

Table 1. Hybrid zone localities, distances from western edge of transect, and numbers of chromosomes sampled (averaged across loci). Locality numbers correspond to those in the map in Figure 1.

Locality	Distance (km)	Number of Chromosomes Sampled
1. Augsburg (A), Germany	2.2	17
2. Augsburg (B), Germany	2.2	11
3. Ebersbach, Germany	39.6	9
4. Kammerberg-Hartl, Germany	45.2	7
5. Kammerberg-Zandt, Germany	45.2	6
6. Appercha, Germany	49.2	27
7. Gesselthausen-Warta, Germany	51.8	105
8. Gesselthausen-Ziigletrum, Germany	51.8	62
9. Giesenbach, Germany	52.4	4
10. Eberspoint, Germany	53.2	18
11. Massenhausen, Germany	53.8	15
12. Neufahrn bei Freising, Germany	55.6	103
13. Pulling-Petryszak, Germany	58.8	12
14. Acherling, Germany	59.2	30

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15. Freising, Germany	61.4	9
16. Tunttenham, Germany	62.2	6
17. Gut Wildschwaig, Germany	64.4	6
18. Rudlfing, Germany	67.2	30
19. Schwaig, Germany	69.2	21
20. Tittenkofen, Germany	77.2	3
21. Sonnendorf, Germany	84.6	37
22. Brundl, Germany	103.4	23
23. Attenham, Germany	130.4	13
24. Simbach, Germany	152.4	27
25. Ranshofen-Holfinger, Austria	155.0	6
27. Branau, Austria	157.4	5
28. Nofing, Austria	160.4	11
29. Aufhausen, Austria	170.0	3
30. Rodham, Austria	170.2	10
31. Leitham-Fuchs, Austria	176.0	23
32. Leitham-Hubinger, Austria	176.4	3

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