**Optimal dose and safety of molnupiravir in patients with early SARS-CoV-2: a phase 1, open-label, dose-escalating, randomised controlled study**

**Short Title: Optimal Dose and Safety of Molnupiravir**

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**Synopsis**

**Objectives**

AGILE is a phase Ib/IIa platform for rapidly evaluating COVID-19 treatments. In this trial (NCT04746183) we evaluated the safety and optimal dose of molnupiravir in participants with early symptomatic infection.

**Methods**

We undertook a dose-escalating, open-label, randomised-controlled (standard-of-care) Bayesian adaptive phase I trial at the Royal Liverpool and Broadgreen Clinical Research Facility. Participants (adult outpatients with PCR-confirmed SARS-CoV-2 infection within 5 days of symptom onset) were randomised 2:1 in groups of 6 participants to 300mg, 600mg and 800mg doses of molnupiravir orally, twice daily for 5 days or control. A dose was judged unsafe if the probability of 30% or greater dose-limiting toxicity (the primary outcome) over controls was higher than 25%. Secondary outcomes included safety, clinical progression, pharmacokinetics and virologic responses.

**Results**

Of 103 participants screened, 18 participants were enrolled between 17 July and 30 October 2020. Molnupiravir was well tolerated at 400, 600 or 800mg doses with no serious or severe adverse events. Overall, 4 of 4 (100%), 4 of 4 (100%) and 1 of 4 (25%) of the participants receiving 300, 600 and 800mg molnupiravir respectively, and 5 of 6 (83%) controls, had at least one adverse event, all of which were mild (≤grade 2). The probability of ≥30% excess toxicity over controls at 800mg was estimated at 0.9%.

**Conclusion**

Molnupiravir was safe and well tolerated; a dose of 800mg twice-daily for 5 days was recommended for Phase II evaluation.

Abstract 235 words

Main text 3134 words

**INTRODUCTION**

In addition to life-saving therapies for COVID-19, there is an urgent need for effective antivirals in order to reduce disease burden, prevent hospitalisation and death and potentially decrease transmission of SARS-CoV-2. Since the natural history of infection is characterised by an early peak in viral load (within the first 5 days of infection),1 any antiviral would be expected to exert most effect when given early in the course of infection. AGILE is a randomised multi-arm, multi-dose, phase Ib/IIa platform in the UK using a seamless Bayesian adaptive design 2 to determine the safety, activity and optimal doseof multiple SARS-CoV-2 candidate therapeutics. Several candidates can be tested simultaneously (potentially sharing control group data) to increase efficiency.

We evaluated molnupiravir (EIDD-2801/MK-4482), for the treatment of COVID-19 in a seamless phase I/II trial. Molnupiravir is rapidly and extensively converted (via host esterases) to the active ribonucleoside analogue 14 ß-d-N4-hydroxycytidine (NHC; EIDD-1931); cytochrome P450 enzymes are not a major route in the metabolism of molnupiravir, and are not affected by the drug through enzyme induction or inhibition. Despite differences in model systems, activity of molnupiravir has consistently been demonstrated *in-vitro* and in animal models. In mice implanted with authentic human lung tissue, a prophylactic dose of 500 mg/kg given 12h prior to inoculation with SARS-CoV-2 and every 12h thereafter dramatically reduced viral plaque forming units at 2 days post inoculation.3 Furthermore, a twice daily 200mg/kg dose (but not 75mg/kg) was also able to reduce pulmonary viral RNA and improve lung histopathology in Syrian Golden Hamsters when initiated at the time of inoculation but with much lower efficacy if initiated ≥24h after infection.4 Finally, molnupiravir significantly reduced viral titres in the nasal swabs and turbinate 4 days after infection in ferrets when given at 5mg/kg twice daily initiated 12h after inoculation or 15mg/kg initiated 36h after inoculation,5 and was able to block transmission between ferrets. Current data warrant investigation of molnupiravir in human patients including studies to define the appropriate dose for a human SARS-CoV-2 antiviral indication.

Molnupiravir has been evaluated in healthy volunteers in single (50-1600mg) and multiple (50-800mg for 5.5 days) ascending oral doses, and was found to be well-tolerated.6 Preliminary data have also been presented from a study in patients with mild-to-moderate SARS CoV2 infection who received 200mg, 400mg or 800mg of molnupiravir twice daily for 5 days or placebo.7 Virus was cultured from nasopharyngeal swabs in only 42.9% of all PCR-positive patients at baseline and of these, culture-negativity was seen in all 47 evaluable subjects receiving molnupiravir (regardless of dose) versus 24% subjects allocated to placebo.

Here we report phase Ib results where we sought to determine the safety and tolerability of multiple ascending doses of molnupiravir in participants with PCR-confirmed SARS-CoV-2 infection who had symptoms within the preceding 5 days which did not require hospitalisation, to recommend a dose for phase II. Secondary objectives included characterising adverse events (AEs), serious adverse events (SAEs), clinical outcomes (FLU-PRO, WHO Ordinal Scale, NEWS2 and mortality) as well as the pharmacokinetics of molnupiravir and its major metabolite EIDD-1931.

**METHODS**

**Study design, Participants and Ethics**

This dose-escalation phase I study (clinicaltrials.gov registration number NCT04746183) was designed as an open label, randomised, controlled Bayesian adaptive trial in adult early infection in the community, coordinated by the National Institute for Health Research (NIHR) Southampton Clinical Trials Unit with participants recruited into the NIHR Royal Liverpool and Broadgreen Clinical Research Facility (UK). Eligible participants were men and women aged ≥18 years with PCR-confirmed SARS-CoV-2 infection who were within 5 days of symptom onset, free of uncontrolled chronic conditions, and ambulant in the community with mild or moderate disease (see eligibility below). Women of childbearing potential and men were required to use two effective methods of contraception, one of which should be highly effective, throughout the study and for 50 days and 100 days thereafter respectively. Any of the following criteria excluded participants from the study: pregnant or breast feeding women, stage 4 (severe) chronic kidney disease, clinically significant liver dysfunction, SpO2 <95% by oximetry or lung disease requiring supplementary oxygen, ALT and/or AST > 5 times upper limit of normal , platelets <50x10‑9/L, experiencing any >= Grade 3 Common Terminology Criteria for Adverse Events (CTCAE v5), previously reported hepatitis C infection, known allergy to any study medication or having received any other experimental agents within 30 days of first dose of study drug (use of other comedications was allowed). All participants provided written informed consent before enrolment. The study protocol was reviewed and approved by the UK Medicines and Healthcare product Regulatory Agency (MHRA)(EudraCT 2020-001860-27) and West Midlands Edgbaston Research Ethics Committee (20/WM/0136).

**Randomisation and masking**

Four sequential molnupiravir dosing tiers were defined *a priori* (300mg, 400mg, 600mg and 800mg BD for 5 days) with participants allocated using permuted blocks (block size 3, with no further stratification factors, generated by NIHR Southampton CTU statisticians) via MEDIDATA RAVE. Randomisation used a 2:1 allocation ratio so that within each cohort, 4 participants were randomly assigned to receive molnupiravir plus standard-of-care and 2 participants (controls) standard-of-care alone. This study was open label in accordance with conventional Phase I design, with a follow-on Phase IIa placebo-controlled study.

**Procedures**

Participants with laboratory-confirmed SARS-CoV-2 infection, or who had an illness compatible with COVID-19 (and who were subsequently confirmed to be positive) were screened against eligibility criteria, including presence and onset of symptoms within the previous 5 days. For safety reasons, in each cohort, the first participant randomised to molnupiravir (sentinel patient) was followed up until for 24 hours before any subsequent participants were randomised. All participants who received molnupiravir received drug after at least a two hour fasting period with a 4 hour period of observation after the first dose.

We utilised a Bayesian adaptive design to accelerate decision making in this phase I study. Briefly, we developed a dose toxicity model 8 which estimates (as a continuous incremental probability) the ‘dose limiting toxicity’ at day 7 of molnupiravir in controls and at each of the four pre-defined dosing tiers (300mg, 400mg, 600mg and 800mg BD)– see figure 2. Tolerability was expressed as an excess in prevalence of dose limiting toxicities of treatment over controls (who also suffered from symptoms of COVID-19 infection) with the model continually refined upon completion of each dosing cycle. Details are provided in Supplement S1. For each cohort, the Safety Review Committee (SRC) reviewed all available safety data including at least 7 days data for each participant in the cohort, and all accrued information on previous cohorts (up to a maximum follow-up of 28 days). This included AE data, vital signs data, ECG data and clinical laboratory evaluations, as well as any emerging data from other studies. Following SRC review, recommendations could be to de-escalate, escalate, remain at the same dose, or continue to phase II. A dose was deemed to be unsafe if there was a ≥25% chance that treatment was associated with a >30% risk of dose-limiting toxicities at day 7. The model recommended the next dose-level according to which level is the most likely to correspond to an increase of a 15-25% in the dose limiting toxicity rate over control. However, the SRC made the ultimate decision whether to accept that the current dose was safe and to dose escalate and could decide to skip a dose if it did not more than double and was deemed safe by the Bayesian model. Once the dose escalation Phase I was complete, the independent Data Monitoring and Ethics Committee reviewed data from the final SRC, along with their recommendations on the recommended phase II dose, to ratify the Recommend Phase 2 Dose.

**Outcomes**

The primary outcome was dose limiting toxicity (DLT) using CTCAE version 5 (grades 3 and above) measured over 7 days and CTCAE grading related to platelets and/or lymphocytes, assessed in all participants, who were randomly assigned and received at least one dose of molnupiravir (unless randomised to control). Secondary outcomes: for safety included AEs, SAEs, physical findings, vital signs and laboratory parameters; for pharmacokinetics included concentrations of molnupiravir and EIDD-1931 in plasma; for clinical included Patient Reported Outcome Measures (FLU-PRO), WHO COVID-19 Ordinal Scale (at days 15,29), NEWS2 (assessed during clinic days 15, 29), mortality (days 15, 29) and time from randomisation to death (up to day 29).

**Pharmacokinetic sampling**

Plasma was sampled at Day 1 and 5 to measure the concentrations of molnupiravir and its major active metabolite EIDD-1931. On each sampling day, 2mL of venous blood was collected pre-dose, at 30 mins, and at 1-, 2- and 4-hours post-dose. All samples were rapidly cooled on wet ice and centrifuged (2000g for 10min) within 30 min of sample collection. Within 10 minutes of completing centrifugation, 150mL of plasma was mixed with 450uL of acetonitrile, vortexed and transferred to a -80oC freezer prior to onward shipping for pharmacokinetic analyses. Drug concentrations were measured using a validated LC-MS/MS assay at Covance Clinical Laboratories, Leeds, UK.

Concentrations of EIDD-1931 in plasma on day 1 and day 5 were described using summary statistics (geometric mean (90% CI), mean, standard deviation, median and range) for each time point.

Key pharmacokinetic (PK) parameters such as area under the concentration-time curve 0-4 h (AUC0-4), maximum concentration (Cmax) and time to maximum concentration (Tmax) were determined by non-compartmental modelling methods (WinNonlin, Phoenix, v. 8.3, Pharsight, Mountain View, CA, USA) on day 1 and day 5 for each dose and summarised descriptively. Accumulation ratios to day 5 were calculated for EIDD-1931 AUC0-4 and Cmax.

**Statistical analysis**

All analyses are reported according to CONSORT 2010 and ICH E9 guidelines on Statistical Principles in Clinical Trials. All enrolled participants were included in both the evaluable population and the safety population for analysis.

The primary endpoint of DLTs up to 7 days post first dose were modelled using a Bayesian dose-toxicity model based on Mozgunov *et al*.8 The relationship between dose and toxicity was modelled using a two-parameter logistic model, where information can be shared across doses; in particular, the DLT rate in controls informs estimates for the active doses. The prior distributions for this model were calibrated to maximise the proportion of correct selection under a range of dose-toxicity scenarios where each dose considered in the study was the optimum one. The toxicity risk in controls was *a priori* assumed to be 10%. Further details are given in Supplement S1.

The dose-toxicity model was updated after every cohort of participants, and the final model is presented as estimated DLT rates for each dose, alongside equal-tail 95% credible intervals. For active doses, we also present estimated additional toxicity above controls, the probability that the DLT rate falls within 15-25% additional toxicity over controls (a pre-determined acceptable target range for toxicity) and the probability of at least 30% additional toxicity over controls (deemed as unacceptably toxic). This is supported by the same information for up to day 29.

Baseline demographics are summarised within each dose (and controls) using descriptive statistics. Clinical endpoints are similarly summarised at days 15 and 29. The sample size was flexible, based on the need for the study to adapt to accruing safety data. Simulations to assess model operating characteristics and to calibrate prior assumed four doses (plus controls), with cohorts of size six capped at a total of 30 participants.

Statistical analysis was undertaken in SAS version 9·4, STATA version 16 and R version 3·6 ·0.

**Role of funding source**

The funder of the study had no role in study design, study execution, data collection, data analysis, or data interpretation. WH and WP were non-voting members of the Safety Review Committee meetings as recommended by the MHRA, but did not participate in the Committee’s decision-making. The statisticians SE, GS and KT had full access to all the data in the study and SK & GG had final responsibility for the decision to submit for publication.

**RESULTS**

Of 103 potential participants (Figure 1) who attended for screening, 58 were excluded (31 had no signs or symptoms of COVID-19, 12 tested negative for SARS-CoV-2 by PCR, 7 had signs or symptoms that began after 5 days of planned first dose, 2 had an uncontrolled comorbidity, 3 did not meet contraceptive requirements, 1 did not meet the age range, 1 did not meet the mild/moderate disease criterion, 22 declined, 5 were screened between dosing cohorts and 1 was unknown). Eligible individuals were randomly assigned within three sequential dose cohorts (300mg, 600mg & 800mg) of 6 participants each (i.e. a total of 18 participants within the phase 1) and dosed in the period between 17 July 2020, and 30 October 2020. The baseline characteristics of participants were similar across all groups (Table 1) with an overall median age of 56, 72% (13/18) female and 33% (6/18) having a WHO COVID ordinal score of 1 (ambulatory mild disease). The median number of days (range) from symptom onset to randomisation and treatment by the 18 participants was 4 (range 1-5).

All molnupiravir participants received at least 1 dose with 3/4 (75%), 4/4 (100%) and 3/4 (75%) completing the full treatment in the 300, 600 and 800mg cohort respectively. One participant on 300mg BD only took 1 of 2 intended tablets for 2 of their treatment doses and one participant on 800mg BD took only two doses on day one, withdrawing from treatment for personal reasons unrelated to the study. The median number of molnupiravir doses received (range) was 10 (8-10), 10 (10-10), 10 (2-10) and median number of days on molnupiravir treatment (and range) was 5.5 (5-6), 5 (5-5), 5 (1-5) for the 300mg, 600mg & 800mg cohorts respectively.

**Primary analysis**

No participants in any cohort experienced a DLT or a grade 3 or above change in lymphocytes or platelets (for those with a normal baseline value) or a 2 or more grade increment in lymphocytes or platelets (for those with grade 2 or 3 at baseline). Following review by the Safety Review Committee (SRC), dose cohort escalation went from 300mg to 600mg (skipping 400mg) and then from 600mg to 800mg. Bayesian model DLT point estimates, 95% credible interval, and the target toxicity level of 20% over the controls are shown in Figure 2. For data up to day 7, the maximum dose (800mg) had an estimated DLT rate of 11·0% (equal-tail 95% credible interval of 1·8 to 30·4%), with estimated 7·4% additional toxicity over controls and a probability of additional toxicity ≥30% over controls of 0.9%. As there were no DLTs recorded up to day 28, the results for day 7 are the same for day 28 and so are not repeated. These data support 800mg BD as the recommended phase II dose.

**Analysis of secondary endpoints**

Adverse events were evenly distributed among the dose cohorts including controls. Overall, 4 of 4 (100%), 4 of 4 (100%) and 1 of 4 (25%) of the participants receiving 300, 600 and 800mg of molnupiravir, and 5 of 6 (83%) controls, had at least one adverse event, all of which were mild (≤ grade 2). Molnupiravir was generally well-tolerated compared with controls, and Table 2 describes the frequencies of events across the groups by system organ class and CTCAE term. No serious adverse events were reported. The most common symptoms were gastrointestinal (diarhoea, nausea), respiratory (cough), central nervous system (loss of smell or taste) and flu-like symptoms.

At day 15 all participants had a WHO ordinal scale of 1 or 2, with a median score (range) of 1·5 (1-2), 1·5 (1-2), 2 (2-2) and 1·5 (1-2) for molnupiravir 300mg, 600mg and 800mg and controls respectively. At day 15 the molnupiravir 300mg, 600mg and controls had a median NEWS2 Score of 0 (range 0-0), with molnupiravir 800mg a median score of 1 (range 0-1). Median O2 saturation (range) was 97 (97-100), 97 (96-99), 99.5 (97-100), 97 (96-99) for 300, 600 and 800mg molnupiravir and controls respectively with median FLU-PRO totals 0·4 (0·2-10), 0·2 (0·1-0·6), 0·1 (0-0·3) and 0·2 (0-0·5) respectively (further details with comparable day 29 endpoints are provided in Supplement 2).

Virology samples were collected for future sequencing and characterisation of variants; PK-PD analysis was not included in this Phase I analysis since sample sizes at each dose are too small to make inferences about antiviral activity.

**Pharmacokinetics**

The prodrug molnupiravir was generally not detectable, or detected at low concentrations only at early timepoints (0·5, 1 hour post-dose), at all 3 doses (Table 3). Plasma concentrations of the nucleoside metabolite EIDD-1931 were detectable, and showed no accumulation between day 1 and day 5. At day 5, geometric mean NHC exposures (%CV) over the first 4 hours of dosing (AUC0-4) in the 300mg (N=4), 600mg (N=4) and 800mg (N=3) dose were 3470 (42·4), 3880 (56·3) and 7880 (39·0) ng.h/mL, with corresponding peak (Cmax) concentrations of 1620 (51·0), 1820 (84·6) and 4180 (28·1) ng/mL. Time to peak plasma concentration was 0·5 – 2·0 hours.

**DISCUSSION**

To study the tolerability and safety of molnupiravir, we enrolled participants who presented within five days of symptoms, and who did not have severe disease, since we judged that the largest public health impact of this antiviral drug would be through deployment in the community for preventing hospitalisation. In untreated SARS-CoV-2 infection, viral load peaks in the first week of illness 1 suggesting that early antiviral treatment may influence disease progression and potentially transmission.

We have established the safety and tolerability of molnupiravir in SARS-CoV-2 infected individuals, alongside a conventional phase I dose ranging study in healthy volunteers (NCT04392219). We have shown that a dose of 800mg bd of molnupiravir is safe and well-tolerated in participants with SARS CoV-2 infection; the plasma concentrations attained are within the target range based on scaling from animal models.3,4 Adverse effects were commonly reported, affecting 9/12 and 5/6 participants on molnupiravir and controls respectively. All were mild (Grade 1-2) and included flu-like and upper respiratory symptoms, headache, myalgia, diarrhoea and nausea which were also consistent with symptomatic COVID-19 disease.

AGILE utilises complex innovative trial design methodology to accelerate early phase evaluation of novel antiviral agents against SARS CoV-2. Our Bayesian approach was selected to optimise statistical efficiency and to accelerate decision-making. Drug safety is not definitively established during phase I and requires large numbers of individuals dosed in phase III or IV. Rather, the AGILE design allowed us to establish (within an accelerated timescale) that a dose of molnupiravir 800mg bd for 5 days was sufficiently safe to progress into our continuation phase IIa placebo-controlled trial (where safety continues to be monitored). Since full reproductive toxicologic datasets were not available at the time of initiation, our study required stringent precautions to avoid pregnancy in participants or their partners.

To our knowledge this is the first published report describing the use of molnupiravir in SARS-CoV-2 infected individuals. We observed comparable exposures of EIDD1931 to healthy volunteers,1 and describe an approach for rapidly estimating a dose-toxicity relationship for phase II evaluation. Whether or not molnupiravir will prove effective in treating COVID-19 will be determined in phase II trials which are currently underway, including our own, but the paucity of potent antiviral agents in the COVID-19 pipeline strongly argues for such accelerated approaches to early phase drug development.

**Data sharing**

The AGILE Trial Steering Committee will consider all reasonable requests by health-care providers, investigators, and researchers to provide anonymised data to address specific scientific or clinical objectives. The AGILE investigators are committed to reviewing requests from researchers for access to clinical trial protocols, de-identified patient-level clinical trial data, and study-level clinical trial data.

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**Transparency declarations**

**Conflicts of interest**

Potential conflicts of interest. WH is a cofounder, owner and advisor of/to Ridgeback Biotherapeutics LP. WP is employed by Ridgeback Biotherapeutics.

Potential conflicts of interest SK has received research funding from ViiV Healthcare, Gilead Sciences, and Merck for the Liverpool HIV Drug Interactions programme and for unrelated clinical studies. Unrelated to the current work, AO is a Director of Tandem Nano Ltd and co-inventor of patents relating to drug delivery. AO has received research funding from ViiV Healthcare, Merck, Janssen and consultancy from Gilead, ViiV and Merck not related to the current work. GG has received funding from Janssen-Cilag, Astra Zeneca, Novartis, Astex, Roche, Heartflow, Celldex, BMS, BionTech, Cancer Research UK, NIHR, British Lung Foundation, Unitaid, GSK for unrelated academic clinical trials and programme funding. WG has received funding from the Wellcome Trust.

No conflict. RF, TF, SE, TJ, RL, ND, LW, OTH, CW, HR, EM, PM, EA, KB, MB, JG, GS, AC, CH, KT, JC, SC, HP, EW, LJ, SY, KM, MR, KF, VS, DL, MJ

**Contributors Statement**

SK, GG, TJ, SE, RF, PM contributed to study design. SK, GG, TJ, SE, GS, KT, PM, HP contributed to data analysis and interpretation. RF led clinical conduct as the principal investigator of the clinical site. TF, LW, RL, MB participated in clinical assessment and data collection. KF participated in the management of pharmacovigilance. SC, EW, MR, LJ contributed to the digital data collection and data management of the trial. MR led set up of the randomisation system. WG, TF, VS, EA, KB, CH contributed to study bioanalysis. AC, ND, EM, OTH, SY, HR, JC, RL, CW, JB, AO, MJ, DGL contributed to study management and execution. KM contributed to monitoring activities. WP, WH contributed preclinical and safety data on molnupiravir. SK, GG, MJ, AO, DGL were involved in primary manuscript writing. All authors contributed to the final version of the manuscript.

**References**

1.Cevik M, Tate M, Lloyd O *et al*. SARS-CoV-2, SARS-CoV, and MERS-CoV viral load dynamics, duration of viral shedding, and infectiousness: a systematic review and meta-analysis. *Lancet Microbe*. 2020; **2**: e13-e22

2.Griffiths G, Fitzgerald R, Jaki T *et al*. AGILE-ACCORD: A Randomized, Multicentre, Seamless, Adaptive Phase I/II Platform Study to Determine the Optimal Dose, Safety and Efficacy of Multiple Candidate Agents for the Treatment of COVID-19: A structured summary of a study protocol for a randomised platform trial. *Trials*. 2020; **21**: 1-3

3.Wahl A, Gralinski L, Johnson C *et al*. SARS-CoV-2 infection is effectively treated and prevented by EIDD-2801. *Nature* 2021; **591**: 451-457.

4.Abdelnabi R, Foo CS, Kaptein SJ *et al*. *Molnupiravir (EIDD-2801) inhibits SARS-CoV2 replication in Syrian hamsters model.*

https://doi.org/10.1101/2020.12.10.419242

5.Cox RM, Wolf JD, Plemper RK. Therapeutically administered ribonucleoside analogue MK-4482/EIDD-2801 blocks SARS-CoV-2 transmission in ferrets. *Nat Microbiol*. 2021; **6**: 11-8

6.Painter WP, Holman W, Bush JA *et al*. Human Safety, Tolerability, and Pharmacokinetics of a Novel Broad-Spectrum Oral Antiviral Compound, Molnupiravir, with Activity Against SARS-CoV-2. *Antimicrob Agents Chemother.* 2021;

https://doi.org/10.1128/AAC.02428-20

7.Painter WP, Sheahan T, Baric R, *et al*. Reduction in infectious SARS-CoV-2 in treatment study of COVID-19 with molnupiravir. *Virtual Conference on Retroviruses and Opportunistic Infections (vCROI) 2021*. Abstract 777

8.Mozgunov P, Jaki T, Paoletti X. Randomized dose-escalation designs for drug combination cancer trials with immunotherapy. *J Biopharm Stat*. 2019; **29**: 359-77

TABLE 1: Baseline characteristics

|  | **MOLNUPIRAVIR** | **MOLNUPIRAVIR** | **MOLNUPIRAVIR** | **Standard Care** |  |
| --- | --- | --- | --- | --- | --- |
|  | **(300mg)** | **(600mg)** | **(800mg)** | **Total** | **Total** |
|  | **(n=4)** | **(n=4)** | **(n=4)** | **(n=6)** | **(n=18)** |
| **Age at consent (years)** |  |  |  |  |  |
| n | 4 | 4 | 4 | 6 | 18 |
| Median  | 56·0 | 43·0 | 39·0 | 59·0 | 56·0 |
| Range | 51·0 to 80·0 | 22·0 to 60·0 | 25·0 to 63·0 | 22·0 to 63·0 | 22·0 to 80·0 |
| **Gender – n (%)** |  |  |  |  |  |
| Male | 1 (25·0%) | 2 (50·0%) | 0 (0·0%) | 2 (33·3%) | 5 (27·8%) |
| Female | 3 (75·0%) | 2 (50·0%) | 4 (100%) | 4 (66·7%) | 13 (72·2%) |
| **Ethnicity**  |  |  |  |  |  |
| White - English / Welsh / Scottish / Northern Irish | 4 (100%) | 4 (100%) | 4 (100%) | 6 (100%) | 18 (100%) |
| **BMI** |  |  |  |  |  |
| n | 4 | 4 | 4 | 6 | 18 |
| Median  | 28·1 | 33·9 | 21·0 | 31·3 | 29·5 |
| Range | 25·6 to 32·7 | 30·0 to 51·1 | 20·4 to 34·0 | 27·2 to 36·2 | 20·4 to 51·1 |
| **WHO Score (day 1) – n (%)** |  |  |  |  |  |
| 1. Ambulatory mild disease, asymptomatic; viral RNA detected | 2 (50·0%) | 0 (0·0%) | 1 (25·0%) | 3 (50·0%) | 6 (33·3%) |
| 2. Ambulatory mild disease, symptomatic; independent | 1 (25·0%) | 4 (100%) | 3 (75·0%) | 3 (50·0%) | 11 (61·1%) |
| 3. Ambulatory mild disease, symptomatic; assistance needed | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 1 (5·6%) |
| **WHO Score (day 1)** |  |  |  |  |  |
|  n | 4 | 4 | 4 | 6 | 18 |
|  Median  | 1·5 | 2 | 2 | 1·5 | 2 |
|  Range | 1 to 3 | 2 to 2 | 1 to 2 | 1 to 2 | 1 to 3 |
| **NEWS2 Score (day 1)** |  |  |  |  |  |
|  n | 4 | 4 | 4 | 6 | 18 |
|  Median  | 0 | 0 | 0.5 | 0 | 0 |
|  Range | 0 to 1 | 0 to 0 | 0 to 1 | 0 to 0 | 0 to 1 |
| **O2 Saturation % (day 1)** |  |  |  |  |  |
|  n | 4 | 4 | 4 | 6 | 18 |
|  Median  | 97·5 | 96·5 | 99·0 | 98·0 | 97·5 |
|  Range | 95·0 to 98·0 | 96·0 to 99·0 | 95·0 to 100·0 | 96·0 to 100·0 | 95·0 to 100 |
| **FLU-PRO total (day 1)** |  |  |  |  |  |
|  n | 3 | 4 | 4 | 6 | 17 |
|  Median  | 0·7 | 0·8 | 1·0 | 0·6 |  |
|  Range | 0·4 to 1·3 | 0·8 to 1·4 | 0·5 to 1·6 | 0·3 to 1·6 | 0·3 to 1·6 |
| Missing from eCRF – n (%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 1 (5·6%) |
| **Time from symptom onset to randomisation (days)2** |  |  |  |  |  |
|  n | 4 | 4 | 4 | 6 | 18 |
|  Median  | 4·0 | 4·0 | 3·5 | 4·0 | 4·0 |
|  Range | 3·0 to 4·0 | 4·0 to 4·0 | 2·0 to 4·0 | 1·0 to 5·0 | 1·0 to 5·0 |

**NOTE:** Percentages are based on the number of patients in the study arm

¹Reason for missing FLU-PRO:

2**NOTE:** Date of randomisation is the same as date of first dose for all patients randomised to molnupiravir

**TABLE 2:** **Overall toxicity Summary by CTCAE version 5 term - Safety population**

| **Characteristic** | **MOLNUPIRAVIR** | **MOLNUPIRAVIR** | **MOLNUPIRAVIR** | **Standard Care** |
| --- | --- | --- | --- | --- |
|  | **(300mg)** | **(600mg)** | **(800mg)** | **Total** |
|  | **(n=4)** | **(n=4)** | **(n=4)** | **(n=6)** |
| Number of patients that experienced at least one AE – n (%) \* | 4 (100%) | 4 (100%) | 1 (25%) | 5 (83·3%) |
| Summary of AEs – n (%) |  |  |  |  |
| **Cardiac disorders** | **0 (0·0%)** | **0 (0·0%)** | **1 (25·0%)** | **0 (0·0%)** |
| Palpitations | 0 (0·0%) | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) |
| **Ear and labyrinth disorders** | **0 (0·0%)** | **0 (0·0%)** | **0 (0·0%)** | **1 (16·7%)** |
| Tinnitus | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| **Eye disorders** | **0 (0·0%)** | **1 (25·0%)** | **0 (0·0%)** | **0 (0·0%)** |
| Blurred Vision | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| **Gastrointestinal disorders** | **2 (50·0%)** | **3 (75·0%)** | **0 (0·0%)** | **2 (33·3%)** |
| Abdominal Pain | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Diarrhea | 2 (50·0%) | 1 (25·0%) | 0 (0·0%) | 1 (16·7%) |
| Dyspepsia | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| Nausea | 1 (25·0%) | 2 (50·0%) | 0 (0·0%) | 1 (16·7%) |
| Oral Dysesthesia | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Vomiting | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| **General disorders and administration site conditions** | **1 (25·0%)** | **0 (0·0%)** | **0 (0·0%)** | **2 (33·3%)** |
| Fatigue | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Flu Like Symptoms | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 2 (33·3%) |
| Non-Cardiac Chest Pain | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| **Infections and infestations** | **1 (25·0%)** | **2 (50·0%)** | **0 (0·0%)** | **0 (0·0%)** |
| Herpes Simplex Reactivation | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Infections And Infestations - Other, Specify1 | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| Thrush | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| **Investigations** | **0 (0·0%)** | **1 (25·0%)** | **0 (0·0%)** | **0 (0·0%)** |
| Alanine Aminotransferase Increased | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| GGT Increased | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| **Musculoskeletal and connective tissue disorders** | **1 (25·0%)** | **0 (0·0%)** | **0 (0·0%)** | **1 (16·7%)** |
| Chest Wall Pain | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Myalgia | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| **Nervous system disorders** | **2 (50·0%)** | **1 (25·0%)** | **0 (0·0%)** | **2 (33·3%)** |
| Anosmia | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Dizziness | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| Dysgeusia | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| Headache | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 2 (33·3%) |
| **Psychiatric disorders** | **0 (0·0%)** | **0 (0·0%)** | **1 (25·0%)** | **0 (0·0%)** |
| Anxiety | 0 (0·0%) | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) |
| **Renal and urinary disorders** | **1 (25·0%)** | **0 (0·0%)** | **0 (0·0%)** | **1 (16·7%)** |
| Chronic Kidney Disease | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| Urine Discoloration | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| **Respiratory, thoracic and mediastinal disorders** | **1 (25·0%)** | **1 (25·0%)** | **0 (0·0%)** | **1 (16·7%)** |
| Cough | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 1 (16·7%) |
| Hoarseness | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| Rhinorrhea | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Sore Throat | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |
| **Unclassified** | **1 (25·0%)** | **1 (25·0%)** | **1 (25·0%)** | **0 (0·0%)** |
| Other - Bilateral Thigh Pain | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) | 0 (0·0%) |
| Other - Loose Stools | 0 (0·0%) | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) |
| Other - Worsening Fatigue | 0 (0·0%) | 1 (25·0%) | 0 (0·0%) | 0 (0·0%) |

Note: Percentages are based on the number of patients in the study arm. CTCAE v5·0 terms are used to classify AEs.

1This AE reported in Other specify free text field as “Chest infection”.

\* NB: All AEs were either Grade 1 or 2

Table 3: Geometric mean (%CV) Pharmacokinetic parameters of EIDD-2801 following single and multiple dose administration of EIDD-2801

|  |  |  |  |
| --- | --- | --- | --- |
|  | **300mg bd** | **600mg bd** | **800 mg bd** |
| **Parameter (Units)**  | **Day 1** | **Day 5** | **Day 1** | **Day 5** | **Day 1** | **Day 5** |
|  | **N=4** | **N=4** | **N=4** | **N=4** | **N=4** | **N=3** |
| Molnupiravir (EIDD2801) |
| AUC0-4 (ng.h/mL)  | NC | NC | NC | NC | NC | NC |
| Cmax (ng/mL)  | 5·76 (13.3) | NC | 25·8 | 9·14c (5·97 – 12·3)d | 8·43 (58·6) | 7·79 (13·9) |
| Tmaxa (h)  | 0·500 (0·500-0·500) | NC | 1·00b (1·00-1·00) | 0·500c (0·500-0·500) | 0·750 (0·500-1·00) | 1·00 (1·00-1·00) |
| NHC (EIDD1931) |
| AUC 0-4 (ng.h/mL)  | 3210 (40·5) | 3470 (42·4) | 4610(33·7) | 3880(56·3) | 9240 (41·0) | 7880 (39·0) |
| Cmax (ng/mL)  | 1490 (29·4) | 1620 (51·0) | 2230(38·2) | 1820(84·6) | 4440 (45·2) | 4180 (28·1) |
| Tmaxa (h)  | 1·50 (1·00-2·00) | 1·00 (1·00-2·00) | 1·50 (1·00-2·00) | 1·00 (1·00-2·00) | 2·00 (1·00–2·00) | 2·00 (1·00-2·00) |

NA = Not applicable, NC = Not calculable,

AUC0-4 = Area under the time-concentration curve (0-4h)

Tmax = Time to peak concentration

Cmax – Peak concentration

a Median (min-max) presented,

b n=1 with quantifiable concentrations out of 4 subjects

c n=2 with quantifiable concentrations out of 4 subjects

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Figure 1: CONSORT diagram

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Figure 2: Primary endpoint - dose toxicity plot up to day 7 (evaluable population)