



Figure S3. Comparison of *rho* regions between Japanese eel and European eel.

(A) Microsynteny of *rho* regions (*lrig1* – *thoc7*). Numbering of syntenies (#1 and #2) and coloring of genes are according to those in Figure 3. (B and C) Comparison of scaffolds around *ds-rho* (B) and *fw-rho* (C) regions, respectively. The horizontal axis is for the scaffolds of Japanese eel and the vertical for those of European eel. The mapped positions of linkage marker in Japanese eel are plotted on horizontal axis.