

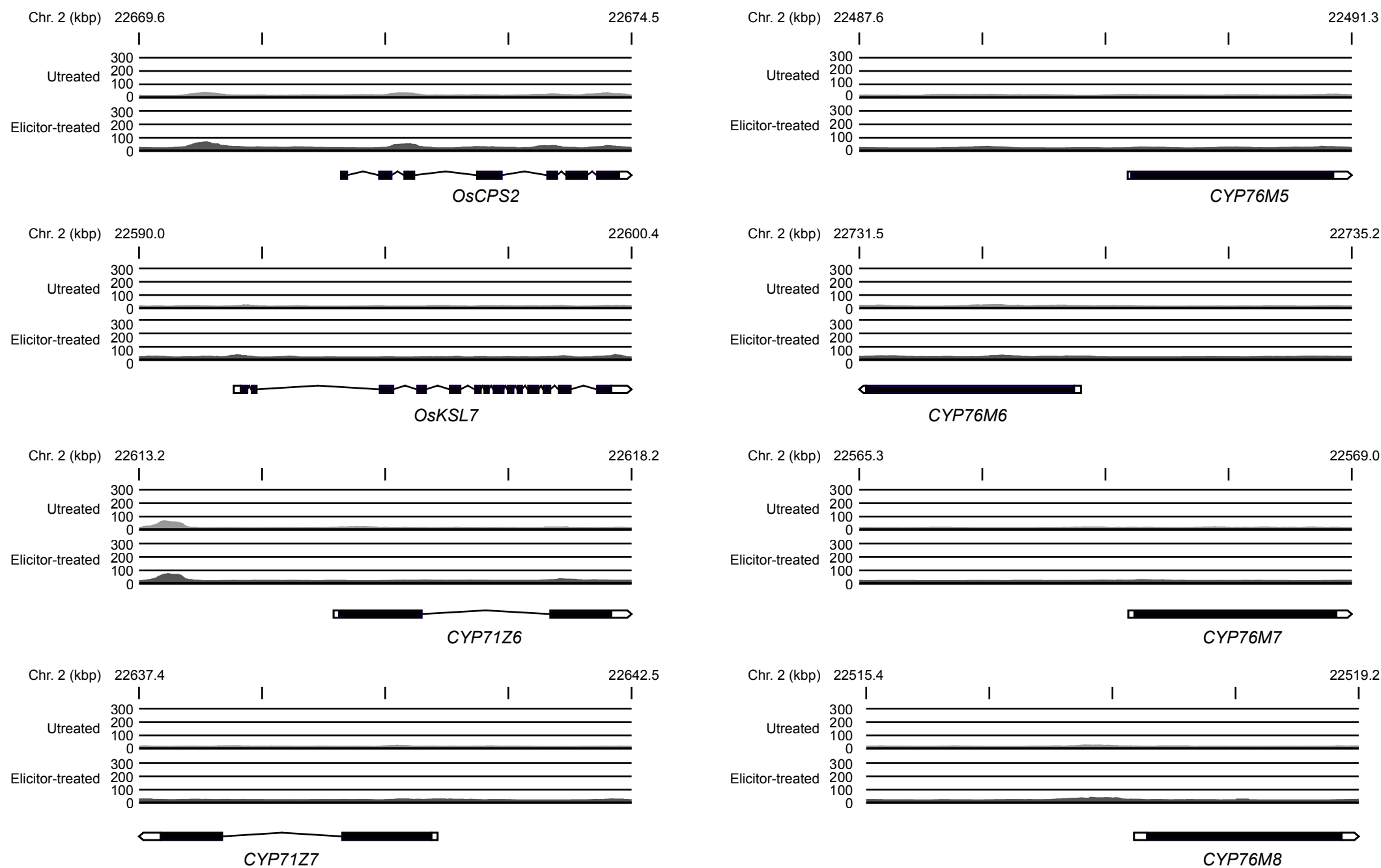


Fig. S5. OsTGAP1-binding regions around phytocassane biosynthetic genes. The mapped ChIP-seq reads in untreated (grey) and elicitor-treated (black) conditions were visualized using Partek Genomics Suite. No binding site was detected in the 2-kbp upstream region of each gene. The gene structure of each phytocassane biosynthetic gene is shown in the bottom row. Open and closed squares indicate untranslated and coding regions, respectively. Lines indicate introns.