EAVE II Data Dictionary

EAVE II Analysts

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1 GP Data Demographics

1.1 Data information

| Description | Baseline GP extract on all patients and their key demographics |
|-------------------|--|
| Datafile | EAVE_demographics.rds |
| Type | Cleaned |
| Data sources | GP data |
| Scripts | NA |
| Update frequency | Bi-monthly to 3 months |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|---------------------------|---------------------------------------|---|-----------|----------------|---|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | GP data | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| ageYear | Age | Age in years | integer | GP data | Min = 0, $Max = 120$ | NA | NA |
| Sex | Sex | Sex at birth | character | GP data | F, M | NA | NA |
| $simd 2020_sc_quintile$ | SIMD Quintile | Scottish Index of Multiple Deprivation (SIMD) Quintile (2020) | integer | GP data | 1-5 | NA | NA |
| ur6_2016 | Urban/Rural classification | Urban/Rural 6-fold classification (2016 classification) | integer | GP data | 1-6 | NA | NA |
| ur6_2016_name | Urban/Rural classification name | Urban/Rural 6-fold classification (2016 classification) | character | GP data | 1 Large Urban Areas, 2 Other urban Areas, 3 Accessible Small Towns, 4 Remote Small Towns, 5 Accessible Rural, 6 Remote Rural | NA | NA |
| hb2019 | NHS Health Board Code | Codes of 2019 NHS Health Boards | character | GP data | \$08000015, \$08000016, \$08000017, \$08000019, \$08000022, \$08000022, \$08000024, \$08000025, \$08000026, \$08000028, \$08000029, \$08000030, \$08000031, \$08000032 | NA | NA |
| hb2019name | NHS Health Board Name | Names of 2019 NHS Health Boards | character | GP data | NHS Ayrshire and Arran, NHS Borders, NHS Dumfries and Galloway, NHS Fife, NHS Forth Valley, NHS Grampian, NHS Greater Glasgow and Clyde, NHS Highland, NHS Lanarkshire, NHS Lothian, NHS Orkney, NHS Shetland, NHS Tayside, NHS Western Isles | NA | NA |

2 GP Diagnostics

2.1 Data information

| Description | Baseline GP extract of all EAVE II risk groups |
|-------------------|--|
| Datafile | cohort_diags.rds |
| Type | Cleaned |
| Data sources | GP data |
| Scripts | NA |
| Update frequency | Bi-monthly to 3 months |
| Update timestamps | 2020-09-11, 2020-09-15 |
| Analyses used | NA |

2.2 Variable information

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-------------|------------------------|---|-----------|----------------|--|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | GP data | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| diag | Diagnosis (GP cluster) | Medical diagnosis, clinical indicators and other health-related characteristics | character | GP data | See Supplementary Information - EAVEII_dd_gp_clusters.xl | NA lsx | NA |

3 ECOSS

3.1 Data information

| Description | Laboratory data on COVID-19 testing |
|-------------------|--|
| Datafile | ECOSSdeduped_linked.rds |
| Type | Cleaned |
| Data sources | ECOSS |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | 2020-12-08, 2020-12-15, 2020-12-22, 2020-12-29, 2021-01-05, 2021-01-19 |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|---------------------|----------------------|---|-----------|----------------|--|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | ECOSS | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| ECOSSID | ECOSS Link Number | Identifier | integer | ECOSS | 8 digits (e.g. 12345678) | NA | NA |
| Sex | Sex | Sex at birth | character | ECOSS | F, M, U (Unknown) | NA | NA |
| AgeYear | Age | Age in years | integer | ECOSS | Min = 0, $Max = 130$ | NA | NA |
| ${\it HealthBoard}$ | NHS Health Board | Name abbreviation of NHS Health Boards | character | ECOSS | AA, BR, DG, FF, FV, GC, GR, HG, LN, LO, OR, SH, TY, UK (Uknown), WI | NA | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|----------------------------|-------------------------------------|--|------------------------|----------------|--|---------------------|-------------------------|
| PostCode LabSpecimenNo | Postcode Number of laboratory | Postcode Number of laboratory | character character | ECOSS ECOSS | 7 digits/letters 11 digits/letters | NA NA | NA NA |
| Specimendate | specimen date of specimen | specimen date of specimen | date | ECOSS | yyyy-mm-dd | NA | NA |
| OriginalOrganism | Original organism | collection Original organism - Coronavirus 2019 positive | character | ECOSS | Coronavirus 2019-nCoV, Coronavirus 2019-nCoV negative | NA | NA |
| NCOV_RESULT | COVID-19 result | or negative Novel Coronavirus result | character | ECOSS | Negative, Positive | NA | NA |
| result SpecimenOrigin | Result Origin of specimen | Result Origin of specimen | integer character | ECOSS ECOSS | 0,1 Community, Drug Service, GP, GUM Clinic, H.M Prison, Hospital, Local Authority EH Dept, NHS- Other, Non-NHS-Other, Occupational Health, Private, Residential Care Homes | NA NA | NA NA |
| ${\bf Ecossdate Received}$ | ECOSS date | date record received by ECOSS | date | ECOSS | yyyy-mm-dd | NA | NA |
| SubLab | Sub lab | Sub lab ID | character | ECOSS | ABD:GHB, AYR:ARL, BOR:ARL, DEE:UNI, DMF:ROY, EDI:RVL, FIF:ARL, GIL:BAN, GLA:GJH, GLA:REV, INV:RAI, LAN:HAR, LAN:LAW, LAN:MON, NHS:COV, ORK:BAL, STI:ROY, WES:LES | NA | NA |
| SourceLab | Source lab | Source lab ID | character | ECOSS | ABD:ALB, ABD:GHB, ABD:ROY, AYR:ARL, BOR:ARL, DEE:UNI, DMF:ROY, DUN:ARL, EDI:ROY, EDI:RVL, EDI:SCH, EDI:UNI, EDI:WGH, FIF:ARL, GIL:BAN, GLA:GAR, GLA:GJH, GLA:NUF, GLA:REV, GLA:ROY, GLA:SCH, GLA:SCS, GLA:SGH, GLA:STR, GLA:UNI, GLA:VIC, GLA:WES, GLA:CLY, INV:RAI, LAN:ALL, LAN:HAR, LAN:LAW, LAN:MON, NHS:UKL, OBA:LAB, ORK:BAL, PAI:RAH, STI:ROY, WES:LES, WLO:BAN | NA | NA |
| CAREHOMEID | Care home ID | Care home identifier | character | ECOSS | NA | NA | NA |
| NRS.Reg.date | NRS registration date | National Records of Scotland registration date | date | ECOSS | yyyy-mm-dd | NA | NA |
| NRS.date.Death | NRS death date | National Records of Scotland death date | date | ECOSS | yyyy-mm-dd | NA | NA |

 $\underline{(continued)}$

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|---------|----------------------|--|---------|-------------------------|--------|---------------------|--|
| death28 | Death within 28 days | Death within 28 days of positive covid test | integer | ECOSS, NRS deaths | 0,1 | Yes | Derived from daily linkage of ECOSS and NRS deaths within PHS. |

4 Health Board Lookup

4.1 Data information

| Description | Lookup of NHS Scotland Healthboards and the total population |
|-------------------|--|
| Datafile | HB2019_pop_lookup.rds |
| Type | Lookup |
| Data sources | PHS lookup file |
| Scripts | NA |
| Update frequency | NA |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|------------|---------------------------|---|-----------|---------------------------|---|---------------------|-------------------------|
| year | Year | Year of the lookup file used to determine the Health Boards | integer | Health board lookup | 2019 | NA | NA |
| hb2019 | Health Board code 2019 | Codes of NHS Health Boards in 2019 | character | Health board lookup | \$08000015, \$08000016, \$08000017, \$08000019, \$08000020, \$08000022, \$08000024, \$08000025, \$08000026, \$08000028, \$08000029, \$08000030, \$08000031, \$08000032 | NA | NA |
| hb2019name | Health Board name 2019 | Names of NHS Health Boards in 2019 | character | Health board lookup | NHS Ayrshire and Arran, NHS Borders, NHS Dumfries and Galloway, NHS Fife, NHS Forth Valley, NHS Grampian, NHS Greater Glasgow and Clyde, NHS Highland, NHS Lanarkshire, NHS Lothian, NHS Orkney, NHS Shetland, NHS Tayside, NHS Western Isles | NA | NA |
| hb2018 | Health Board code 2018 | Codes of NHS Health Boards in 2018 | character | Health board lookup | S08000015, S08000016, S08000017, S08000019, S08000020, S08000022, S08000024, S08000025, S08000026, S08000028, S08000029, S08000030, S08000031, S08000032 | NA | NA |
| hb2014 | Health Board code 2014 | Codes of NHS Health Boards in 2014 | character | Health board lookup | \$08000015, \$08000016, \$08000017, \$08000019, \$08000020, \$08000022, \$08000024, \$08000025, \$08000026, \$08000028, \$08000029, \$08000030, \$08000031, \$08000033 | NA | NA |

| (continued | |
|------------|--|
| | |
| | |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|----------|--------------|--|-----------|-------------------------------------|-----------------|---------------------|-------------------------|
| age | Age | Age in years | integer | Health board | Min= 0, Max= 90 | NA | NA |
| sex | Sex (number) | Sex at birth | integer | lookup Health board lookup | 1, 2 | NA | NA |
| sex_name | Sex | Sex at birth | character | Health board lookup | M, F | NA | NA |
| pop | Population | Number of people by year, Health Board, age and sex | integer | Health board lookup | e.g. 1508 | NA | NA |

5 Severe cases

5.1 Data information

| Description | EAVE II severe cases and outcomes |
|-------------------|---|
| Datafile | cases_severe_dates.rds |
| Type | Derived |
| Data sources | GP data, ECOSS, NRS Deaths, RAPID, SICSAG |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|------------------|---|--|-----------|-------------------------------|-------------------------------------|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | GP data | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| SPECIMENdate | Specimen date | Date of test | date | ECOSS | date in format yyyy-mm-dd | NA | NA |
| CURRENT_POSTCODE | Postcode | Postcode of residence | character | HPS case- control files | Postcode in pc7 format | NA | NA |
| is.case | is a covid case | Flag to indicate person is a covid case (+ve test and/or covid on death certificate) | logic | ECOSS, NRS deaths | TRUE/FALSE (should all be true) | NA | NA |
| DATE_OF_DEATH | Date of death | Date of patients death | date | NRS deaths | date in format yyyy-mm-dd | NA | NA |
| covid_ucod | COVID is underlying cause of death (COD) | Indicator or whether covid is underlying or primary COD on death certificate | logic | NRS deaths | 0,1 | NA | NA |
| covid_cod | Covid listed as any cause of death | Indicator or whether covid is listed as any COD on death certificate | logic | NRS deaths | 0,1 | NA | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|----------------|----------------------------|---|---------|-------------------------|---|---------------------|-------------------------|
| dead28 | Death within 28 days | Indicator of whether patient has died within 28 days of +ve covid Test | logic | ECOSS, NRS deaths | 0,1 | Yes | NA |
| inhosp | In hospital when test done | Indicator whether patient was already admitted to hospital at time of test | logic | ECOSS, RAPID | 0,1 | Yes | NA |
| icu | ICU admission | Indicator of whether patient was admitted to Intensive Care Unit (ICU) | logic | SICSAG | 0,1 | NA | NA |
| hdu | HDU admission | Indicator of whether patient was admitted to High Dependency Unit (HDU) | logic | SICSAG | 0,1 | NA | NA |
| RAPID | rapid flag | flag to indicate person in RAPID dataset | logic | RAPID | 0,1 | Yes | NA |
| ICU_date | ICU admission date | Date admitted to ICU | date | SICSAG | date in format yyyy-mm-dd | NA | NA |
| days_to_icu | Days to ICU | Number of days from test to admission to ICU | numeric | ECOSS, SICSAG | any +ve value | Yes | NA |
| admission.date | Hospital admission date | Date admitted to hospital | date | RAPID | date in format yyyy-mm-dd | NA | NA |
| days_2_hosp | Days to hospital | Number of days from test to admission to hospital | numeric | ECOSS, RAPID | any +ve value (if testedbefore admission, value changed to 0) | Yes | NA |

6 Household identifier

6.1 Data information

| Description | EAVE II Household from CHI registry |
|-------------------|-------------------------------------|
| Datafile | HID_lookup_MarSept2020.rds |
| Type | Derived |
| Data sources | CHILI |
| Scripts | NA |
| Update frequency | 6 months |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|------------------------------------|---|---|-----------|----------------|-------------------------------------|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | GP data | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| Date_address_changed.x | Address change date (March) | Date when address changed (March update) | numeric | CHILI | date as number yyyymmdd | NA | NA |
| March_hid | Household ID (March) | Household ID number (March update) | numeric | CHILI | ID | NA | NA |
| March_hid_count | Number of ID's in household (March) | Number of people in household (March update) | numeric | CHILI | any +ve whole number | NA | NA |
| ${\bf Date_address_changed.y}$ | Address change data (September) | Date when address changed (September update) | numeric | CHILI | date as number yyyymmdd | NA | NA |
| ${\it March_transfer_out_date}$ | Move or transfer date (March) | Date when resident moved out of scotland (March update) | date | CHILI | date | NA | NA |
| Sept_hid | Household ID (September) | Household ID number (September update) | numeric | CHILI | ID | NA | NA |
| Sept_hid_count | Number of ID's in household (September) | Number of people in household (September update) | numeric | CHILI | any +ve whole number | NA | NA |
| Sept_transfer_out_date | Move or transfer date (September) | Date when resident moved out of scotland (September update) | date | CHILI | date | NA | NA |

7 QCOVID deduplicated

7.1 Data information

| Description | EAVE II data for QCOVID validation analysis |
|-------------------|---|
| Datafile | QCOVIDdeduped.rds |
| Type | Bespoke |
| Data sources | GP data, ECOSS, HES, RTDS |
| Scripts | NA |
| Update frequency | NA |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Туре | Data source | Values | Derived variable | Derivation hierarchy |
|-------|--------|---------------|---------|----------------|--------|---------------------|-------------------------|
| rowid | Row ID | UID for batch | integer | n/a | n/a | NA | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|----------------------|--|--|---------|--|------------------|---------------------|-------------------------|
| sex | Sex | Sex | logic | GP data or Patient demo- graphic | 0:female; 1:male | NA | NA |
| age | Age | Age | integer | service GP data or Patient demo- graphic service | 19-100 | NA | NA |
| Q_RX_IMMUNO | Prescribed immunosup- pressants indicator | Have you been prescribed immunosup- pressants prescribed by your GP | logic | GP data | 0:false; 1:true | NA | NA |
| Q_RX_LABA | Anti- leukotriene or long acting beta2-agonists (LABA) indicator | Are you taking anti- leukotriene or long acting beta2-agonists (LABA)? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_RX_PRED | Oral steroid indicator | Have you been prescribed oral steroids by your GP in the last 6 months? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_AF | Atrial fibrillation indicator | Do you have atrial fibrillation? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_CCF | Heart failure indicator | Do you have heart failure? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_asthma | ${ m Asthma}$ indicator | Do you have asthma? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_blood_cancer | Cancer of the blood or bone marrow and are at any stage of treatment indicator | Have you a cancer of the blood or bone marrow such as leukaemia, myelodysplastic syndromes, lymphoma or myeloma and are at any stage of treatment? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_cerebralpalsy | Cerebral palsy indicator | Do you have cerebral palsy? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_chd | Coronary heart disease indicator | Do you have coronary heart disease? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_cirrhosis | Cirrhosis of the liver indicator | Do you have cirrhosis of the liver? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_Congen_HD | Congenital heart disease or had surgery for it in the past indicator | Do you have congenital heart disease or have you had surgery for it in the past? | logic | GP data or HES (OPCS) | 0:false; 1:true | NA | NA |
| Q_DIAG_COPD | Chronic obstructive pulmonary disease (COPD) indicator | Do you have chronic obstructive pulmonary disease (COPD)? | logic | GP data | 0:false; 1:true | NA | NA |
| $Q_DIAG_dementia$ | Dementia indicator | Do you have dementia? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_epilepsy | Epilepsy indicator | Do you have epilepsy? | logic | GP data | 0:false; 1:true | NA | NA |

 $\underline{(continued)}$

| Name | Label | Description | Туре | Data source | Values | Derived variable | Derivation hierarchy |
|---------------------|--|---|---------|----------------|--|---------------------|-------------------------|
| Q_DIAG_fracture | A prior fracture of hip, wrist, spine or humerus indicator | Have you had a prior fracture of hip, wrist, spine or humerus? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_neurorare | Motor neurone disease, multiple sclerosis, myaesthenia, or Huntingtons's Chorea | Do you have motor neurone disease, multiple sclerosis, myaesthenia, or Huntingtons's | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_parkinsons | indicator Parkinson's disease indicator | Chorea? Do you have Parkinson's disease? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_pulm_hyper | Pulmonary hypertension or pulmonary fibrosis indicator | Do you have pulmonary hypertension or pulmonary fibrosis? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_pulm_rare | Cystic fibrosis or bronchiectasis or alveolitis | Do you have cystic fibrosis or bronchiectasis or alveolitis? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_pvd | indicator Peripheral vascular disease indicator | Do you have peripheral vascular disease? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_ra_sle | Rheumatoid arthritis or SLE indicator | Do you have rheumatoid arthritis or SLE? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_respcancer | Lung or oral cancer indicator | Do you have lung or oral cancer? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_SEV_MENT_ILL | Severe mental illness indicator | Do you have severe mental illness? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_sickle_cell_ | Sickle cell disease or severe combined immune deficiency syndromes indicator | Do you have sickle cell disease or severe combined immune deficiency syndromes? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_stroke | A stroke or TIA indicator | Have you had a stroke or TIA? | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_DIABETES_1 | Diabetes indicator | Type 1 diabetes | logic | GP data | 0:none; 1:Type 1; 2:Type 2 | NA | NA |
| Q_DIAG_DIABETES_2 | Diabetes indicator | type 2 diabetes | logic | GP data | 0:false; 1:true | NA | NA |
| Q_DIAG_VTE | A thrombosis or pulmonary embolus indicator | Have you had a thrombosis or pulmonary embolus? | logic | GP data | 0:false; 1:true | NA | NA |
| BMI | BMI | Body Mass Index | numeric | GP data | 0-40 | NA | NA |
| ethnicity | Ethnic group | What is your ethnic group? | integer | GP data | 0-17 | NA | NA |
| homecat | Housing category | what is your housing category - care home or homeless or neither? | integer | GP data | 0: neither; 1: care home; 2: homeless | NA | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|----------------------|---|---|-----------|---|--|---------------------|-------------------------|
| learncat | Learning disability or Down's Syndrome indicator | Do you have a learning disability or Down's Syndrome? | integer | GP data | 0: neither; 1: learning disability; 2: Down's | NA | NA |
| p_marrow6 | Bone marrow or stem cell transplant in the last 6 months indicator | Have you had a bone marrow or stem cell transplant in the last 6 months? | logic | HES (OPCS) | 0:false; 1:true | NA | NA |
| p_radio6 | Radiotherapy in the last 6 months indicator | Have you had radiotherapy in the last 6 months? | logic | Radiotherap Data Set (RTDS) or HES (OPCS) | y0:false; 1:true | NA | NA |
| $p_solidtransplant$ | Solid organ transplant indicator | Have you had a solid organ transplant (lung, liver, somach, pancreas, spleen, heart or thymus)? | logic | HES | 0:false; 1:true | NA | NA |
| renalcat | Kidney disease indicator | Do you have kidney disease? | integer | GP data, HES (OPCS) | 0-5 | NA | NA |
| Postcode | Postcode | What is your postcode? | numeric | QCovid Postcode Table | double (nullable) | NA | NA |
| town | Town | Townsend score | character | QCovid Postcode Table | string (nullable) | Yes | NA |

8 Sgene linked

8.1 Data information

| Datafile Sgene_linked.rds | |
|--|-------|
| 0 = | |
| Type Bespoke | |
| Data sources GP data, ECOSS, RAPID, NRS d | eaths |
| Scripts NA | |
| Update frequency Weekly to fortnightly | |
| Update timestamps 2020-12-30, 2021-01-05, 2021-01-14 | 1 |
| Analyses used NA | |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|--------------|---------------------|--|-----------|----------------|-------------------------------------|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | GP data | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| ECOSSID | ECOSS ID | ECOSS ID | numeric | ECOSS | ECOSSID digits | NA | NA |
| age_year | Age | Age in years | numeric | ECOSS | 0-100 | NA | NA |
| sex | Sex | Sex | character | ECOSS | M/F | NA | NA |
| SpecimenDate | Specimen Date | Date sample taken | date | ECOSS | date in format | NA | NA |

| Name | Label | Description | Туре | Data source | Values | Derived variable | Derivation hierarchy |
|---|---|--|-----------|------------------------------------|---|---------------------|-------------------------|
| SubLab | Sub Lab | location sample processed | character | ECOSS | Lab ID | NA | NA |
| result | COVID result | result of covid test | numeric | ECOSS | 1 (only +ve tests in this file) | NA | NA |
| NRS.Date.Death | Date of NRS death record | Date of death | date | NRS deaths | date in format | NA | NA |
| death28 | Death within 28 days of positive test | Died within 28 days of +ve test | logic | ECOSS, NRS deaths | 0,1 | Yes | NA |
| simd2020 | SIMD Quintile 2020 | Quintile of scottish index of multiple deprivation (from postcode of residence) | numeric | ECOSS | 1-5 | NA | NA |
| admission_date | Admission to hospital date | Day admitted to hospital | date | RAPID | dates | NA | NA |
| discharge_date | Discharge from hospital date | Day discharged from hospital | date | RAPID | dates | NA | NA |
| covid_adm | COVID admission | Flag for covid admission | logic | ECOSS, RAPID | 0,1 | Yes | NA |
| time_test_covid_adm | Time from positive test to admission | Time (days) from +ve test to admission | numeric | ECOSS, RAPID | 0-14 | Yes | NA |
| covid dth | COVID death | Covid death (defined as death within 28 days of +ve test ?check if also includes covid on death certificate >28 days | logic | ECOSS, RAPID | 0,1 | Yes | NA |
| time_test_covid_dth | Time from poistive test to death | Time (days) from +ve test to death | numeric | ECOSS, RAPID | 0-28 | Yes | NA |
| days | Days of | Days of ? | numeric | ECOSS, RAPID | 0+ | Yes | NA |
| true_s_gene_dropout | True S Gene Dropout | Whether S gene positive, S gene negative or weak Sgene positive on PCR test. | character | ??ECOSS - need to check with Chris | Positive S Gene/True S Gene Dropout/Weak Positive/Unknown | NA | NA |
| new | New patient since last linkage | Flag for new compared to last linkage of gene dropouts | logic | ECOSS, RAPID | 0,1 | Yes | NA |

9 Vaccinations

9.1 Data information

| Description | COVID-19 vaccinations from GP system and Turas Vaccination Management Tool (TVMT) |
|-------------------|---|
| Datafile | C19vaccine.rds |
| Type | Cleaned |
| Data sources | GP data, Turas |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

9.2 Variable information

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-----------------|--------------------------|--|-----------|----------------|--|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | derived/ass | sigr EA VE(7)digits (e.g. EAVE1234567) | NA | NA |
| Run_Date | Date of extract | Date extract run | date time | GP | Date time | NA | NA |
| Date_of_birth | DOB | Date of birth | date | GP | Date | NA | NA |
| postcode | Postcode of residence | Postcode (if available) | character | GP | UK postcode | NA | NA |
| occurrence_time | Date of vaccination | Date of vaccination | date | GP | Date | NA | NA |
| type | Vaccine type | Vaccine type | character | GP | AstraZenica, Pfizer, Moderna | NA | NA |
| stage | Dose number | Dose number | numeric | GP | 1 or 2 (a few 0 , 3 or blank) | NA | NA |
| batch | Batch number | Batch number | character | GP | batch numbers of vaccines | NA | NA |
| site | Injection site | Injeciton site | character | GP | RUA (right upper arm), LUA, etc | NA | NA |
| method | Method of delivery | Method of delivery | character | GP | IM (intramuscular) | NA | NA |

10 Adverse events (clotting)

10.1 Data information

| Description | COVID-19 vaccine adverse events for clotting |
|-------------------|--|
| Datafile | AZ_safety |
| Type | Cleaned |
| Data sources | GP data |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-----------------|----------------------------|--|-----------|----------------|--|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | derived/assi | gr EA VE(7)digits (e.g. EAVE1234567) | NA | NA |
| ExtractDate | Date of data extraction | Date of data extraction | date | GP | Date | NA | NA |
| Practice_Number | GP practice number | GP practice number | character | GP | 5 digit code | NA | NA |
| Age | Age | Age at time of event | character | GP | 0-100 | NA | NA |
| Sex | Sex | Sex | character | GP | F, M | NA | NA |
| RegStatus | Registration status | Registration status at practice | character | GP | Registration code | NA | NA |
| RegDate | Registration date | Date registered at practice | date | GP | Date | NA | NA |
| IsDeducted | Registration deducted | Whether deducted (deregistered) | logic | GP | TRUE, FALSE | NA | NA |

| (continued | ١ |
|------------|---|

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|---------------|----------------|---|-----------|-----------------|---|---------------------|--|
| DeductionDate | Deduction date | Date deregistered | date | GP | Date | NA | NA |
| StartDate | Event date | Event date | date | GP | Date | NA | NA |
| Quick_groups | Cluster groups | Event type (each group icludes several read codes) | character | GP / derived | "Arterial_thromb", "cereb_haem", "CVT", "dv_throm", "haem", "haem_gigu" "haem_rb", , "haem_sc", "itp", "itp_gen", "not relevant", , "pulm_emb", "pv_throm", "sub_arach_haem" "SVT", "th_haem" "throm" | NA | For read codes included in groupings see code library: https://argoshare.is.ed.ac.uk |

11 Whole Genome Sequencing (WGS)

11.1 Data information

| Description | COVID-19 genome sequencing |
|-------------------|----------------------------|
| Datafile | genomic_dataset |
| Type | Cleaned |
| Data sources | WGS |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-----------------|-----------------------|---|-----------|----------------|--|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | derived/ass | igræAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| LabSpecimenNo | Lab number | Lab number - links to ECOSS to find CHI (and other info) | character | WGS | NA | NA | NA |
| query | Full Sequence ID | Full Sequence ID using within analysis server to allow for duplicated sequence in case of quality issues | character | WGS | NA | NA | NA |
| SequenceID | Sequence ID Number | Sequence ID number | character | WGS | NA | NA | NA |
| Collection_Date | Collection date | Date sample collected | date | WGS | NA | NA | NA |
| Sequencing_Date | Sequencing date | Date sample sequenced | date | WGS | NA | NA | NA |

 $\underline{(continued)}$

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-----------------------------|---------------------------|---|-----------|----------------|--------|------------------|-------------------------|
| Alignment_Date | Alignment date | The date the sequence is first available to PHS for analysis (used for sample tracking and calculating turn around times) | date | WGS | NA | NA | NA |
| adm2 | County of patient | County level based on patient's outer postcode | character | WGS | NA | NA | NA |
| ${\it HealthBoard}$ | Healthboard of patient | Healthboard as extracted from ECOSS. Typically based on residential postcode if present, otherwise, uses CHI seeding to determine heathboard. | character | WGS | NA | NA | NA |
| ${\bf Variant of Interest}$ | Variant of interest | whether VOC/VOI or not | character | WGS | NA | NA | NA |
| VariantShorthand | Variance shorthand | seems incomplete? - better to use lineage | character | WGS | NA | NA | NA |
| VariantCalling | Variant calling | Based on the PHE VUI/VOC genomic definition, there are thresholds within number of mutation / deletions present within your sequence that match the profile. HR-LQG is high risk low quality. | character | WGS | NA | NA | NA |
| ref_count | Counts of reference bases | Counts of reference bases (not mutations) | integer | WGS | NA | NA | NA |
| alt_count | Counts of mutations | Counts of mutations (given that the mutation is an expected one within the VUI/VOC genomic profile) | integer | WGS | NA | NA | NA |
| $other_count$ | Other counts | Counts of bases that are not reference or expected mutations | integer | WGS | NA | NA | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-----------------|---------------------------------|--|-----------|----------------|--------|---------------------|-------------------------|
| Ambiguous | Counts of ambiguous bases | Counts of bases which are ambiguous (typically due to low sequencing quality) | integer | WGS | NA | NA | NA |
| aa.S.L452R | Specific site mutation | Refers to specific site mutations wt = wild type, other letters are the mutation seen at this site. So L452R means that a site 452 you normally see the amino acid L, but the mutation of interest has an R in that site. Amb means ambiguous. | character | WGS | NA | NA | NA |
| aa.S.E484K | Specific site mutation | see above | character | WGS | NA | NA | NA |
| aa.S.G446V | Specific site mutation | see above | character | WGS | NA | NA | NA |
| Lineage | Lineage | Lineage (if applicable) in format e.g. B.1.1.7 | character | WGS | NA | NA | NA |
| Lineage_version | Lineage version | Date lineages last de- fined/assigned (as lineage deifinition can change over time) | date | WGS | NA | NA | NA |

12 EAVE weights

12.1 Data information

| Description | NA |
|-------------------|----|
| Datafile | NA |
| Type | NA |
| Data sources | NA |
| Scripts | NA |
| Update frequency | NA |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|-------------|---------------------|--|-----------|----------------|--|---------------------|-------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | derived/assi | gr EA VE(7)digits (e.g. EAVE1234567) | NA | NA |
| eave_weight | EAVE II weight | Weight assigned to individual person based on NRS mid-year estimates | numeric | derived | 0-1 | NA | NA |

13 Demographic endpoints

13.1 Data information

| Description | Summary of all outcomes and baseline characteristics |
|-------------------|--|
| Datafile | Cohort_Demog_Endpoints_Dates.rds or Cohort_Demog_Endpoints_Times.rds |
| Type | Derived |
| Data sources | GP data, ECOSS, RAPID, NRS deaths, SICSAG |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|--------------------------|------------------------------|---|-----------|----------------|--|---------------------|------------------------------|
| EAVE_LINKNO | EAVE Link Number | Randomly assigned study index number | character | derived/assi | græAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| Sex | Sex | Sex | character | GP data | F, M | NA | NA |
| ageYear | Age | Age as individual year | integer | GP data | 0-100 | NA | NA |
| $simd2020_sc_quintile$ | SIMD Quintile (2020) | Scottish Index of Multiple Deprivation Quintile (2020) | character | GP data | 1 - High, 2, 3, 4, 5 - Low | Yes | Postcode to SIMD quintile |
| DataZone | Data Zone | Data Zone | character | GP data | Data zone codes | NA | NA |
| ur6_2016_name | Urban Rural Index (2016) | Urban Rural Index (2016) | character | GP data | 1 Large Urban Area, 2 Other Urban Areas, 3 Accessible Small Towns, 4 Remote Small Towns, 5 Accessible Rural, 6 Remote Rural | Yes | Postcode to UR |
| HealthBoard | NHS Scotland health board | NHS Scotland health board | character | GP data | NA | NA | NA |
| Hospital | NHS facility of test | Location where COVID test took place (NHS Facility) | character | ECOSS | NA | NA | NA |
| SpecimenDate | Speciment date | Date of COVID-19 test | date | ECOSS | Date | NA | NA |
| OriginalOrganism | Test result (formal) | Result of COVID-19 test (formal) | character | ECOSS | Coronavirus 2019-nCoV,Coronavirus 2019-nCoV negative | NA | NA |
| result | Test result | Result of COVID-19 test | logic | ECOSS | 0, 1 | NA | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|--|--|--|-----------|--|--------|------------------|-------------------------|
| death28 | Death with 28 days of test | Indicator if individual died within 28 days of COVID-19 test | integer | ECOSS, NRS deaths | 0, 1 | NA | NA |
| SubLab | Sub Lab | Location where sample processed (lab code) | character | ECOSS | NA | NA | NA |
| NRS.Date.Death | Date of death | Date of death | date | $ \begin{array}{c} \text{NRS} \\ \text{deaths} \end{array} $ | Date | NA | NA |
| NRS.Reg.Date | Date of death registered | Date of death registered | date | NRS deaths | Date | NA | NA |
| is.case | COVID-19 case | Indicator if positive COVID case | logic | ECOSS | 0, 1 | NA | NA |
| death_covid | COVID death | Indicator if death was caused by COVID-19 | integer | NRS deaths | 0, 1 | NA | NA |
| $\operatorname{covid}\operatorname{_cod}$ | COVID as any cause of death | COVID as any cause of death | integer | NRS deaths | 0, 1 | NA | NA |
| DATE_OF_DEATH | Date of death | Date of death | date | NRS deaths | Date | NA | NA |
| icu_death | Died in ICU | Indicator if individual died in ICU | logic | SICSAG, NRS deaths | 0, 1 | NA | NA |
| date_icu_death | Date of ICU death | Date of ICU death | date | SICSAG, NRS deaths | Date | NA | NA |
| hosp_covid | Hospitalised with COVID | Indicator if individual was hospitalised with COVID-19 | logic | RAPID | 0, 1 | NA | NA |
| date_hosp_covid | Date of COVID-19 hospital admission | Date of hospital admission for COVID-19 | date | RAPID | Date | NA | NA |
| days_to_death | Days from positive test to death | Days from positive test to death | numeric | ECOSS, NRS deaths | NA | NA | NA |

14 Previous tests

14.1 Data information

| Description | Individuals tested for COVID-19 with a summary of their results |
|-------------------|---|
| Datafile | Tests.rds |
| Type | Cleaned |
| Data sources | ECOSS |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|------------------|--|--------------------------------------|-----------|----------------|-------------------------------------|---------------------|-------------------------|
| EAVE Link Number | Randomly assigned study index number | character | character | derived | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| n_tests | Number of COVID tests | Number of COVID tests | integer | ECOSS | NA | NA | NA |
| n_pos | Number of positive COVID tests | Number of positive COVID tests | integer | ECOSS | NA | NA | NA |
| n_neg | Number of negative COVID tests | Number of negative COVID tests | integer | ECOSS | NA | NA | NA |

15 Previous positive tests

15.1 Data information

| Description | Individuals with a positive COVID-19 test |
|-------------------|---|
| Datafile | Positive_Tests.rds |
| Type | Cleaned |
| Data sources | ECOSS |
| Scripts | NA |
| Update frequency | Weekly to fortnightly |
| Update timestamps | NA |
| Analyses used | NA |

15.2 Variable information

| Name | Label | Description | Type | Data source | Values | Derived variable | Derivation hierarchy |
|------------------|--|--------------------------------------|-----------|----------------|-------------------------------------|---------------------|-------------------------|
| EAVE Link Number | Randomly assigned study index number | character | character | derived | EAVE(7)digits (e.g. EAVE1234567) | NA | NA |
| $specimen_date$ | Speciment date | Date of positive COVID-19 test | date | ECOSS | Date | NA | NA |

16 Supplementary Information

All supplementary information are stored in the Supplementary GitHub folder. This contains the following:

| Name | Description | File |
|----------------------------------|--|--|
| GP Cluster ReadCodes | Links each GP risk group cluster to the GP ReadCodes within the GP IT system | $\overline{EAVEII_dd_gp_clusters.xlsx}$ |
| Data source external information | Provides extra information and links on linked data sources | $EAVEII_dd_external_info.xlsx$ |