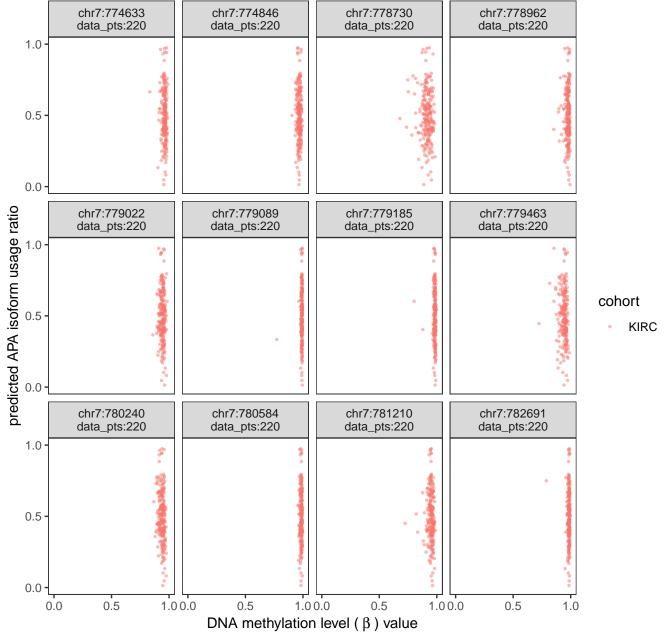
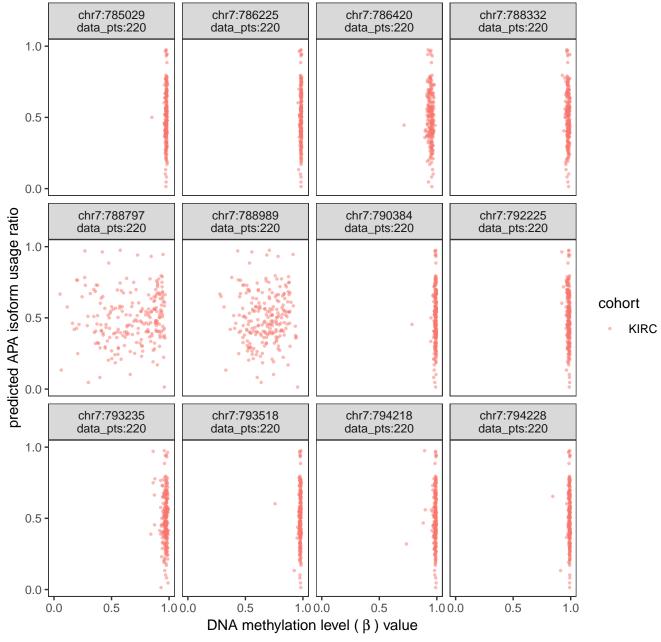
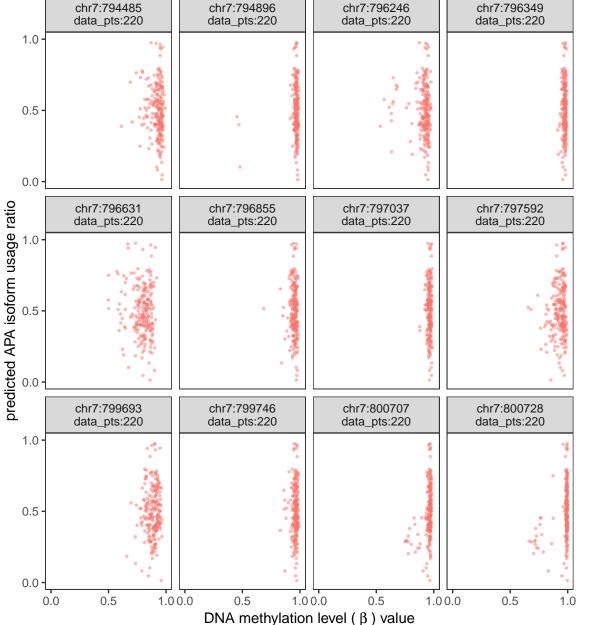
(558)HEATR2[chr7:761,338-834,190], gtop8000, #1/8



(558)HEATR2[chr7:761,338-834,190], gtop8000, #2/8



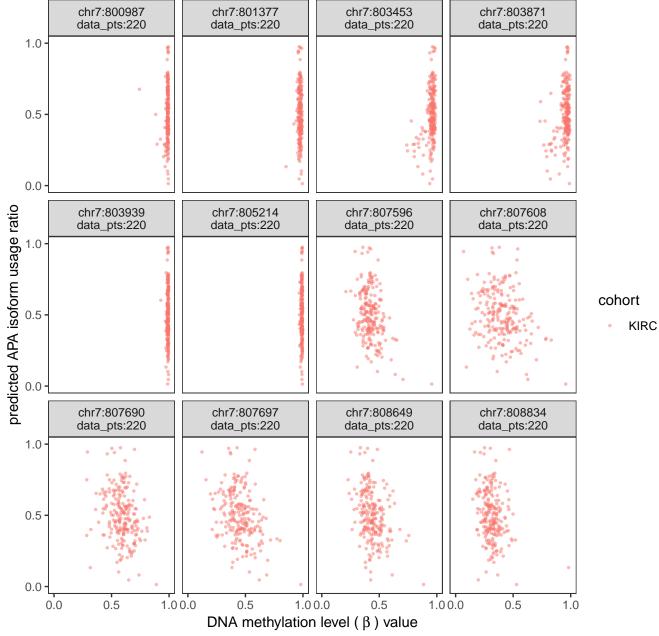
(558)HEATR2[chr7:761,338-834,190], gtop8000, #3/8 chr7:794896 chr7:796246 chr7:794485 data_pts:220 data_pts:220 data_pts:220 1.0



cohort

KIRC

(558)HEATR2[chr7:761,338–834,190], gtop8000, #4/8



(558)HEATR2[chr7:761,338-834,190], gtop8000, #5/8 chr7:809048 chr7:809078 chr7:809109 chr7:810958 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 0.5 0.0 predicted APA isoform usage ratio chr7:811206 chr7:811390 chr7:811491 chr7:811698 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 cohort 0.5 **KIRC** chr7:811786 chr7:813708 chr7:813957 chr7:814391 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 0.5

0.0 1.00.0 0.5 1.00.0 0.5 1.00.0 0.5 0.0 0.5 1.0 DNA methylation level (β) value

(558)HEATR2[chr7:761,338-834,190], gtop8000, #6/8 chr7:814447 chr7:814621 chr7:814701 chr7:814759 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 0.5 0.0 predicted APA isoform usage ratio chr7:815404 chr7:815417 chr7:816960 chr7:817139 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 cohort **KIRC** chr7:817576 chr7:817700 chr7:818287 chr7:818309 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 0.5

0.0 0.5 1.00.0 0.5 1.00.0 0.5 1.00.0 0.5 0.0 1.0 DNA methylation level (β) value

(558)HEATR2[chr7:761,338-834,190], gtop8000, #7/8 chr7:819621 chr7:819747 chr7:818362 chr7:819902 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 0.5 0.0 predicted APA isoform usage ratio chr7:821690 chr7:822036 chr7:822431 chr7:822633 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 cohort **KIRC** chr7:823500 chr7:824325 chr7:824549 chr7:824876 data_pts:220 data_pts:220 data_pts:220 data_pts:220 1.0 0.5

0.0 0.5 1.00.0 0.5 1.00.0 0.5 1.00.0 0.5 0.0 1.0 DNA methylation level (β) value

