

Table 1 - Markers genotyped and used in the linkage analyses by each data collection group.

Marker	JHU 1998	Mayo 1998	Finn 1998	Swed 1998	BC/CA/HI	FHCRC	Finn	France	JHU (GW)	Mayo (GW)	Michigan	Sweden	UK	Ulm	Utah
DXS1216*							10		191		174	67			
DXS6800						186							64		
DXS986							10		191	161	174	67			
DXS990							10		191	161	174	67			
DXS6789						186							64		
DXS1106							10		191	161	174	67			
DXS6797						186							64		
GATA172D05						186							64		
DXS8055							10		191	161	174	67			
DXS1001							10		191	161	174	67			
GATA165B12						186							64		
DXS1047						186	10		191	161	174	67	64		
DXS1192															146
DXS1232	139	123	57							161					
DXS984	139	123	57	41		186	10	66	191	161	174	67		104	146
GATA31E08					100	186							64		
DXS1205	139	123	57							161	153				
DXS1227							10		191	161	174	67			
DXS6751	139	123	57		100					161				104	
DXS6798	139		57												146
DXS8106	139	123	57		100	186		66		161				103	
DXS7127					100								64		
DXS6806	139	123	57	41		186				161					
DXS8043	139	123	57	41			10		191	161	174	67			
MXMAF MA113ZF5			57							161					
DXS1200	139	123	57	41	100	186		66		161	153			104	146
DXS297	139	123	57			186				161					
DXS731										161					

MXMAFM323Y F1										161				
DXS8091	139	123	57				10		191	161	174	67		
AFM136yb10 (MXMAFM136 YB10)	139	123	57							161				146
MXMAFMA107 XF5										161				
DXS1193	139		57	41	100	186					153		103	146
DXS1123										161				
DXS8069	139	123	57	41		186				161				146
DXS8011	139		57											
DXS8103	139	123	57	41		186				161				146
AFMa225xh9 (MXMAFMA22 5XH9)	139	123	57							161				146
MXMAFMA082 XA5										161				
DXS1073							10		191	161	174	67		146

* Yellow-shaded, bolded text indicates markers included in the GWS marker set for each sample and blue-shaded, non-bolded text indicates markers also included in the GWS plus fine-mapping marker set. Yellow-shaded, bolded text of the marker name indicates that it served as a GWS marker for at least one sample, but some of these markers served only as finemapping markers in some densely typed data sets. The first four columns indicate families that were included in the original linkage paper [3]. Numbers in the table represent the number of families genotyped for each marker in each dataset.