

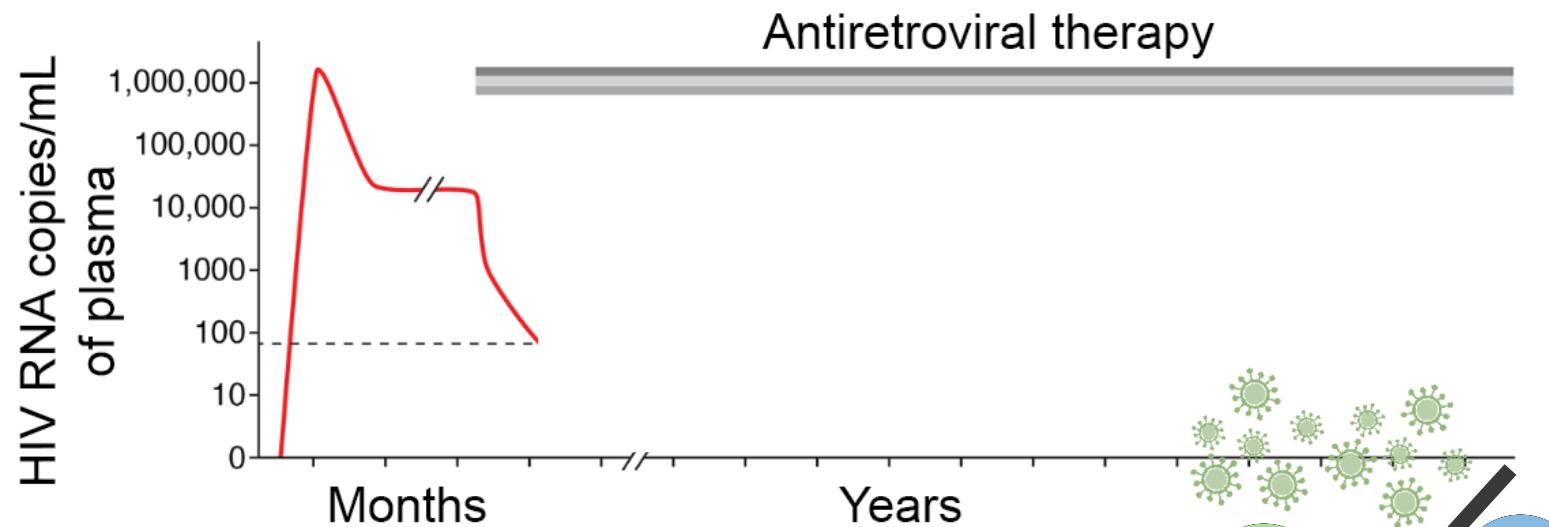


5'Leader-defective proviruses are a common source of persistent viremia despite effective ART

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I have received honoraria from Gilead to present at scientific conferences

Persistent viremia complicates HIV clinical care

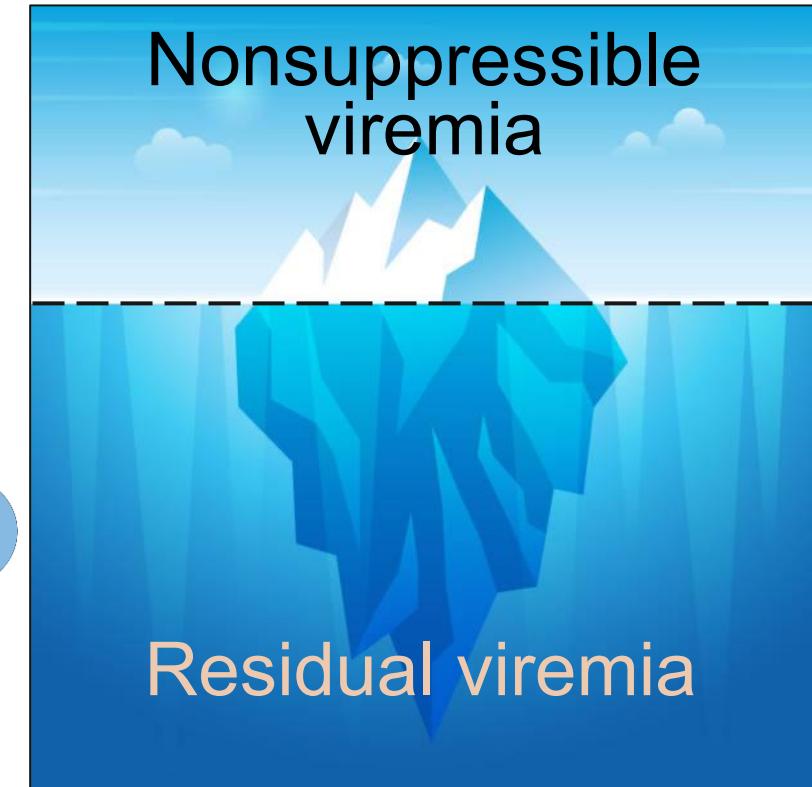
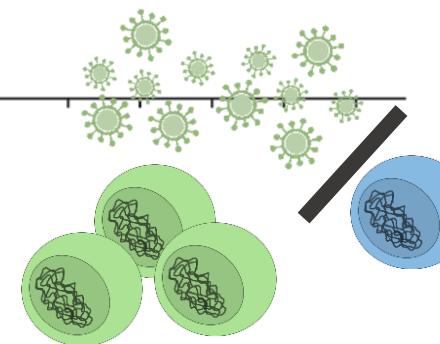


Bob and Janet Siliciano, JCI 2020

Bailey, JV 2006

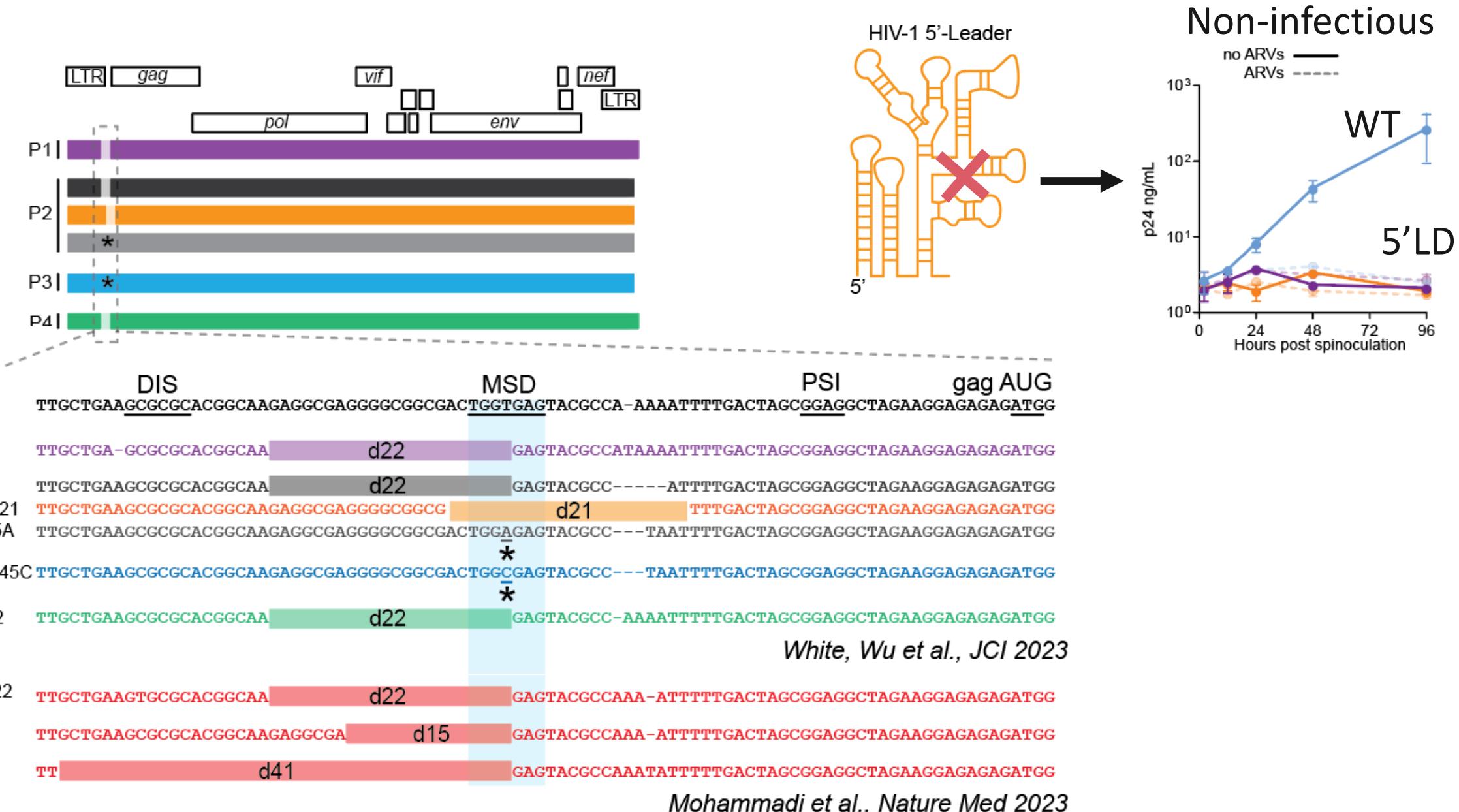
Simonetti, PNAS 2016

Halvas, JCI 2020



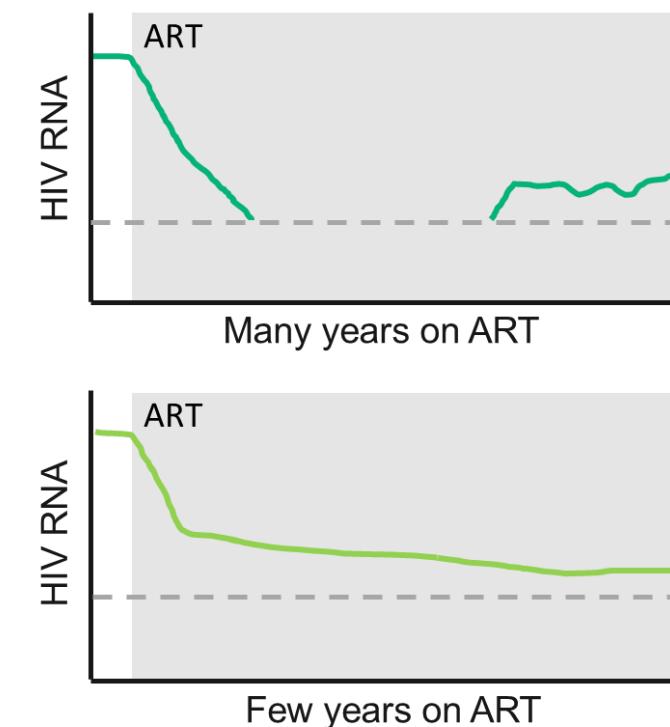
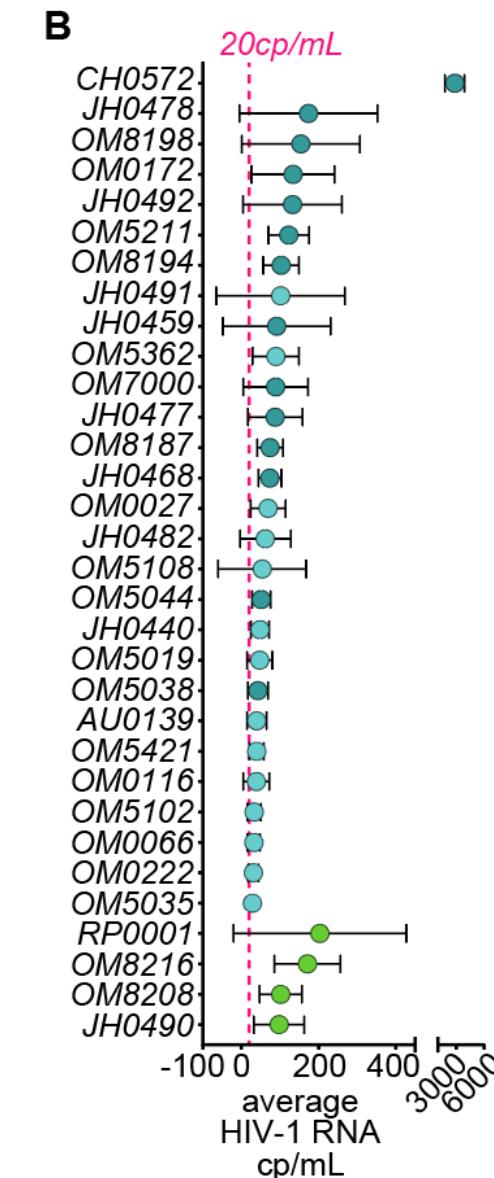
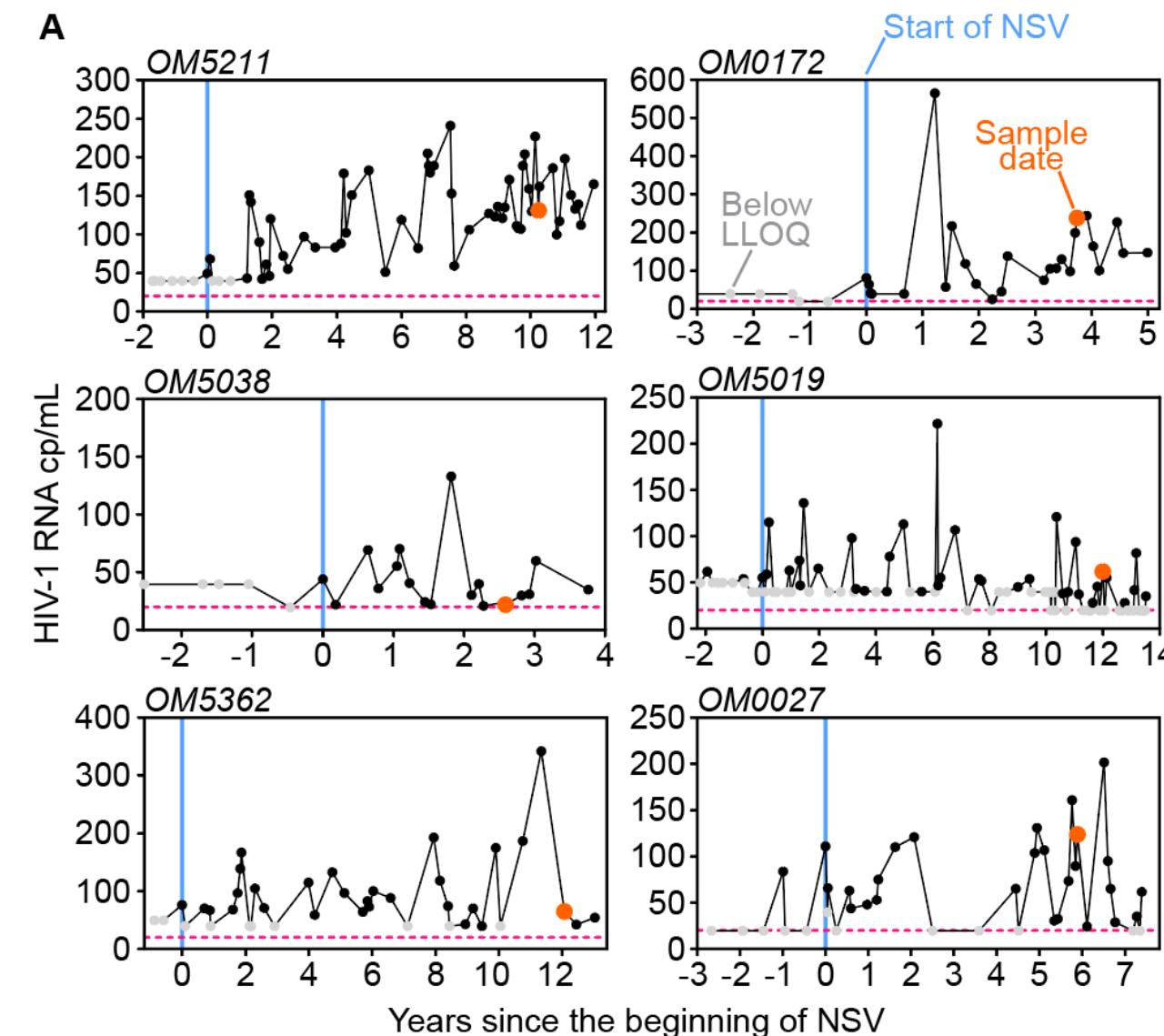
Determining virus and host drivers of persistent viremia can improve clinical care and our understanding of HIV persistence

Proviruses with 5'Leader Defects can cause NSV

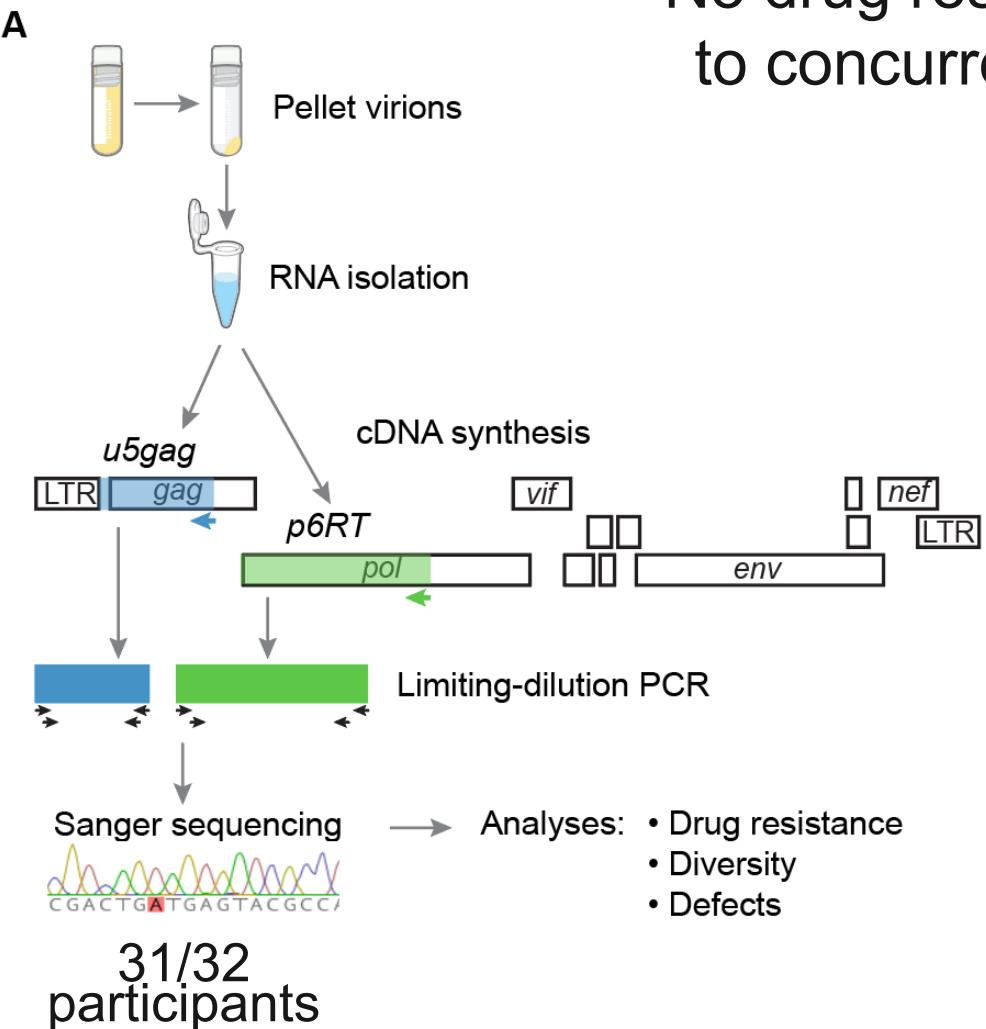


To which extent do 5'L defects contribute to NSV?

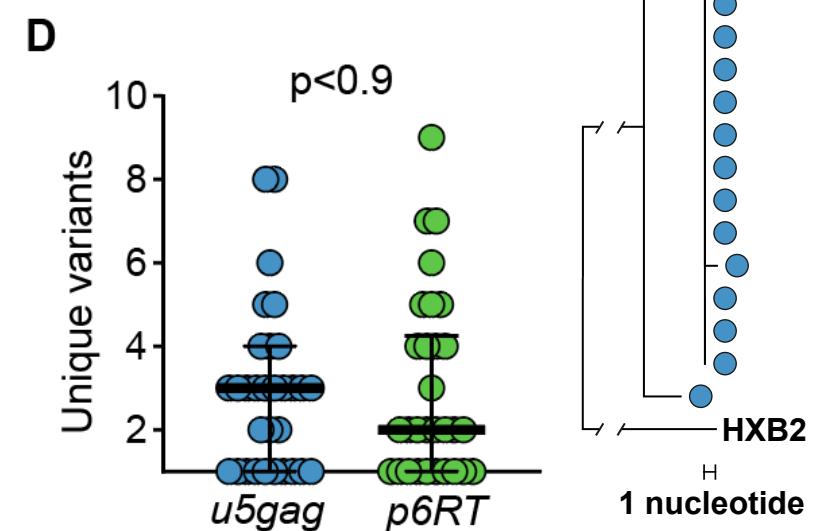
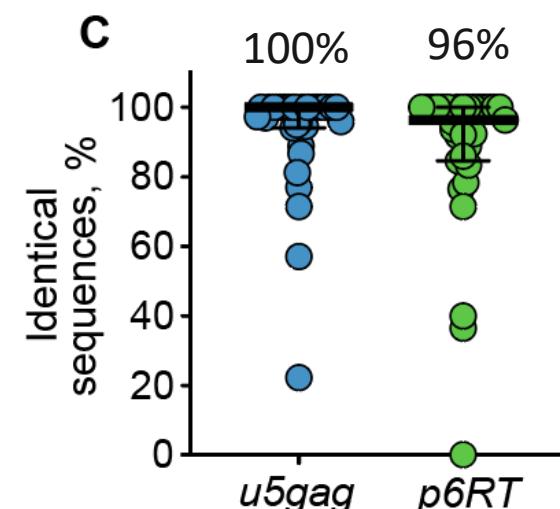
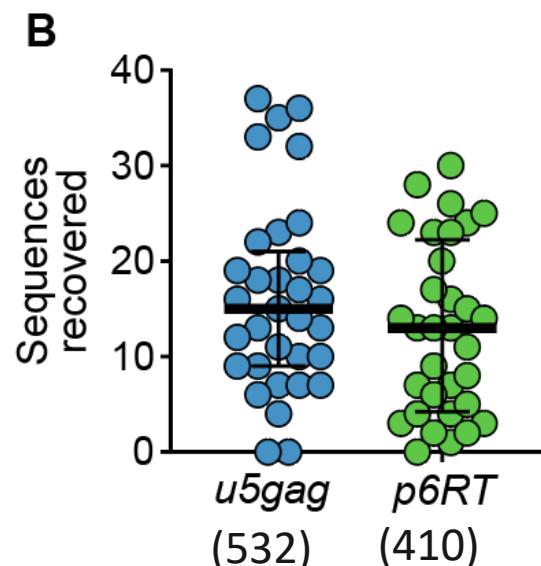
We studied **32 people on ART** with persistent (or intermittent) viremia >20 cp/mL



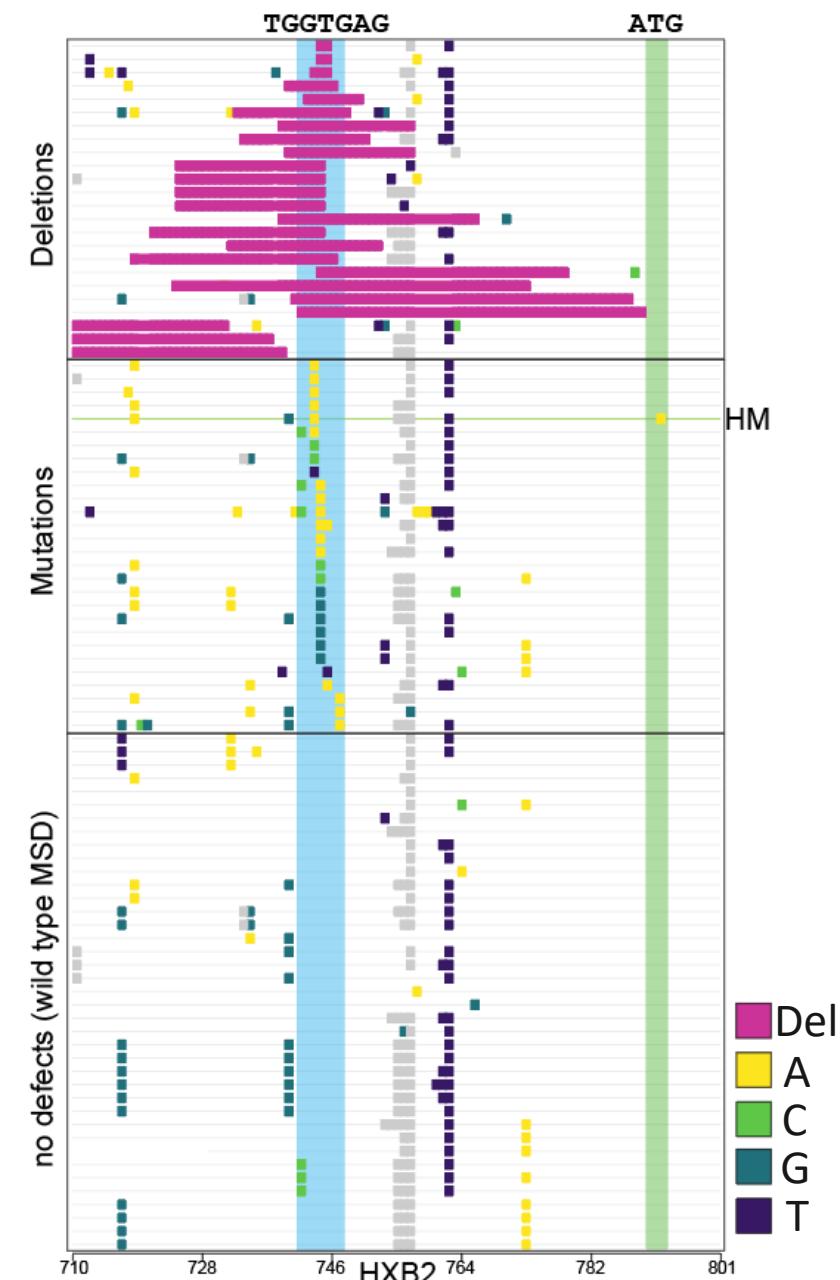
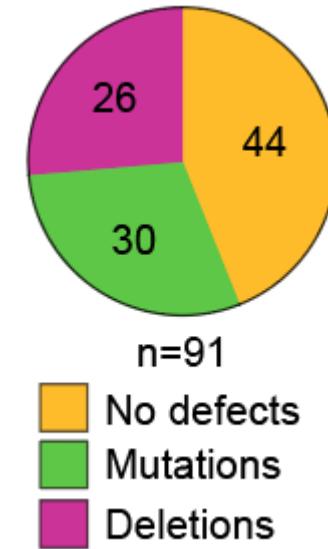
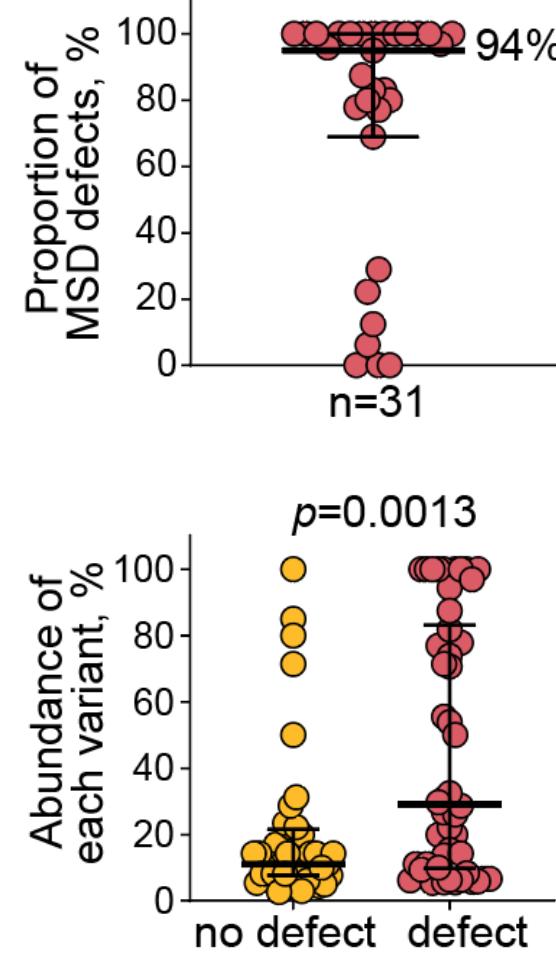
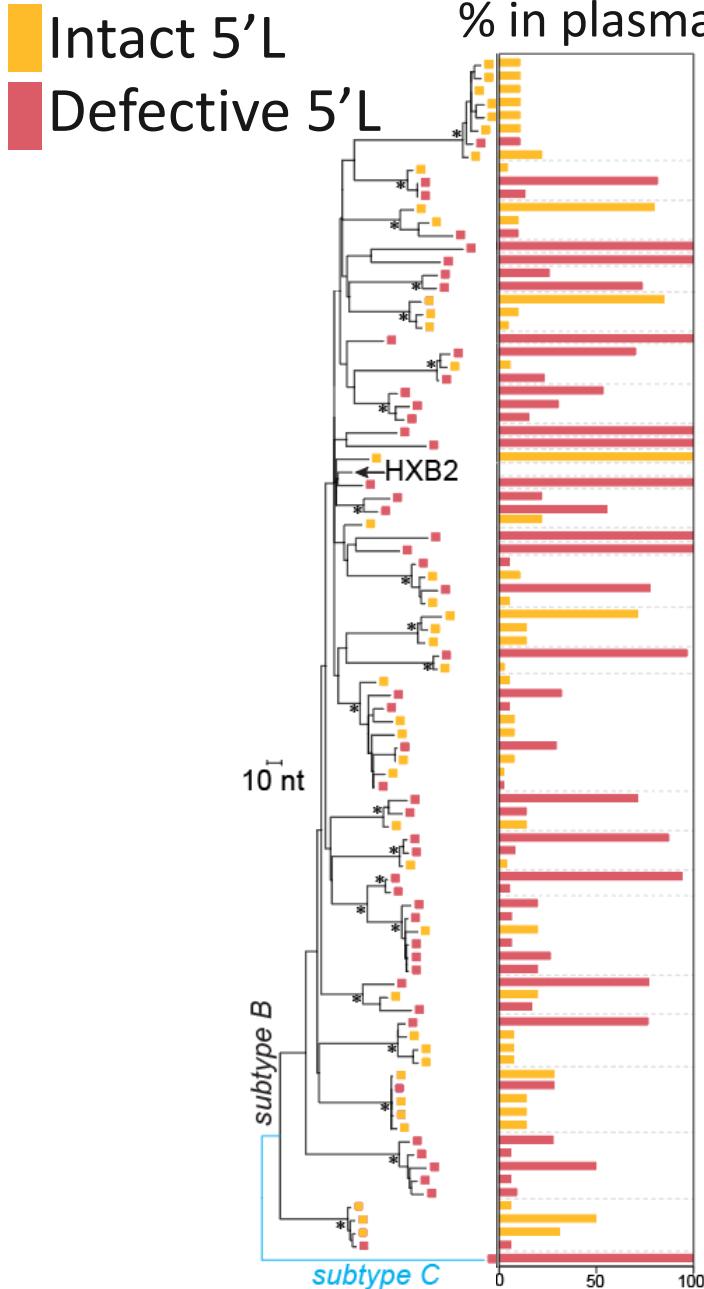
HIV-1 variants contributing to NSV are highly clonal



No drug resistance mutations detected to concurrent PIs, NRTIs, or NNRTIs

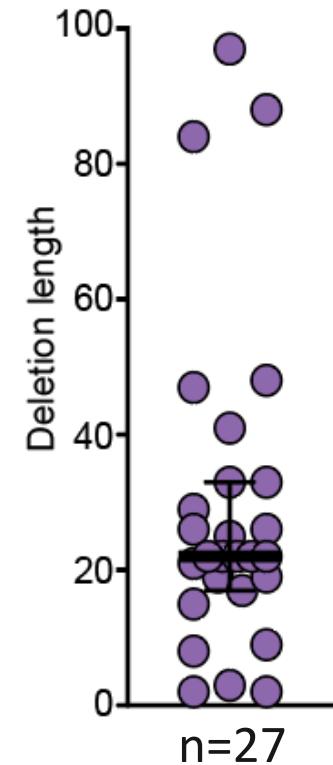
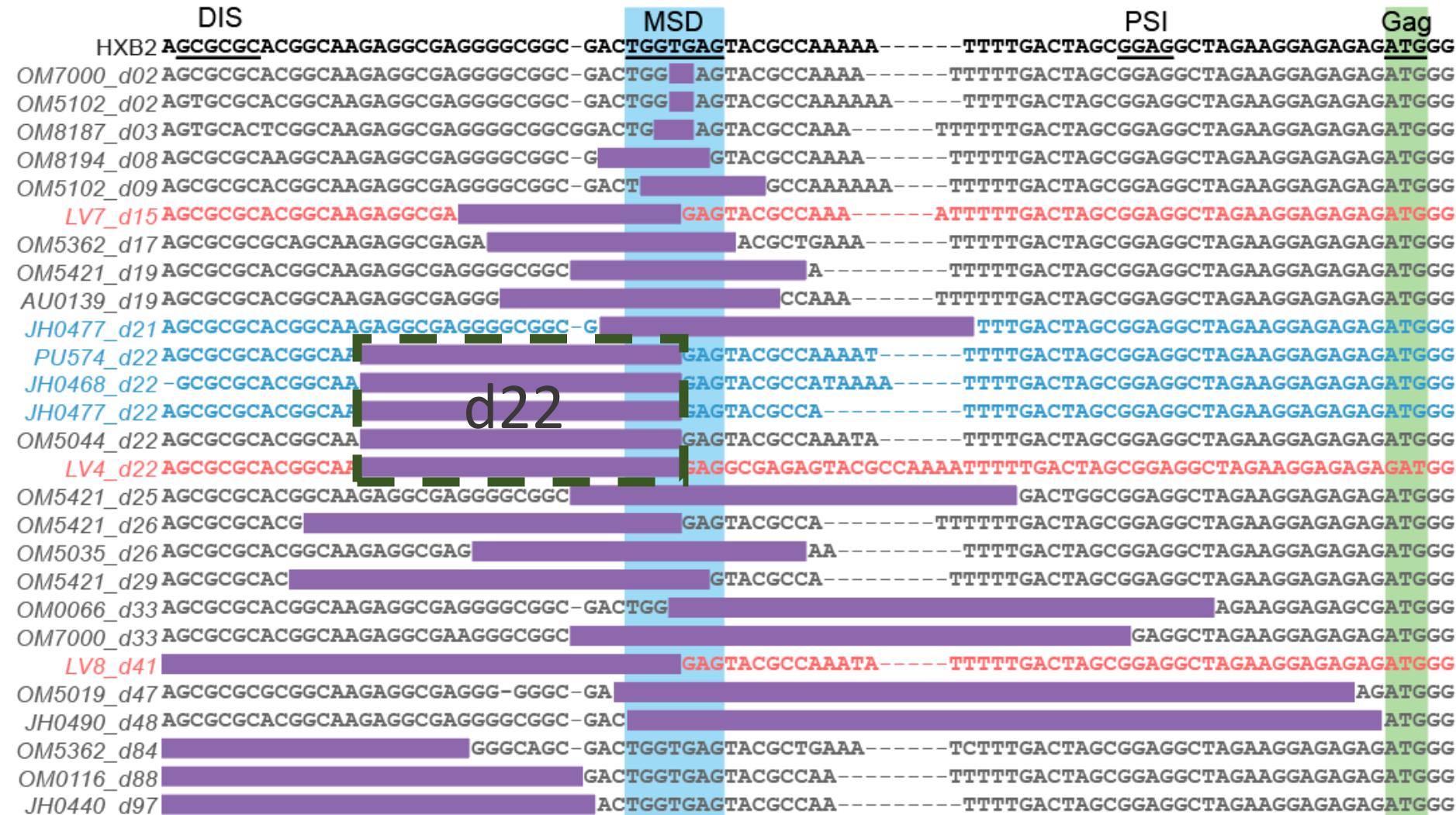


>90% of virus in plasma has 5'Leader defects



Deletions involve the Major Splicing Donor

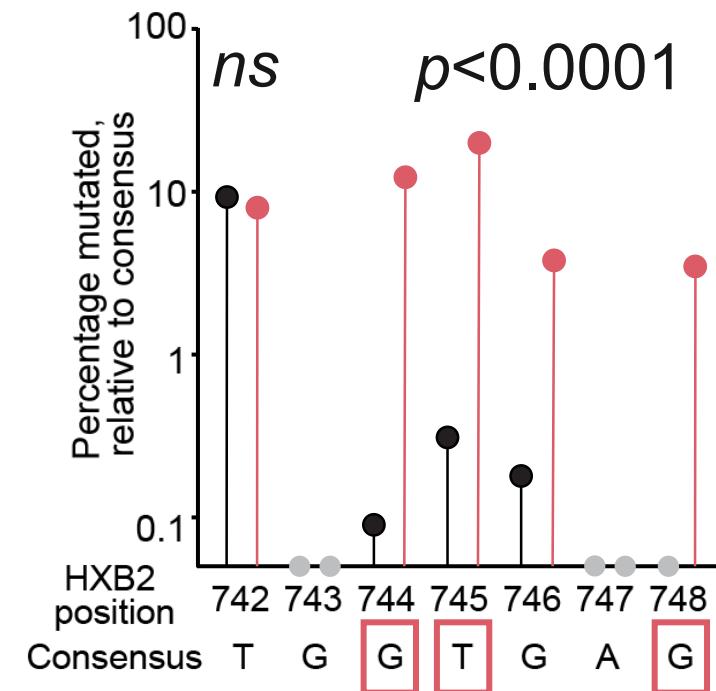
Deletions



Deletion
White, Wu et al.
Mohammadi et al.

Specific MSD mutations are enriched in NSV

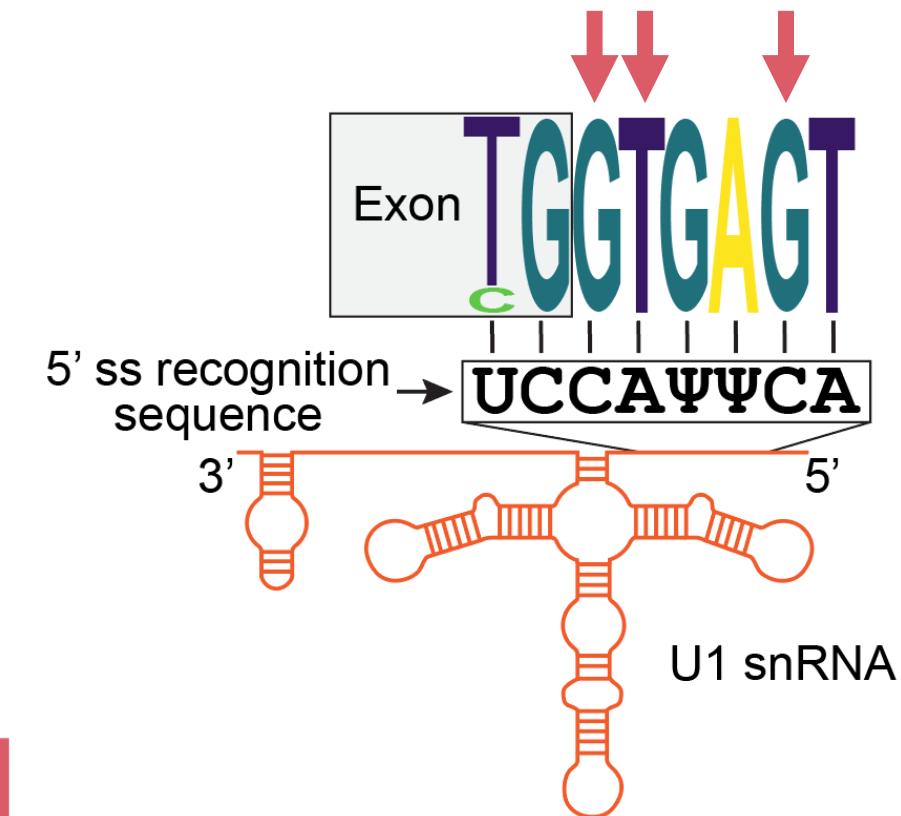
MSD	
HXB2	GCGACTGGT GAGT TACGCC
JH0492_G744A	GCGACTG A TGAGTACGCC
OM5038_G744A	GCGACTG A TGAGTACGCC
OM8194_G744A	GCGACTG A TGAGTACGCC
JH0440_G744A	GCGACTG A TGAGTACGCC
OM0116_G744A	GCG G CTG A TGAGTACGCC
OM5211_G744A	GCGAC G TGAGTACGCC
JH0482_G744C	GCGACTG C TGAGTACGCC
OM5019_G744C	GCGACTG C TGAGTACGCC
OM0222_G744T	GCGACTG T TGAGTACGCC
OM5211_T745A	GCGAC C G A GAGTACGCC
JH0477_T745A	GCGACTGGGAGTACGCC
JH0491_T745A	GCGA AC G G GAGTACGCC
AU0139_T745A	GCGACTG G AAGTACGCC
OM5108_T745A	GCGACTG G GAGTACGCC
OM5421_T745A	GCGACTG G GAGTACGCC
JH0478_T745C	GCGACTGGGAGTACGCC
RP0001_T745C	GCGACTG G GAGTACGCC
JH0440_T745G	GCGACTG G GAGTACGCC
JH0440_T745G	GCGACTG G GAGTACGCC
OM8208_T745G	GCGGCTG G GAGTACGCC
OM7000_T745G	GCGACTG G GAGTACGCC
OM0172_T745G	GCGACTG G GAGTACGCC
OM0172_T745G	GCGACTG G GAGTACGCC
JH0477_G746T	G C A T GGT A GAGTACGCC
AU0139_G746A	GCGACTGGT A AAGTACGCC
JH0440_G748A	GCGACTGGT G AATACGCC
OM5035_G748A	GCG G CTGGT G AATACGCC
OM8208_G748A	GCG G CTGGT G AATACGCC



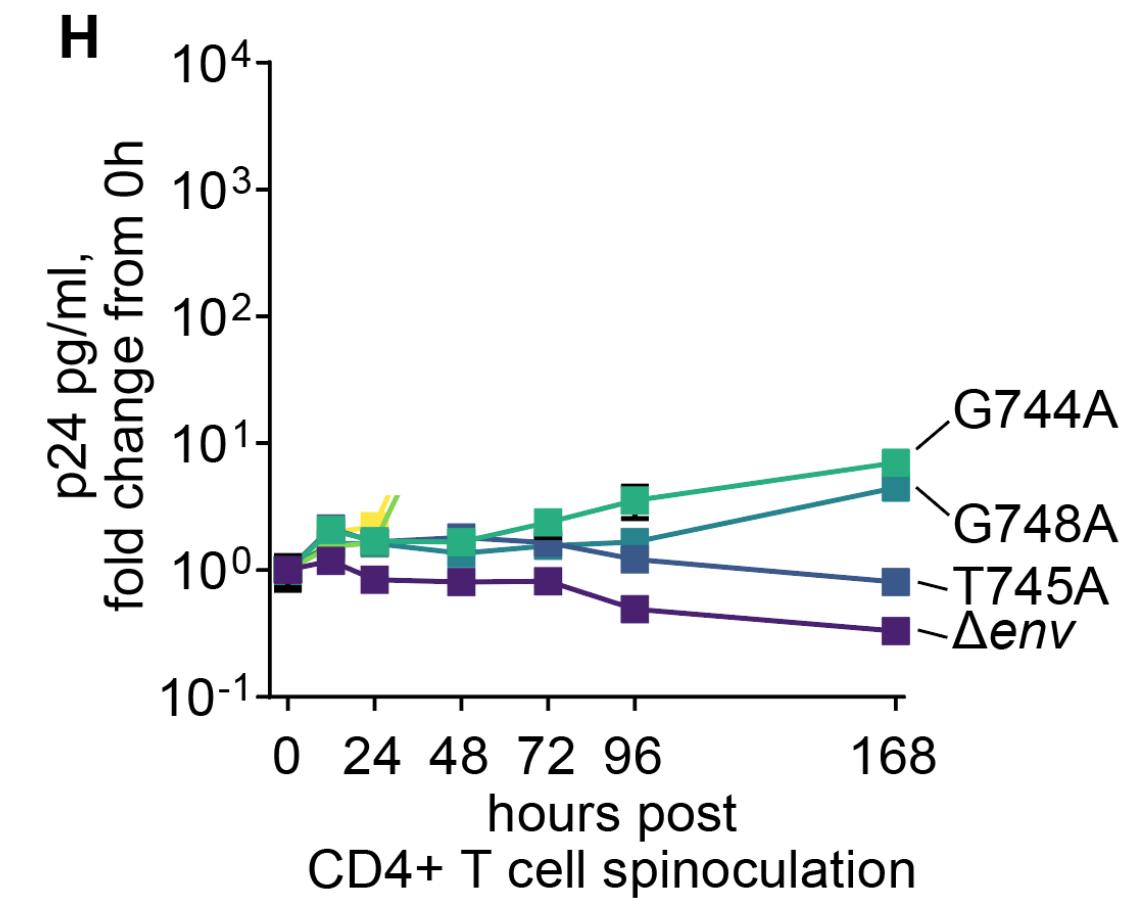
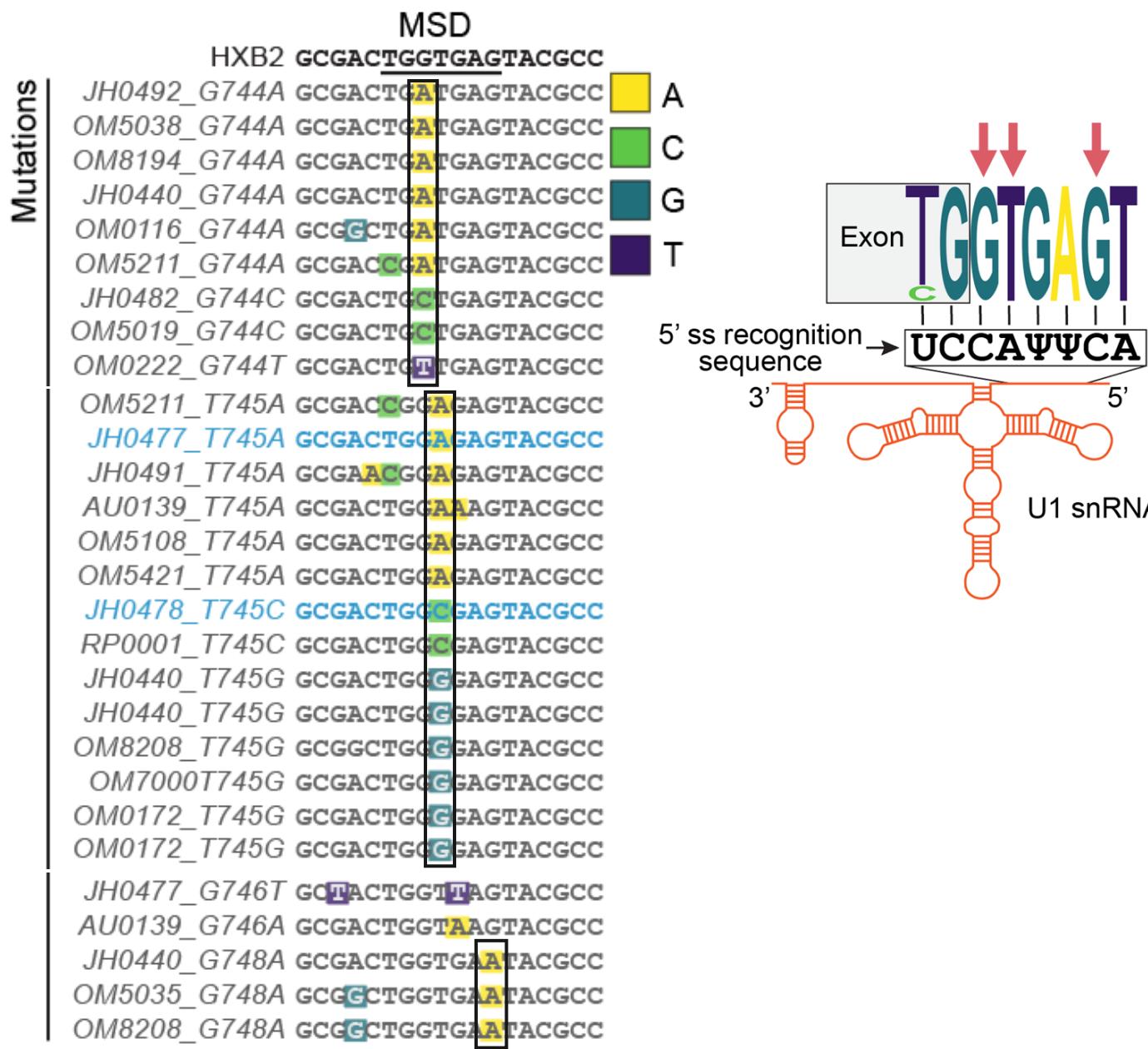
HIV sequence database (plasma RNA, all subtypes) n=2238

χ^2 test

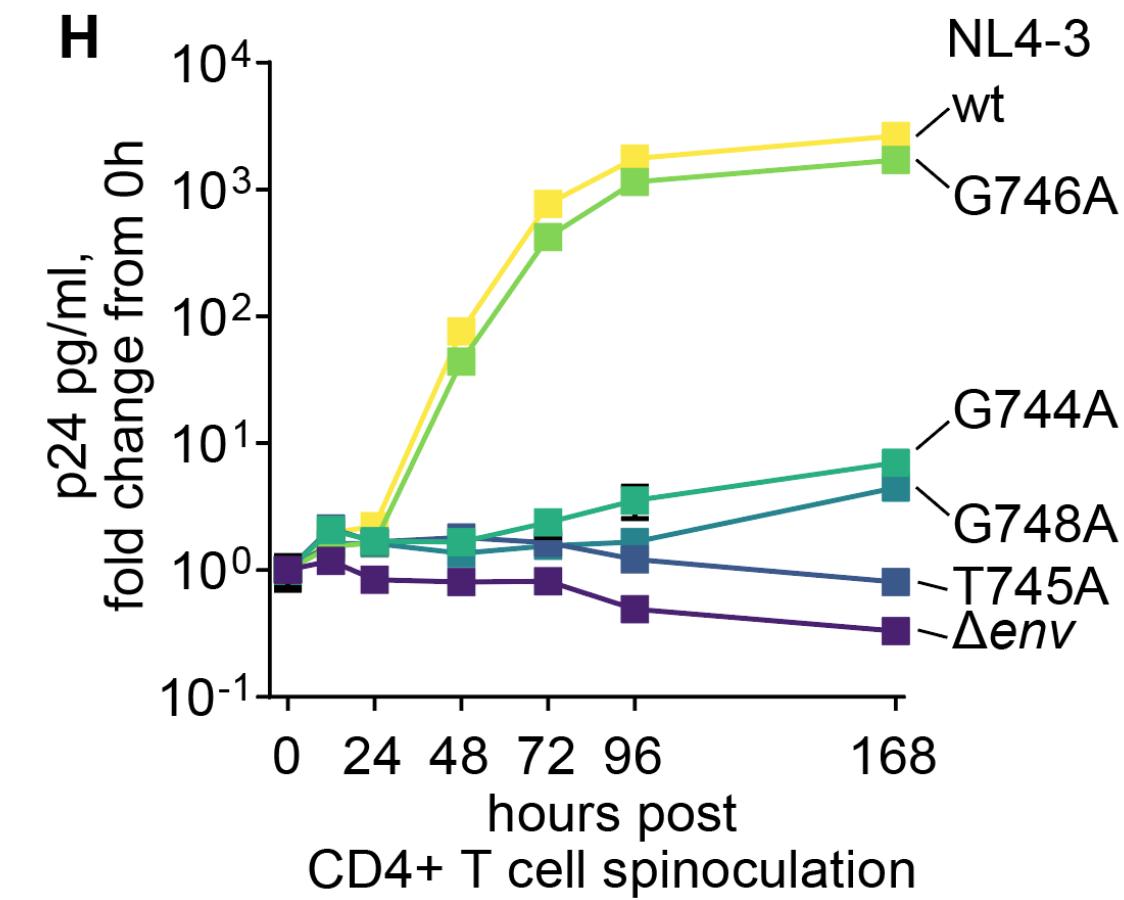
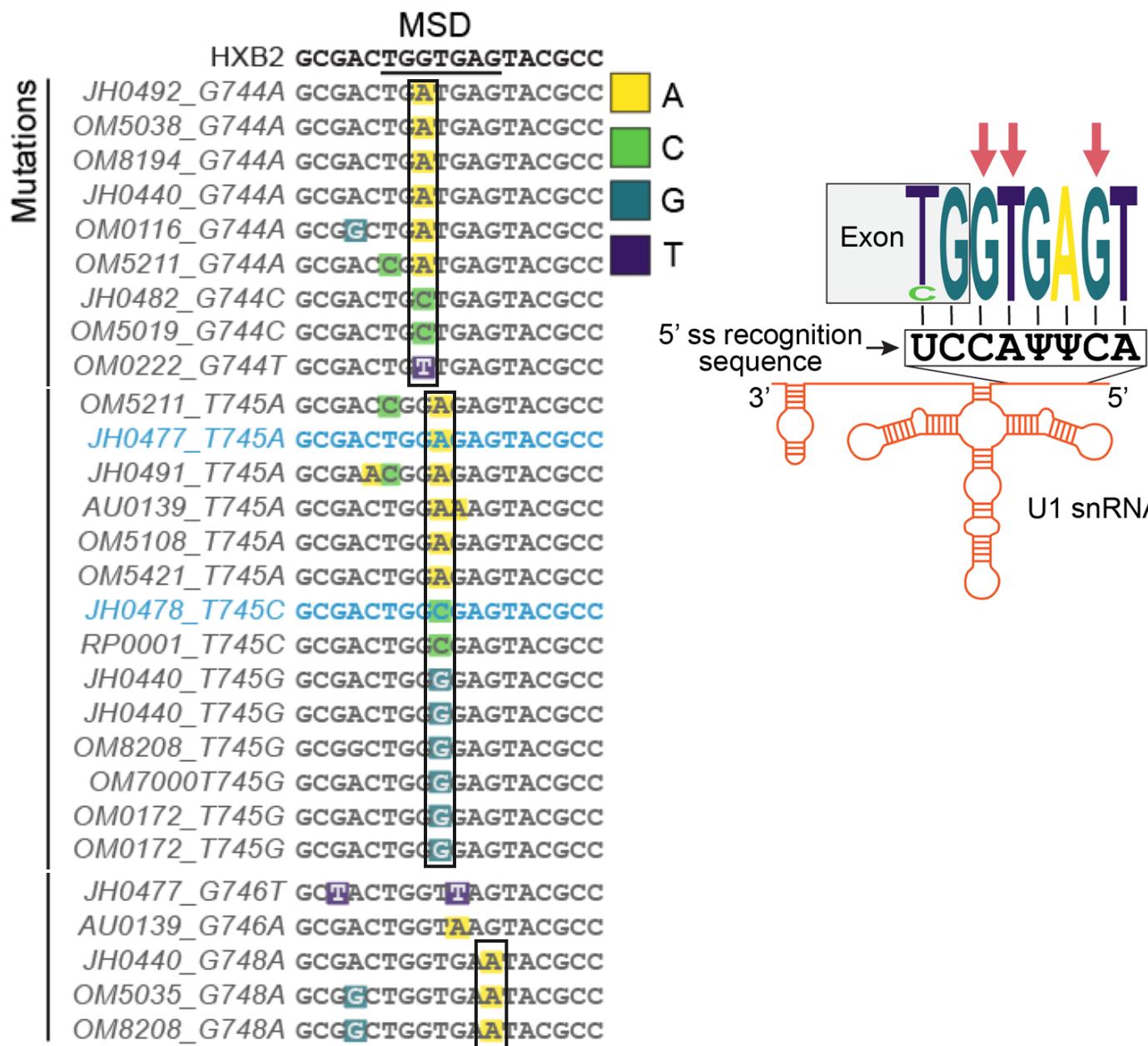
NSV dataset n= 94



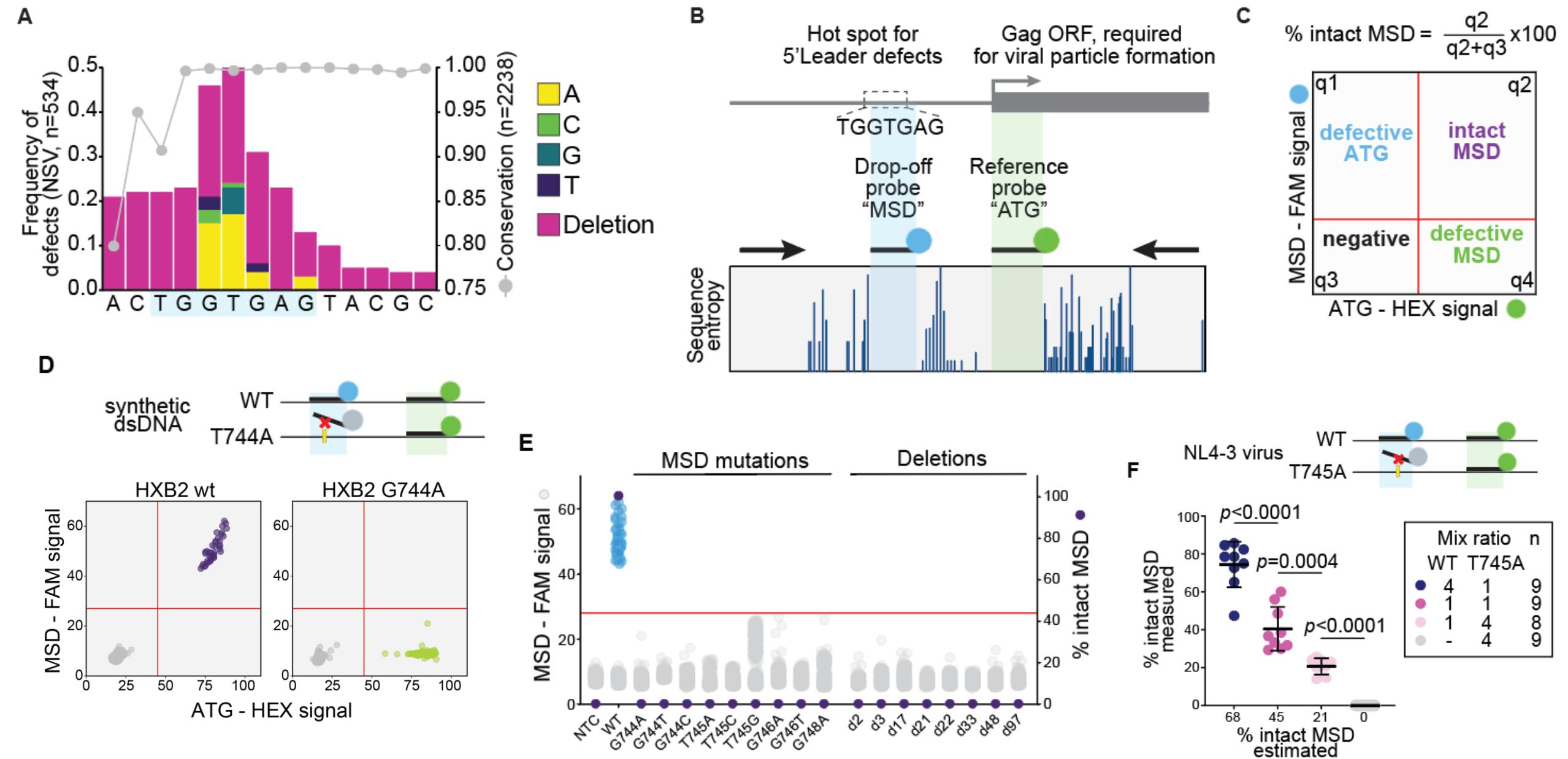
Mutations affect RNA binding to splicing machinery



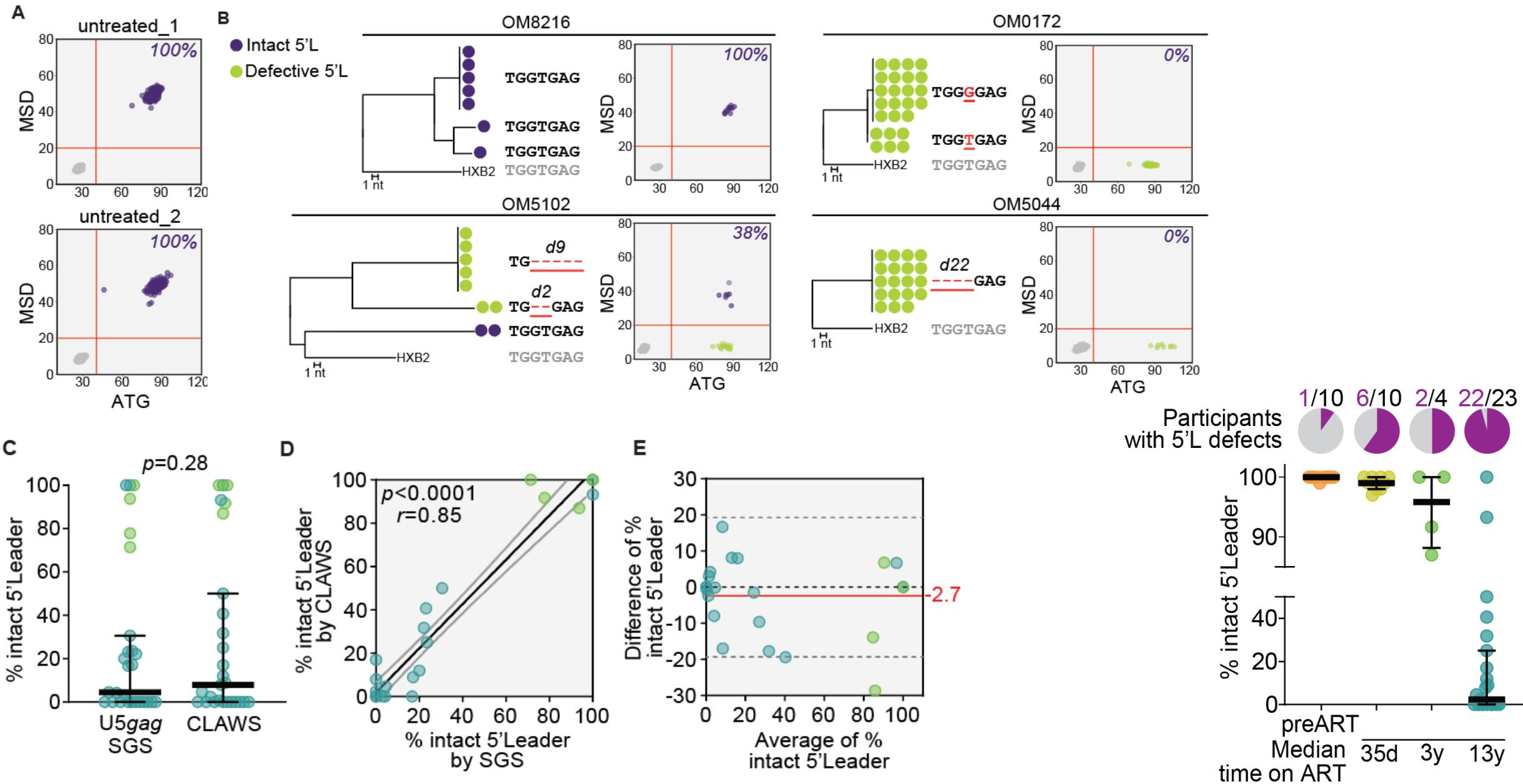
Mutations affect RNA binding to splicing machinery



Capturing 5'L Anomalies Without Sequencing



CLAWS strongly correlates with 5'L sequencing



Conclusions

The new onset of persistent viremia despite no issues in adherence and drug efficacy is driven by virus production from expanded infected clones, and not replication.

Proviruses with small defects in the 5'Leader region are a common cause detectable viremia, complicating ART management.

These defects result in non-infectious virus, due to inefficient HIV splicing

→ differential selection pressure compared to intact proviruses?

→ These results allow a ***better clinical interpretation of detectable viral load***

→ Sequencing data informed the design of a ***simpler, targeted assay*** that can distinguish intact from defective viral RNA

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