Figure S2. A KFRS4/Kyo-specific deletion of approximately 50 kb in length located approximately 50 kb upstream of Ednrβ. RefSeq gene and the capture targets of the TargetEC method are shown in the first and the second tracks, respectively. The results of the peak-calling for PVG/Seac and KFRS4/Kyo are shown in the third and fourth tracks, respectively, followed by the results for the peak-calling for the other 20 strains. The KFRS4/Kyo-specific deletion is indicated by a red rectangle.