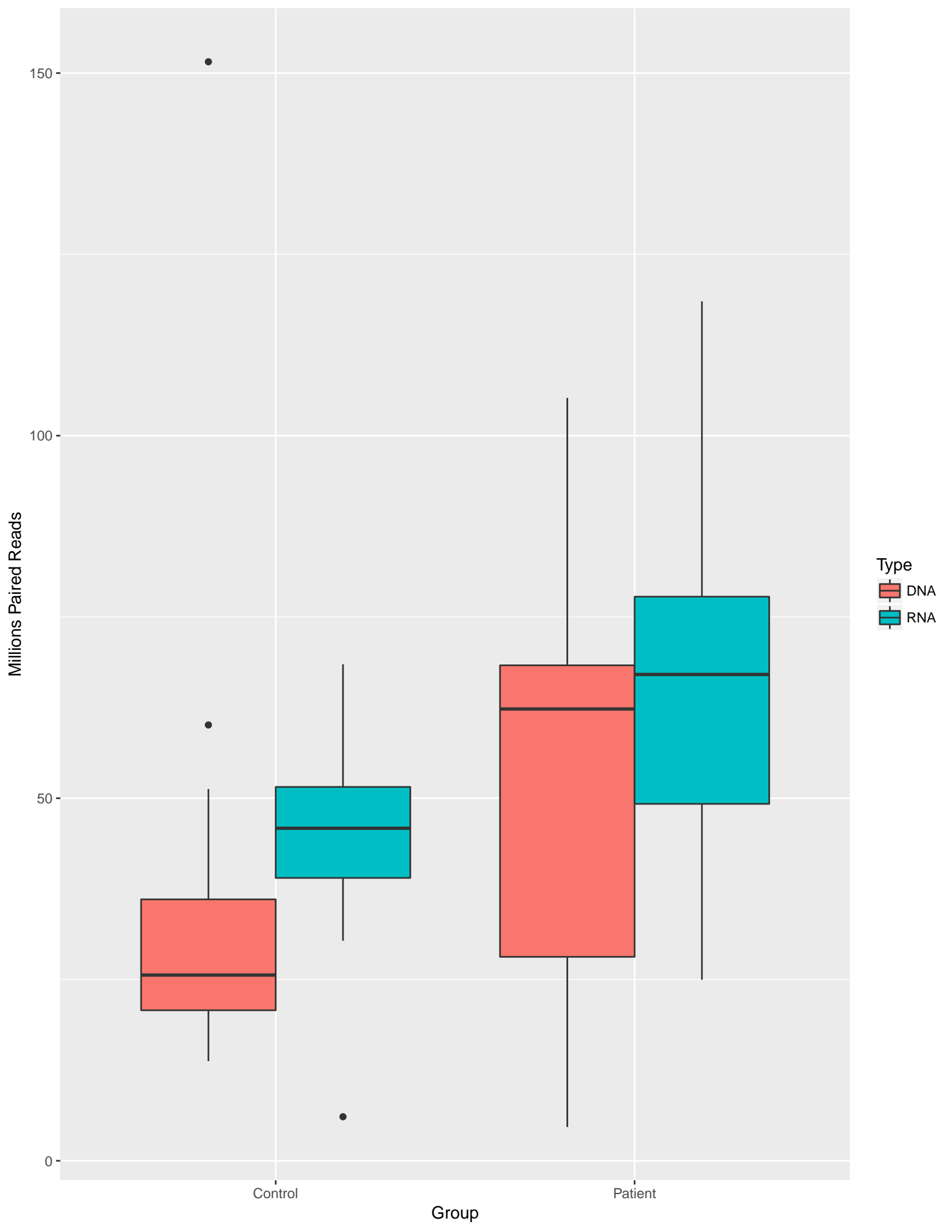


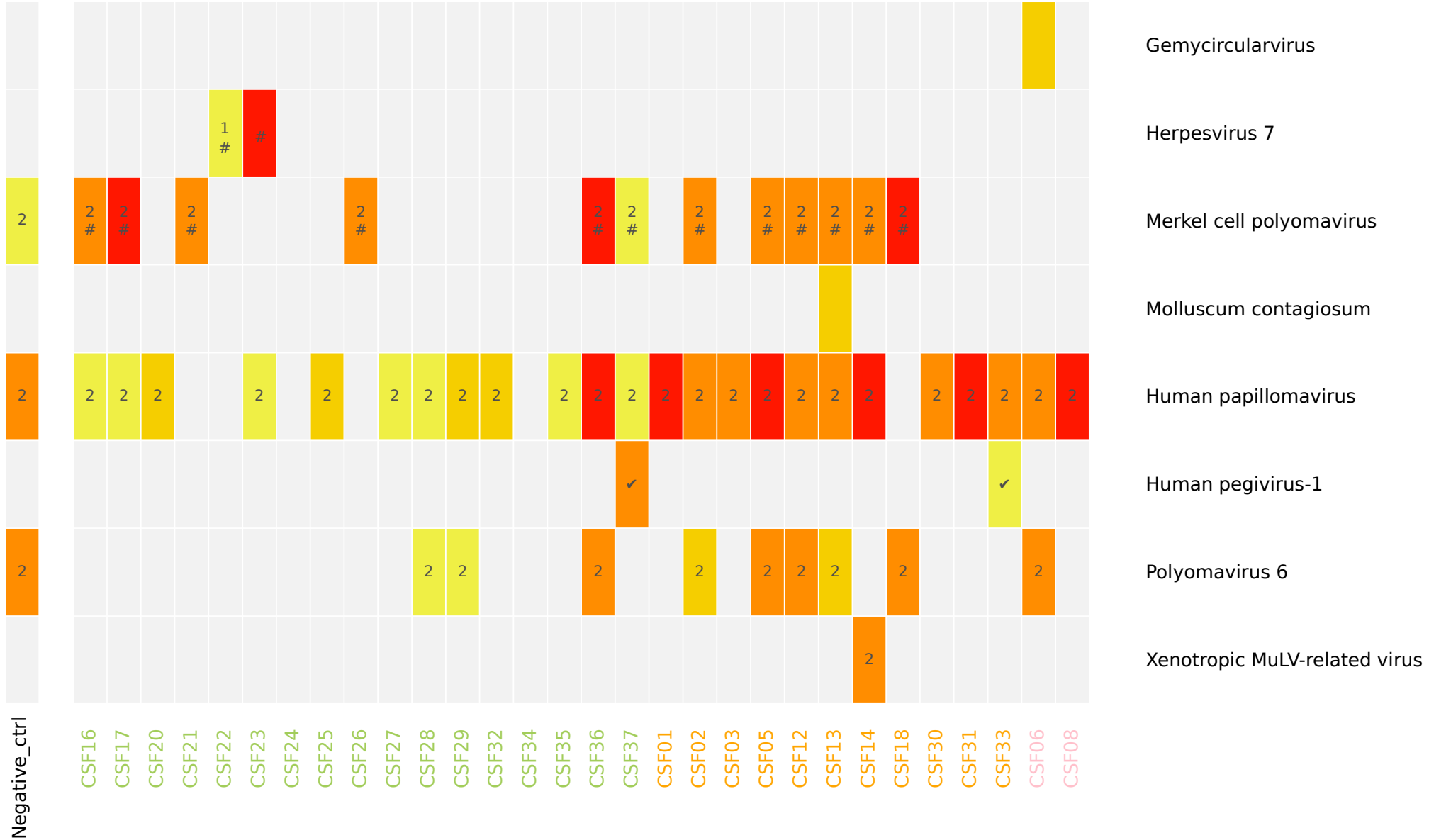
r(RT-)PCR assays	Viruses Detected	Sequence (5' to 3')			Final [uM] Fwd/Rev/ Probe	r(RT-) PCR annealing tempera ture °C	Reference
		Fwd primer	Probe	Rev Primer			
	Torque Teno virus	GTTTTTCYACGCCCGTCC	FAM - ACTCACCTHCGGCACCCGC - BHQ1	CCTTGACTYCGGTGTGTA A	0.9/0.9/0.2	60	Masouridi-Levrat S, et al. Bone Marrow Transplant. 2016;51:440-442.
	Adenovirus	GCCCCAGTGGTCTTACA TGCACATC	FAM - TGCACCAGACCCGGGCTCAGGTA CTCCG A -TAMRA	gvGCCACGGTGGGGTTTC TAAACTT	0.9/0.9/0.2	60	adapted from Verheyen J, et al. Appl Environ Microbiol 2009; 75: 2798-2801.
	Merkel cell polyomavirus	CCCTTTGGAGCAAATTC CA	FAM - CAAAATATCCACAAGCTCAGAAGTGA - TAMRA	CTGACCTCATCAAACATA GAGAA	0.6/0.6/0.2 5	55	Arvia R. et al. Journal of Virological Methods 246 (2017) 15-20
	Human pegivirus-1	GGCGACCGGCCAAAA	FAM - TGACCGGGATTTACGACCTACCAACCCT - TAMRA	CTTAAGACCCACCTATAG TGGTACC	0.9/0.9/0.2	55	Chivero ET et al. The Journal of general virology. 2014;95:1307-19.
	Epstein-Barr virus	CGGAAGCCCTCTGGACT TC	FAM - TGTACACGCACGAGAAATGCGCCT - TAMRA	CCCTGTTTATCCGATGGA ATG	0.2/0.2/0.1	60	Kimura H et al. J Clin Microbiol 1999: 37:132-6
	Human herpes virus 7	Genesig™ Detection Kit, Human Herpesvirus 7			unknown	60	PrimerDesign, England, United Kingdom

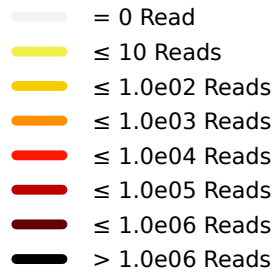


- Groups :
- █ Elective cesarean section
 - █ Elective urologic or digestive surgery
 - █ Elective orthopedic surgery

- = 0 Read
- ≤ 10 Reads
- ≤ 1.0e02 Reads
- ≤ 1.0e03 Reads
- ≤ 1.0e04 Reads
- ≤ 1.0e05 Reads
- ≤ 1.0e06 Reads
- > 1.0e06 Reads

- 1 : Most likely due to index hopping
- 2 : Probable contaminations
- ✓ : Positive by r(RT-)PCR
- # : Undetected by r(RT-)PCR





2 : Probable contaminations
 ✓ : Positive by r(RT-)PCR
 # : Undetected by r(RT-)PCR
 ⌘ : Insufficient volume to be tested by r(RT-)PCR

