

Breed	Recombination Rate	Total Hotspots	Median hotspot width (range)	Mean hotspot rate (range)
Arabian	0.95	31,927	4.61 (1.0-54.1)	2.43 (0.68-152.46)
Belgian	1.39	39,218	4.60 (1.0-49.0)	3.41 (1.15-169.0)
Franches-Montagnes	1.13	38,131	4.48 (1.0-50.9)	2.56 (0.94-277.98)
French Trotter	1.36	45,604	4.30 (1.0-45.98)	3.43 (1.10-122.87)
Icelandic	2.11	38,999	4.90 (1.0-68.0)	5.22 (1.77-198.19)
Lusitano	1.30	36,207	4.70 (1.0-49.1)	3.06 (1.13-125.50)
Maremmano	1.31	40,204	4.70 (1.0-50.1)	3.07 (1.12-129.69)
Morgan	1.14	50,953	3.70 (1.0-37.7)	2.89 (0.93-147.89)
Quarter Horse	0.90	50,814	3.60 (1.0-37.1)	2.45 (0.73-92.41)
Standardbred	1.48	43,069	4.00 (1.0-50.0)	2.45 (0.92-211.31)
Thoroughbred	1.53	21,992	5.44 (1.0-55.2)	3.10 (1.14-171.37)
Welsh Pony	1.07	47,802	3.90 (1.0-42.1)	3.00 (0.76-156.50)

**Supplemental Table S2.** Summary of breed-specific recombination maps. Recombination rate refers to the average genome-wide recombination rate in cM/Mb. Median hotspot width is in kb. Mean hotspot rates are in cM/Mb and represent average rate per Mb across all hotspots combined.