Available samples
N=725

Excluded samples
N=127
The 127 samples from the implementation study in Brazil were excluded for two reasons: 1, they were not genotyped and 2, they were processed by a different laboratory than the rest of samples (results from this sub-study are summarized separately in the manuscript)

Index test
N=598

Positive result n=582

Reference standard positive n=566

Reference standard negative n=16

Negative result n=16

Reference standard positive n=2

Reference standard negative n=14

Target condition (HCV infection) present in all samples

*In addition to clinical sensitivity we determined analytical sensitivities (limits of detection) for every genotype separately. This allows for a more general comparison of sensitivity with existing assays.*