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1. Introduction

One of the challenges facing health care computing is the representation of patient data in a usable form. The typical approach is to encode the information using some standard terms taken from a controlled vocabulary. Applications such as order entry, summary reporting, automated decision support, and data aggregation for clinical research all require recording the data in standard ways [1,2]. This need for controlled vocabulary to support clinical applications has been recognized for decades (see, for example, [3,4,5]). Understandably, health care providers, educators, researchers and policy makers often take for granted the existence of an appropriate standard terminology and assume that it is in routine use. In reality, the lack of a standard for representing patient data is one of the greatest impediments to

Review Paper

Coding Systems in Health Care

Abstract: Computer-based patient data which are represented in a coded form have a variety of uses, including direct patient care, statistical reporting, automated decision support, and clinical research. No standard exists which supports all of these functions. Abstracting coding systems, such as ICD, CPT, DRGs and MeSH fail to provide adequate detail, forcing application developers to create their own coding schemes for systems. Some of these schemes have been put forward as possible standards, but they have not been widely accepted. This paper reviews existing schemes used for abstracting, electronic record systems, and comprehensive coding. It also discusses the remaining impediments to acceptance of standards and the current efforts to overcome them, including SNOMED, the Gabrieli Medical Nomenclature, the Read Clinical Codes, GALEN, and the Unified Medical Language System (UMLS).

Keywords: Controlled Medical Vocabulary, Nomenclature, Taxonomy, Electronic Medical Records, Medical Record Coding, Review

medical computing today [6,7]. The importance of patient data encoding to the medical informatics community is reflected in the recent increase in published literature on the subject. For example, in the newly established Journal of the American Medical Informatics Association, 18 of the 51 papers in the first 8 issues deal with coding of clinical data. A survey of medical informatics conference proceedings, spanning the years 1974 to 1992, showed 8.4% were primarily about coding issues [8]; in the most recent Symposium on Computer Applications in Medical Care (SCAMC) 24 of the 167 papers appeared in the Vocabulary and Nomenclature track, and an additional 24 dealt with applications requiring coded patient data [9].

In this paper, I review the current state of the coding schemes with general suitability for health care applications. First, I will survey the coding schemes which are used for abstracting patient data, as is done for health statistics reporting and reimbursement. Next, I will review the controlled vocabularies which are intended to support coding of detailed patient data, as in comprehensive electronic medical records and automated decision support. I will then report on current efforts to develop comprehensive clinical coding schemes that seek to serve both purposes. Finally, I will close with a summary of the research issues which remain to be addressed.

2. Coding for Medical Record Abstraction

The coding of patient information has been carried out long before the advent of computers. This coding has always been directed at simpli-

fying the data, converting it to a general form which is easier to manipulate. For example, while a patient may have pneumonia that may be caused by any of a variety of organisms, involve different sites in the lungs, be accompanied by any of several different symptoms, and be of varying severity, coding the patient's diagnoses as simply "bacterial pneumonia" allows it to be aggregated with other cases for statistical purposes. If finer granularity is needed, more specific terms can be added to the coding scheme (such as "Gram-negative bacterial pneumonia", "lobar bacterial pneumonia", and "bacterial pneumonia requiring mechanical ventilation"). A set of patient records can be classified with such codes and then retrieved when cases of certain types are needed. Because the coding represents only a simplified synopsis of information extracted from the record, this kind of coding is referred to as abstraction. Record abstraction has been performed since the advent of formal medical records, to allow assessment of incidence of a disease, mortality of a surgical procedure or (in the era of prospective payment) costs for a hospital stay.

481 Pneumococcal Pneumonia

482 Other Bacterial Pneumonia

482.0 Pneumonia due to Klebsiella Pneumoniae

482.1 Pneumonia due to Pseudomonas

482.2 Pneumonia due to Haemophilus Influenzae

482.3 Pneumonia due to Streptococcus

482.4 Pneumonia due to Staphylococcus

482.8 Pneumonia due to Other Specified Bacteria

484 Pneumonia in Infectious Disease Classified Elsewhere

484.3 Pneumonia in Whooping Cough

484.4 Pneumonia in Tularemia

484.5 Pneumonia in Anthrax

Figure 1 - Bacterial Pneumonias Coded in ICD-9. The very extensive set of codes for mycobacterial disease has been omitted for simplicity.

The archetypal coding system for medical record abstraction is the International Classification of Diseases (ICD). Other major coding schemes are usually presented in terms of their compatibility with ICD and their ability to resolve some of ICD's problems with granularity or coverage of a particular domain. ICD was first published in 1893. It has been revised at roughly 10-year intervals, first by the Statistical International Institute and later by the World Health Organization (WHO). The Ninth Edition (ICD-9) was published in 1977 [10], and the Tenth Edition (ICD-10) in 1992 [11]. The coding system consists of a "core" classification of three-digit codes which are the minimum required for reporting mortality statistics to WHO. A fourth digit (in the first decimal place) provides an additional level of detail; usually .0 to .7 are used for more specific forms of the core term, .8 is usually used for an "other" category and .9 for "unspecified". Terms are arranged in a strict hierarchy, based on the digits in the code. For example, bacterial pneumonias are classified as shown in Figure 1. While ICD proper is limited to disease terminology, WHO also provides a set of expansions for different "families" of terms for medical specialty diagnoses, health status, disablements, procedures and reasons for contact with health care providers.

The publication of ICD-9 was immediately followed by publication of criticisms regarding its inadequacy for general coding and specific specialty coverage [12,13,14]. In order to address these and other perceived problems with ICD-9, the United States National Center for Health Statistics published a set of "clinical modifications" to ICD-9, known as ICD-9-CM [15]. While completely compatible with ICD-9, the additions provided an additional level of detail in many places by adding a fifth digit to the code, contact the code of the code of the code of the code of the code, contact the code of the code, contact the code of the code of the code of the code of the code, contact the code of the cod

003 Other Salmonella Infections

003.0 Salmonella Gastroenteritis

003.1 Salmonella Septicemia

003.2 Localized Salmonella Infections

003.20 Localized Salmonella Infection, Unspecified

003.21 Salmonella Meningitis

003.22 Salmonella Pneumonia

003.23 Salmonella Arthritis

003.24 Salmonella Osteomyelitis

003.29 Other Localized Salmonella Infection

003.8 Other specified salmonella infections

003.9 Salmonella infection, unspecified

Figure 2 - Example of "fifth digit" codes the Clinical Modifications of ICD-9 (ICD-9-CM). The four-digit codes are identical to those in ICD-9; the five-digit codes were introduced in ICD-9-CM. Note that Salmonella Pneumonia has been added as a child in the 003 section; it is not included under 482 (Other Bacterial Pneumonia) or 484 (Pneumonia in Infectious Disease Classified Elsewhere).

responding to another level in the hierarchy (see Fig. 2).

Another American creation for the purpose of abstracting medical records has been the Diagnosis-Related Groups (DRGs), developed initially at Yale University for use in prospective payment in the Medicare program [16]. In this case, the coding system is an abstraction of an abstraction: it is applied to lists of ICD-9-CM codes which are themselves derived from medical records. The purpose of DRG coding is to provide a relatively small number of codes for classifying patient hospitalizations while at the same time providing some separation of cases based on severity of illness. The principal motivations for the groupings are factors which affect cost and length of stay. Thus, a medical record containing the ICD-9-CM primary diagnosis of Pneumococcal Pneumonia (481) might be coded with one of eighteen codes (see Figure 3) depending on associated conditions and procedures; additional codes are possible if the pneumonia is a secondary diagnosis.

A more international response to perceived deficiencies in ICD-9 came in the form of the International Classification of Primary Care (ICPC) from the World Organization of National Colleges, Academies and Academic Associations of General Practitioners/Family Physicians (WONCA) [17]. ICPC provides seven axes of terms and a structure to combine them to represent clinical encounters. While the granularity of the terms is generally less than that of other classifications schemes (e.g., all pneumonias are coded as R81), the ability to represent the interactions of the concepts found in a medical record is much greater through the postcoordination of atomic terms (see Figure 4). In postcoordination, the coding is accomplished through the use of multiple codes as needed to describe the data. So, for example, a case of bacterial pneumonia would be

coded in ICPC as a combination of the code R81 and the code for the particular test result which identifies the causative agent. This is in contrast to the precoordination approach, in which every type of pneumonia is assigned its own code (as in Fig. 1).

Professional specialty groups find that general coding schemes are of little use for their purposes and often resort to developing their own coding schemes for medical record abstraction. For example, the American Medical Association developed the Current Procedural Terminology (CPT) in 1966 [18] to provide a precoordinated coding scheme for diagnostic and therapeutic procedures which has since been adopted in the US for billing and reimbursement. Like the DRG codes, CPT codes specify information about the codes which differentiates them based on their cost. For example, there are different codes for pacemaker insertions, depending on whether the leads are "epicardial, by

| Respiratory disease with major chest operating room procedure, no major complication or comorbidity | 75 | |
|---|-----|--|
| Respiratory disease with major chest operating room procedure, minor complication or comorbidity | 76 | |
| Respiratory disease with other respiratory system operating procedure, no complication or comorbidity | 77 | |
| Respiratory infection with minor complication, age greater than 17 | 79 | |
| Respiratory infection with no minor complication, age greater than 17 | 80 | |
| Simple Pneumonia with minor complication, age greater than 17 | 89 | |
| Simple Pneumonia with no minor complication, age greater than 17 | 90 | |
| Respiratory disease with ventilator support | 475 | |
| Respiratory disease with major chest operating room procedure and major complication or comorbidity | 538 | |
| Respiratory disease, other respiratory system operating procedure and major complication or comorbidity | 539 | |
| Respiratory infection with major complication or comorbidity | 540 | |
| Respiratory infection with secondary diagnosis of bronchopulmonary dysplasia | 631 | |
| Respiratory infection with secondary diagnosis of cystic fibrosis | 740 | |
| Respiratory infection with minor complication, age not greater than 17 | 770 | |
| Respiratory infection with no minor complication, age not greater than 17 | 771 | |
| Simple Pneumonia with minor complication, age not greater than 17 | 772 | |
| Simple Pneumonia with no minor complication, age not greater than 17 | 773 | |
| Respiratory infection with primary diagnosis of tuberculosis | 798 | |

Figure 3 - DRG codes assigned to cases of bacterial pneumonia depending on co-occurring conditions and/or procedures (mycobacterial disease is not shown except as a co-occurring condition). "Simple Pneumonia" codes are used when the primary bacterial pneumonia corresponds to ICD-9 codes 481, 482.2, , 482.3 or 482.9 (refer to Figures 1 and 2) and there are only minor or no complications. The remaining ICD-9 bacterial pneumonias (482.0, 482.1, 482.2, 482.4, 482.8, 484, and various other codes such as 003.22 (refer to Figure 2) are coded as "Respiratory Disease" or "Respiratory Infection". Cases in which pneumonia is a secondary diagnosis may also be assigned other codes (such as 798), depending on the primary condition.

| | | Chapter | Chapter . | | |
|----|-----------------------------------|---------|-----------------|--|--|
| | Components | | R - Respiratory | | |
| 1. | Symptoms and complaints | | | | |
| 2. | Diagnostic, screening, prevention | | • | | |
| 3. | Treatment, procedures, medication | | | | |
| 4. | Test results | | | | |
| 5. | Administative | | | | |
| 6. | Other | | | | |
| 7. | Diagnoses, disease | | R81 | | |

Figure 4 - ICPC Coding for Pneumonia. Only one of seventeen chapters (Respiratory System) is shown. Coding a clinical encounter for a patient with pneumonia entails the assignment of the code R81 as the diagnosis and including codes in any of the other six components that can be used to describe the severity and etiology of the case.

thoracotomy" (33200), "epicardial, by xiphoid approach" (33201), "transvenous, atrial" (33206), "transvenous, ventricular" (33207), or "transvenous, AV sequential" (33208). CPT also provides information about the reasons for a procedure. For example, there are codes for arterial punctures for "withdrawal of blood for diagnosis" (36600), "monitoring" (36620), "infusion therapy" (36640), and "occlusion therapy" (75894).

Another successful specialty coding scheme is the American Psychiatric Association's Diagnostic and Statistical Manual of Mental Disorders, published in 1987 in its Revised Third Edition (DSM-III-R) [19]. Publication of the Fourth Edition (DSM-IV) has been coordinated with the development of psychiatric diagnoses in ICD-10 [20]. The DSM nomenclature provides definitions of the disorders including diagnostic criteria. Thus it is used not only for coding patient data but as a tool for actually assigning diagnoses. Each edition of DSM has been coordinated with corresponding editions of ICD. Compatibility between ICD-9 and DSM-III-R was found to be reasonably good [21]; a number of studies have shown that compatibility

between ICD-10 and DSM-IV is variable across its different sections.

Nursing organizations have been extremely active in the development of standard coding systems for abstracting patient records. One review counted a total of 13 separate projects world-wide [22]. Two recent reports analyze the current state of these classification systems (as well as the more general purpose standard coding systems) and describe their shortcomings [23,24,25]. The findings of these authors and others are serving as the basis for the development of an International Classification of Nursing Practice by the International Council of clasNurses.

Another domain with a successful abstracting scheme is in anatomic pathology. Drawing from the New York Academy of Medicine's Standard Nomenclature of Diseases and Operations (SNDO) [26], the College of American Pathologists developed the Standard Nomenclature of Pathology (SNOP) as a multiaxial system for describing pathologic findings [27] through postcoordination of topographic (anatomic), morphologic, etiologic and functional terms. SNOP has been used widely in pathology sys-

tems in the US; its successor, the Systematized Nomenclature of Medicine (SNOMED) has evolved beyond an abstracting scheme toward a comprehensive coding system and is described below.

No review of medical coding schemes would be complete without mention of the Medical Subject Headings (MeSH), maintained by the US National Library of Medicine (NLM) [28]. MeSH is the vocabulary by which the world medical literature is indexed. MeSH arranges terms in a structure that breaks from the strict hierarchy used by most other coding schemes. Terms are organized into hierarchies and may appear in multiple places in the hierarchy (see Figure 5). Although it is not generally used as a direct coding scheme for patient information, it plays a central role in the Unified Medical Language System (described below).

The medical literature is replete with arguments about the pros and cons of the available standards for abstracting medical records. Inadequacies in one coding system may even be blamed on those of another [29], but problems are typically reported when a scheme blurs impor-

tant clinical distinctions through its coarse granularity [30] or because it simply lacks sufficient content to cover the requisite domain [31].

The structure of a controlled vocabulary may also be the source of problems [32]. For example, a strict hierarchical structure precludes the ability to classify terms in two or more ways. By way of illustration, refer to Figure 2, which shows refinement of the ICD-9 term 003.2 Localized Salmonella Infections with the ICD-9-CM term 003.22 Salmonella Pneumonia. This position in the coding hierarchy appears correct, but it ignores the fact that ICD-9-CM (and ICD-9, as shown in Figure 1), classifies such terms under 482 Other Bacterial Pneumonia or 484 Pneumonia in Infectious Disease Classified Elsewhere. Since ICD-9-CM is a strict hierarchy, Salmonella Pneumonia may appear only as a descendent of one of its possible parents (Pneumonia or Localized Salmonella Infections). The structure used by MeSH offers a way to overcome the limitations of a strict hierarchy by allowing multiple contexts; however, as Figure 5 demonstrates, allowing a term to appear in multiple contexts may lead to some ambiguity about its meaning.

3. Coding for Medical **Record Systems**

Abstracting systems are a fact of life for medical record keeping, both for health statistics reporting and, at least in the US, for reimbursement [33]. The relevant question here is: can these systems support computerbased health care systems? When an abstract system fails at its original task (reporting causes of mortality and morbidity) [34], it should not be surprising that it is inappropriate for more strenuous tasks, such as coding a research database [35]. An

even more challenging task is the coding of data in a record in a way that retains sufficient detail for a care provider to use it directly in patient care. Treatment decisions, for example, require more detail than "Pneumonia Due to Other Specified Bacteria" in order to select an appropriate antibiotic. At the same time, coding of detailed data must consider the additional uses for the data, such as case review, summary review, decision support, research, quality assurance and, of course, reporting of mortality and morbidity.

Electronic medical record (EMR) systems typically have the greatest vocabulary requirements, assuming that the data in the record are to be encoded. In general, developers of health care applications have difficulty using existing coding systems. For example, the developers of TMR

(The Medical Record) at Duke University have explicitly rejected standard vocabularies as inappropriate for use in an EMR [36]. They, and others, have resorted to developing their own controlled vocabularies. In some cases, they are created in an ad hoc manner, adding coded terms as needed. In other cases, developers have applied a deliberate methodology to vocabulary development.

One of the most comprehensive EMRs is the HELP System in use at the LDS Hospital in Salt Lake City, Utah [37]. The data in HELP are drawn from most of the hospital departments, cover a wide range of functional types, and are used for a variety of purposes [38]. Almost all of the data in HELP are encoded with the PTXT data dictionary. This dictionary is structured as a strict hierarchy with each term having an eight-byte code in which the first

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Respiratory Tract Diseases
       Lung Diseases
              Pneumonia
```

Bronchopneumonia

Pneumonia, Aspiration

Pneumonia, Lipid

Pneumonia, Lobar

Pneumonia, Mycoplasma

Pneumonia, Pneumocystis Carinii

Pneumonia, Rickettsial

Pneumonia, Staphylococcal

Pneumonia, Viral

Lung Diseases, Fungal

Pneumonia, Pneumocystis Carinii

Respiratory Tract Infections

Pneumonia

Pneumonia, Lobar

Pneumonia, Mycoplasma

Pneumonia, Pneumocystis Carinii

Pneumonia, Rickettsial

Pneumonia, Staphylococcal

Pneumonia, Viral

Lung Diseases, Fungal

Pneumonia, Pneumocystis Carinii

Figure 5 - Partial tree structure for the Medical Subject Headings (MeSH) showing pneumonia terms. Note that terms can appear in multiple locations, although they may not always have the same children, implying that they have somewhat different meanings in different contexts. For example, Pneumonia means "lung inflammation" in one context (line 3) and "lung infection" in another (line 16).

three bytes specify general information about the type of data being stored and the last five define the term's position in the PTXT hierarchy. The system is now commercially available and the PTXT vocabulary is common across the various HELP installations; however, as of this writing PTXT has not been implemented in any other EMRs. Furthermore, while PTXT is used successfully by the on-line decision support capabilities of the HELP system, it has proven difficult to use for a diagnostic expert system developed by the same research group [39].

COSTAR (Computer-Stored Ambulatory Record) [40], developed at the Massachusetts General Hospital, also makes extensive use of a formal, albeit "home grown" controlled vocabulary called the Directory. Like PTXT, the COSTAR Directory is a strict hierarchy with a coding system (in this case, three alpha-numeric digits, plus a check digit and optional modifiers) which provides terms for coding a wide range of information in the record. COSTAR is available from commercial vendors, but can also be obtained in a public domain form that is available from the COSTAR Users Group. A standard Directory is supplied with the software; however, it only specifies the uppermost levels in the hierarchy. It is left to each installation site to flesh out the hierarchy with specific terms for their own institution. There has been no attempt to standardize these individual development efforts.

The Regenstrief Medical Record System (RMRS) at the University of Indiana [41] also uses a coded vocabulary for representing a portion of its data. This particular vocabulary construction task was complicated by the need to coordinate terminologies from four different hospitals. Despite the effort expended

to make RMRS inter-institutional, it remains institution-dependent and has not been adopted for use in other systems.

There is one notable exception to the rule that abstracting systems have failed to support EMRs. Developed at Erasmus University in Rotterdam and now in use in a majority of private practitioners' offices in the Netherlands, the Elias system makes use of the ICPC for coding diagnoses and reasons for encounters [42]. This adoption was not without cost, however. An extensive project was undertaken to translate ICPC to Dutch and to match the ICPC codes with the terms entered by users of the ELIAS system [43]. This project resulted in a greatly enhanced version of ICPC, with a significant addition of index terms and synonyms. Evaluations thus far have shown relatively good general acceptance. Similar success in other settings awaits further work to establish vocabulary standards [44].

All of the aforementioned EMRs make use of coding schemes which, while varying in their domain coverage and richness of detail, all share a fairly simple structure - that of a strict hierarchy. In some cases, synonyms are allowed and in some cases appropriate modifiers are specified. However, the depth of the representation of the vocabularies is generally shallow compared to that invested in other aspects of the systems. The approach at the University of Manchester has been quite different. In the PEN & PAD project (Practitioners Entering Notes and Practitioners Accessing Data), the vocabulary model is based on a semantic net formalism (Structured Meta Knowledge, or SMK) which allows for a variety of vocabularyrelated information to be specified and allows multiple hierarchies [45]. System developers have found that the extra effort made in vocabulary

development ultimately pays off in terms of the ability of the EMR to remain faithful to the description of the original patient care processes it records. The structure of the PEN & PAD vocabulary also provides the flexibility needed to support secondary uses of the data and to adapt the system for uses in a variety of patient care settings and populations,

The Medical Entities Dictionary (MED) used in the Columbia-Presbyterian clinical information system is also based on a semantic network model [46]. This vocabulary integrates terms from national coding schemes with those from local ancillary systems to produce a unified coding scheme that retains the fine granularity from the original coding schemes while accommodating the coarser granularity of a variety of applications making use of the patient data. The semantic network model is useful both for supporting the addition of new terms from ancillary systems [47] and for maintaining currency with changes in the national vocabularies [48].

4. Current Efforts to Develop Medical Coding Systems

The developers of each EMR have dealt with controlled vocabulary in a unique way. The results have been generally satisfactory for supports ing the needs at each site; however, the ability to share the coding scheme for use at other sites has been limited, when it occurs at all. The implication is that other developers may enjoy the same successes but they will, essentially, be required to start from scratch. With several decades of experience in computer-based vocabulary requirements, researchers are now beginning to collaborate to apply their individual experiences to the task of developing general purpose, comprehensive controlled

wocabularies to support health care applications.

The first coding scheme which attempted to provide terms for a broad range of clinical domains was the Systematized Nomenclature of Medicine (SNOMED), from the College of American Pathologists. First published in 1975 and then

revised as SNOMED II in 1979, it has recently been released in a greatly expanded version: the Systematized Nomenclature of Human and Veterinary Medicine - SNOMED International [49]. SNOMED consists of a set of axes (now eleven), each of which serve as a taxonomy for a specific set of concepts (organisms,

diseases, procedures, etc.), containing a total of over 130,000 terms. Coding patient information is accomplished through the postcoordination of terms from multiple axes to represent complex terms, which may be desired but do not exist in SNOMED. For example, although many of the various bacterial pneu-

| DE-10000 Bacterial infectious disease, NOS (L-10000) DE-11205 Pneumonia in anthrax (T-28000)(M-40000) DE-13212 Pneumonia in pertussis (T-28000)(M-40000) DE-13430 Pneumonic plague, NOS (T-28000)(L-1E401)(DE-01750) DE-13431 Primary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13432 Secondary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-14210 Staphylococcal pneumonia (T-28000)(M-4000)(L-25100) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (T-28000)(L-1F701) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to Klebsiella pneumoniae (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-23400) DE-19162 Pneumonia due to Proteus mirabilis (T-28000)(M-40000)(L-16802) | | | |
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| DE-11205 Pneumonia in anthrax (T-28000)(M-40000) DE-13212 Pneumonia in pertussis (T-28000)(M-40000) DE-13430 Pneumonic plague, NOS (T-28000)(L-1E401)(DE-01750) DE-13431 Primary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13432 Secondary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13510 Pneumococcal pneumonia (T-28000)(M-4000)(L-25116) DE-13934 Salmonella pneumonia (T-28000)(M-4000)(L-25116) DE-14120 Staphylococcal pneumonia (T-28000)(M-4000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-17701) DE-15716 Pittsburg pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (T-28000)(M-40000)(L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-23400) | | | |
| DE-13212 Pneumonia in pertussis (T-28000)(M-40000) DE-13430 Pneumonic plague, NOS (T-28000)(L-1E401)(DE-01750) DE-13431 Primary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13432 Secondary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-13934 Salmonella pneumonia (T-28000)(L-17100) DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15014 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(M-40000) DE-15716 Pittsburg pneumonia (T-28000)(L-1F701) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) < | DE-10000 | Bacterial infectious disease, NOS | (L-10000) |
| DE-13430 Pneumonic plague, NOS (T-28000)(L-1E401)(DE-01750) DE-13431 Primary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13432 Secondary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-13934 Salmonella pneumonia (T-28000)(L-17100) DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (T-28000)(L-1F701) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-11205 | | (T-28000)(M-40000) |
| DE-13431 Primary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13432 Secondary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-13934 Salmonella pneumonia (T-28000)(L-17100) DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-13212 | Pneumonia in pertussis | (T-28000)(M-40000) |
| DE-13431 Primary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13432 Secondary pneumonic plague (T-28000)(L-1E401)(DE-01750) DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-13934 Salmonella pneumonia (T-28000)(L-17100) DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-13430 | Pneumonic plague, NOS | (T-28000)(L-1E401)(DE-01750) |
| DE-13510 Pneumococcal pneumonia (T-28000)(M-40000)(L-25116) DE-13934 Salmonella pneumonia (T-28000)(L-17100) DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | | Primary pneumonic plague | (T-28000)(L-1E401)(DE-01750) |
| DE-13934 Salmonella pneumonia (T-28000)(L-17100) DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-13432 | Secondary pneumonic plague | (T-28000)(L-1E401)(DE-01750) |
| DE-14120 Staphylococcal pneumonia (T-28000)(L-24800) DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-13510 | Pneumococcal pneumonia | (T-28000)(M-40000)(L-25116) |
| DE-14213 Pneumonia due to Streptococcus (T-28000)(M-40000)(L-25100) DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-13934 | Salmonella pneumonia | (T-28000)(L-17100) |
| DE-14817 Tuberculous pneumonia (T-28000)(M-40000)(L-21801) DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-14120 | Staphylococcal pneumonia | (T-28000)(L-24800) |
| DE-15104 Pneumonia in typhoid fever (T-28000)(M-40000) DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) Mebsiella pneumoniae (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-14213 | Pneumonia due to Streptococcus | (T-28000)(M-40000)(L-25100) |
| DE-15613 Haemophilus influenzae pneumonia (T-28000)(L-1F701) DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) Klebsiella pneumoniae (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-14817 | | (T-28000)(M-40000)(L-21801) |
| DE-15716 Pittsburg pneumonia (L-20402) DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to (L-16001) Klebsiella pneumoniae (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-15104 | Pneumonia in typhoid fever | (T-28000)(M-40000) |
| DE-15810 Mycoplasma pneumonia (T-28000)(L-22018) DE-19110 Bacterial infection due to Klebsiella pneumoniae (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-15613 | | (T-28000)(L-1F701) |
| DE-19110 Bacterial infection due to Klebsiella pneumoniae (L-16001) | DE-15716 | Pittsburg pneumonia | (L-20402) |
| Klebsiella pneumoniae (L-16001) DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-15810 | | (T-28000)(L-22018) |
| DE-19111 Pneumonia due to Klebsiella pneumoniae (T-28000)(M-40000)(L-16001) DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | DE-19110 | Bacterial infection due to | |
| DE-19151 Pneumonia due to Pseudomonas (T-28000)(M-40000)(L-23400) | | | (L-16001) |
| | DE-19111 | Pneumonia due to Klebsiella pneumoniae | (T-28000)(M-40000)(L-16001) |
| DE-19162 Pneumonia due to Proteus mirabilis (T-28000)(M-40000)(L-16802) | DE-19151 | Pneumonia due to Pseudomonas | (T-28000)(M-40000)(L-23400) |
| | DE-19162 | Pneumonia due to Proteus mirabilis | (T-28000)(M-40000)(L-16802) |
| DE-19204 Pneumonia due to E.coli (T-28000)(M-40000)(L-15602) | DE-19204 | Pneumonia due to E.coli | (T-28000)(M-40000)(L-15602) |
| DE-21611 Ornithosis with pneumonia (T-28000)(M-40000)(L-2A902) | DE-21611 | Ornithosis with pneumonia | (T-28000)(M-40000)(L-2A902) |
| DE-21704 Pneumonia in Q fever (T-28000)(M-40000) | DE-21704 | Pneumonia in Q fever | (T-28000)(M-40000) |
| DE-3632A AIDS with bacterial pneumonia (T-28000)(L-34800)(L-10000) | DE-3632A | AIDS with bacterial pneumonia | (T-28000)(L-34800)(L-10000) |
| DE-3632B AIDS with pneumococcal pneumonia (T-28000)(L-34800)(L-25100) | DE-3632B | | (T-28000)(L-34800)(L-25100) |
| DE-36333 AIDS with pneumonia, NOS (T-28000)(M-40000)(L-34800) | DE-36333 | AIDS with pneumonia, NOS | (T-28000)(M-40000)(L-34800) |
| D2-50100 Bronchopneumonia, NOS (T-26000)(M-40000) | D2 50100 | Dranch announce in NOC | (T-26000)(M-40000) |
| D2-50100 Bronchopneumonia, NOS (T-26000)(M-40000) D2-50104 Peribronchial pneumonia (T-26090)(M-40000) | 1 | | , , , , |
| 1 | ł | | |
| D2-50110 Hemorrhagic bronchopneumonia (T-26000)(M-40790) D2-50120 Terminal bronchopneumonia (T-26000)(M-40000) | 1 | | |
| D2-50130 Pleurobronchopneumonia (T-26000)(M-40000) (T-26000)(M-40000) | { | | |
| D2-50130 Pleuropneumonia (1-26000)(M-40000) T-26000)(M-40000) | 1 | | · · · · · · · · · · · · · · · · · · · |
| D2-50140 Pneumonia, NOS (T-28000)(M-40000) | | | the state of the s |
| | 1 | | , , , |
| D2-50142 Catarrhal pneumonia (T-28000)(M-40000) D2-50150 Unresolved pneumonia (T-28770)(M-40000) | | | |
| D2-50150 Unresolved pheumonia (1-28000)(M-40000) (T-28000)(M-40000) | | | |
| D2-50300 Aspiration pneumonia, NOS (T-28000)(M-40000)(G-C001)(F-29200) | | | |
| D2-61020 Gangrenous pneumonia (T-28000)(M-40700) (T-28000)(M-40700) | 1 | | |
| D8-72532 Infective pneumonia acquired prenatally, NOS | | | (1-20000)(111-40/00) |
| Intective photonicina acquired prenatany, 1405 | - 0 12332 | intective pholinoma acquired prematany, 1405 | |

Figure 6 - SNOMED international codes for pneumonia. The first set of terms are those from the Disease axis which are included under the Bacterial Infectious Disease hierarchy (excluding several veterinary diseases). "NOS" stands for "Not Otherwise Specified". The codes shown on the right are the SNOMED codes which, when taken together, are the equivalent of the precoordinated bacterial pneumonia terms. For example, "Pneumococcal pneumonia " (DE-13510) is the precoordination of the terms "Lung, NOS" (T-28000), "Inflammation, NOS" (M-40000), and "Streptococcus pneumoniae" (L-25116). The second set of terms shows some of the other pneumonia terms in SNOMED which could be coupled with specific Living Organism terms to allow postcoordinated coding of concepts not found explicitly in SNOMED.

Respiratory Disorder

Infection of the Lower Respiratory Tract and Mediastinum
Acute Lower Respiratory Tract Infection
Pneumonia

Bacterial Pneumonia

Actinomycotic Pneumonia
Haemophilus Influenzae Pneumonia
Legionnaires Disease
Pneumococcal Pneumonia
Pneumonic Plague

Prieumonic Plague
Primary Pneumonic Plague
Secondary Pneumonic Plague
Salmonella Pneumonia
Typhoid Pneumonia
Staphylococcal Pneumonia
Meningococcal Pneumonia

Figure 7 - Bacterial pneumonia in the Read Clinical Codes. Additional infections can be coded by using Bacterial Pneumonia with one of the prescribed modifiers (Bacteria). Some of these terms also appear in other hierarchy locations; for example, Meningococcal Pneumonia also appears under Meningococcal Infection (which is under Bacterial Disease). However, Bacterial Pneumonia is not listed under Bacterial Disease, nor is Actinomycotic Pneumonia under Actinomycotic Infection, although Pulmonary Actinomycosis does appear. Unlike MeSH, when a term appears in multiple places (such as Pneumonic Plague, which also appears under Plague), its children must appear as well.

monia terms seen in other terminologies are in SNOMED (see Fig.6), additional terms can be constructed by pairing a generic pneumonia term with a bacteria term taken from the Living Organism axis.

Despite its long history and extensive efforts to provide the codes needed for coding in EMRs, SNOMED has not been widely embraced. The latest version goes a long way toward addressing past complaints about missing terms; however, the structure of previous versions, also found to be an impediment to use, has persisted in SNOMED International. The main problem with using SNOMED for coding patient information is that it is too expressive. Because there are few rules about how the postcoordination coding should be done, the same expression might be represented differently by different coders. For example, "acute appendicitis" can be coded as a single disease

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term, as a combination of a modifier ("acute") and a disease term ("appendicitis"), or as a combination of a modifier ("acute"), a morphology term ("inflammation") and a topography term ("vermiform appendix"). Each of these codings is correct, yet there is no formal way, in SNOMED, to know they have equivalent meaning. Such freedom of expression may be welcome to those who must encode human utterances, but it is frustrating to system developers who must make sure that their applications can recognize medical concepts.

One proposed solution to this redundant coding problem is the representation of the semantics of SNOMED expressions in a formal way that would allow different surface forms to be recognizable as equivalent [50]. For example, if the disease term "acute appendicitis" was formally represented as equivalent to the combination of a modifier

term and a disease term, and the disease term "appendicitis" was formally represented as a combination of a morphology term and a topography term, then the three coding schemes for "acute appendicitis" would be computationally equivalent. Such equivalence would permit the development of rules for consistent coding and/or sophisticated retrieval of patient data. The SNOMED developers have embraced this approach and work is now under way to formalize the semantics in SNOMED to make it meet the needs of EMRs [51].

The Read Clinical Codes are a set of codes designed specifically for use in coding electronic medical records. Developed privately in the 1980's [52,53], the first version was adopted by the British National Health Service in 1990. Version 2 was developed to meet the needs of hospitals for cross-mapping their data to ICD-9. Version 3 [54] was developed to support not only medical record summarization, but to support patient care applications directly. While previous versions of the Read Codes were organized in a strict hierarchy, Version 3 made an important step by allowing terms to have multiple parents in the hierarchy; that is, the hierarchy became that of a directed acyclic graph. Figure 7 shows the hierarchy for bacterial pneumonia. Version 3.1 added the ability to make use of term modifiers through a set of templates for combining terms in specific, controlled ways so that both precoordination and postcoordination is used. Finally, the NHS has undertaken a series of "terms projects" which are expanding the content of the Read Codes to assure that the terms needed by practitioners are represented in the Codes [55].

At about the same time, the Gabrieli Medical Nomenclature was described in the US [56]. This sys-

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4-3-3-2-1-7-1 Pneumonia
4-3-3-2-1-7-1-3 Causes of Pneumonia
4-3-3-2-1-7-1-3-1 Bacterial Pneumonia
4-3-3-2-1-7-1-3-1-1 Presumed Bacterial Pneumonia
4-3-3-2-1-7-1-3-1-2 Streptococcus Pneumonia
4-3-3-2-1-7-1-3-1-3 Staphylococcus Aureus Pneumonia
4-3-3-2-1-7-1-3-1-3-1 Staphylococcal Pneumonia
4-3-3-2-1-7-1-3-1-4 Streptococcus Pyogenes Pneumonia
4-3-3-2-1-7-1-3-1-5 Neisseria Meningitidis Pneumonia
4-3-3-2-1-7-1-3-1-6 Branhamella Catarrhalis Pneumonia
4-3-3-2-1-7-1-3-1-7 Hemophilus Influenzae Pneumonia
4-3-3-2-1-7-1-3-1-8 Klebsiella Pneumonia
4-3-3-2-1-7-1-3-1-9 Escherichia Coli Pneumonia
4-3-3-2-1-7-1-3-1-10 Serratia Species Pneumonia
4-3-3-2-1-7-1-3-1-11 Enterobacteria Species Pneumonia
4-3-3-2-1-7-1-3-1-12 Proteus Species Pneumonia
4-3-3-2-1-7-1-3-1-13 Pseudomonas Aeruginosa Pneumonia
4-3-3-2-1-7-1-3-1-14 Pseudomonas Capacia Pneumonia
4-3-3-2-1-7-1-3-1-15 Pseudomonas Multiphilia Pneumonia
4-3-3-2-1-7-1-3-1-16 Pseudomonas Pseudoalcaligenes Pneumonia
4-3-3-2-1-7-1-3-1-17 Actinobacter Species Pneumonia
4-3-3-2-1-7-1-3-1-18 Legionella Species Pneumonia
4-3-3-2-1-7-1-3-1-19 Anaerobic Microbial Pneumonia
4-3-3-2-1-7-1-3-1-19-1 Fusobacterium Species Pneumonia
4-3-3-2-1-7-1-3-1-19-2 Bacteroides Species Pneumonia
4-3-3-2-1-7-1-3-1-19-3 Peptostreptococcus Species Pneumonia
4-3-3-2-1-7-1-3-1-19-4 Microaerophilic Streptococcus Pneumonia
4-3-3-2-1-7-1-3-1-20 Actinomyces Pneumonia
4-3-3-2-1-7-1-3-1-21 Nocardia Species Pneumonia
4-3-3-2-1-7-1-3-1-22 Mycoplasma Pneumonia
4-3-3-2-1-7-1-3-1-23 Coxiella Burnetti Pneumonia
4-3-3-2-1-7-1-3-1-24 Chlamydia Psittaci Pneumonia
4-3-3-2-1-7-1-3-1-25 Chlamydia Trachopmatis Pneumonia
4-3-3-2-1-7-1-3-1-26 Pseudomonas Pseudomallei Pneumonia
4-3-3-2-1-7-1-3-1-27 Paturella Pneumonia
4-3-3-2-1-7-1-3-1-28 Francisella Pneumonia
4-3-3-2-1-7-1-3-1-29 Yersinia Pestis Pneumonia
4-3-3-2-1-7-1-3-1-30 Bacillis Anthracis Pneumonia
4-3-3-2-1-7-1-3-1-31 Brucella Species Pneumonia
4-3-3-2-1-7-1-3-1-32 Chlamydial Pneumonia
4-3-3-2-1-7-1-3-1-33 Mycobacterial Pneumonia
4-3-22-1 Bacterial Disease
4-3-22-1-1 Bacteriogenic Pneumonia
4-3-22-1-1-2 Pneumococcus Pneumonia
4-3-22-1-1-3 Staphylococcal Pneumonia
4-3-22-1-1-3-1 Primary Staphylococcal Pneumonia
4-3-22-1-1-3-2 Secondary Staphylococcal Pneumonia
4-3-22-1-1-4 Streptococcal Pneumonia
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Figure 8 - Bacterial pneumonia coded in the Gabrieli (ASTM) Medical Nomenclature. Sixteen descendants of Mycobacterial pneumonia not shown. Some terms appear in multiple locations (e.g., Staphylococcal Pneumonia, which has additional descendants in one context). Note that Bacterial Pneumonia and Bacteriogenic Pneumonia are not considered synonymous and have different descendants. Similarly, Streptococcus Pneumonia (4-3-3-2-1-7-1-3-1-2) and Streptococcal Pneumonia (4-3-22-1-1-4) are not considered synonymous. Additional bacterial pneumonias can be found elsewhere in the hierarchy, such as Listerial Pneumonia (4-3-22-1-29-6-1), Staphylococcus Aureus Pneumonia in a Granulocytopenic Host (4-3-3-2-1-7-1-1-1-2), its child Staphylococcus Epidermidis Pneumonia in a Granulocytopenic Host, and Staphylococcus Pneumonia in Children (16-10-5-7-2-14-1-3).

tem, first developed at the University of Buffalo, was adopted for use in a proprietary system. It consists of a single, large hierarchy which contains successively more complex expressions as one moves down through the hierarchy. The aim of this system is to take precoordination to the extreme, providing a code for each utterance that might be found in a medical record (see Figure 8). Although initially available as a commercial product, the developers have used it as the basis for nomenclature work under the American Society for Testing and Materials (ASTM an international standards organization based in the US) [57]. The ASTM is currently working to move this nomenclature through the standards development process.

In Europe, a consortium of universities, agencies and vendors, with funding from the Advanced Informatics in Medicine initiative (AIM), has formed the GALEN project to develop standards for representing coded patient information [58]. GALEN is developing a reference model for medical concepts using a formalism based on the SMK of PEN & PAD. The reference model is intended to allow representation of patient information in a way that is independent of the language being recorded and independent of the data model used by an EMR system. The GALEN developers are working closely with the Technical Committee on Medical Informatics (TC251) of the Comité Europeén de Normalisation (CEN) to develop the content that will populate the reference model with actual terms.

A collaborative effort is currently under way between ASTM (LOINC) [59] and CEN (EUCLIDES) [60] to develop the reference model and content for the domain of laboratory test names. The standard specifies structured coded semantic information about each test, such as the sub-

stance measured and the analytical method used. Rather than establish a vocabulary for use in laboratory systems, this standard is aimed at providing a vocabulary into which local laboratory terms can be mapped for exchange with other institutions.

The Canon Group [61] has experimented with the use of conceptual graphs as a form of concept representation. Using this approach, they have experimented with collaborative vocabulary development. The development work thus far has resulted in a reference model and content for the domain of chest radiograph reports which can serve a variety of purposes, including natural language processing, predictive data entry and automated decision support [62].

For some time, the NLM has been developing the Unified Medical Language System (UMLS) [63] to serve a number of controlled vocabulary needs [64]. Included in the UMLS is the Metathesaurus, which contains concepts, and the UMLS Semantic Net, which provides information about how the semantic classes of concepts can be interrelated. The

Bacterial pneumonia Pneumonia, Lobar Pneumonia, Staphylococcal Pneumonia, Streptococcal Pneumonia due to Streptococcus Pneumonia in anthrax Pneumonia, anthrax Bronchopneumonia Pasteurellosis, Pneumonic Salmonella Pneumonia Other bacterial Pneumonia Pneumonia due to Klebsiella Pneumoniae Pneumonia due to other specified bacteria Pneumonia in whooping cough Pneumonia due to Pseudomonas

Figure 9 - Pneumonia concepts in the Unified Language Systems (UMLS) Metathesaurus.

Pneumonia due to Hemophilus

influenzae (H. influenzae)

concepts in the Metathesaurus are drawn from established controlled vocabularies, such as MeSH, ICD. 9-CM, and SNOMED. Information about each concept includes the preferred form of the concept in the various source vocabularies, synonyms and lexical variants of the concepts, and information about relationships between specific concepts (Figures 9 and 10). Various uses for the UMLS have been described, including the coding of patient data. However, the NLM has acknowledged that the UMLS does not serve clinical encoding well. This is largely due to the fact that the source vocabularies do not themselves serve this function. The NLM is now developing ways in which the UMLS can be enhanced to support the coding of clinical data and has enlisted the help of a large number of researchers (including most of the Canon Group) to provide input for and evaluation of this UMLS expansion.

Finally, vocabulary servers have become a research issue in their own right. The servers are intended to provide open, distributed health care systems with information about upto-date vocabulary content. Groups working on vocabulary servers include GALEN [65], the NLM [66], the University of Utah [67], and Stanford University [68].

5. Research Issues

The preceding discussions of standard codes for abstraction, codes for electronic medical records, and current research efforts supports my opening statement that no accepted standard exists for coding patient information. In the past, the tendency for developers to create their own coding schemes, rather than adopt an existing one, may have been due to the "Not Invented Here" phe

Bacterial pneumonia

Source: CSP93/PT/2596-5280; DOR27/DT/U000523; ICD91/PT/482.9; ICD91/IT/482.9

Parent: Bacterial Infections; Pneumonia; Influenza with Pneumonia

Child: Pneumonia, Mycoplasma

Narrower: Pneumonia, Lobar; Pneumonia, Rickettsial; Pneumonia, Staphylococcal; Pneumonia due to Klebsiella

Pneumoniae; Pneumonia due to Pseudomonas; Pneumonia due to Hemophilus influenzae (H. influenzae)

Other: Klebsiella Pneumoniae, Streptococcus Pneumoniae

Pneumonia, Lobar

Source: ICD91/IT/481; MSH94/PM/D011018; MSH94/MH/D011018; SNM2/RT/M-40000; ICD91/PT/481; SNM2/PT/

D-0164; DXP92/PT/U000473; MSH94/EP/D011018; INS94/MH/D011018; INS94/SY/D011018

Synonym: Pneumonia, diplococcal

Parent: Bacterial Infections; Influenza with Pneumonia

Broader: Bacterial Pneumonia; Inflammation

Other: Streptococcus Pneumoniae
Semantic: inverse-is-a: Pneumonia

has-result: Pneumococcal Infections

Pneumonia, Staphylococcal

Source: ICD91/PT/482.4; ICD91/IT/482.4; MSH94/MH/D011023; MSH94/PM/D011023; MSH94/EP/D011023; SNM2/

PT/D-017X; INS94/MH/D011023; INS94/SY/D011023

Parent: Bacterial Infections; Influenza with Pneumonia

Broader: Bacterial Pneumonia

Semantic: inverse-is-a: Pneumonia; Staphylococcal Infections

Pneumonia, Streptococcal

Source: ICD91/IT/482.3

Other: Streptococcus Pneumoniae

Pneumonia due to Streptococcus

Source: ICD91/PT/482.3

ATX: Pneumonia AND Streptococcal Infections AND NOT Pneumonia, Lobar

Parent: Influenza with Pneumonia

Pneumonia in Anthrax

Source: ICD91/PT/484.5; ICD91/IT/022.1; ICD91/IT/484.5

Parent: Influenza with Pneumonia

Broader: Pneumonia in other infectious diseases classified elsewhere

Other: Pneumonia, Anthrax

Pneumonia, Anthrax

Source: ICD91/IT/022.1; ICD91/IT/484.5

Other: Pneumonia in Anthrax

Figure 10 - Some of the information available in the UMLS about selected pneumonia concepts. Concept preferred names are shown in italics. Sources are identifiers for the concept in other vocabularies. Synonyms are names other than the preferred name. ATX is an associated MeSH expression which can be used for Medline searches. The remaining fields (Parent, Child, Broader, Narrower, Other and Semantic) show relationships between concepts in the Metathesaurus. Note that concepts may or may not have hierarchical relations to each other through Parent/Child, Broader/Narrower, and Semantic (is-a/inverse-is-a) relations. Note also that Pneumonia, Streptococcal and Pneumonia due to Streptococcus are

nomenon. However, as systems have become larger and begin to address more comprehensive domains (such as EMRs), developers are quite willing to take advantage of existing standards. Their continued inability to do so points to failure on the part of controlled vocabularies to meet their needs. The developers of the locabularies, on the other hand, have continued to be surprised at this resistance to use. The source of the

problem is that the vocabularies are created for specific purposes and often have characteristics which limit their usefulness for other purposes. The standards developers are investing considerable effort to address this problem. In the process, a variety of issues has come to light. Some of these are internal issues, dealing with the structure and content of the vocabularies themselves, and others are external issues, dealing with the

relationships between vocabulary developers and vocabulary users.

Internal Issues

The first basis on which vocabularies are judged is their content. A user cannot adopt a coding scheme if it does not have the ability to express the necessary concepts. The vocabulary domains are moving targets as medical knowledge grows with new terms to add and old ones

to discard. The developers of the comprehensive vocabularies devote substantial energy into expanding their content. This usually involves development of committees and interaction with professional specialty groups to provide input. As a result, the large vocabularies being built today seem to be coming close to having the content needed.

One place where vocabularies have run into trouble has been the codes they use to represent terms. In many cases, the codes are designed to reflect the position of the term in the hierarchy. There is a certain elegance to this approach; however, in the real world of medical terminology, this elegance breaks down. If the code has a limited number of positions or digits, then the depth of the hierarchy is limited. If the positions in the code are limited to a fixed number of characters, then the breadth of the hierarchy is limited. These limitations can adversely affect vocabulary content, since some domains become too full to allow additional terms, requiring the use of catch-all "Other" terms. In addition, multiple hierarchies (see below) cannot be accommodated with a single code.

Vocabulary developers are addressing the coding issue by divesting the unique identifiers for the terms from their hierarchical positions. Among the comprehensive coding systems, only SNOMED continues to use a hierarchy-based unique identifier. The remainder either provide hierarchical information as semantic links or they allow tree addresses which can be of arbitrary length and breadth.

A related issue is the need for medical terms to be organized in multiple classes. If a vocabulary permits only a single hierarchy, it will invariably be the one that meets the developer's view of the world. When this view differs from the user's view, the user may look elsewhere for a coding scheme. For example, users may wish to be able to access patient diagnoses based on location or on etiology. This becomes awkward when the user, for example, wants to identify all patients with bacterial pneumonia but the coding scheme scatters the codes as in ICD, with some in the Pneumonia class, and others in the various bacterial disease classes.

Most vocabulary developers have recognized the need to accommodate multiple classes and allow them. This has been simplified by the departure from the use of hierarchical codes. In systems such as Read, GALEN and UMLS, hierarchies are represented as links between parents and children, so multiple hierarchies are simply the result of multiple links. In systems which use tree addresses, such as MeSH and the Gabrieli Nomenclature, the solution is simply to allow terms to have multiple tree addresses. Still to be resolved are the issues of variation of meaning and variation of children across different hierarchical addresses for the same term.

Researchers are realizing, though, that allowing multiple classification was the easy part. As the structures of the vocabularies become more powerful and complex, the task of where to place a term becomes as important as what term to place [69]. New techniques are being explored by several groups to take advantage of the semantic information included about the terms, either as frames, semantic nets, or conceptual graphs. One of these techniques is automated term subsumption, long used in artificial intelligence research, in which the attributes of the term define its location. For example, if the ICD-9-CM term "Salmonella pneumonia" included attributes that identify it as being caused by Salmonella and occurring in the lung, it might be possible to automatically assign it as a child of both of the desired parents.

A continuing controversy in vocabulary development revolves around the choice between precoordination and postcoordination. On one hand, a precoordinated term like "Salmonella pneumonia" is probably a useful concept and more natural than the combination "Salmonella"+"Pneumonia" On the other hand, precoordination can easily lead to combinatorial explosion as all permutations of all modifiers are appended to terms in order to have a preassigned code for the composite. Attempting to choose one or the other approach is probably not feasible. Terms which seem reasonably atomic to one user of the vocabulary will seem to some other user to be a precoordination of smaller concepts, Precoordinated terms will often be found to be missing some minute detail, requiring the addition of a modifier, turning it into a postcoordination. The reality is that vocabularies which do not allow postcoordination are usually too limiting, while those that allow postcoordination always have a healthy collection of precoordinated terms. The use of conceptual graphs, as described in the appendicitis example in SNOMED, may accommodate both approaches while allowing equivalence between a precoordinated term and a postcoordinated phrase to be recognized.

External Issues

Once vocabularies are created, continuity needs to be maintained. Besides the issues related to how to include new terms (described above), there are epistemologic issues related to identifying new terms for inclusion and marking old ones for deletion. Monitoring usage of terms, such as is done by the National Library of Medicine for MeSH [70] will be important for determining what users need. Changes will include the addition of new terms, the addition of new classes.

or aggregations of terms, the addition of an existing term to an existing class, identification of a particular type of (semantic) relationship between two terms, and the addition of entirely new types of relationships.

The development of mechanisms for responding to needs for additions will be crucial for the success of any controlled vocabulary, since the lack of necessary terms in a standard coding scheme will merely push system developers to create their own coded terminologies. Any vocabulary that is interested in meeting user needs would do well to follow the lead of the NLM, which requests UMLS users to submit suggestions for changes via electronic mail [71].

An important part of maintaining a vocabulary is the communication of changes to the users. The traditional method has been to convene a committee of experts periodically to review the current version of a vocabulary and prescribe changes. This approach seems to result in updates measured in years and decades. However, for many applications, this is inadequate. For example, if a new drug goes on the market, or a new test can be ordered from the laboratory, waiting a year - or even a day - is too long if the new term is encountered and needs to be coded immediately. Users need to get changes as soon as they are available. This issue is being addressed in the various projects to develop vocabulary servers. Such servers will facilitate the dissemination of changes from the central authority and also provide a link back to the authority to recommend changes when they are seen, rather than waiting for the next standards-setting group to meet.

6. Conclusion

The application of computers to medicine has accelerated the breadth of uses and depth of detail needed

for the representation of patient data. Legacy abstracting systems were recognized as inadequate for applications such as electronic medical records and automated decision support, but simply expanding their content has not solved the problem. Today, research into medical data representation is livelier than ever, as formal computer science techniques are being applied to large, real-world domains. Local solutions have shown great promise for the application builders who have had the resources needed for vocabulary development. For those who do not have such resources, current efforts to develop thoughtful solutions at national and international levels are under way.

Acknowledgements

I thank my collaborators on the InterMed project, sponsored by the National Library of Medicine, and the members of the Canon group for their stimulating discussions on vocabulary issues. I also thank Leslie Juceam and George Hripcsak for their contributions to the editing of the manuscript.

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