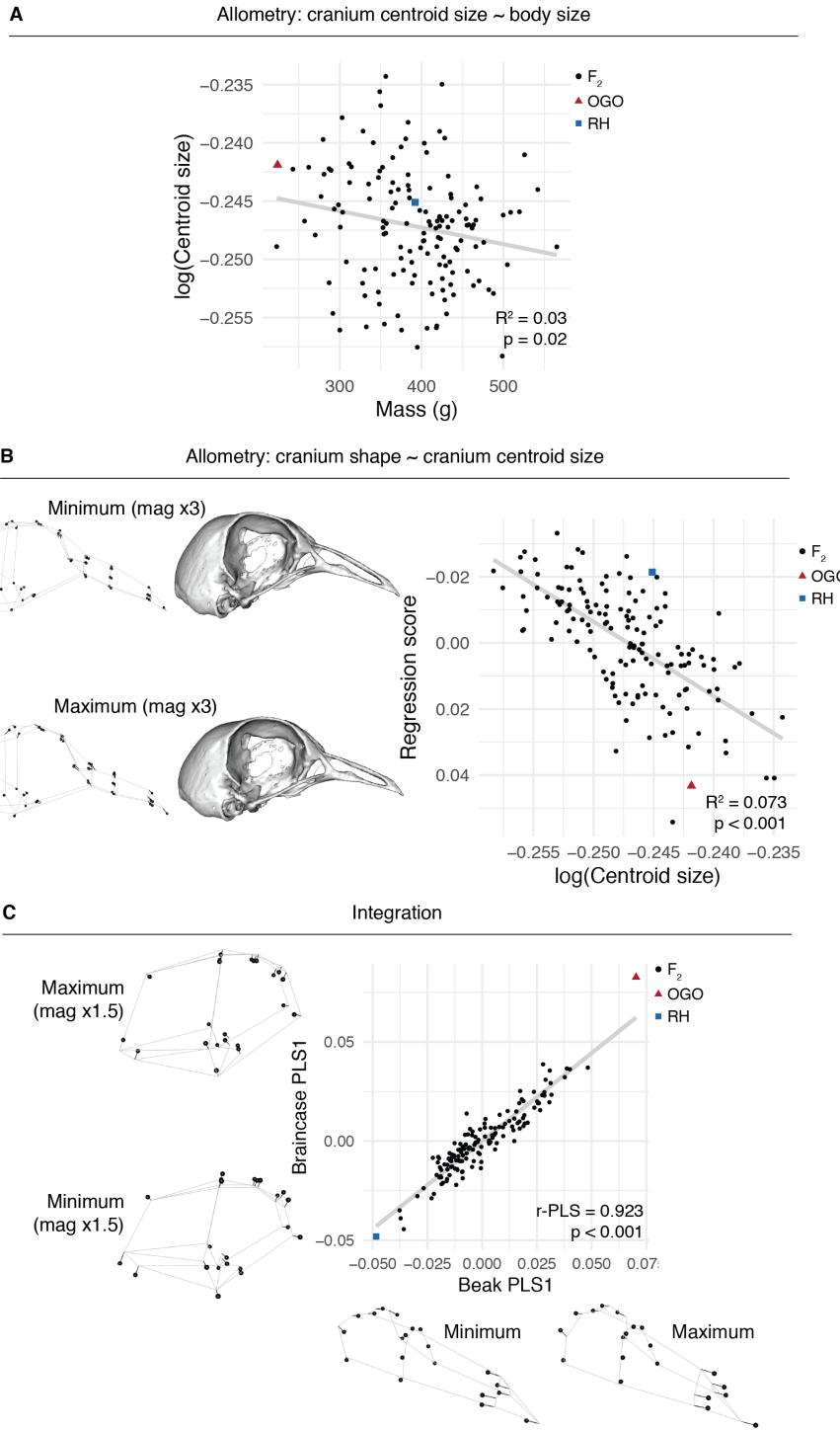
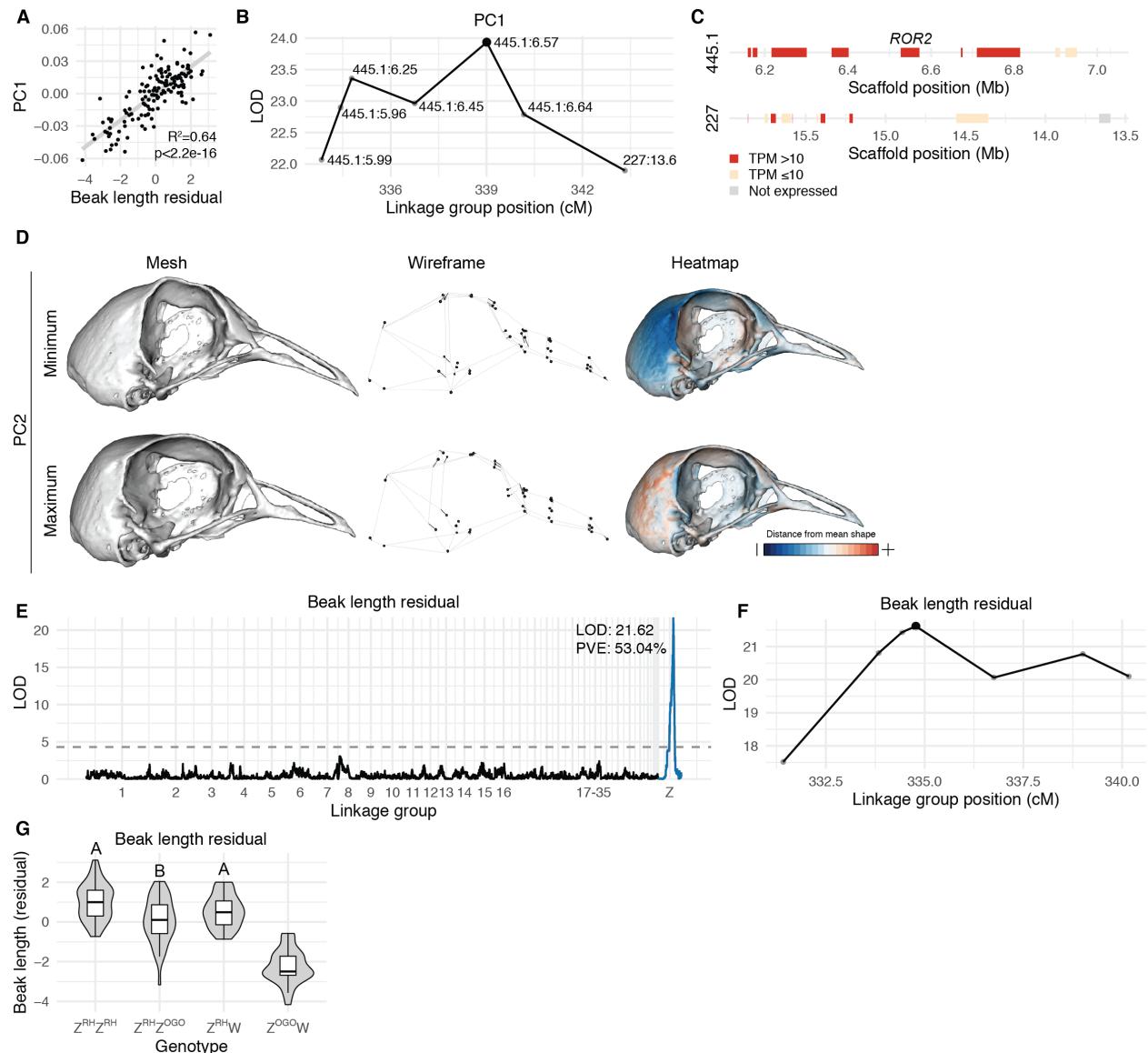


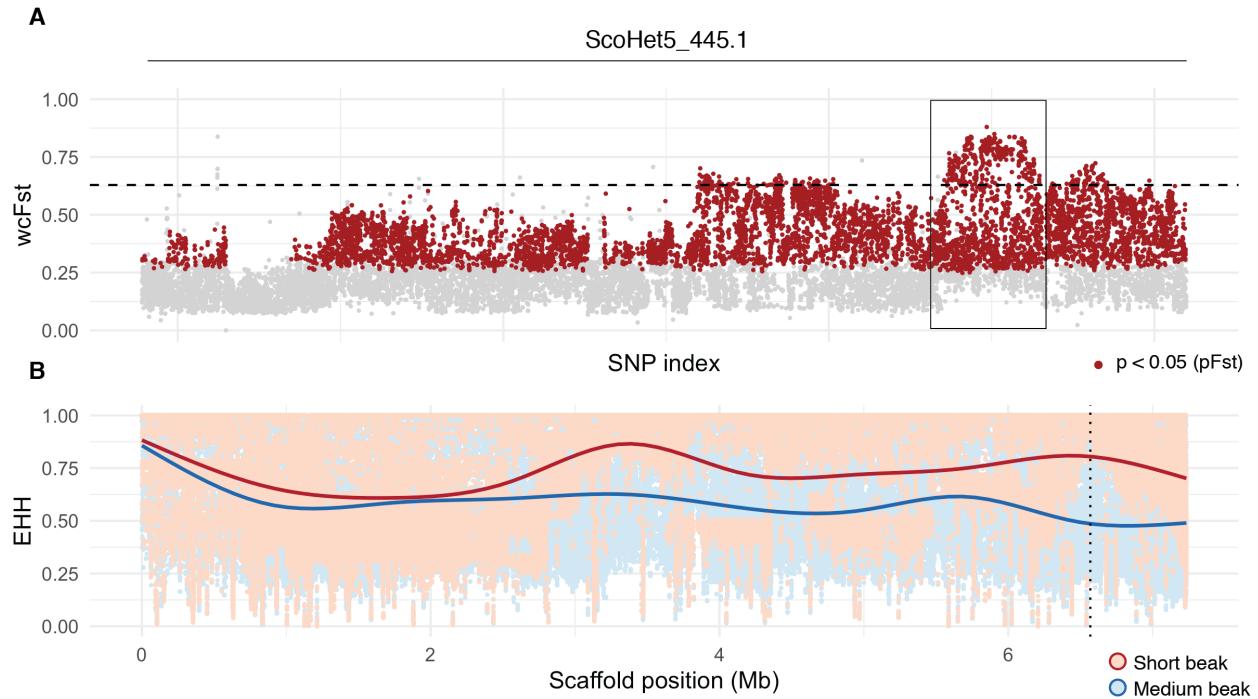
Figure S1. Pigeon craniofacial landmark atlas, related to Figures 1-2. Landmarks are indicated by blue discs.



Supplemental Figure 2. Allometry and integration in the RH x OGO cross, related to Figures 1-2. (A) Cranium centroid size ~ body size (mass) linear regression. (B) Cranium shape ~ centroid size linear regression. (B) Beak vs. braincase PLS1 shapes. Minimum and maximum shapes are depicted as wireframes and/or warped meshes along corresponding axis.



Supplemental Figure 3. Geometric morphometric and QTL analysis of craniofacial skeleton shape in RH x OGO cross, related to Figures 1-2. (A) Scatter plot of beak length residual vs. PC1 score for all RH x OGO F₂ individuals. (B) LOD support interval for PC1 QTL peak on Z. Markers in interval are denoted with dots and labeled with genomic scaffold name (ScoHet5_445.1 or ScoHet5_227) and position (in Mb); black dot indicates QTL peak marker (ScoHet5_445.1:6.57-Mb), which was used to estimate QTL effects. (C) Genes in PC1 QTL interval, color-coded by mRNA expression level in facial primordia derived from HH29 RH embryos. *ROR2* is located directly under QTL peak. (D) Minimum and maximum PC2 shapes depicted as warped mesh (left), wireframe showing landmark displacement (center), or heatmap indicating regional shape changes (right). For mesh and wireframe models, shape change is magnified 1.5x to aid visualization. (E) Genome-wide QTL scan for beak length using residuals from beak length ~ body mass linear regression. (F) LOD support interval is nearly identical to PC1 QTL interval (displayed in Figure 2D). Genes in interval on ScoHet5_445.1 are displayed at bottom and color-coded by expression level. (G) Plot of QTL effects, using peak marker highlighted by black dot in (F).



Supplemental Figure 4. Allele frequency differentiation (F_{ST}) and extended haplotype homozygosity (EHH) on scaffold ScoHet5_445.1 in short and medium/long beak pigeons, related to Figure 3. (A) F_{ST} on ScoHet5_445.1. Boxed region indicates ~293-kb peak region displayed in Figure 3D-E. (B) EHH on ScoHet5_445.1. Smoothed lines represent local regression fitting; dotted vertical line indicates position of ScoHet5_445.1:6568443.

49	Junction of palatine and pterygoid	Left	Skull	Bilateral
50	Junction of palatine and pterygoid	Right	Skull	Bilateral
51	Coronoid process of mandible	Left	Mandible	Bilateral
52	Coronoid process of mandible	Right	Mandible	Bilateral
53	Lateral edge of basioccipital	Left	Skull	Bilateral
54	Lateral edge of basioccipital	Right	Skull	Bilateral
55	Ventral point on mandible	Left	Mandible	Bilateral
56	Ventral point on mandible	Right	Mandible	Bilateral
57	Dorsal orbital rim	Left	Skull	Bilateral
58	Dorsal orbital rim	Right	Skull	Bilateral
59	Midline of frontal bone	Midline	Skull	Midline
60	Ventral point on mandible	Left	Mandible	Bilateral
61	Ventral point on mandible	Right	Mandible	Bilateral
62	Dorsal point on mandible	Left	Mandible	Bilateral
63	Dorsal point on mandible	Right	Mandible	Bilateral
64	Ventral point on maxilla	Left	Beak	Bilateral
65	Ventral point on maxilla	Right	Beak	Bilateral
66	Ventral point on premaxillary	Left	Beak	Bilateral
67	Ventral point on premaxillary	Right	Beak	Bilateral
68	Back of skull	Midline	Skull	Midline
69	Dorsal point on mandible	Left	Mandible	Bilateral
70	Dorsal point on mandible	Right	Mandible	Bilateral
71	Lateral edge of nasal bone	Right	Beak	Bilateral
72	Lateral edge of nasal bone	Left	Beak	Bilateral
73	Supraoccipital bone	Midline	Skull	Midline

