

Norovirus GII isolate NORO_200_25_11_2015, complete genome

GenBank: MH218661.1

[FASTA Graphics](#)

[Go to:](#)

LOCUS MH218661 7538 bp RNA linear VRL 11-JUN-2018

DEFINITION Norovirus GII isolate NORO_200_25_11_2015, complete genome.

ACCESSION MH218661

VERSION MH218661.1

KEYWORDS .

SOURCE Norovirus GII

ORGANISM [Norovirus GII](#)

Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes;
Picornavirales; Caliciviridae; Norovirus.

REFERENCE 1 (bases 1 to 7538)

AUTHORS Brown,J.R., Roy,S., Shah,D., Williams,C.A., Williams,R., Dunn,H.,
Hartley,J., Harris,K. and Breuer,J.

TITLE Norovirus transmission dynamics in a paediatric hospital using full
genome sequences

JOURNAL Clin. Infect. Dis. (2018) In press

PUBMED [29800111](#)

REMARK Publication Status: Available-Online prior to print

REFERENCE 2 (bases 1 to 7538)

AUTHORS Brown,J.R., Roy,S., Shah,D., Williams,C.A., Williams,R., Dunn,H.,
Hartley,J., Harris,K. and Breuer,J.

TITLE Direct Submission

JOURNAL Submitted (17-APR-2018) Division of Infection and Immunity,
University College London, 90 Gower St, London, London WC1E 6BT,

United KingdomCOMMENT ##Assembly-Data-START##

Assembly Method :: CLC Genomics Workbench v. 10.1

Sequencing Technology :: Illumina

##Assembly-Data-END##FEATURES Location/Qualifiers

source 1..7538

/organism="Norovirus GII"

```
/mol_type="genomic RNA"
/isolate="NORO_200_25_11_2015"
/host="Homo sapiens"
/db_xref="taxon:122929"
/country="United Kingdom"
/collection_date="25-Nov-2015"
/note="genotype: GII.P7_GII.7"
gene      5..5098
          /gene="ORF1"
CDS      5..5098
          /gene="ORF1"
          /codon_start=1
          /product="nonstructural polyprotein"
          /protein_id="AWR17643.1"
          /translation="MKMASNDATAAFGSQNSVNDSNNATPSNKEEVGAFSNIKVGFKK
MLGAVPKGTKAPSSNQNCPTVKIGPKTLTIPPEPPNGEDTVQFDAKSETIIGLPDLTT
VQNEHENNPYTPPPLSEREYRPATEPLPGTILEMWDGEFYHYSVYVSDGKALGVHKPP
AAISLATIELTPISLYWRPVYTPNYLVSPDTLKGLAGEKFPYAFSNNCYNFCCWVLD
LNDTWLNRRCISRTTGFFKPYQSWNRKPLPTVDDGKIKKVANAILCALGSLFSKPIKD
LLGKLKPLNLLHLLASCDWTFAGIVETVILMAELFNVFWTPPDVSNFIASLIGDFELQ
GPEDLAVELVPVVMGGIGMVLGFTAEEKIGRMLSSAASTLRACKDLGNYALDILKLMK
WFFPKKEEKAEMETLRAIEDAVLDMEAIGNNHLTTLLKDKDSLTA FMKTL DLEEEKAR
KLSTKSSSPDIVGTINAILARIAAARSL LHKAKEEMFSRIRPVVVMISGRPGIGKTHM
ARHLAKSIANTMSGDQRVGLVPRNGVDHWDAYRGERVVLWDDYGMGSPVKDALTLQEL
ADTCPVTLNCDRIENKGKMFDSDVIIITTNLVNPAPLDYVNF EACSRRVDFLVYAESP
EIEKVKRDFPGQPDMMWKDHF KPDFS HIKLTLAPQGGFDKNGNTPHGKGTMRSLTQGSL
TARVAGLVHERRDEFQLQGNDLQTYNFD TNRVSAFRKLAADNKYGIMETMRVGAALKS
VKTLEDLKAALRDVKFNECEIIYKNSKYRVSSNGKGSVSDKVEDNASQTVNEVHSAL
LRLRQARARYYISCFQDLVYTLMQVAGASFVVSRI SRFCWERWVKPTETQEVNEPEK
EVAQGGWEIEPKDTEPEGKKGKNKKGRGKKHTAFSSKGLSDEEYDEFKRIKEERNKY
SIEEYLQDRDRYEEVAVARATEEDFCEEEEA KIRQRI FRPTRKQRKEERGVLGLVTG
SDIRKRRPDDFQPKGNLWADDTRSVDYNERLD FEAPPSVWSRIVPLGTGWGFWVSSNL
LITTHVLPKGIKELFGVEIKQIQVHKSGEFCRFRFPPIRPDVTGLVLEEGAPEGTV
```

CSVLVKRPTGEMIPLAVRMGTHASMKIQGRTVGGQMGMILLTGANAKNMDLGTGPGDCG
CPYIYKRGNDIVVAGVHTAAARGGNTVICATQGQDGEAVLEGNEGLGTYCGAPILGPG
KAPKLSTKTKFWRSSPDALPPGTYEPAYLGGKDPVEKGPSLQQVMRDQLRPFTEPRG
KPPRPVLEEAKKTVMNVLEQTIDPAKPWSYSQACASLDKTTSSGSPHHVKKNDHWNG
ESFTGFLADQASKANLMEYQAKHVQPVYTAALKDELVKTDKIYKKIKRLLWGSDLGT
MIRCARAFGGLMDSMKASCIALPCRVMNMNEDGPIIFDKHSKYRYHYDADYSRW DST
QQRSILSAAMEVMVRFSAEPELAQVVAEDLLAPSQLDVGDFVISVQEGLPSPGVPCTSQ
WNSIAHWILTLSAMAESVGLSPDVVQAHSCFSFYGDDEIVSTDINLDPMKLTQKLREY
GLVPTRPDKTEGPLVITEDLTGLTFLRRSIARDPAGWFGKLDQDSILRQLYWTRGPNH
ENPYESMVPHSQRATQLMALLGEASLHGPQFYKKVSKMVINEIKSGGLEFYVPRQEAM
FRWMRFSDLSTWEGDRNLAPEGVNEDGVE"

gene

5079..6701

/gene="ORF2"

CDS

5079..6701

/gene="ORF2"

/codon_start=1

/product="VP1"

/protein_id="[AWR17641.1](#)"

/translation="MKMASNDAAPSNDGAAGLVPEINNEVMPLPVPAGASLATPVVGQ
QNIIDPWIRNNFVQAPAGEFTVSPRNSPGEILLDLELGPENPYLAHLARMYNGHAGG
MEVQIVLAGNAFTAGKIIFAAIPPGFPYENLSPSQITMCPHVIIDVRQLEPVLLPMPD
IRNNFFHYNQGNPKLRLVAMLYTPLRANNSGDDVFTVSCRVLTKPSPDFEFTFLVPP
TVESKTKQFTLPILKISEMTNSRFPVPVEMMYTARNENQVVQPQNGRVTLDGELLGTT
PLLAVNICKFKGEVIAKNGDVRSYRMDMEITNTDGTPIIDPTEDTPGPIGSPDFQGILF
GVASQRNKSEQNPATRAHEANINTGGDQYAPKLAQVKFFSESQDFEVHQPTVFTPVGV
AGDPSHPFRQWVLPTRYGGHLTNTHLAPAVAPLFPGEQILFFRSQIPSSGGHELGYMD
CLVPQEWVQHIFYQEAATAQSEVALIRFINPDTGRVLFKAKLHKQGFTITVAHTGDNPIV
MPPNGYFRFEAWVNQFYSLAPVGTGNGRRRIQ"

gene

6701..7474

/gene="ORF3"

CDS

6701..7474

/gene="ORF3"

/codon_start=1

/product="VP2"
/protein_id="AWR17642.1"
/translation="MASAFFANLAGGVASNGISSLINAGANAINQKVEFDYNKQLQNA
SFSDKEMLQAQIEATKRLQQEMIDIKQGVLTAAAGFSPTDAARGSVGAPMTKLVDWSG
TRYWAPGAMRTTGYSGTFTHLPPMQSASYVNNQNRQLFKVDPSPSSSSGSSSVSVKTQ
STALSGTSRVDSSSQSTLSTRPSSRTSDWVRSQNERLSPFMNGALQTAYVTPSSRAS
SVSTVSTVPKEVLDSWTPMFNTHRQPLFAHLRRRGESQI"ORIGIN

1 gtgaatgaag atggcgtcta acgacgctac cgctgccttt ggcagtcaaa actctgtcaa
61 tgacagtaac aacgccaccc cttctaacaa agaagagggt ggtgcattct ccaacattaa
121 agttggattt aagaaaatgc tgggcgccgt acccaaaggt accaaagcgc ccagcagtaa
181 tcaaaattgc cccacagtta aaatcgggcc taaaacgtta acaattcctc ctgagcccc
241 aaatggtgaa gacaccgtac aatttgatgc aaagtcagaa actataattg ggctaccaga
301 cttgacaacg gtgcaaaacg aacacgaaaa caatccatac actgtccctc cattaagtga
361 gagggagtac aggccggcca ctgaaccact tcctggcaca atactggaga tgtgggatgg
421 tgagttttac cattactccg tgtacgtcag tgatggcaaa gctctggggg tccacaaacc
481 accggcgggt ataagcctcg cgacgataga gctcaccccc atatcactct actggaggcc
541 cgtttacacc cccaactact tggctctctc agacacatta aagggcctcg ctgggtagaa
601 gttcccctac acggccttca gtaataactg ttacaacttc tgctgttggg tgcttgatct
661 caacgacacg tggcttaata gaagatgcat ttccagaact accggcttct tcaaaccata
721 ccagtcttgg aatagaaaac ctctcccaac tgttgatgat ggaaagatca aaaagggtggc
781 caatgcaatc ctctgtgcac tcggttcgct gttttcaaaa ccaatcaaag acctattagg
841 caaactcaag ccattgaatc tactacactt gctcgcgtcc tgtgattgga cgtttgacgg
901 catagtggag acagtcattc taatggcgga acttttcaac gtattctgga cccaccaga
961 tgtctctaata tttatagcct ccctgatagg tgattttgaa ttgcaagggc ctgaagacct
1021 ggctgtagag ctcgccccg tggttatggg tggcataggg atggttctcg gcttcaccgc
1081 tgagaagata ggccgcatgc tgctgtccgc tgcatacaac ctgcgggcgt gcaaagacct
1141 agggaaactac gcccttgaca tactcaaatt ggtcatgaaa tggttcttcc caaagaagga
1201 agagaaggct gagatggaaa ccttgagggc gattgaagat gccgtccttg atatggaagc
1261 tatagggaac aaccacctca cgacccttct caaggacaaa gacagcctca cagctttcat
1321 gaagaccttg gatctagaag aggagaaggc gagaaaactg tccactaaat catcttcacc
1381 ggacatagtc ggcaccatca acgcatatt agctaggata gctgctgcta gatccctact
1441 acataaggcc aaagaggaga tgttcagtag gatcagacca gtggtcgtca tgatctcagg
1501 cagacctggc attggaaaaa cccacatggc gaggcacttg gctaagagca tcgccaatc

1561 catgagtggc gatcagagag tcggactcgt tccacgcaac ggtgtcgatc actgggacgc
1621 ctacagggga gagaggggtg tcttgtggga tgactacggc atggggagcc ctgttaaaga
1681 cgccctgaca ctgcaagagt tggccgacac atgtccagtc actttaaatt gtgataggat
1741 tgagaataag ggaaagatgt ttgatagcga tgtcatcata atcacgacca acctagtcaa
1801 ccccgacccc ctcgactatg tgaacttcga ggcgtgttct aggagagttg acttcctggt
1861 ctatgcagag tcaccagaaa ttgaaaaggt caagagagac ttccctggcc agcctgacat
1921 gtggaaagac cattttaaac cagacttttc ccatattaag ttgactctag cccccaggg
1981 tgggtttgat aagaatggca atacccaca cggcaagggt accatgcggt ccttaacca
2041 ggggtcccta actgcgagag tcgcaggcct tgtccatgag aggagggacg aattccagct
2101 ccaaggaaat gacctccaaa catacaactt tgacaccaac agagtctccg catttagaaa
2161 attggctgca gataacaagt atgggatcat ggagacgatg agagttggtg cagcgctcaa
2221 gagtgtgaag accttggaag atcttaaagc tgcattgagg gatgtgaaat tcaatgagtg
2281 tgaaataatt tacaaaaact ctaagtaccg agtctcttcc aacggcaaa gctcggtttc
2341 tgttgacaaa gttgaagaca atgcttccca gactgtgaac gaggtgcatt cagcacttct
2401 tagactcagg caggcaaggg ccagatacta catcagctgc ttccaagacc tcgtttacac
2461 tctcatgcag gttgcagggg catcattcgt cgtcagtagg atctccagaa ggttctgctg
2521 ggaaaggtgg gtcaaaccga ccgagaccca ggaagtgaat gagcccgaaa aggaagtggc
2581 ccaagggtga tgggaaattg agcccaagga cacagaacca gaaggcaaga agggcaagaa
2641 caagaaagga aggggcaaga aacacacagc cttctctagc aaaggcctga gtgatgaaga
2701 gtatgatgag ttcaagagaa tcaaggaaga aaggaacgga aagtactcca ttgaggaata
2761 cctgcaggac cgagaccgct attatgaaga ggtggcagtt gctcgggcaa cagaggagga
2821 cttctgcgaa gaggaggaag ccaagataag acaaagaatt ttccgtccga caaggaaaca
2881 gaggaaggag gaaaggggcg tgcttggtt ggtcactggc tcagacatca gaaagaggag
2941 acccgatgac ttccaaccaa agggcaacct gtgggcagat gacaccagaa gtgtggacta
3001 caatgagagg cttgattttg aagcacctcc gagtgtttgg tcaagaatag tccattagg
3061 cactggttgg gggttttggg tttcatccaa ctttctgac acaacaacac acgtcctgcc
3121 taaagggatc aaggaactct ttggagtga aatcaagcaa atccaagttc ataagtctgg
3181 agagttttgc aggttttaggt ttccaagacc catcaggcca gatgtcacag gactcgtgtt
3241 ggaggaaggc gccccagagg gcaactgtctg ttctgtactc gtaaaaagac ccacaggtga
3301 aatgatcccc cttgcagtga ggatgggcac gcacgcattc atgaaaatac agggcaggac
3361 cgttgggtgg caaatgggga tgctcctcac aggagcaaac gcgaagaaca tggatctcgg
3421 tactggctct ggtgactgcg gttgcccta catctacaaa cgtggcaacg acatagtgtg
3481 cgcgggtggt cacaccgcag cagcccgggg gggcaacact gttatatgcg ccaccaagg

3541 gcaagacggg gaagcagtc ttgagggaaa tgaggacctt ggcacctact gtggcgcccc
 3601 aattctgggt cctggcaagg cgcccaaact cagcacgaag accaagttct ggcgctcatc
 3661 accagatgcc ttgccacctg gcacatacga acctgcttac ctaggaggca aagaccctag
 3721 ggtggaaaaa ggtccctccc tgcagcaagt catgagagac cagttgagac cttcacaga
 3781 acccaggggc aaaccgccc gacctgcagt cctagaggaa gccaaaaaga cagtgatgaa
 3841 tgttttagaa caaaccattg accccgcaa gccatggctc tactcccagg catgtgcctc
 3901 acttgacaaa accacctcta gtggtagtcc ccaccacgtt aagaaaaatg accactggaa
 3961 tggggagtcc ttcactggtc cccttgcala tcaggcatct aaagccaacc tcatgtatga
 4021 acaggccaaa catgtgcagc ccgtgtatac ggcagcgctt aaagatgagc ttgttaagac
 4081 tgacaagatc tacaagaaga taaagaagag gtcctatgg gggtcggacc ttggcacaat
 4141 gatcagggtc gccagggctt ttggtggtct catggacagc atgaaggcaa gttgtatagc
 4201 tcttccttgc aggggtggaa tgaacatgaa tgaagatggc cccatcatat ttgacaaaca
 4261 ctctaagtac aggtaccact atgatgctga ctattctagg tgggactcaa cccaacagag
 4321 gagcattctc tctgccgcta tggaagtaat ggtgcggtt tctgctgaac cagaactggc
 4381 acaagtgggt gcagaggacc ttttggtccc tagccagctt gacgttggcg actttgtcat
 4441 ctcagtccag gaggtctac catcaggagt cccatgcaca tcacaatgga attcaatagc
 4501 gcaactggatc ctgactttga gtgcaatggc agaagtgtca ggtctctcac cagatgttgt
 4561 ccaagctcac tctgctttt cattttacgg ggacgatgag atcgtcagta ctgatatcaa
 4621 tcttgacccc atgaagttga cacagaaact cagggagtat ggtctggtcc ccaccggcc
 4681 tgataaaact gaaggccccc tcgttataac tgaagacctc accggcctga cattcctccg
 4741 taggtcgatt gtcggggacc cagctggttg gtttggaata ctggaccaag actcaattct
 4801 cagacagcta tactggacaa gggggcctaa ccatgagaac ccgtatgaga gcatgggtcc
 4861 tcattctcag cgggccacac agctcatggc cttcttgggt gaggcctcgc tgcatggctc
 4921 ccagttttac aagaaagtta gcaagatggt catcaatgag attaagagcg gtggtctgga
 4981 gttttatgtg ccagacaag aggccatgtt caggtggatg agattctctg acctcagcac
 5041 atgggagggc gatcgcaatc ttgctcccga aggtgtgaat gaagatggcg tcgaatgacg
 5101 cagctccatc taatgatggt gcagcaggcc tcgtgccaga gatcaacaat gaggtcatgc
 5161 cccttgagcc cgtggctggt gcatcgctgg cgacaccagt tgcggggcaa caaaatataa
 5221 ttgatccctg gataagaaat aattttgtac aagctcctgc aggtgagttt acagtttccc
 5281 ccagaaattc ccctggtgaa attttacttg atttggagtt gggaccagag ttgaaccct
 5341 atctcgctca tttggccgt atgtacaatg ggcacgctgg tggcatggaa gtgcaaattg
 5401 tgttggtggt gaacgcgttc acagctggca agatcatatt tgctgccatc cctccagggt
 5461 tccatacga aaatttgtca cttcacaaa ttacaatgtg ccccatgtt ataattgatg

5521 ttagacaatt agaaccagtc ctattaccta tgccagatat tagaaataat ttcttccatt
5581 acaatcaggg caatgatcca aaattgaggt tggtagctat gctctacact cctttaaggg
5641 ctaacaattc tggatgatgat gtgttcacag tttctttagt ggttctcact aaaccgtcac
5701 ctgattttga attcacattt ttggttcccc ctacagttga atccaagacc aagcaattca
5761 cattgcccatt cctcaaaaata tcagagatga ccaactcaag attcccagtg ccggtggaaa
5821 tgatgtacac agcaaggaat gagaatcaag ttgtccagcc acagaatggg agggtcacac
5881 tcgatggcga actattgggc accacaccct tgttggctgt caacatctgt aaattcaagg
5941 gagaggatcat agctaagaat ggagatgtga gatcctatag gatggacatg gaaatcacca
6001 aactgatgg caccccaatt gaccctactg aagatacacc tgggtccatt ggatcacctg
6061 attttcaagg tatactcttt ggcgttgcca gtcaacgcaa caagagttag caaaatcctg
6121 ccacgagagc ccacgaagcc aacatcaata ctggtgggga ccaatatgcc cccaaattgg
6181 cccaagttaa attcttctca gaggccaggt atttcgaggt gcatcagccc acagttttca
6241 cccagtttgg tgtggcaggt gacccttccc accctttcag acagtgggtc ctgcctcgat
6301 atggtggaca tctcaccaac aatacccact tggctccagc agtggcccct ctattcccag
6361 gtgagcagat tctgttcttt agatcacaaa tacctagttc aggcggccat gaattgggtt
6421 acatggactg ccttgtccca caagaatggg tocaacactt ctatcaagaa gctgccacag
6481 ctcagtctga ggtggcactc ataaggttca tcaaccaga cactggtaga gtcctcttgc
6541 aggccaagct acacaagcaa ggctttataa ctgtggctca cactgggtgac aaccgaattg
6601 ttatgccacc aaatgggtat ttcagatttg aagcctgggt caatcagttt tactcactcg
6661 ccccggtggg aactgggaat gggcgtagga ggatccaata atggctagtg ccttctttgc
6721 aaatttggca gggggcggtg cctccaatgg aattagttca ctcataaatg ctggggctaa
6781 tgctattaac caaaaggttg agtttgatta taataaaca ctccaaaatg catccttctc
6841 ccacgataag gaaatgctcc aagcccaat tgaagccact aaacgcttgc aacaagagat
6901 gattgacata aaacaagggg tgctgacggc tgcgggattc tctccacgg atgccgagag
6961 gggttctgtc ggcgcgcaa tgacaaaatt agttgattgg agtggcacta gatattgggc
7021 tcttgggtgc atgaggacaa caggctactc agggaccttc actcacttgc ctctatgca
7081 atctgcttca tatgttaata atcagaatag acaacttttt aaagtggatc cttctccctc
7141 ttcttcaagt ggttcttctg tttcttctgt taaaacacaa tcaactgtc tctcaggcac
7201 ttcaaggggt gactcttcat ctacgtcaac cctgtcgacc aggcgagct caaggacatc
7261 tgattgggtg cgtagccaaa atgaaagatt gagccccctt atgaatggag cattgcaaac
7321 agcgtatgta acgccacat ctagcagggc atcaagtgtc agcacagtct cgaccgtgcc
7381 gaaagaggtt ttggactcct ggacgcctat gtttaacact cacaggcagc cgctcttcgc
7441 ccacctgcgt aggcgagggg agtcacagat ttagtgaaaa gattaattag gaaagtttga

7501 ttagttttct ttctattat tctttaatt ttctttt

//