

# Human Parechovirus A conventional nested genotyping RT-PCR ["Harvala assay"; 2008-2015] Version 2

Ian Mackay

## Abstract

I and my team used this assay between 2008-2015; we dubbed it the "Harvala assay". It produces an amplicon that includes the 3' end of VP3 and the 5' end of VP1, spanning the junction.

*In silico* sequence alignments indicated the highly degenerate oligonucleotides could theoretically detect at least HPeV 1-7, 17 and 18.

**Citation:** Ian Mackay Human Parechovirus A conventional nested genotyping RT-PCR ["Harvala assay"; 2008-2015]. **protocols.io**

[dx.doi.org/10.17504/protocols.io.krrcv56](https://dx.doi.org/10.17504/protocols.io.krrcv56)

**Published:** 11 Nov 2017

## Guidelines

- This protocol assumes the user is familiar with working in a laboratory, with PCR, the thermocycler and software used to run it
- This protocol should be re-evaluated if being used with different reagents, if the oligonucleotide sequences are changed or if the cycling conditions are altered

## Protocol

### Oligonucleotides...

#### Step 1.

Round	Name	5'-3' oligonucleotide sequence
Round 1. RT-PCR	HPeV_VP3/1_OS	GAYAATGCTATMTAYCAWATYTGTA
Round 1. RT-PCR	HPeV_VP3/1_OAS	ACWGTRAARATRTCHACATTSATDG
Round 2. nPCR	HPeV_VP3/1_IS	TTYTCMACTGGATGMGGAARAC
Round 2. nPCR	HPeV_VP3/1_IAS	DGGYCCATCATCYTGWGCTGA

OS-outer sense; OAS-outer antisense; IS-inner sense; IAS-inner antisense

## Reagents...

### Step 2.



#### REAGENTS

SensiFAST Probe no ROX one-step kit BIO-76005 by [Bioline](#)

MyTaq HS DNA Polymerase BIO-21113 by [Bioline](#)

## Reaction setup...

### Step 3.

Below is the reaction setup for a single RT-PCR reaction.

Ideally, this work is conducted in a laboratory separate to any space used to *perform* PCR, molecular cloning or the analysis of high concentration DNA.

This volume has been used in 0.1-0.2ml tubes or various other connected tube configurations such as 100-place rings.

Multiply this according to the number of reactions you will need, remembering to include a positive control and at least two non-template controls (NTCs)

You may also need to allow some extra volume, depending on the method used to pipette mix into tubes for the run. For example, some robot-loaded tubes can require two reaction 'dead volumes'.

### Round 1: RT-PCR

<b>Reagent (stock concentration)</b>	<b>Vol (μL) / reaction</b>	<b>Final concentration</b>
Nuclease free water	7.28	
AN345_panHPEV/LV (200pmol/ul [200uM])	0.06	600nM
AN344_panHPEV/LV (200pmol/ul [200uM])	0.06	600nM
SensiFast OneStep Mix(2x)	10	1X
RNase inhibitor	0.4	
RT/Taq (6U/mL)	0.2	1X
Template extract RNA	2	
Final volume	20μl	

### Round 2: nPCR

<b>Reagent (stock concentration)</b>	<b>Vol (<math>\mu</math>L) / reaction</b>	<b>Final concentration</b>
Nuclease free water	12.424	
AN345_panHPeV/LV (200pmol/ul [200uM])	0.038	380nM
AN344_panHPeV/LV (200pmol/ul [200uM])	0.038	380nM
MyTaq Reaction Buffer (5X)	4	1X
MgCl <sub>2</sub> (25mM)	1.4	
MyTaq HS DNA Polymerase (5U/uL)	0.1	1X
Round 1 amplicon	2	
Final volume	20 $\mu$ l	

#### Cycling conditions...

##### Step 4.

This assay has been optimized and validated for use with a RotorGene 6000 or RotorGene Q thermal cyclers.

The cycling conditions for the one-step RT-PCR and the nested PCR (nPCR) are as follow:

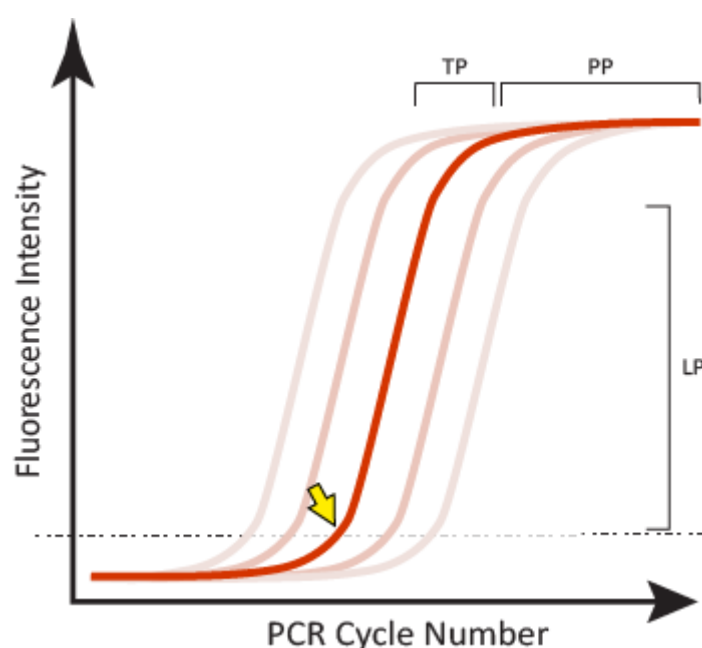
<b>Round 1: RT-PCR</b>			
45°C	20min		
94°C	2min		
95°C	30s		
60°C	30s		40X
72°C	105s		
72°C	7min		
4°C	$\infty$		

<b>Round 2: nPCR</b>			
94°C	1min		
94°C	30s		
50°C	30s		40X
72°C	105s		
72°C	7min		
4°C	$\infty$		

**Step 5.**

The definition used for a satisfactory positive result from a real-time fluorogenic PCR should include each of the following:

1. A **sigmoidal curve** – the trace travels horizontally, curves upward, continues in an exponential rise, curves and reaches a horizontal plateau phase
2. A **suitable level of fluorescence** intensity as measured in comparison to a positive control (y-axis)
3. A defined threshold ( $C_T$ ) value which the fluorescent curve has clearly exceeded (Fig.1 arrow) and which sits early in the log-linear phase
4. A flat or non-sigmoidal curve or a curve that crosses the threshold with a  $C_T$  value  $>40$  cycles is considered a negative result
5. No template controls (NTCs; water instead of specimen extract) should not produce a curve



**Figure 1.** Examples of satisfactory sigmoidal amplification curve shape when considering an assay's fluorescent signal output. The crossing point or threshold cycle ( $C_T$ ) is indicated (yellow arrow); it is the value at which fluorescence levels surpass a predefined (usually set during validation, or arbitrary) threshold level as shown in this normalized linear scale depiction. LP-log-linear phase of signal generated during the exponential part of the PCR amplification; TP-a slowing of the amplification and accompanying fluorescence signal marks the transition phase; PP-the plateau phase is reached when there is little or no increase in fluorescent signal despite continued cycling.