ISAC APPLICATION FORM PROTOCOLS FOR RESEARCH USING THE CLINICAL PRACTICE RESEARCH DATALINK (CPRD)

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PRJ	Study Title PRJ2282: CHESS: CPRD-COPD Hawthorn Effect Study in Salford: A UK cohort study to characterise patients enrolled in the Salford Lung Study and to evaluate a potential Hawthorne effect								
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	Affiliation (full		<u>in Health Data S</u> s)	cience, uni	versity or	Mancheste	r, <u>Matthewis</u>	<u>sperrin@manc</u>	<u>mester.ac.uk</u>
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9.	Has this proto	col beer	n peer reviewed b	by another	Committe	e?			
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	Reviewing Committee: The protocol was reviewed by GSK Worldwide Epidemiology Protocol Review Forum (GSK WWEpi PRF).								
Committee review process: GSK WWEpi PRF review and approval is required for all studies conducted under the sponsorship of GSK Worldwide Epidemiology and is a binding procedure within GSK under SOP_72625(3.0). The									
review committee is chaired by the Vice-President (VP) of WWEpi (or their delegate) and the committee is composed of									
Hea	GSK staff from the following roles: 2 Epidemiologists; 1 Observational Data Analyst; 1 Medically Qualified Member; 1 Health Outcomes Advisor (optional); 1 Statistician and 1 Secretary. All members (excluding secretary) are expected to								
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11. This study is intended for:				
Publication in peer reviewed journals Presentation at company/institutional meetings	=	itation at s	scientific conference	

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19. Does this stu	udy involve linking	to patient <i>identif</i>	<i>īable</i> data from oth	ner sources	?	
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22. Experience/e	expertise available	2				
actively involved	in the proposed r	esearch, including	analysis of data a	and interpre		esearchers
Previous	s GPRD/CPRD Stu	dies	Publications usir	ng GPRD/CP	PRD data	
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		thin the research t			Yes	No
All applicants hav	ve advanced statis	stical expertise – e	.g. all have PhDs <u>c</u>	or equivaler	nt with statistical compor	ient.
Is experience of handling large data sets (>1 million records) available within the research team?						
If yes, please outline level of experience The study will be based within the Centre for Health Informatics at the University of Manchester, which has substantial infrastructure and experience for handling large data sets. Both University of Manchester and GSK already have CPRD						

licer	licences and applicants have been using CPRD actively.					
	Is UK primary care experience available within the research team? If yes, please outline level of experience All applicants are actively researching in primary care.					
23.	References relating to your study					
Plea	Please list up to 3 references (most relevant) relating to your proposed study.					
	New JP, Bakerly ND, Leather D, Woodcock A. Obtaining real-world evidence Apr 26 e pub ahead of print.	e: the Salford Lung Study.	Thorax. 2014			
2.	van Staa T-P, Goldacre B, Gulliford M, Cassell J, Pirmohamed M, Taweel A, randomised trials using routine electronic health records: putting them to the 10.1136.	• •	-			
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PROTOCOL CONTENT CHECKLIST

In order to help ensure that protocols submitted for review contain adequate information for protocol evaluation, ISAC have produced instructions on the content of protocols for research using CPRD data. These instructions are available on the CPRD website (www.cprd.com/ISAC). All protocols using CPRD data which are submitted for review by ISAC must contain information on the areas detailed in the instructions. IF you do not feel that a specific area required by ISAC is relevant for your protocol, you will need to justify this decision to ISAC.

Applicants must complete the checklist below to confirm that the protocol being submitted includes all the areas required by ISAC, or to provide justification where a required area is not considered to be relevant for a specific protocol. Protocols will not be circulated to ISAC for review until the checklist has been completed by the applicant.

Please note, your protocol will be returned to you if you do not complete this checklist, or if you answer 'no' and fail to include justification for the omission of any required area.

	Included in protocol?		
Required area	Yes	No	If no, reason for omission
Lay Summary (max.200 words)			
Background			
Objective, specific aims and rationale			
Study Type Descriptive Hypothesis Generating Hypothesis Testing			
Study Design	\boxtimes		
Sample size/power calculation (Please provide justification of sample size in the protocol)			
Study population (including estimate of expected number of relevant patients in the CPRD)			
Selection of comparison group(s) or controls			
Exposures, outcomes and covariates Exposures are clearly described Outcomes are clearly described			
Use of linked data (if applicable)	\boxtimes		
Data/ Statistical Analysis Plan There is plan for addressing confounding There is a plan for addressing missing data			
Patient/ user group involvement †			This is a purely observational study, and quality of life will not be evaluated.
Limitations of the study design, data sources and analytic methods			
Plans for disseminating and communicating study results			

Voluntary registration of ISAC approved studies:

Epidemiological studies are increasingly being included in registries of research around the world, including those primarily set up for clinical trials. To increase awareness amongst researchers of ongoing research, ISAC encourages voluntary registration of epidemiological research conducted using MHRA databases. This

[†] It is expected that many studies will benefit from the involvement of patient or user groups in their planning and refinement, and/or in the interpretation of the results and plans for further work. This is particularly, but not exclusively true of studies with interests in the impact on quality of life. Please indicate whether or not you intend to engage patients in any of the ways mentioned above.

will not replace information on ISAC approved protocols that may be published in its summary minutes or annual report. It is for the applicant to determine the most appropriate registry for their study. Please inform the ISAC secretariat that you have registered a protocol and provide the location.

PRJ2282: CHESS: CPRD-COPD Hawthorn Effect Study in Salford: A UK cohort study to characterise patients enrolled in the Salford Lung Study and to evaluate a potential Hawthorne effect

Lay Summary

The Salford Lung Study (SLS) is an evaluation of a new drug, RELVAR, taking place in Salford, UK. RELVAR is a drug that treats chronic obstructive pulmonary disease (COPD). The study is unique in that it is taking place in routine care, so it is hoped that the results will show how effective the drug is in the 'real world'.

However, one issue with a study of this kind is that the fact that the study is taking place raises awareness of COPD in both patients and clinicians taking part. This could lead to the 'Hawthorne effect', in which patients in both the group receiving RELVAR, and the group receiving usual care, do better than the general population.

In this study we aim to compare outcomes for patients in the SLS with other COPD patients in the UK who are not in the study, to see whether the Hawthorne effect occurs and its severity. We will also assess whether the COPD patients in the SLS are similar to the rest of the UK. The conclusions will help us understand the extent to which the findings of the SLS can be generalised to the rest of the UK.

Background

COPD is a chronic obstructive disease of the airways associated with a significant social and healthcare burden [1, 2, 3]. Most patients with COPD are managed in primary care, as reflected in recent UK guidelines, which are specifically targeted at primary care physicians [4]. The major goals of treatment are to relieve symptoms, improve activity/exercise tolerance, prevent and treat exacerbations, reduce mortality risk and improve health status. However, despite such guidelines, COPD remains under-diagnosed and under-treated; variations in treatments, standards of care and adherence to guidelines have been reported across different geographical regions [2, 5, 6, 7, 8].

Large computerised patient databases provide a useful source of real life observational data, and the General Practice Research Database (GPRD) has been successfully used to generate descriptive epidemiology data in COPD [9, 10, 11, 12] from a large group of UK primary care practices. GPRD has also been used previously for descriptive drug utilization studies for prescription medications in respiratory diseases [16, 17, 18]. Historically, the limitations of the GPRD were a time gap between data capture and availability for the researcher and limited links to other healthcare databases, although these are currently being addressed with the development of the Clinical Practice Research Datalink (CPRD) and in recent Phase 4 pragmatic clinical trials [12]. The use of electronic medical record (EMR) data in health research is a key objective in the Department of Health's national research strategy [13].

The SLS is an ongoing Phase IIIB pragmatic trial comparing a new once daily ICS/LABA fixed dose combination (RELVAR: fluticasone furorate + vilanterol) in which patients are identified by EMR, enrolled in their GP office, randomized to

RELVAR or standard of care (SOC) maintenance therapy via their neighbourhood pharmacy, and followed for safety and effectiveness via linked primary care/secondary routine care with the primary endpoint of COPD moderate/severe exacerbations over 12 months [14]. MHRA and NICE provided joint advice for the SLS protocol and were supportive of the design to generate "real world" evidence which will demonstrate the value of the medicine against the most relevant standard of care.

Although the SLS will give evidence on the relative effectiveness of RELVAR compared with SOC, the SOC may be prone to the Hawthorne effect, which may distort the effect size.

The Hawthorne effect (also referred to as the observer effect) is a type of <u>reactivity</u> in which individuals improve or modify an aspect of their behaviour in response to their awareness of being observed. The original "Hawthorne effect" studies at the <u>Hawthorne Works</u> in Chicago, USA between 1924 and 1933 suggested that the novelty of being research subjects and the increased attention from such could lead to temporary increases in workers' productivity [15].

Additionally, Salford may not be representative of general UK population; hence the prognostic profiles, and potential effect of RELVAR in terms of the outcomes may differ in the general target COPD population compared with Salford.

Both of these issues mean that extrapolation of the results of the SLS to the wider UK population would be subject to major caveats. This proposal aims to explore, and assess how severe the caveats need to be.

Objectives

Co-primary objectives:

PO1: To characterize the patients enrolled in the SOC arm of the SLS compared with the UK population of COPD patients (using CPRD linked data), including the distribution of deprivation levels, to evaluate the extent to which the SLS participants are representative of the UK patient population targeted for RELVAR. The comparator set will be specified on two bases: firstly, overall, and secondly, the subset fulfilling the protocol inclusion/exclusion criteria.

PO2: To compare the rate of COPD exacerbation over the 12 months in the SOC arm of the SLS compared with the SOC recorded in the CPRD linked data, in order to detect a potential Hawthorne effect as a result of potentially increased attention paid by GPs and nurses to patients with COPD during the SLS study period.

PO3: To compare the rate of severe pneumonia over the 12 months in the SOC arm of the SLS compared with the SOC recorded in the CPRD linked data. Severe pneumonia is a safety signal being assessed in the SLS.

Secondary objectives:

SO1: To make comparisons between the SLS Standard of care and the CPRD cohort on the following health care utilisation (HCU) endpoints: GP visits, hospital admissions, mortality and adherence.

SO2: To evaluate other definitions of COPD exacerbation in SOC from CPRD.

SO3: Self-controlled comparison of COPD and other HCU endpoints in Salford before and after SLS commenced.

Study Type

The study will primarily be descriptive.

Study Design

This will be an observational COPD cohort study that will utilize the CPRD linked data and the Salford EHR system to compare selected cohorts with SLS.

For SLS, the reference/index date is study entry. This will be matched in the CPRD cohort by the following algorithm:

- 1. Draw up a long-list of potentially eligible individuals in the CPRD linked data (patients who would be eligible at some point during the SLS recruitment phase).
- 2. For each individual:
 - a. Randomly sample an entry date from full list of SLS entry dates.
 - b. If patient is eligible at that entry date, then they will be included, otherwise, they will be excluded.

First, the COPD populations in both Salford and in the wider CPRD cohort (excluding Greater Manchester area) will be compared. Second, we will focus on comparisons between all patients enrolled in the SLS versus the larger CPRD defined COPD cohort "eligible for RELVAR prescription" (target patient population per label). Third, we will compare the COPD exacerbation response between eligible CPRD population, and the Salford population in SLS receiving SOC, in a multilevel model – i.e. evaluating Hawthorne effect.

Sample Size

The target sample size for the number of COPD patients enrolled in the SLS is 2,800. A preliminary feasibility count using the February 2015 version of the CPRD GOLD database identified 265,076 patients with a Read code for COPD in their clinical or referral files (see Annex 1 for codelist), with at least one day of up-to-standard registration aged 40+, and eligible for linkage to Hospital Episode Statistics (HES), ONS mortality data and deprivation data. This is a large number of cases and as the study is primarily descriptive, a power calculation is not required.

Data Linkage

Linkages to inpatient HES data, Index of Multiple Deprivation (IMD) data and ONS mortality data will be used in this study. HES will be used to identify two of the primary endpoints - hospital admissions for COPD exacerbation and severe pneumonia. IMD data will be used adjust for confounding due to deprivation. ONS mortality data will be used to censor at death. The entire study will be undertaken among linked practices only.

Fully linked data will only be available up to a certain date when analyses are undertaken (currently March 2014), and as such, primary analyses will be restricted to include SLS enrolled patients up to that date. Subsequent analyses will be conducted once linkage is available for the entire recruitment period.

Study Population

Two cohorts will be produced. First, a CPRD cohort, using linked primary care, medication, Hospital Episode Statistics, and socio-economic data, according to the following inclusion criteria:

- 1. Diagnosis of COPD before index date
- 2. Aged \geq 40 at index date.
- 3. Alive, and registered with a GP, at index date.
- 4. Not registered with a GP in the Greater Manchester area.

Second, a Salford cohort will be constructed using the Salford Integrated Record (SIR). The SIR is a comprehensive primary and secondary care database detailing healthcare contacts, diagnostic tests and prescriptions of all patients registered with a GP in Salford, UK. Patients will be included according to the following criteria:

- 1. Diagnosis of COPD before index date.
- 2. Aged \geq 40 at index date.
- 3. Alive, and registered with a GP, at index date.

Restricted cohorts will then be constructed in both the Salford and CPRD populations, based on the inclusion/exclusion criteria and study period for the SLS:

- 1. Patients with documented GP diagnosis of COPD, and currently receiving maintenance therapy
- 2. Male or female subjects aged ≥40 years of age at index date
- 3. Patients who have a history of treatment with systemic/oral corticosteroids, antibiotics (in association with GP contact) and/or hospitalisation for at least one COPD exacerbation in the 3 years prior to index date.
- 4. Current COPD Therapy: all patients currently receiving either:
 - inhaled corticosteroid (ICS) alone or in combination with a long acting bronchodilator (this could be a fixed dose combination or an ICS/LABA provided in two separate inhalers, or ICS and LAMA),
 - or long-acting bronchodilator therapy alone (e.g. tiotropium or salmeterol, or the use of two bronchodilators i.e. LABA/LAMA),
 - or "triple therapy" i.e. ICS/LABA plus a Long Acting Muscarinic Antagonist (LAMA)

Finally, the third data source, the SLS, will be used as-is. The SLS is described in [14]. In brief, it is a pragmatic trial, carried out in Salford, UK, to evaluate the relative effectiveness of RELVAR compared with SOC.

Subjects meeting any of the following criteria will be excluded from the restricted cohorts:

- 1. Patients with any life threatening condition or uncontrolled/clinically significant disease
- 2. Patients with unstable COPD: Patients with an exacerbation (defined by treatment with oral corticosteroids and/or antibiotic or hospital discharge listing COPD) with an onset within 2 weeks of index date.
- 3. Chronic user of oral corticosteroids: Subjects who are considered to be a chronic user of oral corticosteroids for respiratory or other indications
- 4. In the Salford population only, those patients who are entered in the SLS and randomised to the RELVAR arm.

Exposures

This is a binary comparison of COPD patients enrolled in the SLS and COPD patients in the CPRD cohort. Hence the primary exposure of interest is whether a patient is enrolled in SLS (yes/no). A third grouping, COPD patients in Salford (who are not in SLS) will also be examined.

Outcomes

Primary outcomes/endpoint:

- Rate of COPD exacerbation: Moderate/severe COPD exacerbations will be identified using an algorithm combining GP visits, prescriptions for oral corticosteroids and/or antibiotics, or hospital admission, as defined using information from the ongoing study being conducted by Jenny Quint et al. (collaborative project between London School of Hygiene and Tropical Medicine and GSK). The rate of exacerbation during the 12 month follow-up will be calculated and compared with the SLS rate in the standard of care arm; if technically possible, exacerbation rates for the 12 months prior to index date (matched enrolment date) will also be compared.
- Pneumonia: To be defined as per the codelist used in SLS see Annex 2 and 3

Secondary outcomes/endpoints:

- Healthcare utilisation: All GP visits/encounters and all hospital admissions during the 12 month follow-up.
- Adherence to index prescription: Defined as percent days covered (PDC) and medication possession ratio (MPR) will also be calculated for the matched cohort, as well as discontinuation, switching medicine or adding on other medicines, to be compared with the SLS SOC arm.
- Deaths: All cause, pneumonia death, COPD-attributed death during the 12 month follow-up. For the CPRD, deaths will be determined using Office of National Statistics (ONS) linked mortality data.
- Other definitions of COPD exacerbation: Other definitions will be described as per the outputs of Jenni Quint's study.

Covariates

Covariates will include:

- demographic variables (sex, age, IMD)
- current exposure to LAMA, LABA and ICD
- comorbidities (heart failure, myocardial infarction, stroke, depression, anxiety, asthma, pneumonia, gastro-oesophageal reflux and peptic ulcer disease, Charlson score (COPD will be removed from score), disability)
- markers of COPD severity (previous COPD exacerbation, FEV1 % predicted, FEV1/FVC ratio, GOLD stage, MRC Dyspnoea score)
- co-medications and vaccinations
- lifestyle behaviours (smoking status, BMI)

Analysis

All analyses will be conducted using SAS.

For PO1, distributions of the confounders and effect modifiers will be tabulated, summarised as proportions in each category for binary and categorical variables, and means/medians and standard deviations for continuous variables. Graphical visualisations will also be produced to aid interpretation (for example, boxplots to characterise age distributions in each population, stacked bar charts to visualise deprivation by population). This will be done for a series of the derived populations to separate out true differences in demographics in Salford and differences that arise as a consequence of data quality issues etc.

There are a range of subsets and derivations of the data sources that will be considered for this study, listed here for clarity.

CPRD: all CPRD COPD patients

CPRD-GM: CPRD COPD patients registered in practices in Greater

Manchester

CPRD-xGM: CPRD COPD patients registered in non-Greater Manchester

practices

CPRD-GM-IC: CPRD COPD patients registered in practices in Greater

Manchester meeting the SLS inclusion criteria

CPRD-xGM-IC: CPRD COPD patients registered in non-Greater Manchester

practices meeting the SLS inclusion criteria

SLS-E: SLS – all eligible. Not all of these are enrolled (some decline)

SLS: SLS – all enrolled
SLS-SOC: SLS – SOC arm only
SIR: SIR – all COPD patients

SIR-IC: SIR - COPD patients meeting SLS inclusion criteria only

The following comparisons will be of interest:

- CPRD-GM v CPRD-xGM: to give an indication of true demographic difference from the same data source.
- CPRD-GM-IC v CPRD-xGM-IC: as above, but restricted to patients meeting the inclusion criteria.
- SIR v CPRD-GM: to give an indication of differences arising as a consequence of selection bias of CPRD practices, and through data quality issues etc.

- SIR-IC v CPRD-GM-IC: as above, but restricted to patients meeting the inclusion criteria.
- SIR-IC v SLS-E: to give an indication of recruitment bias and physician researcher bias (at the approach stage).
- SIR-IC v SLS: to give an indication of recruitment bias (at recruitment stage).
- SLS v CPRD-xGM-IC: to indicate the difference between trial recruited, and those meeting the inclusion criteria outside of Salford. This is the key comparison for addressing PO1.

We will then move on to explicit modelling of regional variation of the characteristics within the CPRD cohort to ascertain whether the characteristics observed within Salford are unusual by comparison with CPRD-xGM-IC. We will take local authority regional level (anonymised) as the comparable unit to the study region for SLS. SLS will be considered unusual on a given demographic if an appropriately chosen summary statistic for that demographic (mean for continuous variables) falls outside the 2.5-97.5 percentile range.

Missing data in confounders and effect modifiers will be imputed using multiple imputation for the primary analysis, with complete case analysis included as sensitivity analysis.

For PO2, we will commence with exploratory analyses, similar to described above, to explore the distributions of the primary and secondary endpoints. The Hawthorne effect will be evaluated in two different ways.

Firstly, for descriptive purposes, we will measure the prevalence of the endpoints in a series of subgroups. For example, we will compare the COPD exacerbations in CPRD-xGM-IC with SLS, stratified by SES, gender, etc.

Secondly, we will take a multilevel modelling approach. For this we will combine the SLS and CPRD into one dataset (retaining an indicator of SLS membership). The hierarchies of the model will be patient -> GP practice -> local authority region (with SLS members being treated as a single distinct region) -> strategic health authority region. Strategic health authorities (population threshold of 1 million) are non-anonymised (named) regions. Local authority regions are below the population threshold so an anonymised LA marker will be available.

We will include all confounders and effect modifiers as covariates, with outcomes corresponding to the primary and secondary study outcomes (a separate model for each). Important fixed effects at the local authority level (for example, existence of community teams) will be incorporated into the model if these can be ascertained. As above, confounders and effect modifiers will be imputed using multiple imputation for the primary analysis, with complete case analysis included as sensitivity analysis.

A final model will be selected via backward selection using AIC (Akaike Information Criterion). We will then examine the random effect of the SLS region in the context of the random effects for the other regions. Similar to the above, if the random effect of the SLS region falls outside the 2.5-97.5 percentile range, we will conclude that SOC SLS behaves unusually compared with the rest of the UK, and hence evidence of a Hawthorne effect.

PO3, SO1 and SO2 (which pertain to comparing other endpoints and sensitivity analyses of endpoint definition) will be carried out using the same approach as for PO2.

SO3 makes explicit the possible change in outcomes at commencement of SLS. We will compare outcome rates within Salford before and after the commencement of SLS, in a self-controlling case design. We will do the same thing within the CPRD cohort to control for UK-wide secular trends. This acts as sensitivity analysis to support PO2 (using controls distinct in time rather than in geography).

The QC of analysis will be performed by GSK, in accordance with GSK Standard Operating Procedures (SOPs) and Guidance Documents, specifically the SOP_52213 (4.0): Conducting Quality Control Review of Worldwide Epidemiology Study Results. Wherever feasible, all statistical programming will be independently reviewed by a second analyst, with oversight by a senior statistician.

Limitations

The Hawthorne effect can only be evaluated for the SOC comparison. This does not give definite evidence about whether the prognostic or predictive effect of RELVAR would differ in the general population. This information could only truly be obtained following use of RELVAR in the general population.

There is no direct metric by which 'representativeness' of the Salford cohort can be measured. Early explorations of this will be made in a companion project outside the scope of this protocol.

While GP data are the same, some data (hospital validated COPD diagnoses, pharmacy data) for participants in SLS are collected using a different mechanism to CPRD GOLD. Hence any differences (either in representativeness or treatment response) observed between the SLS and non-SLS cohorts could be attributed to differences in data quality and the data collection mechanism. This will be mitigated by an additional comparison of SLS data with CPRD GOLD data from within Greater Manchester.

Plans for Dissemination

This work is targeted 1) internally at GSK, 2) regulators and 3) the wider scientific community; in order to understand the SLS in wider context. Results will be disseminated externally primarily by manuscripts, including a submission to Thorax in Q2 2016 based on interim results, and followed by a subsequent manuscript containing the final analyses of the data (planned for Q4 2016). In addition, we will present the results of the study at international respiratory conferences as appropriate. The study protocol and results will be posted to GSK Clinical Study Register as per GSK SOPs.

References

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Annex 1 : Read codes to be used to identify COPD

Read Code	Read Term
66YB.00	Chronic obstructive pulmonary disease monitoring
66YB000	Chronic obstructive pulmonary disease 3 monthly review
66YB100	Chronic obstructive pulmonary disease 6 monthly review
66Yd.00	COPD accident and emergency attendance since last visit
66YD.00	Chronic obstructive pulmonary disease monitoring due
66Ye.00	Emergency COPD admission since last appointment
66Yf.00	Number of COPD exacerbations in past year
66Yg.00	Chronic obstructive pulmonary disease disturbs sleep
66Yh.00	Chronic obstructive pulmonary disease does not disturb sleep
66YI.00	COPD self-management plan given
66Yi.00	Multiple COPD emergency hospital admissions
66YL.00	Chronic obstructive pulmonary disease follow-up
66YL.11	COPD follow-up
66YM.00	Chronic obstructive pulmonary disease annual review
66YS.00	Chronic obstructive pulmonary disease monitoring by nurse
66YT.00	Chronic obstructive pulmonary disease monitoring by doctor
8BMW.00	Issue of chronic obstructive pulmonary disease rescue pack
8BP8.00	Antibiotic therapy for acute pulmonary exacerbation
8CE6.00	Chronic obstructive pulmonary disease leaflet given
8CR1.00	Chronic obstructive pulmonary disease clini management plan
8H2R.00	Admit COPD emergency
9h52.00	Excepted from COPD quality indicators: Informed dissent
9kf00	COPD - enhanced services administration
9kf0.00	COPD patient unsuitable for pulmonary rehab - enh serv admin
9kf0.11	COPD patient unsuitable for pulmonary rehabilitation
9kf1.00	Refer COPD structured smoking assessment - enhanc serv admin
9kf1.11	Referred for COPD structured smoking assessment
9kf2.00	COPD structured smoking assessment declined - enh serv admin
9kf2.11	COPD structured smoking assessment declined
90i00	Chronic obstructive pulmonary disease monitoring admin
90i0.00	Chronic obstructive pulmonary disease monitoring 1st letter
9Oi1.00	Chronic obstructive pulmonary disease monitoring 2nd letter
9Oi2.00	Chronic obstructive pulmonary disease monitoring 3rd letter
9Oi3.00	Chronic obstructive pulmonary disease monitoring verb invite
9Oi4.00	Chronic obstructive pulmonary disease monitor phone invite
H300	Chronic obstructive pulmonary disease
Н311	Chronic obstructive airways disease
H3000	Bronchitis unspecified
H3011	Chest infection - unspecified bronchitis
H3012	Recurrent wheezy bronchitis
H300.00	Tracheobronchitis NOS
H301.00	Laryngotracheobronchitis
H302.00	Wheezy bronchitis
H30z.00	Bronchitis NOS
H3100	Chronic bronchitis
H310.00	Simple chronic bronchitis
H310000	Chronic catarrhal bronchitis

H310100	Smokers' cough
H310z00	Simple chronic bronchitis NOS
H311.00	Mucopurulent chronic bronchitis
H311000	Purulent chronic bronchitis
H311100	Fetid chronic bronchitis
H311z00	Mucopurulent chronic bronchitis NOS
H312.00	Obstructive chronic bronchitis
H312000	Chronic asthmatic bronchitis
H312011	Chronic wheezy bronchitis
H312100	Emphysematous bronchitis
H312200	Acute exacerbation of chronic obstructive airways disease
H312300	Bronchiolitis obliterans
H312z00	Obstructive chronic bronchitis NOS
H313.00	Mixed simple and mucopurulent chronic bronchitis
H31y.00	Other chronic bronchitis
H31y000	Chronic tracheitis
H31y100	Chronic tracheobronchitis
H31yz00	Other chronic bronchitis NOS
H31z.00	Chronic bronchitis NOS
H3200	Emphysema
H320.00	Chronic bullous emphysema
H320000	Segmental bullous emphysema
H320100	Zonal bullous emphysema
H320200	Giant bullous emphysema
H320300	Bullous emphysema with collapse
H320311	Tension pneumatocoele
H320z00	Chronic bullous emphysema NOS
H321.00	Panlobular emphysema
H322.00	Centrilobular emphysema
H32y.00	Other emphysema
H32y000	Acute vesicular emphysema
H32y100	Atrophic (senile) emphysema
H32y111	Acute interstitial emphysema
H32y200	MacLeod's unilateral emphysema
H32yz00	Other emphysema NOS
H32z.00	Emphysema NOS
H3600	Mild chronic obstructive pulmonary disease
H3700	Moderate chronic obstructive pulmonary disease
H3800	Severe chronic obstructive pulmonary disease
H3900	Very severe chronic obstructive pulmonary disease
H3A00	End stage chronic obstructive airways disease
H3y11	Other specified chronic obstructive pulmonary disease
H3y0.00	Chronic obstruct pulmonary dis with acute lower resp infectn
H3y1.00	Chron obstruct pulmonary dis with acute rower resp infection.
H3z00	Chronic obstructive airways disease NOS
H3z11	Chronic obstructive pulmonary disease NOS
11,72,11	chrome obstructive pullionary disease 1105

Annex 2: ICD-10 codes to be used to identify pneumonia

ICD-10 code	ICD-10 description
A06.5	Amoebic lung abscess
A15*	Respiratory TB bacteriologically and histologically
	confirmed
A16*	Respiratory TB not confirmed bacteriologically or
	histologically
A19*	Miliary tuberculosis
A20.2	Pneumonic plague
A21.2	Pulmonary tularaemia
A22.1	Pulmonary anthrax
A31.0	Pulmonary mycobacterial infection
A42.0	Pulmonary actinomycosis
A43.0	Pulmonary nocardiosis
A48.1	Legionnaires' disease
B01.2	Varicella pneumonia
B05.2	Measles complicated by pneumonia
B20.6	HIV disease resulting in Pneumocystis carinii pneumonia
B25.0	Cytomegaloviral pneumonitis
B37.1	Pulmonary candidiasis
B38*	Coccidioidomycosis
B39*	Histoplasmosis
B40*	Blastomycosis
B41.0	Pulmonary paracoccidioidomycosis
B42.0	Pulmonary sporotrichosis
B44*	Aspergillosis
B45.0	Pulmonary cryptococcosis
B46.0	Pulmonary mucormycosis
B58.3	Pulmonary toxoplasmosis
B59*	Pneumocystosis
B67.1	Echinococcus granulosus infection of lung
J10.0	Influenza with pneumonia, influenza virus identified
J11.0	Influenza with pneumonia, virus not identified
J12*	Viral pneumonia, not elsewhere classified
J13*	Pneumonia due to Streptococcus pneumoniae
J14*	Pneumonia due to Haemophilus influenzae
J15*	Bacterial pneumonianot elsewhere classified
J16*	Pneumonia due to other infectious organisms NEC
J17*	Pneumonia in diseases classified elsewhere
J18*	Pneumoniaorganism unspecified
J85*	Abscess of lung and mediastinum
J86.0	Pyothorax with fistula
J86.9	Pyothorax without fistula

Annex 3: Read codes to be used to identify pneumonia

A022200 Salmonella pneumonia A054.00 Amoebic lung abscess A1000 Primary tuberculous infection A100.00 Primary tuberculous complex A101.00 Tuberculous pleurisy in primary progressive tuberculosis A102.00 Primary tuberculous infection NOS A1100 Pulmonary tuberculous infection NOS A1101 Lung tuberculosis A1111 Lung tuberculosis A111.00 Infiltrative lung tuberculosis A111.00 Nodular lung tuberculosis A111.00 Tuberculosis of lung with cavitation A113.00 Tuberculosis of bronchus A114.00 Tuberculous fibrosis of lung A115.00 Tuberculous pneumonia A116.00 Tuberculous pneumonia A117.00 Tuberculous pneumonia A117.00 Tuberculous pneumonia A117.00 Tuberculous pneumonia A112.00 Other specified pulmonary tuberculosis A1200 Other respiratory tuberculosis A1200 Tuberculous pleurisy A12000 Tuberculous pleurisy A12000 Tuberculous pleurisy A12000 Tuberculous pleurisy A12000 Tuberculous pleurisy NOS A121.00 Tuberculous pleurisy NOS A121.00 Tuberculosis of intrathoracic lymph nodes A121100 Tuberculosis of intrathoracic lymph nodes A121100 Tuberculosis of mediastinal lymph nodes A121200 Tuberculosis of intrathoracic lymph nodes A121200 Tuberculosis of intrathoracic lymph nodes A121200 Isolated tracheal or bronchial tuberculosis A122200 Isolated tracheal or bronchial tuberculosis A122400 Tuberculosis of lung, confirmed by culture only A12400 Tuberculosis of lung, confirmed by culture only A12400 Tuberculosis of lung, confirmed by unspecified means TB intrathoracic lymph nodes confirm bact histologically A124700 Tuberculosis of lung, confirmed by culture only A124700 Tuberculosis of lung, confirmed by culture only A12400 Tuberculosis of lung, confirmed by	Read Code	Read Term
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A124700 Primary respiratory TB confirm bact and histologically	A124600	· · · · · · · · · · · · · · · · · · ·
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A123.00 Respiratory 1 D not confirmed bact of histologically	A125.00	Respiratory TB not confirmed bact or histologically
A125000 Tuberculosis of lung, bacteriologically & histolog'y neg	A125000	<u> </u>
A125100 Tuberculosis lung bact and histological examin not done	A125100	Tuberculosis lung bact and histological examin not done

A125200	Prim respiratory TB without mention of bact or hist confirm
A125X00	Resp TB unspcf,w'out mention/bacterial or histol confrmtn
A12y.00	Other specified respiratory tuberculosis
A12y000	Tuberculosis of mediastinum
A12y100	Tuberculosis of nasopharynx
A12y200	Tuberculosis of nasal septum
A12y300	Tuberculosis of nasal sinus
A12yz00	Other specified respiratory tuberculosis NOS
A1800	Miliary tuberculosis
A180.00	Acute miliary tuberculosis
A180000	Acute miliary tuberculosis of a single specified site
A180100	Acute miliary tuberculosis of multiple sites
A18y.00	Other specified miliary tuberculosis
A18z.00	Miliary tuberculosis NOS
A203.00	Primary pneumonic plague
A205.00	Pneumonic plague, unspecified
A221.00	Pulmonary anthrax
A221.11	Woolsorters' disease
A310.00	Pulmonary mycobacterial infection
A310000	Pulmonary mycobacterium avium-intracellulare infection
A391.00	Pulmonary actinomycosis
A39y000	Pulmonary nocardiosis
A54x400	Herpes simplex pneumonia
A551.00	Postmeasles pneumonia
A730.00	Ornithosis with pneumonia
A785000	Cytomegaloviral pneumonitis
A789300	HIV disease resulting in Pneumocystis carinii pneumonia
A789311	HIV disease resulting in Pneumocystis jirovecii pneumonia
AB24.00	Candidiasis of lung
AB24.11	Pneumonia - candidal
AB300	Coccidioidomycosis
AB30.00	Primary pulmonary coccidioidomycosis
AB31.00	Primary extrapulmonary coccidioidomycosis
AB32.00	Coccidioidal meningitis
AB33.00	Other progressive coccidioidomycosis
AB33000	Coccidioidal granuloma
AB3z.00	Coccidioidomycosis NOS
AB400	Histoplasmosis
AB40.11	American histoplasmosis
AB40100	Histoplasma capsulatum with meningitis
AB40200	Histoplasma capsulatum with retinitis
AB40300	Histoplasma capsulatum with pericarditis
AB40500	Histoplasma capsulatum with pneumonia
AB40600	Acute pulmonary histoplasmosis capsulati
AB40700	Chronic pulmonary histoplasmosis capsulati
AB40800	Disseminated histoplasmosis capsulati
AB41200	Histoplasma duboisii with retinitis
AB41300	Histoplasma duboisii with pericarditis
AB41400	Histoplasma duboisii with endocarditis
AB41500	Histoplasma duboisii with pneumonia

AB42.00	Pulmonary histoplasmosis
AB4z200	Histoplasmosis with retinitis
AB4zz00	Unspecified histoplasmosis NOS
AB50.00	Blastomycosis
AB50.13	North American blastomycosis
AB50000	Cutaneous blastomycosis
AB50100	Primary pulmonary blastomycosis
AB50200	Disseminated blastomycosis
AB50300	Blastomycosis liver
AB50z00	Blastomycosis NOS
AB51.13	South American blastomycosis
AB52.11	Keloidal blastomycosis
AB62.00	Chromoblastomycosis
AB62100	Phaeomycotic brain abscess
AB62200	Subcutaneous phaeomycotic abscess and cyst
AB62X00	Chromomycosis, unspecified
AB63.00	Aspergillosis
AB63000	Invasive pulmonary aspergillosis
AB63100	Tonsillar aspergillosis
AB63200	Disseminated aspergillosis
AB63300	Allergic bronchopulmonary aspergillosis
AB63400	Pulmonary aspergillus disease
AB63500	Aspergilloma
AB63600	Aspergillus bronchitis
AB63X00	Aspergillosis, unspecified
AB65000	Pulmonary cryptococcosis
AC21.00	Lung echinococcus granulosus
AD63.00	Pneumocystosis
AE00.00	Late effects of respiratory tuberculosis
Ayu1000	[X]Other resp tubercul, confirmed bacteriologicly+histologicly
Ayu1100	[X]Resp tuberculos unspecfd, confirmed bacteriolog+histologicly
Ayu1300	[X]Resp TB unspcf, w'out mention/bacterial or histol confrmtn
Ayu1800	[X]Other miliary tuberculosis
Ayu1900	[X]Miliary tuberculosis, unspecified
AyuEK00	[X]Other forms of aspergillosis
AyuEU00	[X]Other pulmonary aspergillosis
AyuJ400 F501B00	[X]Sequelae of respiratory and unspecified tuberculosis
H2000	Chronic otitis externa due to aspergillosis Viral pneumonia
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H2011	Chest infection - viral pneumonia Pneumonia due to adenovirus
H200.00 H201.00	
	Pneumonia due to respiratory syncytial virus
H202.00	Pneumonia due to parainfluenza virus
H203.00	Pneumonia due to human metapneumovirus
H20y.00	Viral pneumonia NEC
H20y000 H20z.00	Severe acute respiratory syndrome
H20z.00 H2100	Viral pneumonia NOS
	Lobar (pneumococcal) pneumonia Chast infaction pneumococcal pneumonia
H2111	Chest infection - pneumococcal pneumonia
H2200	Other bacterial pneumonia

H2211	Chest infection - other bacterial pneumonia
H220.00	Pneumonia due to klebsiella pneumoniae
H221.00	Pneumonia due to pseudomonas
H222.00	Pneumonia due to haemophilus influenzae
H222.11	Pneumonia due to haemophilus influenzae
H223.00	Pneumonia due to streptococcus
H223000	Pneumonia due to streptococcus, group B
H224.00	Pneumonia due to staphylococcus
H22y.00	Pneumonia due to other specified bacteria
H22y000	Pneumonia due to escherichia coli
H22y011	E.coli pneumonia
H22y100	Pneumonia due to proteus
H22y200	Pneumonia - Legionella
H22yX00	Pneumonia due to other aerobic gram-negative bacteria
H22yz00	Pneumonia due to bacteria NOS
H22z.00	Bacterial pneumonia NOS
H2300	Pneumonia due to other specified organisms
H2311	Chest infection - pneumonia organism OS
H230.00	Pneumonia due to Eaton's agent
H231.00	Pneumonia due to mycoplasma pneumoniae
H232.00	Pneumonia due to pleuropneumonia like organisms
H233.00	Chlamydial pneumonia
H23z.00	Pneumonia due to specified organism NOS
H2400	Pneumonia with infectious diseases EC
H2411	Chest infection with infectious disease EC
H240.00	Pneumonia with measles
H241.00	Pneumonia with cytomegalic inclusion disease
H242.00	Pneumonia with ornithosis
H243.00	Pneumonia with whooping cough
H243.11	Pneumonia with pertussis
H244.00	Pneumonia with tularaemia
H246.00	Pneumonia with aspergillosis
H247000	Pneumonia with candidiasis
H247100	Pneumonia with coccidioidomycosis
H247z00	Pneumonia with systemic mycosis NOS
H24y.00	Pneumonia with other infectious diseases EC
H24y000	Pneumonia with actinomycosis
H24y100	Pneumonia with nocardiasis
H24y200	Pneumonia with pneumocystis carinii
H24y300	Pneumonia with Q-fever
H24y400	Pneumonia with salmonellosis
H24y500	Pneumonia with toxoplasmosis
H24y600	Pneumonia with typhoid fever
H24y700	Pneumonia with varicella
H24yz00	Pneumonia with other infectious diseases EC NOS
H24z.00	Pneumonia with infectious diseases EC NOS
H2500	Bronchopneumonia due to unspecified organism
H2511	Chest infection - unspecified bronchopneumonia
H2600	Pneumonia due to unspecified organism
H2611	Chest infection - pnemonia due to unspecified organism

H260.00	I ober proumonie due to unerceified organism
H260000	Lobar pneumonia due to unspecified organism Lung consolidation
H261.00	<u>e</u>
H262.00	Basal pneumonia due to unspecified organism
H263.00	Postoperative pneumonia
	Pneumonitis, unspecified
H2711	Influence with anoumenie
H270.00	Influenza with pneumonia
H270.11	Chest infection - influenza with pneumonia
H270000	Influenza with bronchopneumonia
H270100	Influenza with pneumonia, influenza virus identified
H270z00	Influenza with pneumonia NOS
H2800	Atypical pneumonia
H2B00	Community acquired pneumonia
H2C00	Hospital acquired pneumonia
H470.11	Aspiration pneumonia
H470312	Aspiration pneumonia due to vomit
H471000	Lipoid pneumonia (exogenous)
H501600	Pyothorax
H5300	Abscess of lung and mediastinum
H530.00	Abscess of lung
H530000	Single lung abscess
H530100	Multiple lung abscess
H530200	Gangrenous pneumonia
H530300	Abscess of lung with pneumonia
H530z00	Abscess of lung NOS
H531.00	Abscess of mediastinum
H53z.00	Abscess of lung and mediastinum NOS
H540000	Hypostatic pneumonia
H540100	Hypostatic bronchopneumonia
H564.00	Bronchiolitis obliterans organising pneumonia
H564.11	Cryptogenic organising pneumonia
H56y000	Endogenous lipoid pneumonia
H56y100	Interstitial pneumonia
H571.00	Rheumatic pneumonia
Hyu0800	[X]Other viral pneumonia
Hyu0A00	[X]Other bacterial pneumonia
Hyu0B00	[X]Pneumonia due to other specified infectious organisms
Hyu0D00	[X]Pneumonia in viral diseases classified elsewhere
Hyu0H00	[X]Other pneumonia, organism unspecified
SP13100	Other aspiration pneumonia as a complication of care