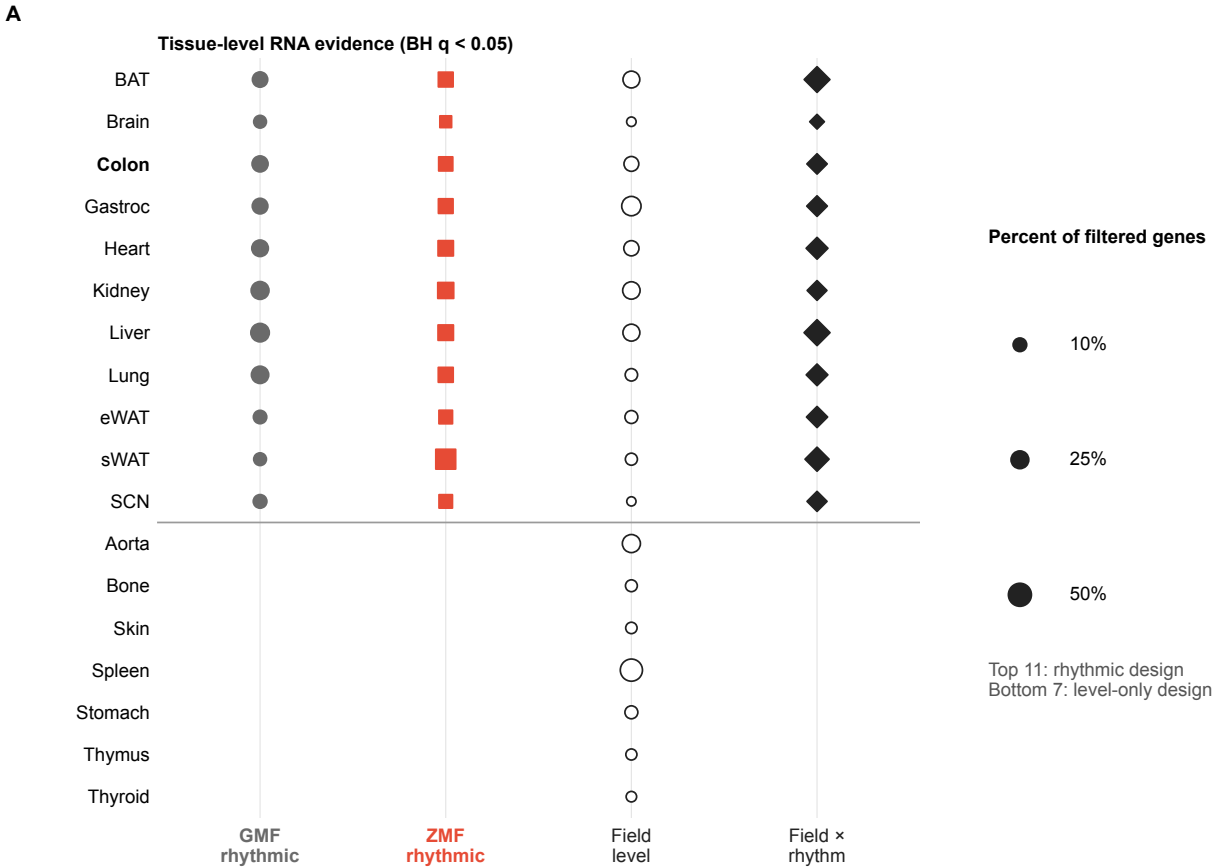
A
Tissue-level RNA evidence (BH q < 0.05)
BAT
Brain
Colon
Gastroc
Heart
Kidney
Liver
Lung
eWAT
sWAT
SCN
Aorta
Bone
Skin
Spleen
Stomach
Thymus
Thyroid
GMF rhythmic
ZMF rhythmic
Field level
Field × rhythm
Percent of filtered genes
10%
25%
50%
Top 11: rhythmic design
Bottom 7: level-only design