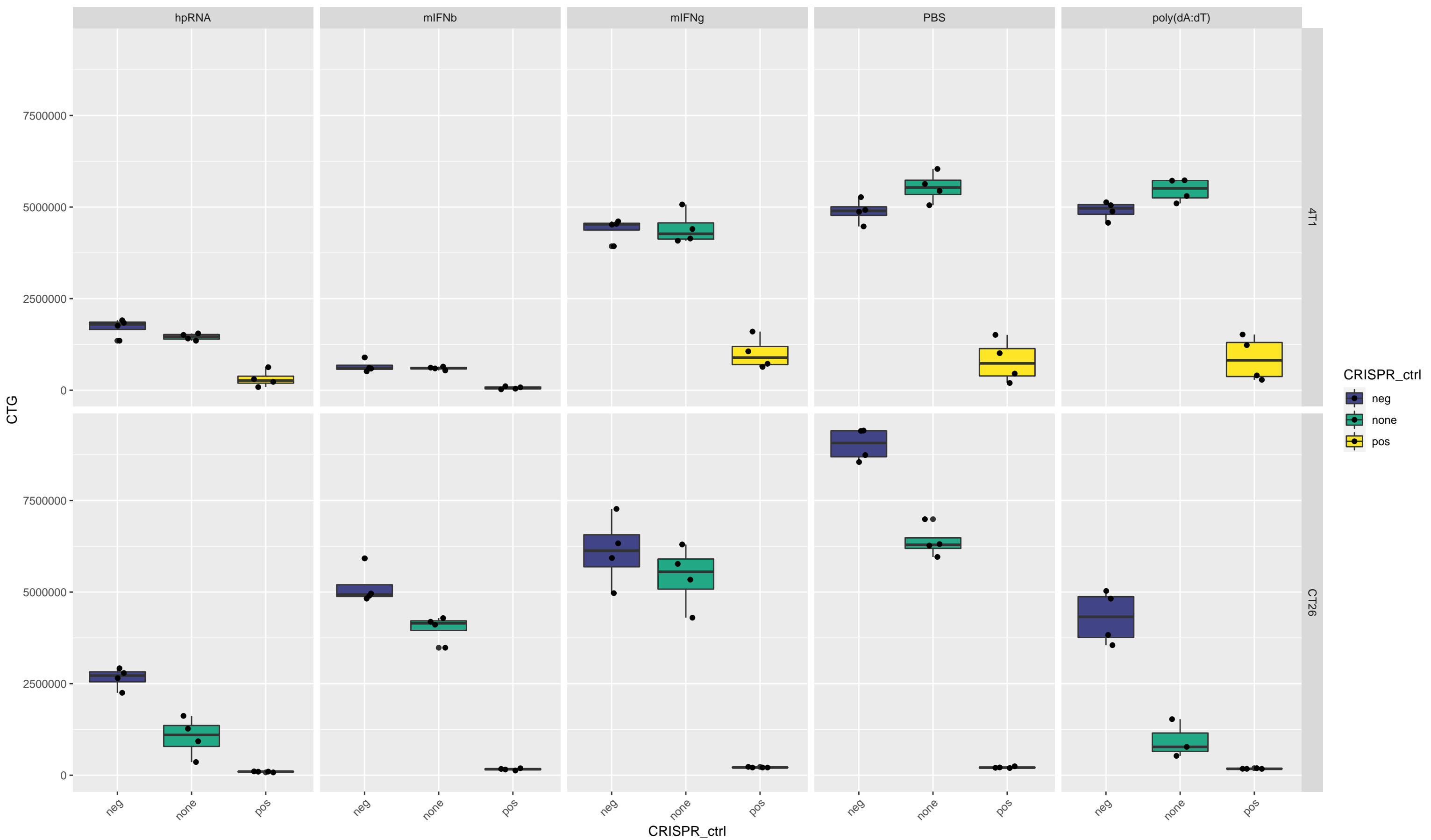
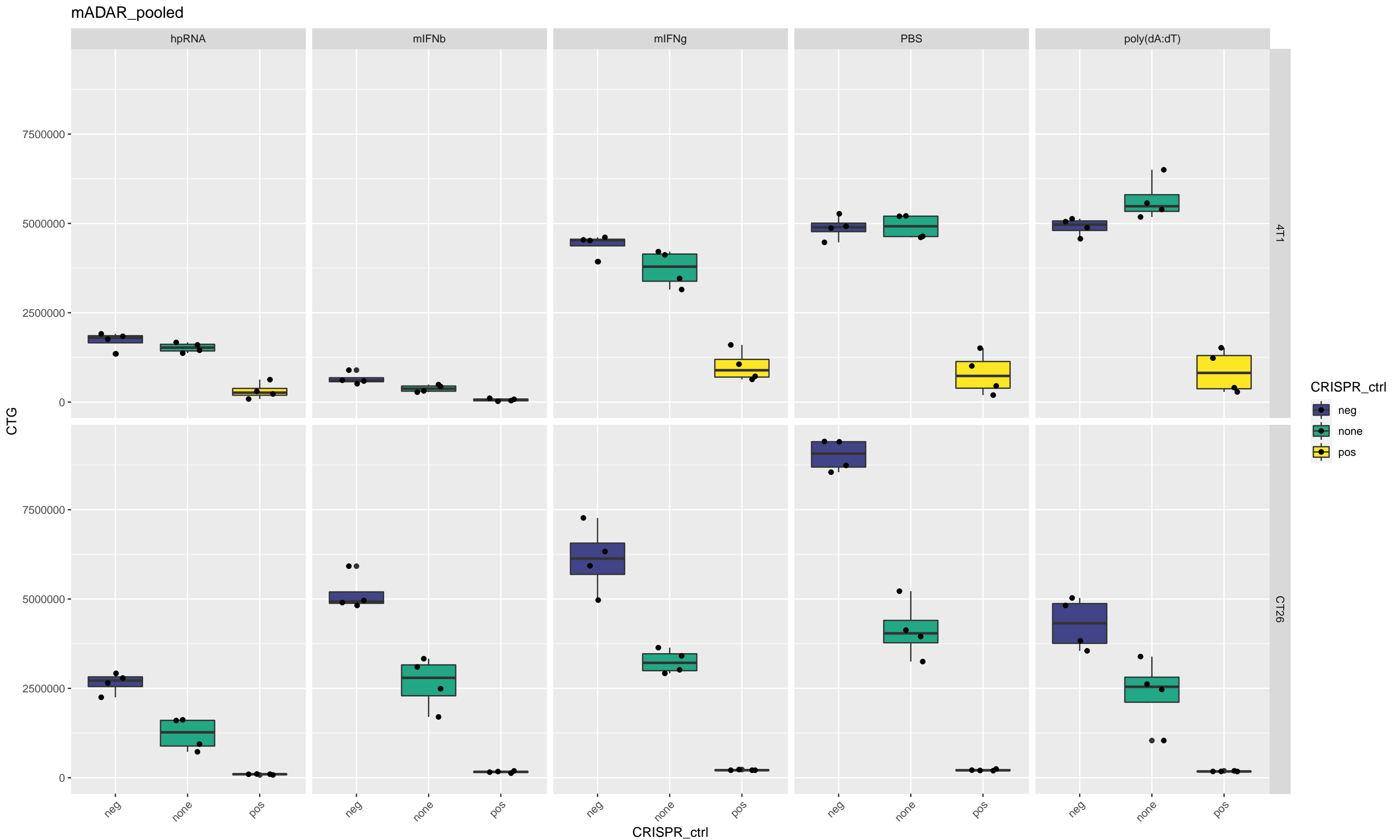
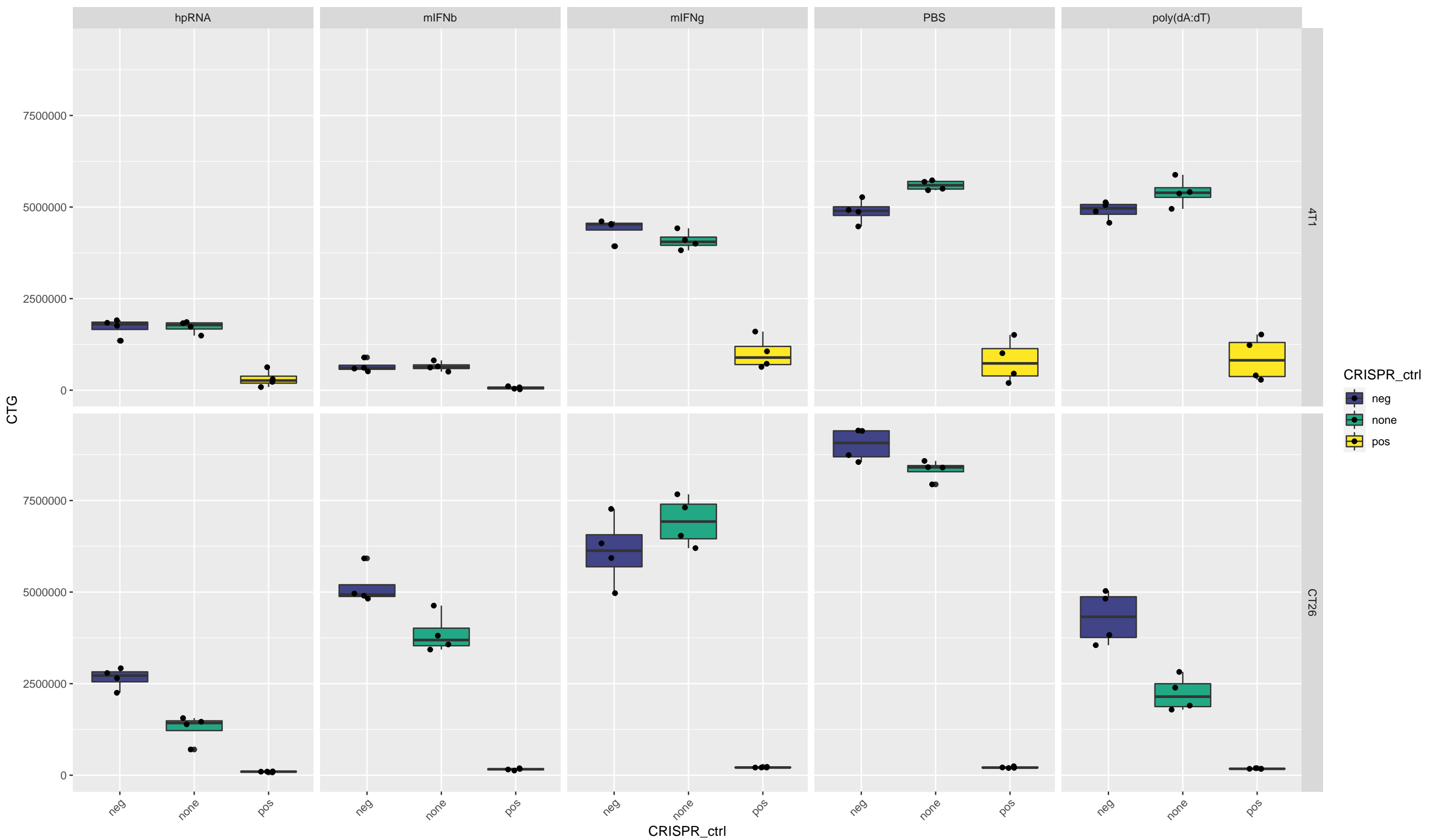


mPARP7\_pooled





mTREX1\_pooled

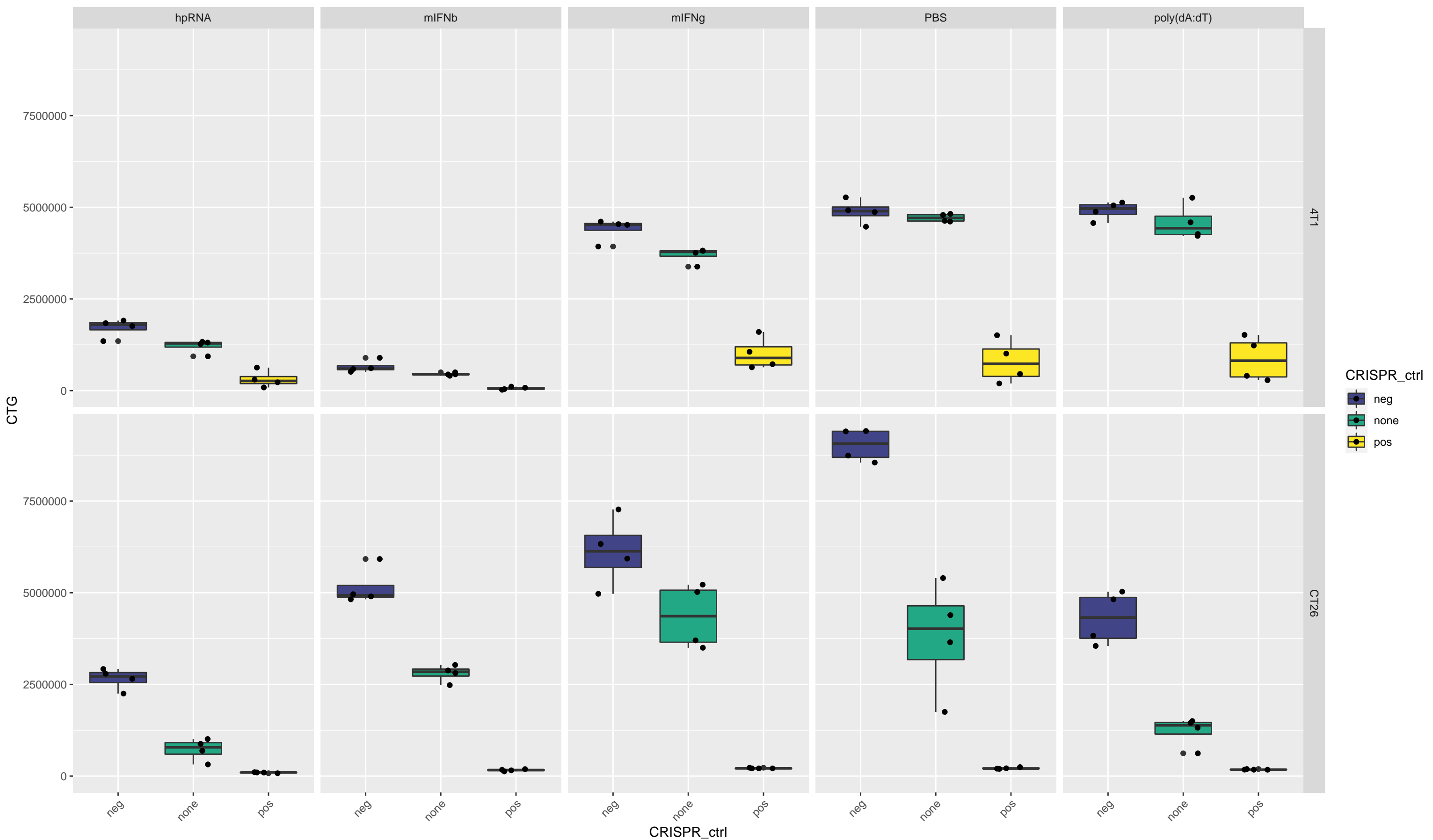


The figure displays a 2x5 grid of box plots comparing the distribution of AT1 and CT26 across three categories (neg, none, pos) for five different methods (hpRNA, mIFNb, mIFNg, PBS, poly(dA:dT)). The y-axis represents a numerical value, and the x-axis represents the categories. The top row is labeled 'AT1' and the bottom row is labeled 'CT26'. The columns are labeled 'hpRNA', 'mIFNb', 'mIFNg', 'PBS', and 'poly(dA:dT)'. Each plot shows a box plot with a median line, a box representing the interquartile range, and whiskers extending to the minimum and maximum values. Individual data points are overlaid on the box plots.

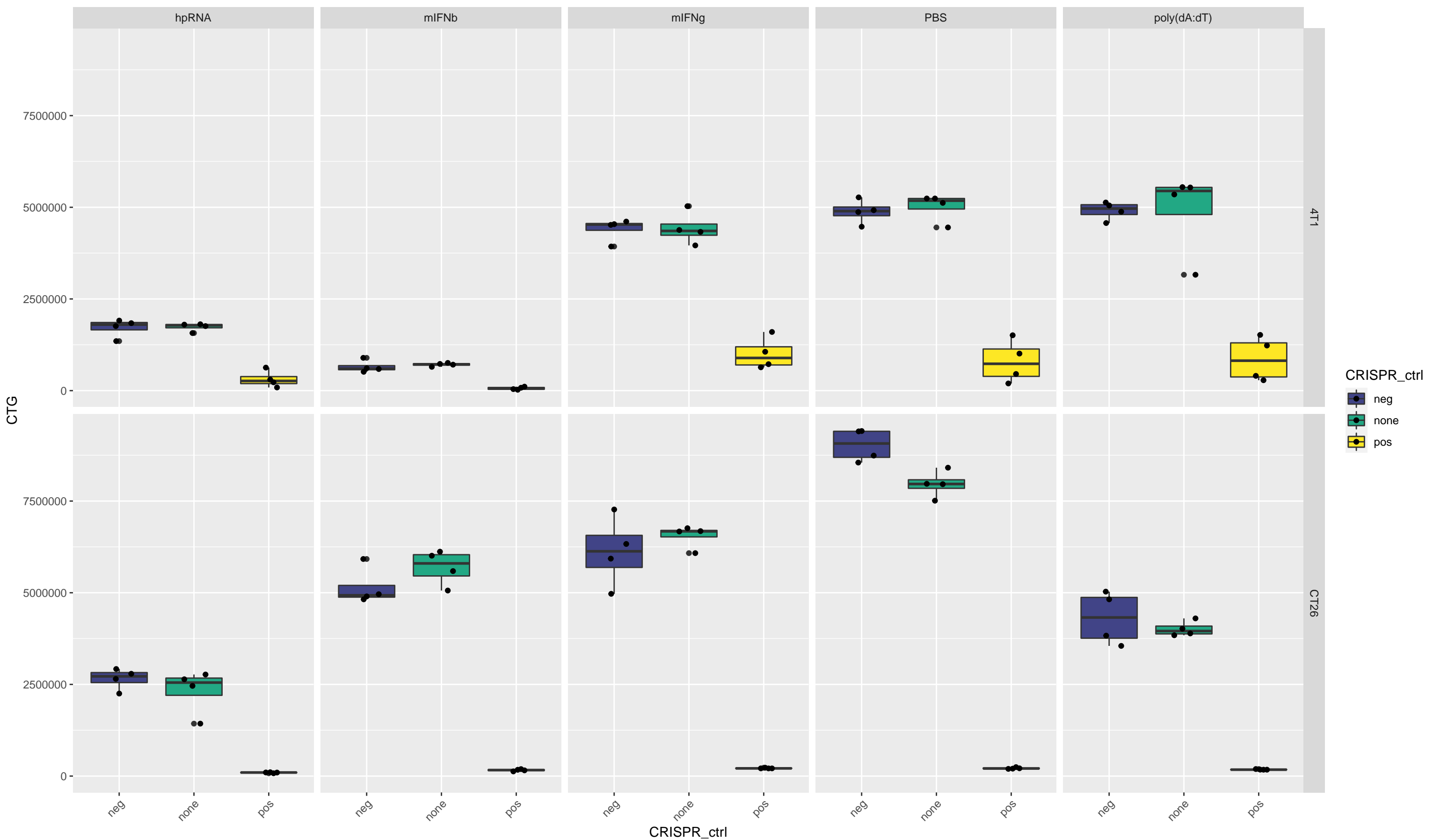
Method	Category	AT1 Median	AT1 IQR	AT1 Min	AT1 Max	CT26 Median	CT26 IQR	CT26 Min	CT26 Max
hpRNA	neg	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	none	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	pos	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
mIFNb	neg	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	none	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	pos	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
mIFNg	neg	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	none	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	pos	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
PBS	neg	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	none	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	pos	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
poly(dA:dT)	neg	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	none	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8
	pos	~0.5	~0.4 - 0.6	~0.2	~0.8	~0.5	~0.4 - 0.6	~0.2	~0.8

Experiment ID: 20201218\_JM  
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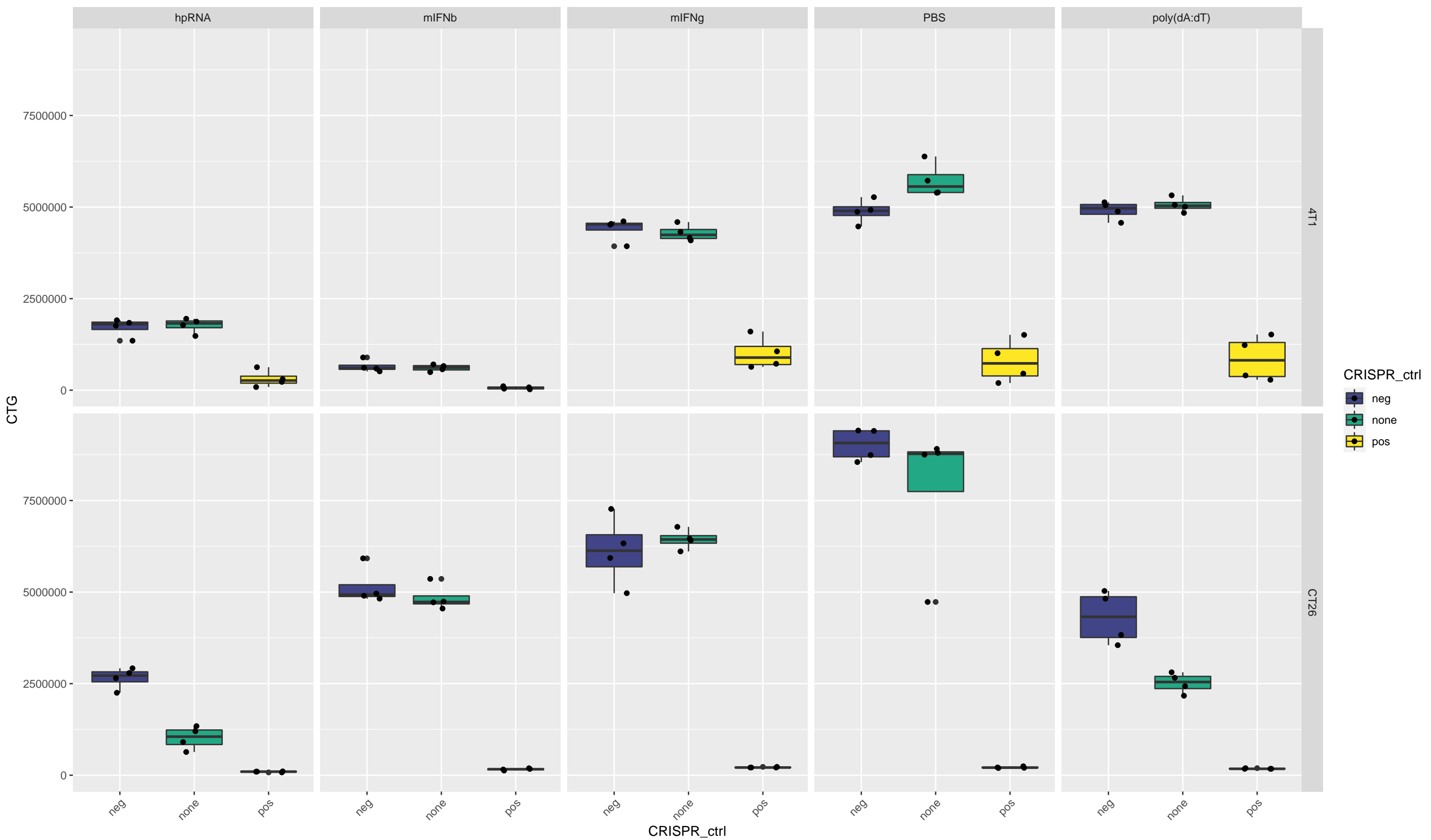
mDHX9\_pooled



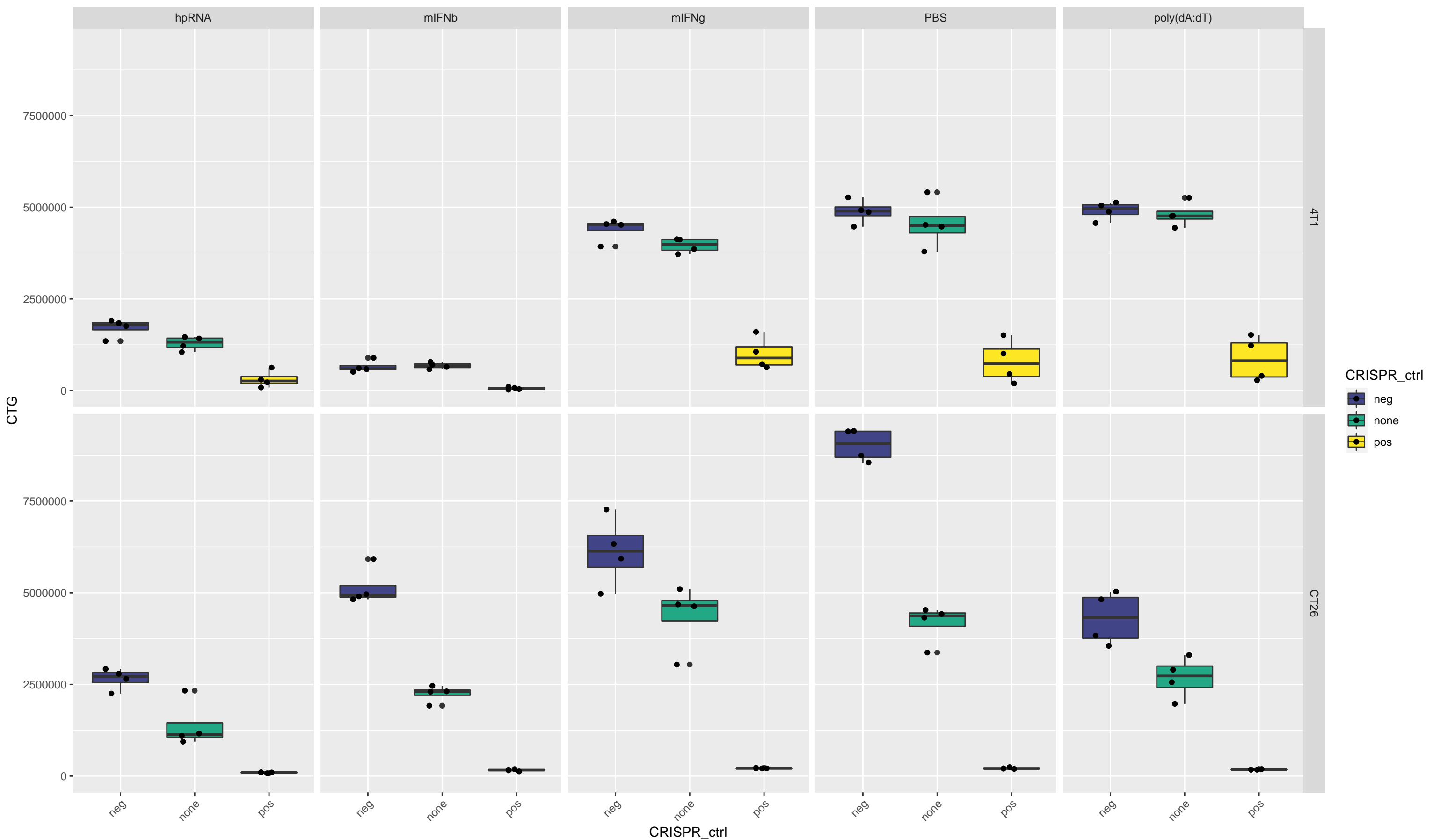
mDDX5\_pooled



## mRNASEH2A\_pooled

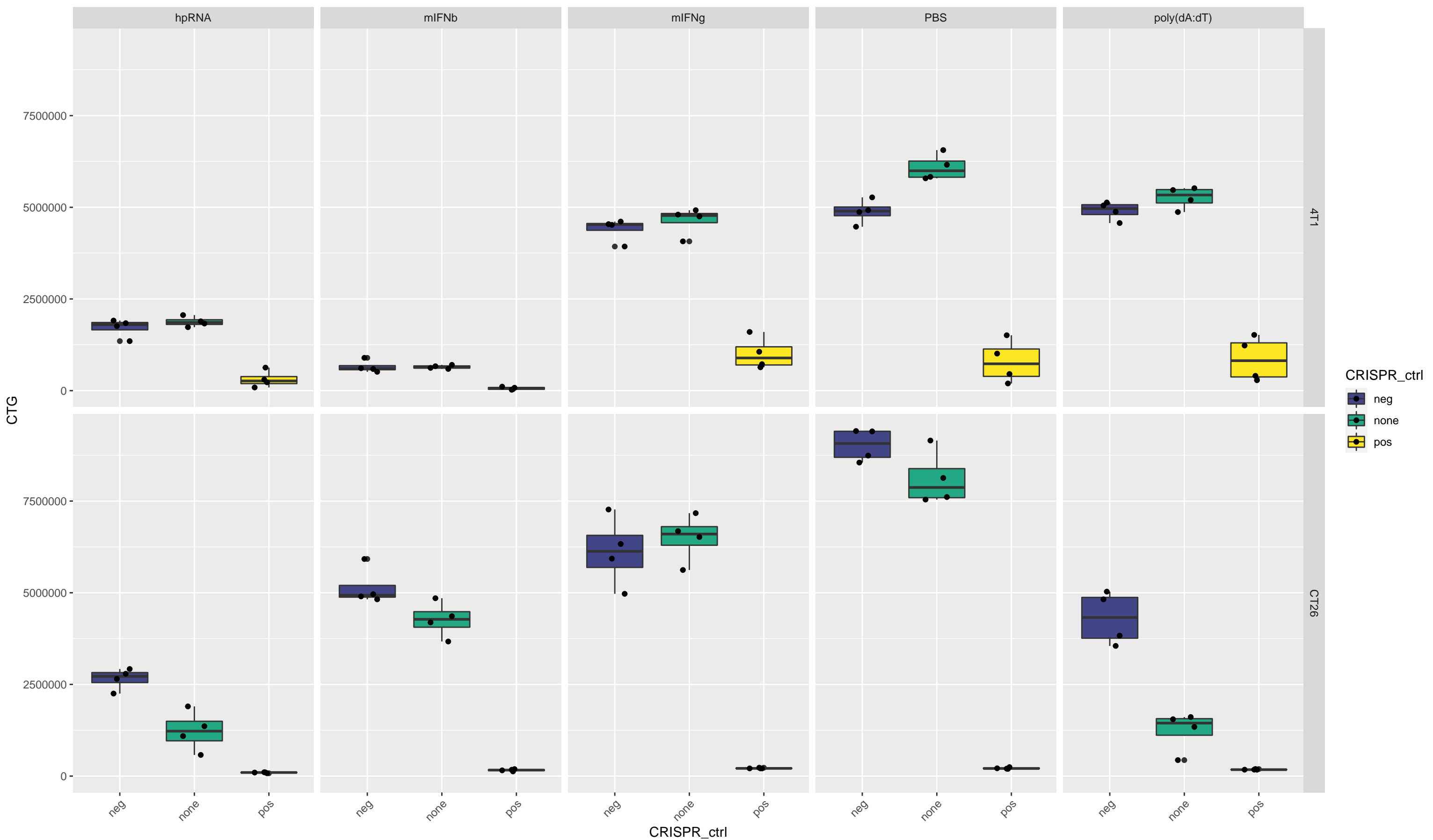


mDDX3X\_pooled

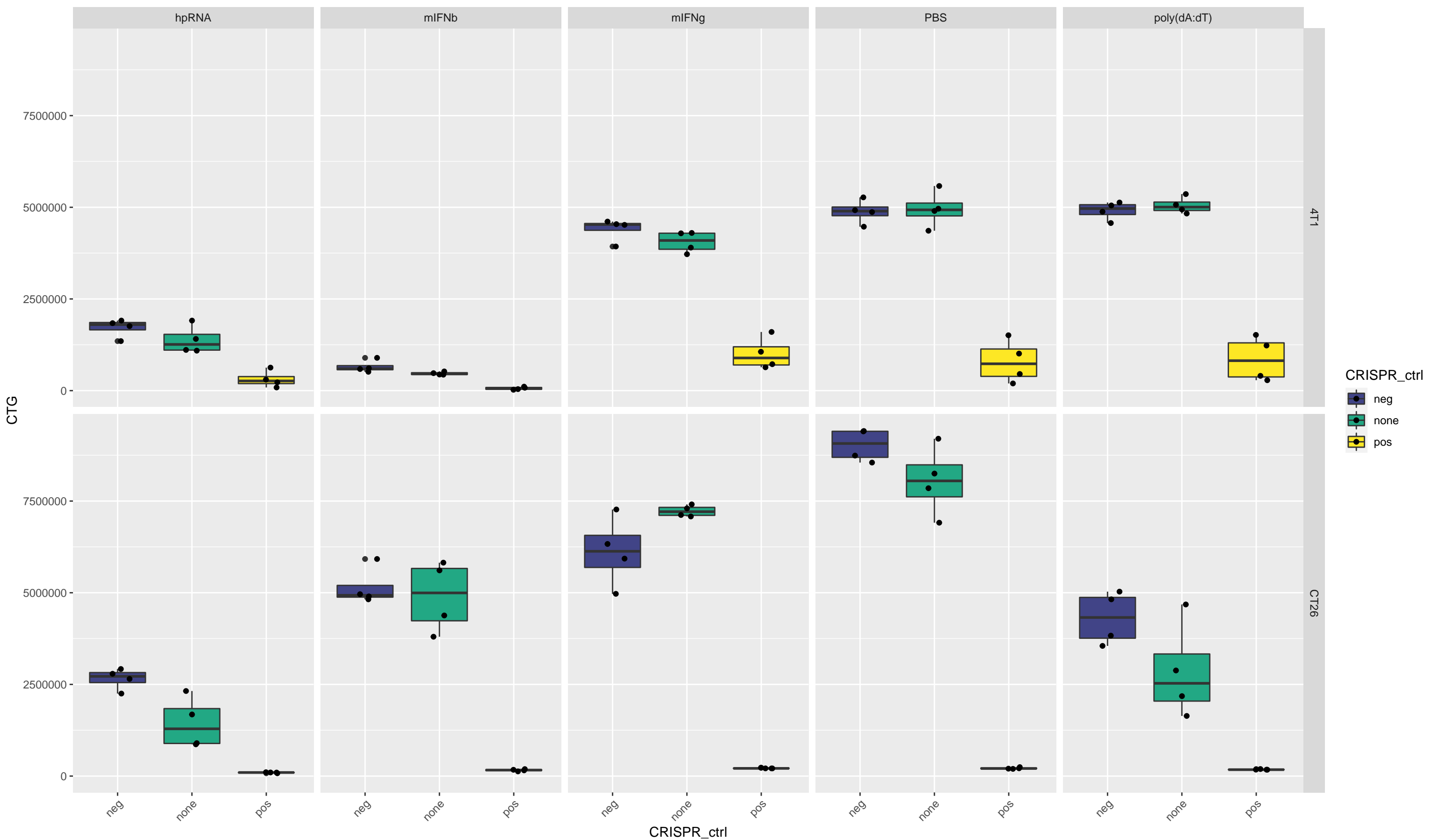


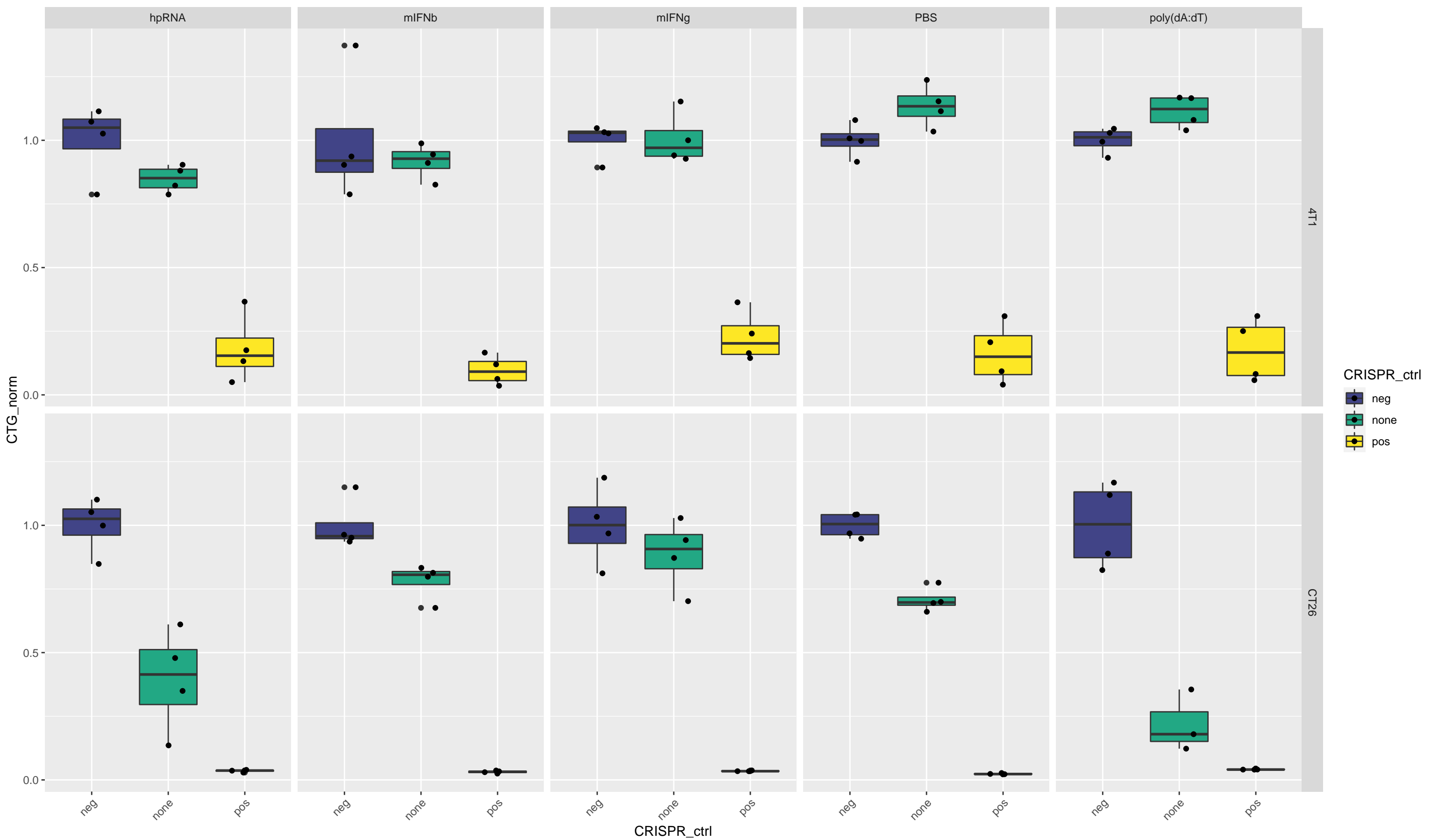


# mPARP14\_pooled

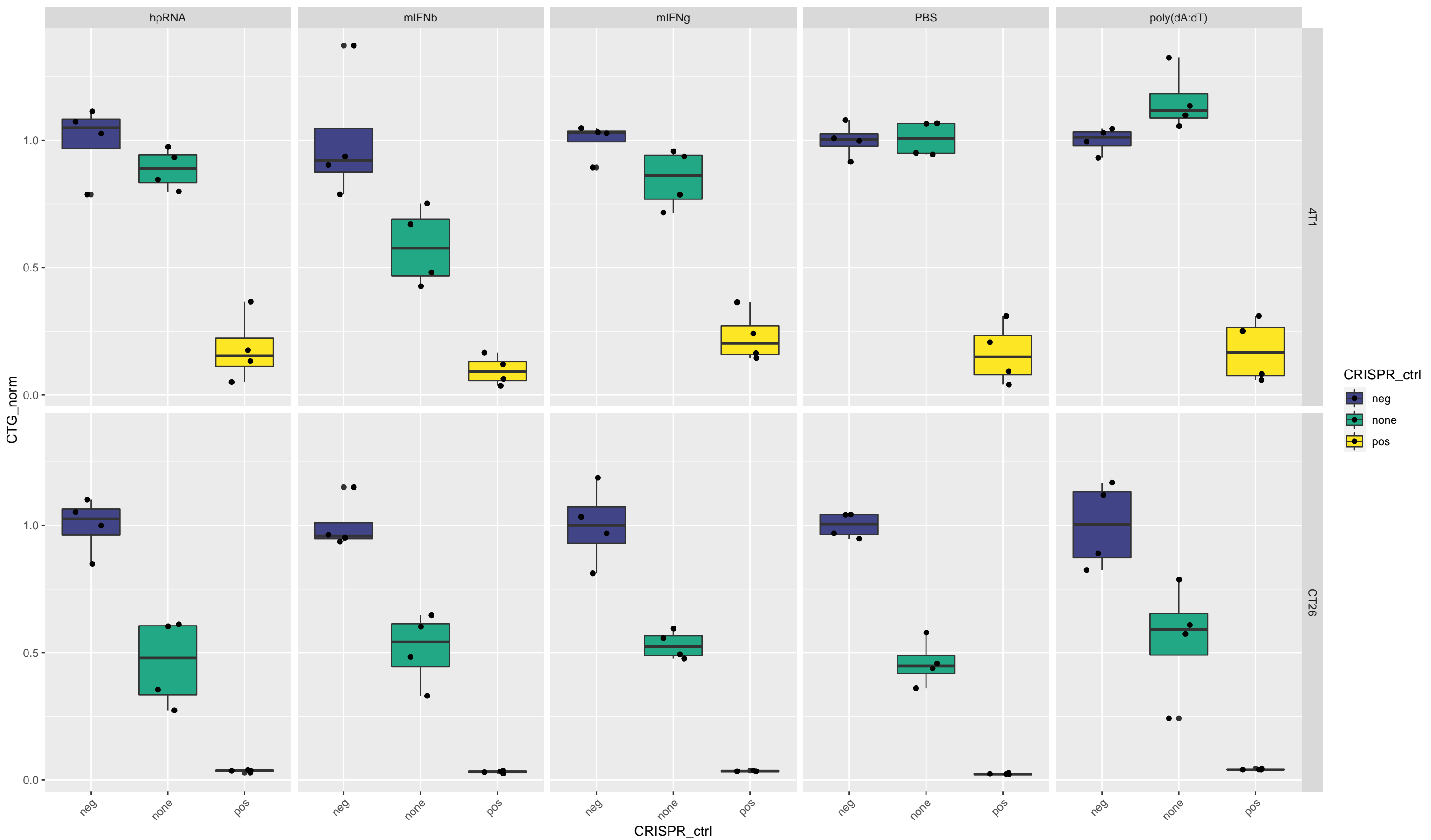


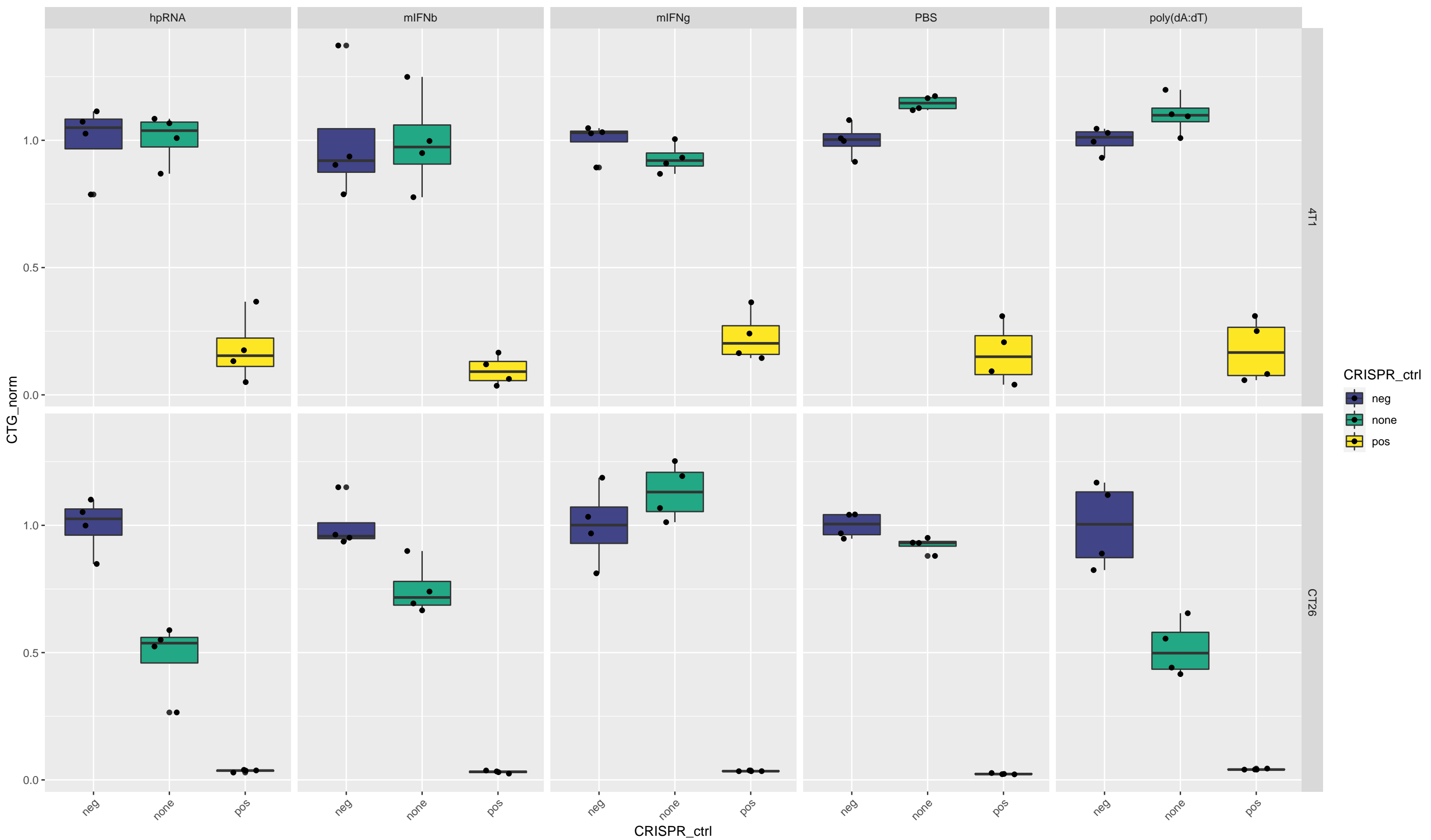
mDDX6\_pooled

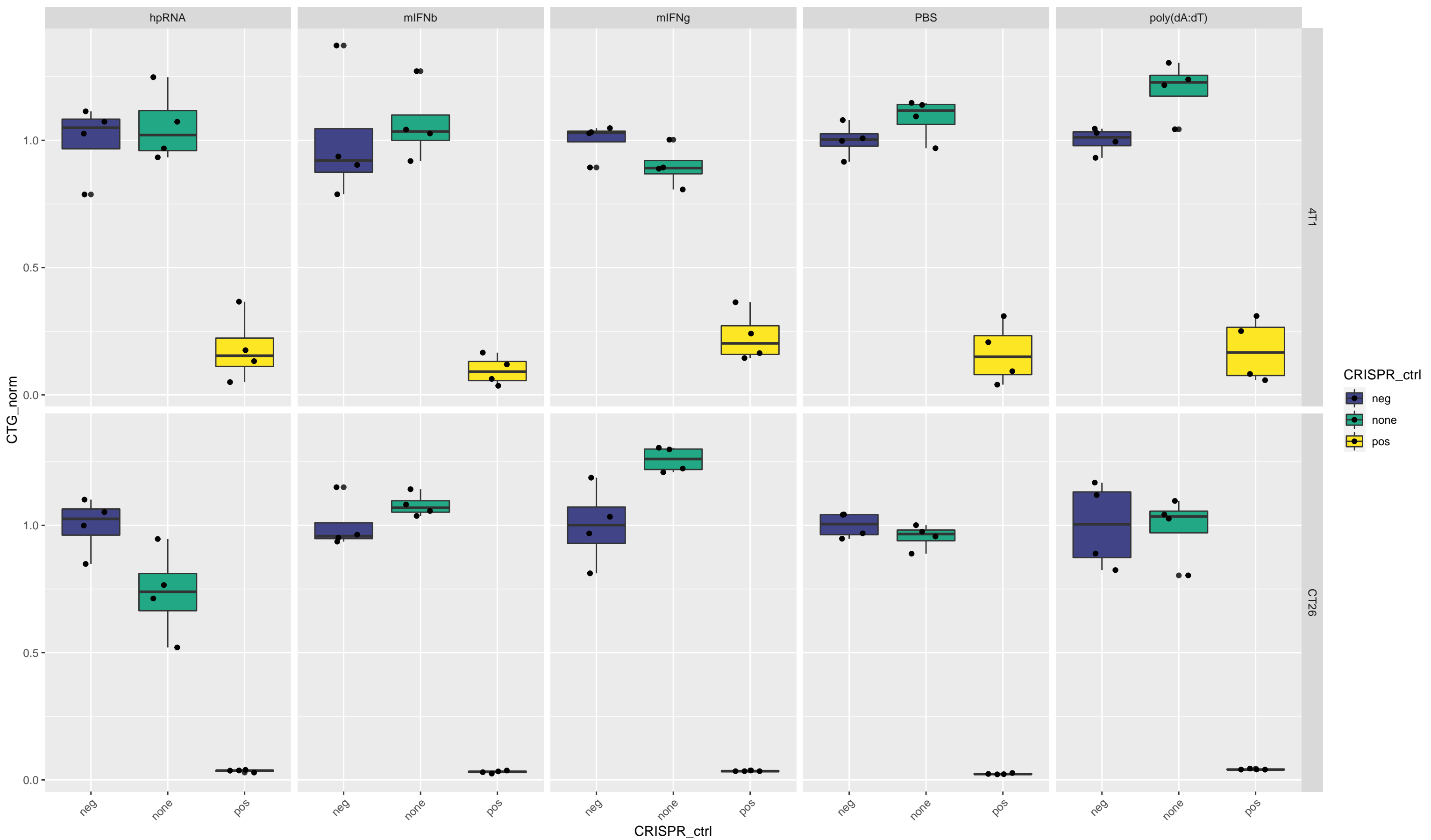


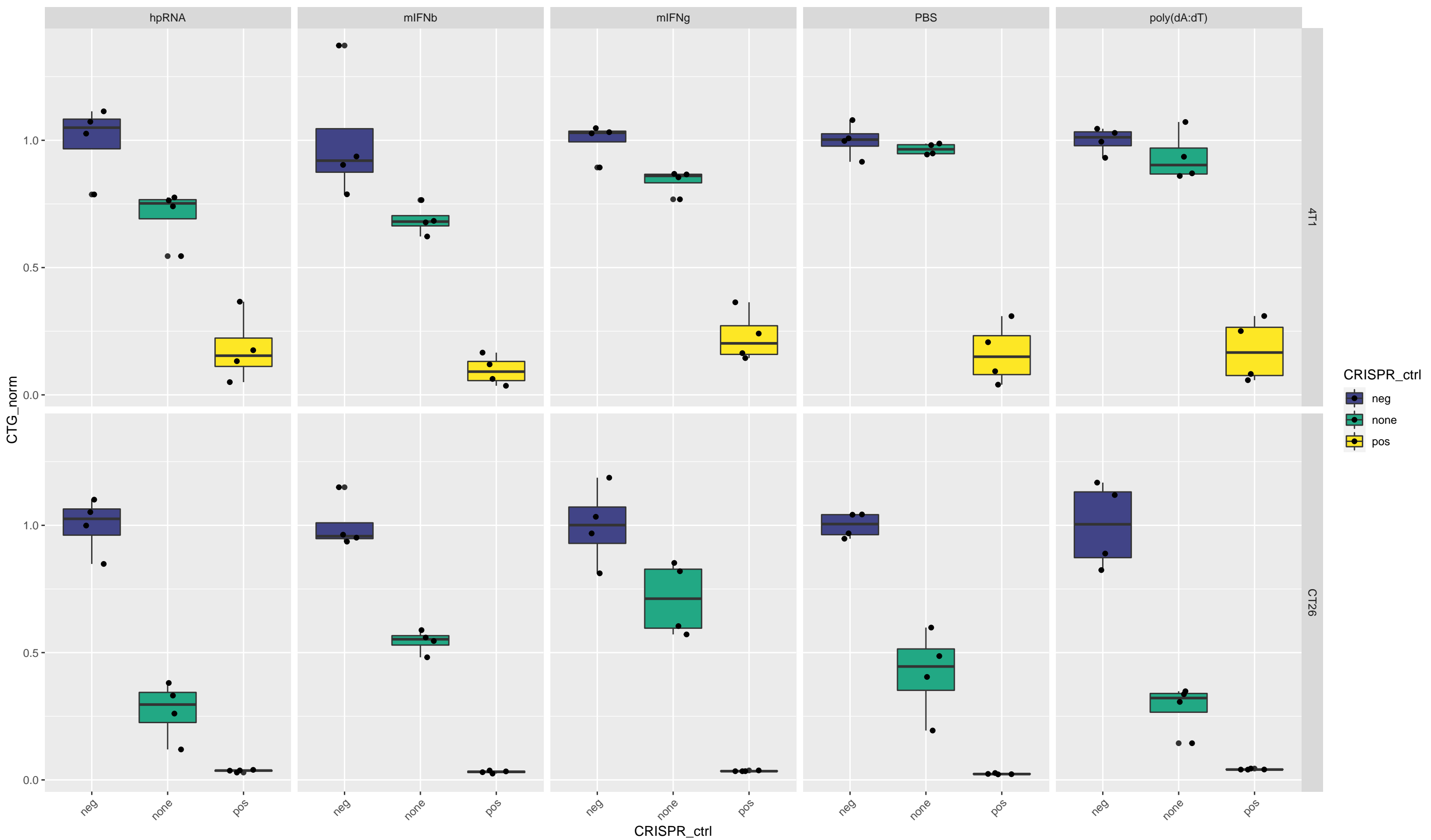


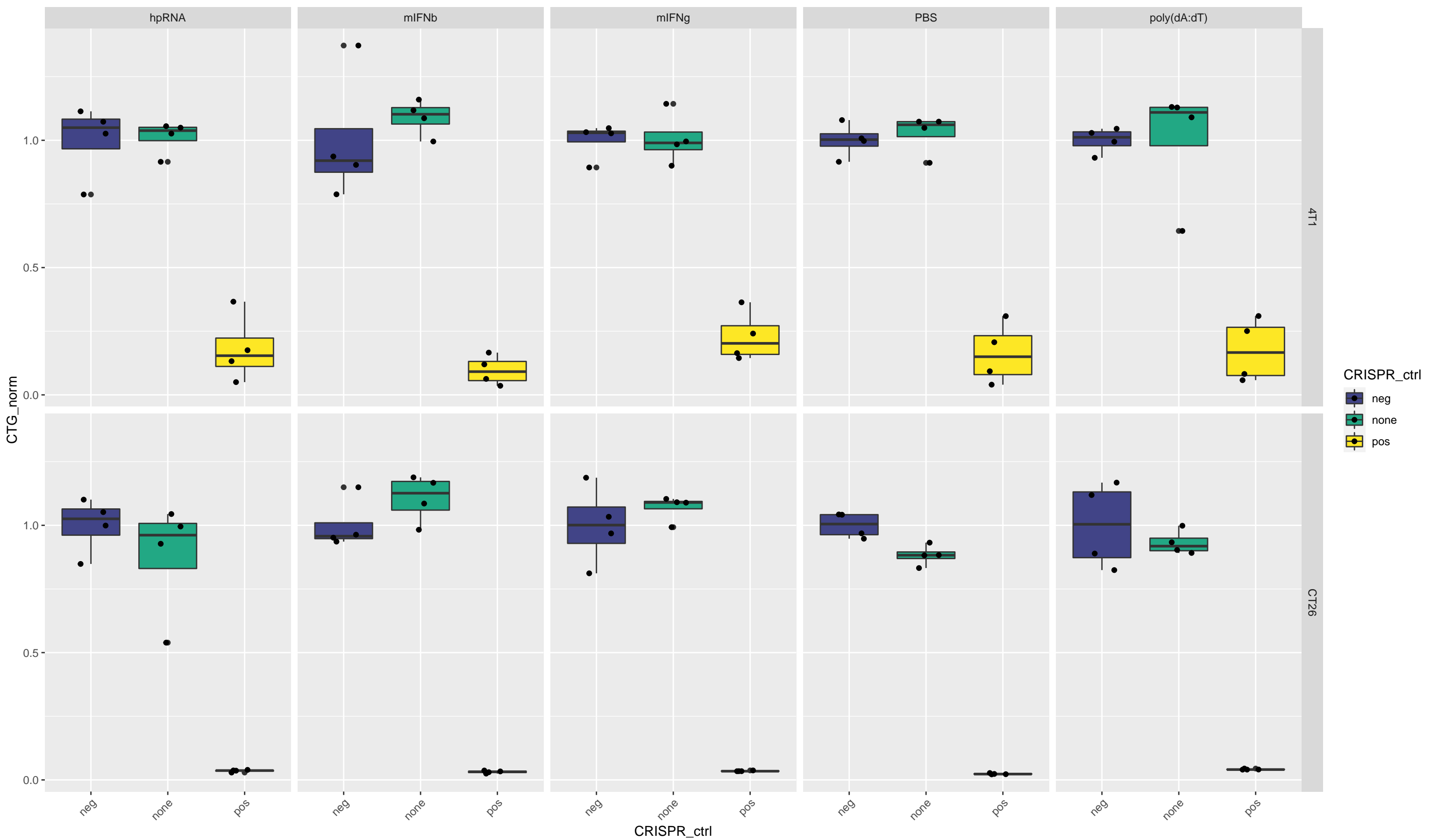
mADAR\_pooled



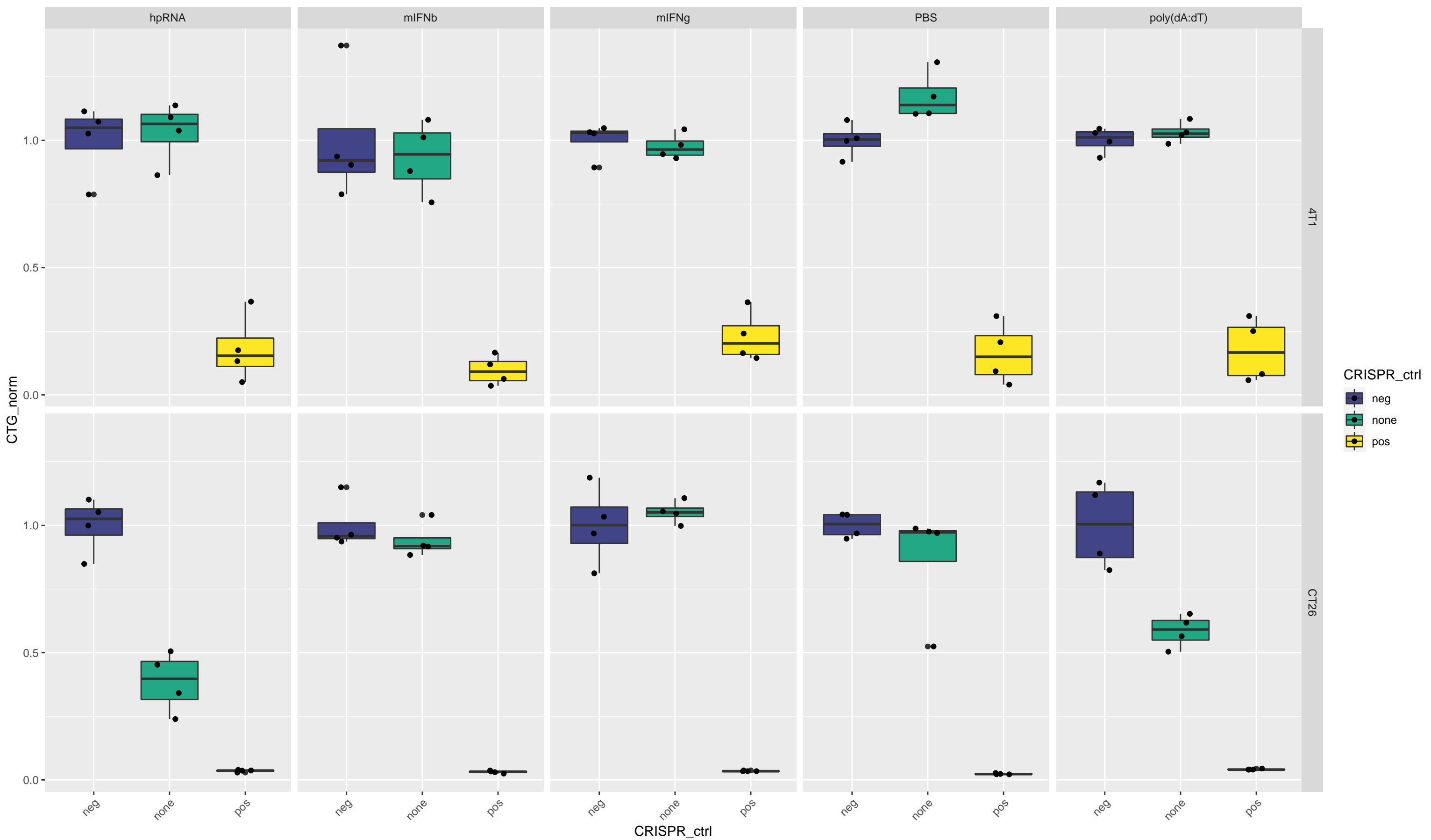


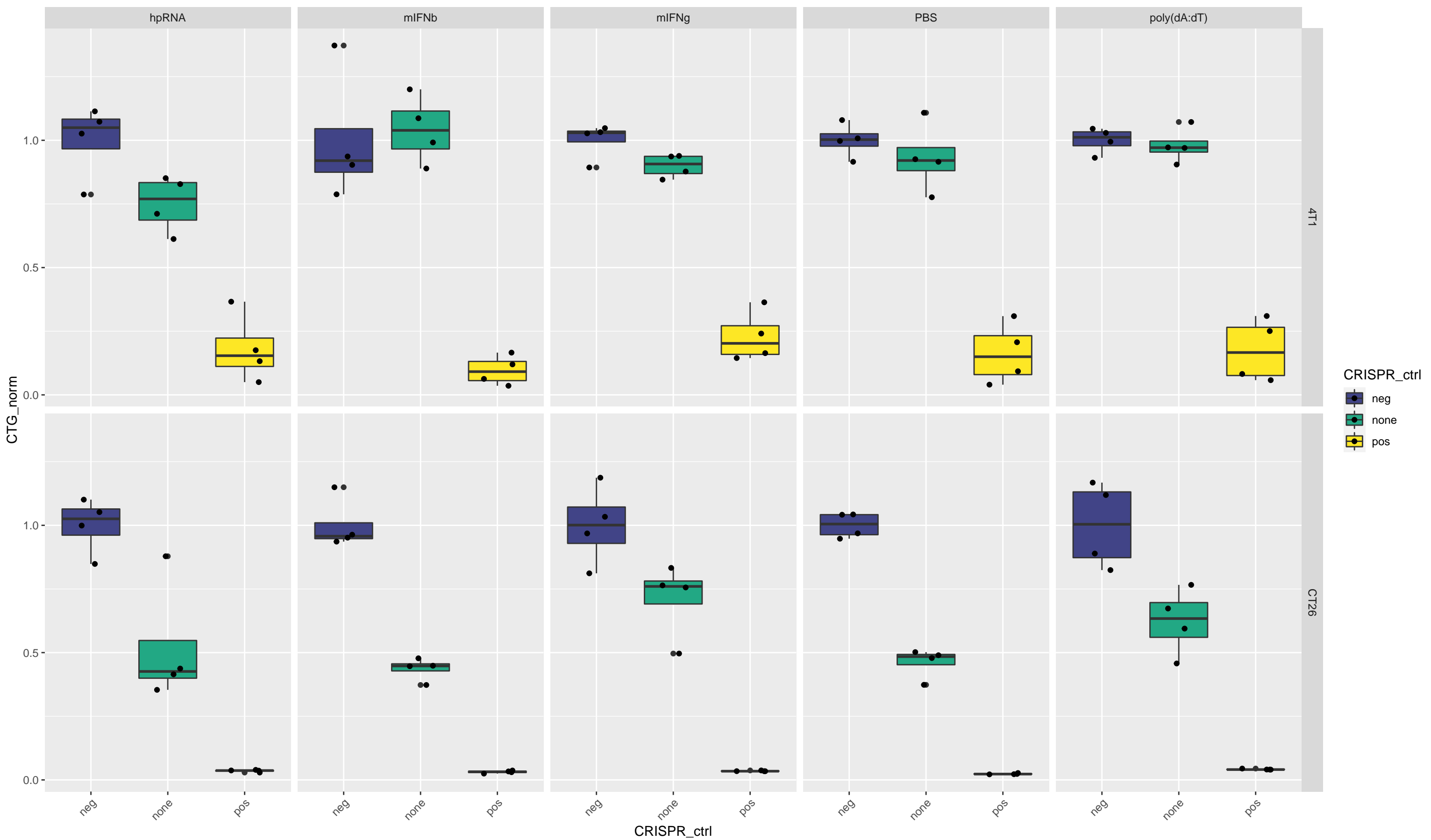












mPARP14\_pooled

