356	0.98626	0.84376
176	C09C-CTT-007-SAT	9	92.6	0.1623	0.6347	0.06378	0.29096			0.91858	0.3653	0.96708	0.5426
177	1A31-F-176-SER	9	95.6	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.
178	NBS-19-sat	9	100.2	0.95312	0.72018	0.90014	0.78914			0.99992	0.4197	1	0.92606
179	NBS-25-ser	9	102.5	0.99998	0.9893	0.99994	0.70212			0.54914	0.17822	0.9104	0.13442
180	C09C-TCC-123-SER	9	106.1	0.70168		0.65638	0.86894			0.76852	0.58888	0.70896	0.13106
181	E35/M59-F-183-SAT	9	108.1	0.38018	0.70582	0.01736	0.90742			0.26104	0.01842	0.7562	0.33128
182	E35/M59-F-096-SAT	9	113.1	0.48452	0.85478	0.06432	0.68264			0.69168	0.18572	0.97866	0.14086
183	C09C-ATA-007-SAT	9	122.5	0.1908	0.76596	0.03828	0.32934			0.8092	0.13652	0.92706	0.67066
184	C09C-AAG-004-SER	9	127.7	0.7229	0.68844	0.59126	0.48468			0.18776	0.16018	0.2488	0.27734
185	C09C-ACT-007-SAT	9	133.4	0.00986	0.84532	0.00048	0.04264			0.99014	0.15468	0.99858	0.8998
186	C09C-ACT-002-SER	9	137	0.07468	0.30256	0.17338	0.06206			0.87718	0.51384	0.69716	0.80972
187	C09C-ATT-011-SAT	u	u	0.16024	0.1111	0.2142	0.58966			0.92136	0.94644	0.7858	0.254
188	E44/M48-F-212-SER	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.
189	1A31-F-074-SAT	u	u	0.03926	0.01466	0.1939	0.42384			0.9974	0.9543	0.88958	0.8633
190	1A31-F-244-SAT	u	u	0.997	0.59188	0.97914	0.99318			0.14716	0.1342	0.3289	0.13044
191	1A31-F-323<N>-P2	u	u	0.76826	0.30302	0.43786	0.94656			0.81542	0.72102	0.8947	0.11408
192	1A31-F-402<N>-P2	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.
193	1A36-F-148<N>-P2	u	u	0.45044	0.04196	0.87182	0.44078			1	0.97966	0.99564	0.93962
194	1A36-F-246-SAT	u	u	0.01552	0.03336	0.10996	0.13608			0.9999	0.93106	0.97552	0.85572
195	1A36-F-299<N>-P2	u	u	0.99936	0.91674	0.99702	0.87806			0.01648	0.14598	0.02414	0.11424
196	1A36-Fb-126.70<N>-P2	u	u	0.0002	0.00124	0.03342	0.0591			1	0.97898	0.99546	0.938
197	E35/M59-F-086<F>-P2	u	u	0.28526	0.0422	0.7901	0.26748			0.99998	0.97846	0.99518	0.94122
198	E35/M59-F-204<F>-P2	u	u	0.96322	0.18142	0.92742	0.9938			0.35152	0.2918	0.7502	0.038
199	E35/M59-F-212-SAT	u	u	0.12998	0.52732	0.0296	0.3109			0.8866	0.25458	0.94808	0.62652

Number (#)	Marker name	Linkage group	cM	All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21		All fields (=Fig. 2) N = 90		Amsterdam n = 29	Sijbekarspel n = 40	Aachen n = 21
				<i>L. serriola</i>	<i>L. serriola</i>					<i>L. serriola</i>	<i>L. serriola</i>			
Distortion towards preference for:														
200	E44/M48-F-328-SAT	u	u	0.00004	0.05784	0.02086	0.00004			0.96892	0.63756	0.72706	0.96456	
201	E44/M48-F-403<F>-P2	u	u	1	1	1	1	1		0.00002	0.00002	0.00002	0.00008	
202	E44/M49-F-231<F>-P2	u	u	1	1	1	1	1		0.00002	0.00002	0.00002	0.00002	
203	E44/M49-F-250<F>-P2	u	u	1	1	1	1	1		0.00002	0.00002	0.00002	0.00002	
204	E44/M49-F-337-SAT	u	u	0.73638	0.11338	0.55966	0.98628			0.95708	0.6613	0.95282	0.4388	
205	E45/M48-F-084<N>-P2	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
206	E45/M48-F-103-SAT	u	u	0.89436	0.92314	0.19816	0.90892			0.27818	0.08718	0.43774	0.39944	
207	E45/M48-F-140-SAT	u	u	1	0.9962	0.99848	0.9858			0.98436	0.8662	0.6991	0.92512	
208	E45/M48-F-146<F>-P2	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
209	E45/M48-F-159<F>-P2	u	u	0.7666	0.84114	0.68954	0.27314			0.99998	0.98014	0.99536	0.9372	
211	1A36-F-434<N>-P1	u	u	1	0.9596	0.99874	0.99418			0.09976	0.499	0.05506	0.11154	
212	E44/M49-F-150<N>-P1	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
213	E44/M49-F-447<F>-P1	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
214	NBS-03-sat	u	u	1	0.9985	1	0.99838			0.00046	0.00498	0.32056	0.00036	
215	NBS-05-sat	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
216	NBS-27-sat	u	u	1	1	1	1			0.00002	0.00538	0.0685	0.00002	
217	NBS-36-sat	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
218	NBS37-sat	u	u	m.d.	m.d.	m.d.	m.d.			m.d.	m.d.	m.d.	m.d.	
221	C09C-AAT-097-SAT	u	u	0.21938	0.36288	0.08842	0.69902			0.70908	0.476	0.84486	0.14818	
222	C09C-AAT-153-SAT	u	u	0.9999	0.99672	0.98276	0.88494			0.00004	0.001	0.00772	0.05022	
223	C09C-AAT-180-SAT	u	u	0.00074	0.69908	0.00138	0.00074			0.99976	0.61438	0.99656	0.99694	
224	C09C-AAT-208-SAT	u	u	0.00002	0.0161	0.00002	0			1	0.9839	1	0.99992	
225	C09C-ATA-003-SAT	u	u	0.99048	0.57446	0.8832	0.99666			0.01704	0.28518	0.06498	0.01112	
226	C09C-ATT-001-SAT	u	u	1	0.6688	1	0.9999			0.00002	0.19978	0.00002	0	
227	C09C-CTT-003-SAT	u	u	0.46296	0.93688	0.67188	0.00122			0.45236	0.02748	0.22154	0.9942	
228	C09C-CTT-009-SAT	u	u	0.06292	0.11056	0.34942	0.06584			0.90532	0.79472	0.52418	0.8457	
229	C09C-GTG-700-SAT	u	u	0.78644	0.22268	0.3099	0.99574			0.21356	0.65268	0.57196	0.00084	
230	C09C-GTT-238-SAT	u	u	0.0382	0.07604	0.05708	0.47356			0.9618	0.85418	0.94292	0.3566	
231	C09-C-GTT-440-SAT	u	u	0.82486	0.73834	0.87318	0.14088			0.17514	0.1551	0.12682	0.72958	
232	C09C-TCC-088-SAT	u	u	0.01166	0.45956	0.18262	0.00014			0.97972	0.39628	0.72284	0.99932	
233	C09C-TCC-146-SAT	u	u	0.047	0.00514	0.79498	0.01886			0.92576	0.98312	0.12328	0.94144	
234	C09C-TCC-199-SAT	u	u	0.28328	0.734	0.1526	0.21486			0.64138	0.15994	0.75838	0.6341	
235	C09C-TCC-640-SAT	u	u	0.87604	0.44852	0.69796	0.94636			0.12396	0.39906	0.20148	0.05364	