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AN EXPLORATION OF DEEP LEARNING IN CONTENT-BASED
MUSIC INFORMATICS

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There once was a time I thought it was some happy accident that I kept finding myself in exactly the right place. Be it college, career, or concert, it always felt like gravity, quietly inevitable, pulling me into the things I should do, the places I should go, the opportunities I should pursue. But now, on the backside of this arc, I can see quite plainly that I always had the direction wrong; it was never a pull, but a *push*. Come to think of it, for as long as I can seem to recall, I have been lucky enough to be surrounded by those who would inspire, encourage, force, drag, and challenge me to always be better. I am nothing short of blessed to be graced by the company and influence of such wonderful human beings, and will try my best to thank as many of you as I can here.

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“What’s it matter? Does it matter,
If we’re all matter when we’re done,
When the sky is full of zeros and ones?”

-Andrew Bird, *Masterfade*

CHAPTER I

INTRODUCTION

It goes without saying that we live in the Age of Information, our day to day experiences awash in a flood of data. As a society, we buy, sell, consume and produce information in unprecedented quantities. Given the accelerating rate at which information is created, one of the fundamental challenges facing the modern world is simply making sense of all this data. The quintessential response to this obstacle is embodied by Google, whose collective *raison d'être* is the organization and indexing of the world's information. To appreciate the value and reach of this technology, one only needs to imagine how difficult it would be to browse the Internet without a search engine.

Understandably, a variety of specialized disciplines have formed under the auspices of developing systems to help people navigate and understand massive amounts of information. Coalescing around the turn of the century, music informatics is one such instance, drawing from several diverse fields including electrical engineering, music psychology, computer science, machine learning, and music theory, among others. Now encompassing a wide spectrum of application areas and the kinds of data considered—from audio and text to album covers and online social interactions—music informatics can be broadly defined as the study of information related to, or is a result of, musical activity.

At a high level, tackling this problem of “information overload” in music is captured by a simple, general analogy: how exactly *does* one find a needle in

a haystack? To answer this question, any system, computational or otherwise, must solve two related problems: first, it is necessary to describe the intrinsic qualities of the item of interest, e.g. a needle is metal, sharp, thin, etc; and second, it is necessary to evaluate the extrinsic relationships between items to determine relevance. A piece of hay is certainly not a needle, for example, but is a pin close enough? Along what criteria might we gauge similarity, or classify objects into groups? Emphasizing the distinction, *description* focuses on absolute representation, whereas *comparison* is concerned with relative associations.

To date, the most successful approaches to large-scale information systems leverage human-provided signals to achieve rich content descriptions. Building on top of robust representations simplifies the problem greatly, and good progress has been made toward the development of useful applications. For example, the Netflix Prize challenge¹ —an open contest to find the best system for automatically predicting a user’s enjoyment of a movie— was built exclusively on movie ratings contributed by a large collection of other users. Similarly, Google’s *PageRank* algorithm associates websites based on how users have linked different pages together, thus facilitating the process of traversing the Internet (Page, Brin, Motwani, & Winograd, 1999).

While this strategy of leveraging manual content description has proven successful in large-scale music recommendation, such as Pandora Radio², its application to more general music information problems is fundamentally limited, manifesting in three related ways. First, human-provided information

¹<http://www.netflixprize.com/>

²<http://www.pandora.com/>

commonly used in such systems —clicks, likes, listens or shares— are easily captured from, or as a natural by-product of, a user’s listening to music. It is one thing to obtain a “thumbs up” for a song; it is quite another to ask that same user to provide a chord transcription of it. Second, manual music description may require a high degree of expertise or effort to perform. The average music listener is not truly capable of transcribing chords from a sound recording, whether or not she possesses the time or willingness to attempt it. Finally, even given the skill, motivation, and infrastructure to manually describe music, this approach cannot scale to *all* music content, now or in the future. The Music Genome Project¹, for example, has resulted in the manual annotation of some 1M commercial recordings, at a pace of 20-30 minutes per track; the iTunes Music Store, however, now offers over 43M tracks worldwide². To illustrate how vast this discrepancy is, consider the following: even assuming the lower bound of 20 minutes, it would still take one sad individual *1,600 years* of non-stop annotation to close that gap. More importantly, this only considers commercial music recordings, neglecting amateur or unpublished content from websites like YouTube³ or Soundcloud⁴, the addition of which makes this goal even more insurmountable. Given the sheer impossibility for humans to meaningfully describe all recorded music, now and in the future, truly scalable music information systems will require good automatic systems to perform this task.

Thus, the development of computational systems to describe music sig-

¹<https://www.pandora.com/about/mgp>

²According to <http://www.apple.com/itunes/music/>, accessed 20 April, 2015.

³<https://www.youtube.com/>

⁴<https://soundcloud.com/>

nals, a flavor of *computer audition* referred to as content-based music informatics, is both a valuable and fascinating problem. In addition to facilitating the search and retrieval of large music collections, automatic systems capable of expert-level music description are invaluable to users who are unable to perform the task themselves, e.g. music transcription. Notably, this problem is also very much unsolved, and given an apparent deceleration of progress, some in the field of music informatics have begun to question the efficacy of traditional research methods. Simultaneously, in the related fields of computer vision and automatic speech recognition, a branch of machine learning, referred to as *deep learning*, has shown great performance in various domains, toppling many long-standing benchmarks. On closer inspection, one recognizes considerable conceptual overlap between deep learning and conventional music signal processing systems, further encouraging this promising union.

Synthesizing these observations, this study explores deep learning as a general approach to the design of computer audition systems for music description applications. More specifically, the proposed research method proceeds thusly: first, methods and trends in content-based music informatics are reviewed in an effort to understand why progress in this domain may be decelerating, and, in doing so, identify possible deficiencies in this methodology; standard approaches to music signal processing are then reformulated in the language and concepts of deep learning, and subsequently applied to classic music informatics problems; finally, the behavior of these deep learning systems is deconstructed in order to illustrate the advantages and challenges inherent to this paradigm.

1 Scope of this Study

This study explores the use of deep learning in the development of systems for automatic music description. Consistent with the larger body of machine perception research, the work presented here aims to computationally model the relationship between stimuli and observations made by an intelligent agent. In this case, “stimuli” are digital signals representing acoustic waves, e.g. sound, “observations” are semantic descriptions in a particular namespace, e.g. timbre or harmony, and the agent being modeled is an intelligent human, e.g. an expert music listener. In practice, the namespace of descriptions considered is constrained to a particular task or application, such as instrument recognition or chord estimation.

Furthermore, if the relationship between stimuli and observation is not a function of the agent, this mapping is said to be “objective”. Objective relationships are those that are true absolutely by definition, such as the statement “A C Major triad consists of the pitches C, E, and G.” Elaborating, all sufficiently capable agents should always produce the same output given the same input. Discrepancies between observations of the same stimuli are understood as one or more of these perspectives being erroneous, resulting from either simple error, bias, or a deficiency of knowledge. For objective relationships, the *quality* of a model is determined by how often it is able to produce “right” answers, often referred to as “ground truth”, to the questions being asked.

Conversely, input-output relationships that *are* a meaningful function of the agent are said to be “subjective”. In contrast to the objective case, which is fundamentally concerned with *facts*, a subjective observation is ultimately an *opinion*. As such, an opinion can only be true or false insofar as it is

held by a competent agent. This is embodied, for example, in the statement “That sounds like a saxophone.” Whether or not the stimuli originated from a saxophone is actually irrelevant; a rational agent has made the observation, and thus it is in some sense valid. Assessing the quality of a computational model at a subjective task must therefore take one of two slightly different formulations. The first transforms a subjective problem to an objective one by considering the perspective of single agent as truth, and thus the quality of a model is a function of how well it can mimic the behavior of that *one* agent. Alternatively, the other approach attempts to determine whether or not a model makes observations on par with other competent agents. In this view, a computational system’s capacity to perform some intelligent task is measured by its ability to convince humans that it is competent (or not) in human ways, e.g. the Turing test (Turing, 1950).

The notions of, and inherent conflict between, objectivity and subjectivity in audition and music perception are central to the challenge posed by the computational modeling of it. Arguably most facets of music perception are subjective and vary in degree from task to task. However, while subjective evaluation might be better suited toward measuring the quality or usability of some computational system, the human involvement required by such assessments make them prohibitively costly in both time and money to conduct with any regularity. As a result, conventional research methodology in engineering and computer science greatly prefers *quantitative* evaluation as a proxy to *qualitative* responses collected from human subjects. Typically quantitative methods proceed by collecting some number of input-output pairs from one or more human subjects beforehand, and treating this data sample as objective

truth. Thus, regardless of whether or not a given task is indeed objective, it is a significant simplification in methodology to treat it as one.

This is all to say that the validity and quality of a music description is often determined by an objective fitness measure, not necessarily out of correctness but rather tractability. Therefore, any quantitative measure is only valid insofar as the assumption of objectivity is as well.

2 Motivation

The proposed research is primarily motivated by two complementary observations: one, large scale music signal processing systems are becoming necessary to help humans navigate and make sense of an ever-increasing volume of music information; two —and, more notably, the specific problem this work seeks to address— the conventional research tradition in content-based music information retrieval is yielding diminishing returns, despite many research areas remaining unsolved.

In the most immediate sense, the proposed research will develop systems to tackle various applications in music informatics. This will at least serve to explore an alternative approach to conventional problems in the field. Based on preliminary results, there is good reason to believe that deep learning may in fact push the state of the art in some, if not most, applications in automatic music description. Sufficiently advanced systems could be deployed in end-user applications, such as navigating music libraries or computer-aided composition and performance.

A thorough exploration and successful extension of deep learning to music signal processing has the potential to encourage a broader study of these

methods. The impacts of such a development could be far reaching, but there are two of particular note. First and foremost, drawing attention to a promising, but otherwise uncharted, research area opens new opportunities for fresh ideas and perspectives. Additionally, deep learning automatically optimizes a system to the data on hand, accelerating research and simplifying the overall design problem. Therefore these methods yield flexible systems that can easily adapt to new data as well as new problems, allowing researchers to seek out novel, exciting applications.

Beyond the scope of music informatics, deep learning research in the context of a different domain, with its own unique challenges, is likely to produce discoveries beneficial to the broad audience of computer science and information processing. One such area where this is likely to occur is in the handling of time-domain signals and sequences. Computer vision, the field in which most breakthroughs in deep learning have occurred, has invested considerable effort in the study of static, 2D images. Certainly some have extended these techniques to image sequences and video, but this is far more the exception than the rule. Other sequential data, such as natural language, in the form of text, speech signals, and motion capture data have also seen a deal of study in deep learning circles. The tradition of music signal processing draws heavily from digital signal theory, a field of study with a considerable focus on an analytical understanding of time.

Therefore, this work offers several potential contributions, both theoretical and practical, to a diverse audience, spanning users of technology, music informatics, and the deep learning community on the whole.

3 Dissertation Outline

Chapter II reviews the current state of affairs in music informatics research, providing context for this work.

Chapter III surveys the body of literature in deep learning, outlining core concepts and definitions.

Chapter IV explores the application of deep learning toward the development of objective timbre similarity spaces.

Chapter V considers the application of deep learning toward automatic chord estimation, as a means to both improve the state of the art and better understand the task at hand.

Chapter VI extends this chord estimation efforts to directly estimate human-readable representations in the form of guitar tablature.

Chapter VII documents the software contributions resulting from this study, contributing to the greater cause of reproducible research efforts.

Chapter VIII concludes this thesis, summarizing the work presented and offering perspectives for future work and outstanding challenges.

4 Contributions

The primary contributions of this dissertation are listed below:

- **Demonstrates an objective approach to the development of timbre similarity embeddings.** The proposed approach extends previous efforts in using pairwise training of deep architectures by relaxing constraints on the output space and generalizing the use of margins as a

ratio, rather than an absolute parameter; in addition to realizing a far more discriminative instrument embedding than a shallow comparison system overall, the margin ratio improves performance slightly over the original pairwise training approach.

- **Advances the state of the art in large vocabulary automatic chord estimation, while illustrating methodological limitations in the current formulation of the task.** Comprehensive error analysis is performed both by comparing two state of the systems against the reference chord transcriptions, as well as the proposed system against another dataset with multiple annotators. The insight gleaned from this study is used to offer perspective on future directions for the task at large.
- **Leveraged traditional chord transcriptions to develop an automatic guitar chord estimation system, which directly maps music audio to fingerings on a fretboard.** Not only does this approach improve some measures over the other large-vocabulary chord estimation system presented here, but it provides a user-friendly interface for both learning and soliciting feedback on system errors.
- **Contributes, in whole or part, to several open source projects to facilitate future efforts.** In addition to a suite of tools that may help serve the larger research community, this includes a framework to reproduce the experimental results and analysis contained herein.

5 Associated Publications by the Author

This thesis covers much of the work presented in the publications listed below:

5.1 Peer-Reviewed Articles

- Humphrey, E. J., Bello, J. P., and LeCun, Y. (2013) “Feature learning and deep architectures: new directions for music informatics.” *Journal of Intelligent Information Systems*. 41 (3), 461–481.

5.2 Peer-Reviewed Conference Papers

- Humphrey, E. J., Salamon, J. Nieto, O., Forsyth, J., Bittner, R. M., and Bello, J. P. “JAMS: A JSON Annotated Music Specification for Reproducible MIR Research.” *Proceedings of the 15th International Society of Music Information Retrieval (ISMIR)*, Taipei, Taiwan, October 2014.
- Raffel, C., McFee, B., Humphrey, E. J., Salamon, J. Nieto, O., Liang, D., and Ellis, D. P. W. “mir eval: A Transparent Implementation of Common MIR Metrics” *Proceedings of the 15th International Society of Music Information Retrieval (ISMIR)*, Taipei, Taiwan, October 2014.
- Humphrey, E. J. and Bello, J.P. “From Music Audio to Guitar Tablature: Teaching Deep Convolutional Networks to Play Guitar.” *Proceedings of the International Conference on Acoustic Signals and Speech Processing (ICASSP)*, Florence, Italy, May 2014.
- Humphrey, E. J., Nieto, O., and Bello, J. P. “Data Driven and Discriminative Projections for Large Scale Cover Song Identification.” to appear in *Proceedings of the 14th International Society of Music Information Retrieval (ISMIR)*, Curitiba, Brazil, November 2013.
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CHAPTER II

CONTEXT

From its inception, many fundamental challenges in music informatics, and in particular those that focus on music audio signals, have received a considerable and sustained research effort from the community. Referred to here as automatic music description, this area of study is based on the premise that if a human expert can experience or observe some musical event from an audio signal, it should be possible to make a machine respond similarly. As the field of music informatics continues into its second decade, there are a growing number of resources that comprehensively review the state of the art in music signal processing across a variety of different application areas (A. Klapuri & Davy, 2006; Casey, Veltkamp, et al., 2008; Müller, Ellis, Klapuri, & Richard, 2011), including melody extraction, chord estimation, beat tracking, tempo estimation, instrument identification, music similarity, genre classification, and mood prediction, to name only a handful of the most prominent topics.

After years of diligent effort however, many well-worn problems in content-based music informatics lack satisfactory solutions and remain unsolved. Observing this larger research trajectory at a distance, it would seem progress is decelerating, if not altogether stalled. For example, a review of recent MIREX* results motivates the conclusion quantitatively, as shown in Figure 1. The

*Music Information Retrieval Evaluation eXchange (MIREX): <http://www.music-ir.org/mirex/>

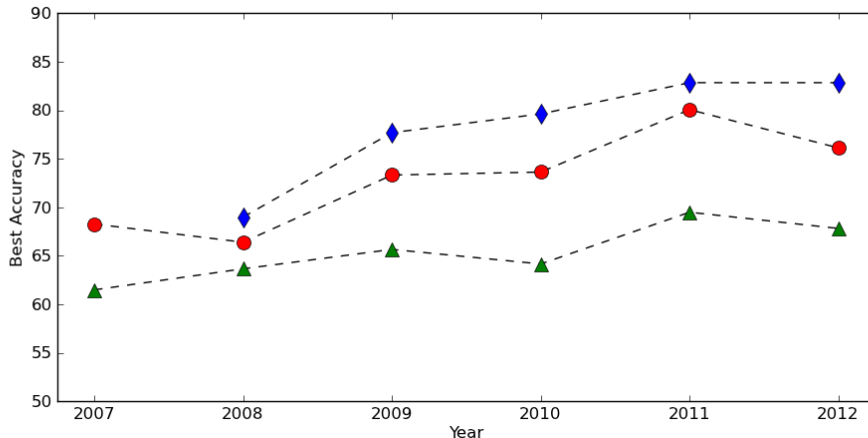


Figure 1: *Losing Steam*: The best performing systems at MIREX since 2007 are plotted as a function of time for Chord Estimation (blue diamonds), Genre Recognition (red circles), and Mood Prediction (green triangles).

three most consistently evaluated tasks for more than the past half decade — chord estimation, genre recognition, and mood prediction— are each converging to performance plateaus below satisfactory levels. Fitting an intentionally generous logarithmic model to the progress in chord estimation, for example, estimates that continued performance at this rate would eclipse 90% in a little over a decade, and 95% some twenty years after that; note that even this trajectory is quite unlikely, and for only this one specific problem (and dataset). Attempts to extrapolate similar projections for the other two tasks are even less encouraging. Furthermore, these ceilings are pervasive across many open problems in the discipline. Though single-best accuracy over time is shown for these three specific tasks, a wider space of MIREX tasks exhibit similar, albeit more sparsely sampled, trends.

Recent research has additionally demonstrated that when state-of-the-art algorithms are used in more realistic conditions, i.e. larger datasets, performance degrades substantially (Bertin-Mahieux & Ellis, 2012). Others have

gone as far as to challenge the very notion that any progress has been made at all, due to issues of problem formulation and validity (Sturm, 2014b). While the truth of the matter likely falls somewhere between “erroneous results” and “sound science”, these varied observations encourage a critical reassessment of content-based music informatics. Does content *really* matter, especially when human-provided information has proven to be more useful than representations derived from the content itself (Slaney, 2011)? If so, what can be learned by analyzing recent approaches to content-based analysis (Flexer, Schnitzer, & Schlueter, 2012)? Do applications in content-based music informatics lack adequate formalization and rigorous validation (Sturm & Collins, 2014)? Is the community considering all possible approaches to solve these problems (Humphrey, Bello, & LeCun, 2012)?

Building on the premise that automatic music description is indeed valuable, this chapter is an attempt to answer the remainder of these questions. Section 1 critically reviews conventional approaches to content-based analysis and identifies three major deficiencies of current systems: the sub-optimality of hand-designing features, the limitations of shallow architectures, and the short temporal scope of conventional signal processing. Section 2 then introduces the ideas of deep architectures and feature learning in terms of music signal processing, two complementary approaches to system design that may alleviate these issues, and surveys the application of these methods in this domain. Finally, Section 3 summarizes the concepts covered herein, and discusses why it is critical point in time for the music informatics community to consider alternative approaches.

1 Reassessing Common Practice in Automatic Music Description

Despite a broad spectrum of application-specific problems, the vast majority of music signal processing systems adopt a common two-stage paradigm of feature extraction and semantic interpretation. Leveraging substantial domain knowledge and a deep understanding of digital signal theory, researchers carefully architect signal processing systems to capture useful signal-level attributes, referred to as *features*. These signal features are then provided to a pattern recognition machine for the purposes of assigning semantic meaning to observations. Crafting good features is a particularly challenging subproblem, and it is becoming standard practice amongst researchers to use precomputed features* or off-the-shelf implementations†, focusing instead on increasingly more powerful pattern recognition machines to improve upon prior work. While early research mainly employed simple classification strategies, such as nearest-neighbors or peak-picking, recent work makes extensive use of sophisticated and versatile techniques, e.g. Support Vector Machines (Mandel & Ellis, 2005), Bayesian Networks (Mauch & Dixon, 2010a), Conditional Random Fields (Sumi, Arai, Fujishima, & Hashimoto, 2012), and Variable-Length Markov Models (Chordia, Sastry, & Sentürk, 2011).

This trend of squeezing every bit of information from a stock feature representation is suspect because the two-tier perspective hinges on the premise that *features are fundamental*. Such representations must be realized in such a way that the degrees of freedom are informative for a particular task; features are said to be *robust* when this is achieved, and *noisy* when variance

* Million Song Dataset

† MIR Toolbox, Chroma Toolbox, MARSYAS, Echonest API

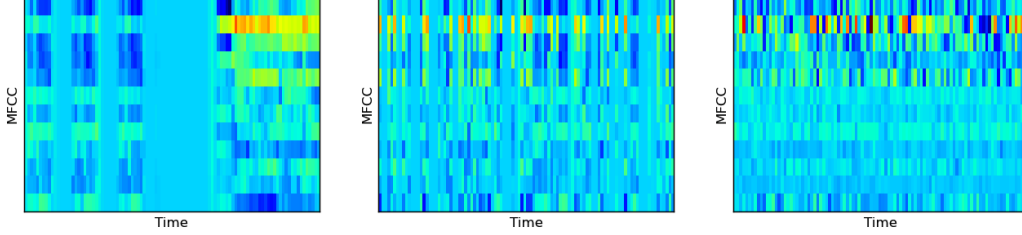


Figure 2: *What story do your features tell?* Sequences of MFCCs are shown for a real music excerpt (left), a time-shuffled version of the same sequence (middle), and an arbitrarily generated sequence of the same shape (right). All three representations have equal mean and variance along the time axis, and could therefore be modeled by the exact same distribution.

is misleading or uninformative. The more robust a feature representation is, the simpler a pattern recognition machine needs to be, and vice versa. It can be said that robust features *generalize* by yielding accurate predictions of new data, while noisy features can lead to the opposite behavior, known as *over-fitting* (Bishop, 2006).

The substantial emphasis traditionally placed on feature design demonstrates that the community tacitly agrees, but it is a point worth illustrating. Consider the scenario presented in Figure 2. Conceptually, the generic approach toward determining acoustic similarity between two music signals proceeds in three stages: short-time statistics are computed to characterize acoustic texture, e.g. Mel-Frequency Cepstral Coefficients (MFCCs); the likelihood that a feature sequence was drawn from one or more probability distributions is measured, e.g. a Gaussian Mixture Model (GMM); and finally, a distance is computed between these representations, e.g. KL-divergence, Earth mover’s distance, etc. (Berenzweig, Logan, Ellis, & Whitman, 2004). Importantly, representing time-series features as a probability distribution discards temporal structure. Therefore, the three feature sequences shown—a real excerpt, a

shuffled version of it, and a randomly generated one with the same statistics—are identical in the eyes of such a model. The audio that actually corresponds to these respective representations, however, will certainly not *sound* similar to a human listener.

This bears a significant consequence: any ambiguity introduced or irrelevant variance left behind in the process of computing features must instead be resolved by the pattern recognition machine. Previous research in chord estimation has explicitly shown that better features allow for simpler classifiers (Cho, Weiss, & Bello, 2010), and intuitively many have spent years steadily improving their respective feature extraction implementations (Lyon, Rehn, Bengio, Walters, & Chechik, 2010; Müller & Ewert, 2011). Moreover, there is ample evidence these various classification strategies work quite well on myriad problems and datasets (Bishop, 2006). The logical conclusion to draw from this observation is that underperforming automatic music description systems are more likely the result of deficiencies in the feature representation than the classifier applied to it.

It is particularly prudent then, to examine the assumptions and design decisions incorporated into feature extraction systems. In music signal processing, audio feature extraction typically consists of a recombination of a small set of operations, as depicted in Figure 3: splitting the signal into independent short-time segments, referred to as blocks or frames; applying an affine transformation, generally interpreted as either a projection or filterbank; applying a point-wise nonlinear function; and pooling across frequency or time. These operations can be, and often are, repeated in the process. For example, MFCCs are computed by filtering a signal segment at multiple frequencies on a Mel-scale (affine transform), taking the logarithm (a nonlinearity), and apply-

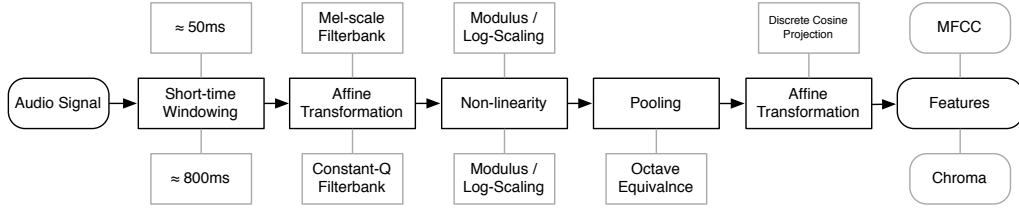


Figure 3: *State of the art*: Standard approaches to feature extraction proceed as the cascaded combination of a few simpler operations; on closer inspection, the main difference between chroma and MFCCs is the parameters used.

ing the Discrete Cosine Transform (another affine transformation). Similarly, chroma features are produced by applying a constant-Q filterbank (affine transformation), taking the complex modulus of the coefficients (non-linearity), and summing across octaves (pooling).

Considering this formulation, there are three specific reasons why this approach might be problematic. First, though the data-driven training of classifiers and other pattern recognition machines has been standard for over a decade in music informatics, the parametrization of feature extractors —e.g. choice of filters, non-linearities and pooling strategies, and the order in which they are applied— remains, by and large, a manual process. Both feature extraction and classifier training present the same basic problem: there exists a large space of possible signal processing systems and, somewhere in it, a configuration that optimizes an objective function over a dataset. Though the music informatics community is privileged with a handful of talented researchers who are particularly adept at exploring this daunting space, crafting good features can be a time consuming and non-trivial task. Additionally, carefully tuning features for one specific application offers no guarantees about relevance or versatility in another scenario. As a result, features developed for one task

are used in others for which they were not specifically designed. The caveat of repurposing features designed for other applications is that, despite potentially encouraging results, they have yet to be optimized for this new use case. Good features for chord estimation may blur out melodic contours, for example, and this information might be particularly useful for structural analysis. In fact, recent research has demonstrated that better features than MFCCs exist for *speech recognition* (Mohamed et al., 2011), the very task for which they were designed, so it is reasonable to assume that there are better musical features as well. The conclusions to draw from this are twofold: continuing to manually optimize a feature representation is not scalable to every problem, and the space of solutions considered may be unnecessarily constrained.

Second, these information processing architectures can be said to be *shallow*, i.e. incorporating only a few non-linear transformations in their processing chain. Sound is a complex phenomena, and shallow processing structures are placed under a great deal of pressure to accurately characterize the latent complexity of this data. Feature extraction can thusly be conceptualized as a function that maps inputs to outputs with an order determined by its *depth*; for a comprehensive discussion on the merits and mathematics of depth, we refer the curious reader to (Bengio, 2009). Consider the example in Figure 4, where the goal is to compute a low-dimensional feature vector (16 coefficients) that describes the log-magnitude spectrum of a windowed violin signal. One possible solution to this problem is to use a *channel vocoder* which, simply put, low-pass filters and decimates the spectrum, producing a piece-wise linear approximation of the envelope. It is clear, however, that with only a few linear components we cannot accurately model the latent complexity of the data, obtaining instead a coarse approximation. Alternatively, the *cepstrum*

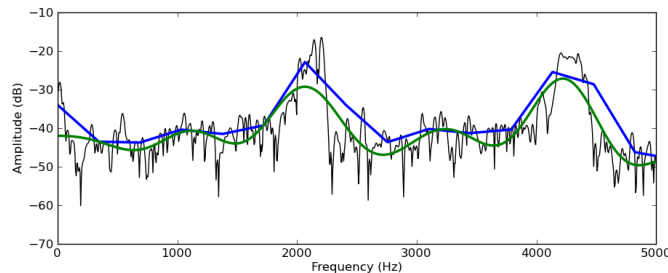


Figure 4: *Low-order approximations of highly non-linear data:* The log-magnitude spectra of a violin signal (black) is characterized by a channel vocoder (blue) and cepstrum coefficients (green). The latter, being a higher-order function, is able to more accurately describe the contour with the same number of coefficients.

method transforms the log-magnitude spectrum before low-pass filtering. In this case, the increase in depth allows the same number of coefficients to more accurately represent the envelope. Obviously, powerful pattern recognition machines can be used in an effort to compensate for the deficiencies of a feature representation. However, shallow, low-order functions are fundamentally limited in the kinds of behavior they can characterize, and this is problematic when the complexity of the data greatly exceeds the complexity of the model.

Third, short-time signal analysis is intuitively problematic because the vast majority of our musical experiences do not live in hundred millisecond intervals, but at least on the order of seconds or minutes. Conventionally, features derived from short-time signals are limited to the information content contained within each segment. As a result, if some musical event does not occur within the span of an observation—a motif that does not fit within a single frame—then it simply cannot be described by that feature vector alone. This is clearly an obstacle to capturing high-level information that unfolds over longer durations, noting that time is extremely, if not fundamentally, important to how music is perceived. Admittedly, it is not immediately obvious how

to incorporate longer, or even multiple, time scales into a feature representation, with previous efforts often taking one of a few simple forms. *Shingling* is one such approach, where a consecutive series of features is concatenated into a single, high-dimensional vector (Casey, Rhodes, & Slaney, 2008). In practice, shingling can be fragile to even slight translations that may arise from tempo or pitch modulations. Alternatively, *bag-of-frames (BoF)* models consider patches of features, fitting the observations to a probability distribution. As addressed earlier with Figure 2, bagging features discards temporal structure, such that any permutation of the feature sequence yields the same distribution. The most straightforward technique is to ignore longer time scales at the feature level altogether, relying on post-filtering *after* classification to produce more musically plausible results. For this to be effective though, the musical object of interest must live at the time-scale of the feature vector or it cannot truly be encoded. Ultimately, none of these approaches are well suited to characterizing structure over musically meaningful time-scales.

1.1 A Concise Summary of Current Obstacles

In an effort to understand why progress in content-based music informatics may be plateauing, the standard approach to music signal processing and feature design has been reviewed, deconstructing assumptions and motivations behind various decisions. As a result, three potential areas of improvement are identified. So that each may be addressed in turn, it is useful to succinctly restate the main points of this section:

- **Hand-crafted feature design is neither scalable nor sustainable:**

Framing feature design as a search in a solution space, the goal is to

discover the configuration that optimizes an objective function. Even conceding that some gifted researchers might be able to achieve this on their own, they are too few and the process too time-consuming to realistically solve every feature design challenge that will arise.

- **Shallow processing architectures struggle to describe the latent complexity of real-world phenomena:** Feature extraction is similar in principle to compactly approximating functions. Real data, however, lives on a highly non-linear manifold and shallow, low-order functions have difficulty describing this information accurately.
- **Short-time analysis cannot naturally capture higher level information:** Despite the importance of long-term structure in music, features are predominantly derived from short-time segments. These statistics cannot capture information beyond the scope of its observation, and common approaches to characterizing longer time scales are ill-suited to music.

2 Deep Learning: A *Slightly* Different Direction

Looking toward how the research community might begin to address these specific shortcomings in modern music signal processing, there is an important development currently underway in computer science. *Deep learning* is riding a wave of promise and excitement in multiple domains, toppling a variety of long-standing benchmarks (Krizhevsky, Sutskever, & Hinton, 2012; G. Hinton et

al., 2012), while slowly permeating the public lexicon (Brumfiel, 2014; Markoff, 2012). Despite all the attention, however, this approach to solving machine perception problems has yet to gain significant traction in content-based music informatics. Before attempting to formally define deep learning, though, it is useful to break down the ideas behind the very name itself and develop an intuition as to why this area is of particular interest.

2.1 Deep Architectures

It was previously shown that deeper processing structures are better suited to characterize complex data. Such systems can be difficult to design, however, as it can be challenging to decompose an abstract music intelligence task into a logical cascade of operations. That said, the evolution of tempo estimation systems is a perfect example of a deep signal processing structure that developed naturally in the due course of research.

The high-level design intuition behind a tempo tracking system is relatively straightforward and, as evidenced by various approaches, widely agreed upon. First, the occurrence of musical events, or onsets, are identified, and then the underlying periodicity is estimated. The earliest efforts in tempo analysis tracked symbolic events (Dannenberg, 1984), but it was soon shown that a time-frequency representation of sound was useful in encoding rhythmic information (Scheirer, 1998). This led to in-depth studies of onset detection (Bello et al., 2005), based on the idea that “good” impulse-like signals, referred to as *novelty functions*, would greatly simplify periodicity analysis. Along the way, it was also discovered that applying non-linear compression to a novelty function produced noticeably better results (A. P. Klapuri, Eronen, & Astola, 2006). Various periodicity tracking methods were simultaneously ex-

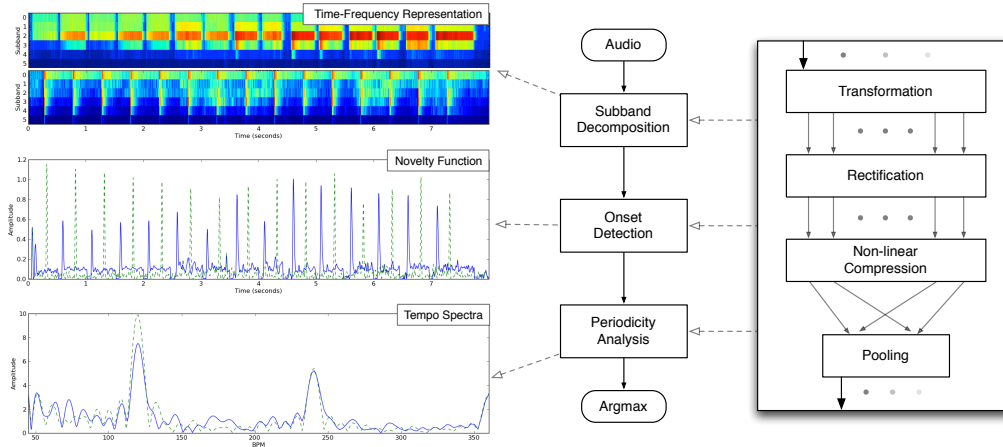


Figure 5: *A complex system of simple parts*: Tempo estimation has, over time, naturally converged to a deep architecture. Note how each processing layer absorbs a different type of variance —pitch, absolute amplitude, and phase— to transform two different signals into nearly identical representations.

plored, including oscillators (Edward & Kolen, 1994), multiple agents (Goto & Muraoka, 1995), inter-onset interval histograms (Dixon, 2007), and tuned filterbanks (Grosche & Müller, 2011).

Reflecting on this lineage, system design has, over time, converged to a deep learning architecture, minus the learning, where the same processing elements —filtering and transforms, non-linearities, and pooling— are replicated over multiple processing layers. Interestingly, as shown in Figure 5, visual inspection demonstrates why it is particularly well suited to the task of tempo estimation. Consider two input waveforms with little in common but tempo; one, an ascending D Major scale played on a trumpet, and the other, a delayed series of bass drum hits. It can be seen that, at each layer, a different kind of variance in the signal is removed. The filterbank front-end absorbs rapid fluctuations in the time-domain signal, spectrally separating acoustic events. This facilitates onset detection, which provides a pitch and timbre

invariant estimate of events in the signal, reducing information along the frequency dimension. Lastly, periodicity analysis eliminates shifts in the pulse train by discarding phase information. At the output of the system, these two acoustically different inputs have been transformed into nearly identical representations. Therefore, the most important lesson demonstrated by this example is how invariance can be achieved by distributing complexity over multiple processing layers.

As mentioned, not all tasks share the same capacity for intuition. Multi-level wavelet filterbanks, referred to as scattering transforms, have also shown promise as a general deep architecture for audio classification by capturing information over not only longer, but also multiple, time-scales (Andén & Mallat, 2011). Recognizing MFCCs as a first-order statistic, this second-order system yielded better classification results over the same observation length while also achieving convincing reconstruction of the original signals. The authors demonstrate their approach to be a multi-layer generalization of MFCCs, and exhibit strong parallels to certain deep network architectures, although the parameterization here is not learned but defined. Perhaps a more intriguing observation to draw from this work though is the influence a fresh perspective can have on designing deep architectures. Rather than propagating all information upwards through the structure, the system keeps summary statistics at each timescale, demonstrating better performance in the applications considered.

2.2 Feature Learning

In traditional music informatics systems, features are tuned manually, leveraging human insight and intuition, and classifiers are tuned automatically,

leveraging an objective criterion and numerical optimization. For this reason, the quality of hand-crafted features is a crucial aspect of system design, as numerical optimization occurs downstream of manual feature design. Many are well aware of the value inherent to good representations, and feature tuning has become a common, if tedious, component in music informatics research. One such instance where this has occurred is in the tuning of chroma features. Developed by Fujishima around the turn of the century (Fujishima, 1999), the last decade and a half has seen consistent iteration and improvement on the same basic concept; estimate the contribution of each pitch class over a short-time observation of audio. Though initially devised for chord estimation, chroma features have been used in a variety of applications, such as structural segmentation (Levy, Noland, & Sandler, 2007) or version identification (Salamon, Serra, & Gómez, 2013).

The fundamental goal in computing chroma features is to consolidate the energies of each pitch class according to a particular magnitude frequency representation. One of the simplest ways to do so, given in Figure 6-(a), shows the averaging of pitch classes in a constant-Q filterbank, e.g. frequencies are spaced like the keys of a piano. Later developments found that weighting the contributions of each frequency with a Gaussian window led to better performance, as shown in Figure 6-(b) (Cho, 2014). This improvement still took time to develop, further motivating the notion that other simple modifications remain undiscovered. That said, this knowledge is attained by maximizing a known objective measure, such as classification accuracy in a chord estimation task. Reflecting, this begs an obvious question: perhaps the parameters of a chroma estimation function could instead be *learned* via numerical optimization?

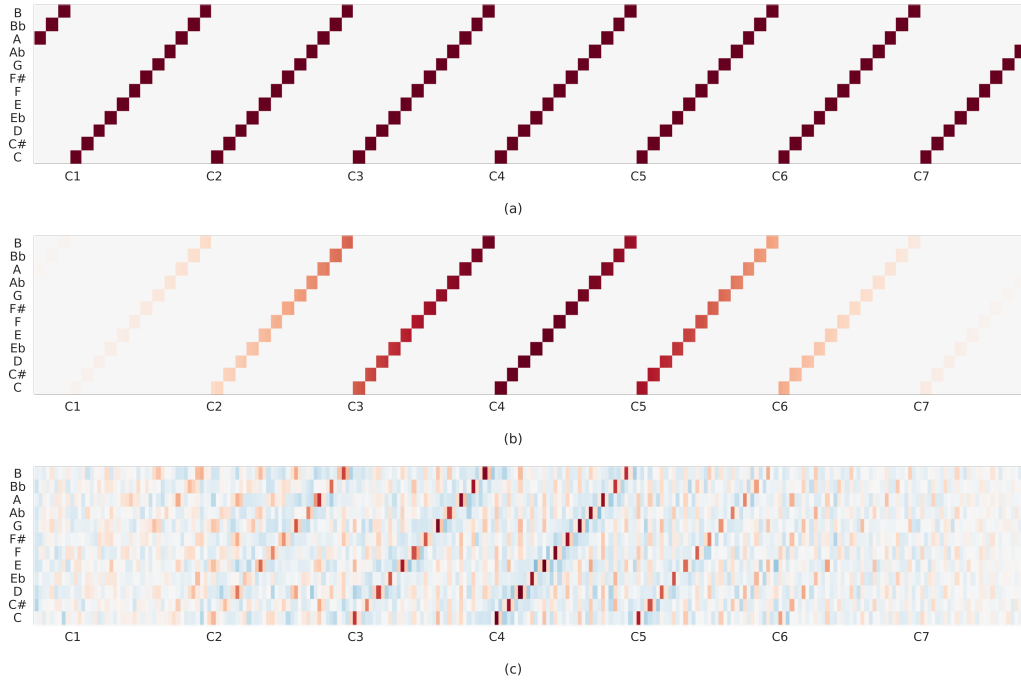


Figure 6: Various weight matrices for computing chroma features, corresponding to (a) uniform average, (b) Gaussian-weighted average, and (c) learned weights; red corresponds to positive values, blue to negative.

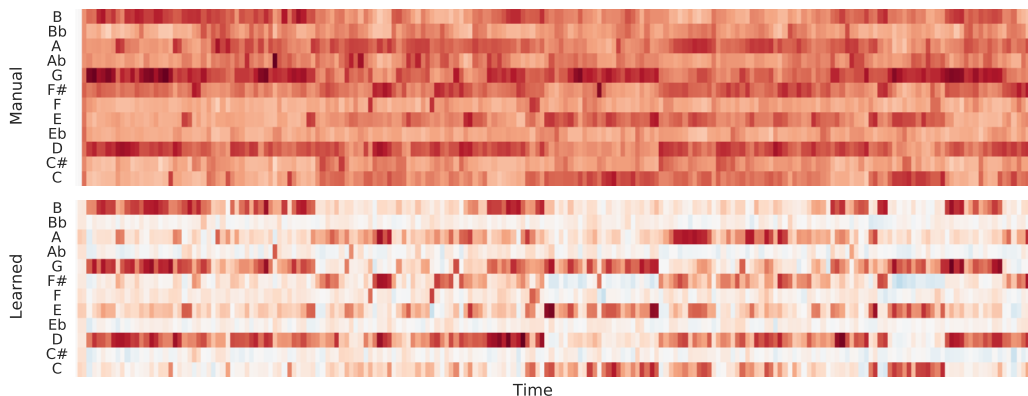


Figure 7: Comparison of manually designed (top) versus learned (bottom) chroma features.

Using the same general equation, a linear dot product between pitch spectra and a weight matrix, the mean-squared error is minimized between estimated chroma features and idealized “target” chroma features. Reference chord transcriptions are used as an information source for the target chroma, producing binary templates from the chord labels. The resulting weight matrix is illustrated in Figure 6-(c), and exhibits three significant behaviors. First, the positive contributions to each pitch class are clearly seen at the octaves, as to be expected. Second, the learned features corroborate the idea that the octave contributions should be weighted by a windowing function, and the one here looks vaguely Gaussian. Third, and most importantly, the learned weights exhibit a small amount of suppression around each octave, shown in blue. Similar to a Ricker wavelet (Vaidyanathan, 1993), negative sidebands serve to diminish wideband regions of energy, like those that found in percussion.

The chroma features obtained by these last two methods, (b) and (c), are shown in Figure 7. The noise floor on the learned chroma features is much higher than the hand-crafted ones, as a direct result of the negative suppression

in the learned weights. While the idea of adjacent pitch energy suppression is novel, it is important to recognize a few things about this example. Most importantly, it is curious to consider what other design aspects “learning” might help tease out from data. The function considered here, a linear dot product, is very constrained, and “better” features are likely possible through more complex models. Additionally, it is possible to directly inspect the learned weights because the system is straightforward; more complex models, however, will make this process far more difficult.

2.3 Previous Deep Learning Efforts in Music Informatics

While far from widespread, an increasing number of researchers have begun investigating deep learning to challenges in content-based music informatics. The most common form of deep learning explored for music applications focuses on such models to single frames of a music signal for genre recognition (Hamel, Wood, & Eck, 2009), mood estimation (Schmidt & Kim, 2011), note transcription (Nam, Ngiam, Lee, & Slaney, 2011), and artist recognition (Dieleman, Brakel, & Schrauwen, 2011). Meanwhile, the earliest instance of modern deep learning methods applied to music signals is found in the use of convolutional networks for the detection of onsets (Lacoste & Eck, 2007). More recently, convolutional networks have also been explored for genre recognition (Li, Chan, & Chun, 2010), instrument similarity (Humphrey, Glennon, & Bello, 2011), chord estimation (Humphrey, Cho, & Bello, 2012; Humphrey & Bello, 2012), onset detection (Schluter & Bock, 2014), and structural segmentation (Ullrich, Schlüter, & Grill, 2014). Recursive neural networks, a powerful, if troublesome, model for sequential data, have also found success in chord transcription (Boulanger-Lewandowski, Bengio, & Vincent, 2013) and

polyphonic pitch analysis (Sigtia et al., 2014). Additionally, predictive sparse decomposition (PSD) and other methods inspired by sparse coding have also seen a spike in interest (Henaff, Jarrett, Kavukcuoglu, & LeCun, 2011; Nam, Herrera, Slaney, & Smith, 2012), but while they make use of learning, neither of these systems are particularly deep. Regardless, it is worthwhile to note that many, if not all, of these works have attained state of the art performance on their respective tasks, often in the first application of the method to the area.

3 Discussion

Recognizing slowing progress across various application areas in content-based music informatics, this chapter has attempted to develop an understanding as to why this might be the case. Revisiting common approaches to the design of music signal processing systems revealed three possible shortcomings: manual feature design cannot scale to every problem the field will need to solve; many architectures are too shallow to adequately model the complexity of the data; and there currently are not many good answers for handling longer time-scales.

In looking to other related disciplines, it seems deep learning may help address some, if not all, of these challenges. Deep architectures are able to distribute complexity across multiple processing layers, thus being able to model more complex data. Feature learning, on the other hand, allows for the automatic optimization of known objective functions, making it easier to discover signal-level characteristics relevant to a given task faster. In fact, a handful of previous deep learning efforts within music informatics have already begun to demonstrate the promise of such methods.

Notably, these are crucial observations to make now for a variety of reasons. From a practical standpoint, many in the research community are investing considerable effort in the curation of datasets. While some, like (Bittner et al., 2014), have gone to great lengths to clear licenses for the source audio signals, most use commercial recordings and thus sharing the original content is problematic. As a result, it is becoming standard practice to apply a respected, but non-invertible, feature extraction algorithm over the original audio content and share the extracted statistics with the community, e.g. the Million Song Dataset (Bertin-Mahieux, Ellis, Whitman, & Lamere, 2011). While these efforts are commendable, such datasets are ultimately limited by the feature extraction algorithm employed.

CHAPTER III

DEEP LEARNING

Deep learning descends from a long and, at times, rocky history of artificial intelligence, information theory, and computer science. The goals of this chapter are two-fold: Section 2 first offers a concise summary of the history of deep learning in three parts, detailing the origins, critical advances, and current state of the art of neural networks. Afterwards, a formal treatment of deep learning is addressed in three parts: Section 2.1 introduces the architectural components of deep networks; Section 2.2 introduces the process of automatic learning, covering the design of loss functions and basic theory of gradient-based optimization; and Section 2.3 outlines various tricks of the trade and other practical considerations in the use of deep learning. Finally, the concepts introduced in this chapter are briefly summarized in Section 3.

1 A Brief History of Neural Networks

Despite the recent wave of interest and excitement surrounding it, the core principles of deep learning were originally devised halfway through the 20th century, grounded in mathematics established even earlier. As a direct descendant of neural networks —computational models with an ambitious moniker burdened by a tumultuous past— the very mention of deep learning often elicits several warranted, if suspicious, questions: What’s the difference? What’s changed? Why do they suddenly work *now*? Thus, before diving into a formal

review of the deep learning and its various components, it is worthwhile to contextualize the research trajectory that has led to today.

1.1 Origins (pre-1980)

For Western Europe and those in its sphere of influence, the Age of Enlightenment marked a golden era of human knowledge, consisting of great advances in many diverse fields, such as mathematics, philosophy, and the physical sciences. Long had humanity contemplated the notions of consciousness and reasoning, but here brilliant thinkers began to return to and explore these concepts with resolve. From the efforts of scholars like Gottfried Leibnitz, Thomas Hobbes, and George Boole, formal logic blossomed into its own mathematical discipline. In doing so, it became possible to symbolically express the act of reasoning, whereby rational thought could be described by a system of equations to be transformed or even solved.

It was this critical development —the idea of logical computation— that encouraged subsequent generations to speculate on the apparent feasibility of artificial intelligence. And, coinciding with the advent of electricity in the 20th century, mathematicians, philosophers, and scientists of the modern era sought to create machines that could *think*. While the space of relevant contributions is too great to enumerate here, there were a handful of breakthroughs that would prove integral to the field of computer science. In 1936, Alan Turing devised the concept of a “universal machine”, which would lead to the proof that a system of binary states, e.g. true and false, could be used to perform *any* mathematical operation (Turing, 1936). Only shortly thereafter, Claude Shannon demonstrated in his *master’s* thesis that Boolean logic could be implemented in electrical circuits via switches and relays, forming the basis of

the modern computer (Shannon, 1938). Shortly thereafter, in 1943, McCulloch and Pitts constructed the first artificial neuron, a simple computational model inspired by discoveries in neuroscience (McCulloch & Pitts, 1943). By coarse analogy to a biology, an artificial neuron “fires” when a weighted combination of its inputs eclipse a given threshold:

$$f(\mathbf{x} \mid \mathbf{w}) = h(\mathbf{w}^T \cdot \mathbf{x})$$

$$h(y) = \begin{cases} 1 : y \geq 0 \\ 0 : y < 0 \end{cases}$$

Importantly, as shown in Figure ??, it was demonstrated that such a model could be used to reproduce Boolean operations, such as AND or OR. Given the clear application to the field of computational logic, artificial neurons only further encouraged the pursuit of artificially “intelligent” machines.

On its own, an artificial neuron is only a general processing structure, and the parameters it takes will specify the precise behavior of the model. Arriving at these parameters, however, was nontrivial and required manual derivation. Thus, in 1958, Frank Rosenblatt’s invention of the *Perceptron* algorithm significantly altered how artificial neurons were conceived (Rosenblatt, 1958). Building upon the work of McCulloch and Pitts, the algorithm, given in 1, offered an automated method of “learning” the parameters necessary to achieve binary classification over a collection of data:

Algorithm 1 Find the optimal parameters for a Perceptron over a collection of data.

```

1: procedure FITPERCEPTRON( $\mathbf{x}, \mathbf{y}, \eta, n_{max}$ )
2:    $\mathbf{w} \leftarrow \mathbf{0}_{(D+1,2)}$ 
3:    $\mathbf{e} \leftarrow \mathbf{1}_N$ 
4:    $n \leftarrow 1$ 
5:   while  $|\mathbf{e}| > 0$  and  $n < n_{max}$  do
6:      $\mathbf{z} \leftarrow f(\mathbf{x}|\mathbf{w})$ 
7:      $\mathbf{e} \leftarrow \mathbf{z} - \mathbf{y}$ 
8:      $\mathbf{w} \leftarrow \mathbf{w} + \eta(\mathbf{e}^T \cdot \mathbf{x})^T$ 
9:      $n \leftarrow n + 1$ 
10:  end while
11:  Return  $\mathbf{w}$ 
12: end procedure

```

The perceptron algorithm requires four inputs: a matrix of observations, $\mathbf{x} \in \mathcal{R}^{N \times D}$, corresponding to N samples with D dimensions; a vector of binary class assignments, \mathbf{y} , in the set $\{0, 1\}$; an learning rate parameter, η ; and an iteration limit, n_{max} . Initializing the algorithm, the weights, \mathbf{w} , are set to a matrix of zeros, shaped $(D + 1, 2)$, an error, \mathbf{e} , is set to a vector of ones, and the iteration counter, n , starts from 1. Then, in an iterative manner, binary outputs, \mathbf{z} , are computed from the perceptron function, $f()$, given in Eq. ??, the error is computed as the difference between class predictions and targets, and the weights are updated with a scaled version of the incorrectly classified inputs. Note that the error vector, \mathbf{e} is only non-zero where the predicted values are wrong, and thus the algorithm naturally terminates when all datapoints are classified correctly. Alternatively, execution is halted once

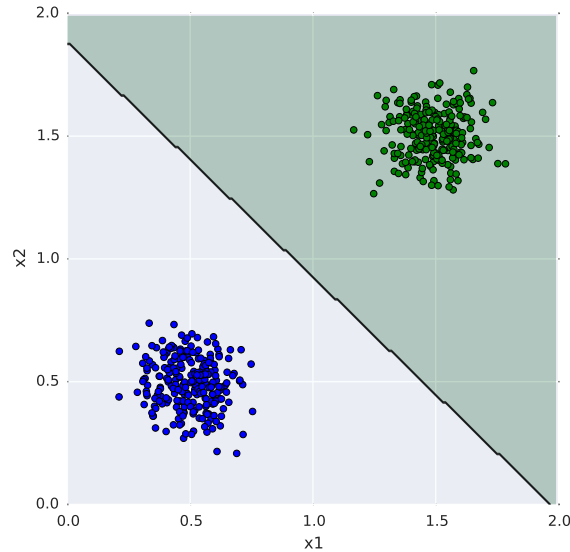


Figure 8: Linearly separable data classified by a trained perceptron.

a fixed number iterations is reached. An example of a perceptron’s stopping condition is given in Figure 8. Here, a perceptron has separated two classes of data, drawn from different Gaussian distributions. As the algorithm proceeds, the total error decreases until a decision boundary is found that correctly classifies the observed data.

Once implemented in hardware, using a slew of potentiometers, motors, and photocells, Rosenblatt’s “Mark I Perceptron” drew considerable attention from the press and the research community alike. The *New York Times* was quick to publish ambitious claims as to the promise this breakthrough held for artificial intelligence and the speed at which subsequent advances would be realized, much to the eventual chagrin of the AI community (Olazaran, 1996). However, the Perceptron was not without limitations nor critics. In their book, *Perceptrons*, published in 1969, Minsky and Papert demonstrated that the model is rather limited in the kinds of behavior it can actually achieve

(Minsky & Papert, 1969). For example, perceptrons are unable to reproduce the logic of an exclusive-or (XOR), and thus can only classify *linearly separable* data, the condition where a single straight line can be drawn between two classes.

This was a critical limitation for researchers in the field of neural computation; if a perceptron could not perform basic logic operations, how could it be expected to reason? The answer, as it would turn out, could be found by transforming how the XOR function is expressed symbolically. Rearranging terms, an equivalent function can be rewritten as the disjunction (OR) of two complementary conjunctions (AND):

$$p \oplus q = (p \wedge \neg q) \vee (\neg p \wedge q) \quad (1)$$

While it is true that a single Perceptron cannot achieve the XOR operation directly, a combination of *three* can: two are used to perform each AND operation and corresponding negation, while a third performs the OR operation. Considering the scenario in Figure 9, this condition can now be easily separated by a *multilayer* perceptron (MLP). Therefore, arbitrarily complex functions could be obtained by cascading simpler non-linear operations, leading to a class of functions that would come to be known as *neural networks*. These models are so versatile, in fact, it would later be shown by the *universal approximation theorem* that a neural network is actually able to model *any* continuous function, within some error tolerance (Cybenko, 1989; Hornik, 1991).

Despite this promising observation, neural network research languished through the closing decades of the 20th century, suffering a considerable drop in funding support and, as a result, interest. While the representational power

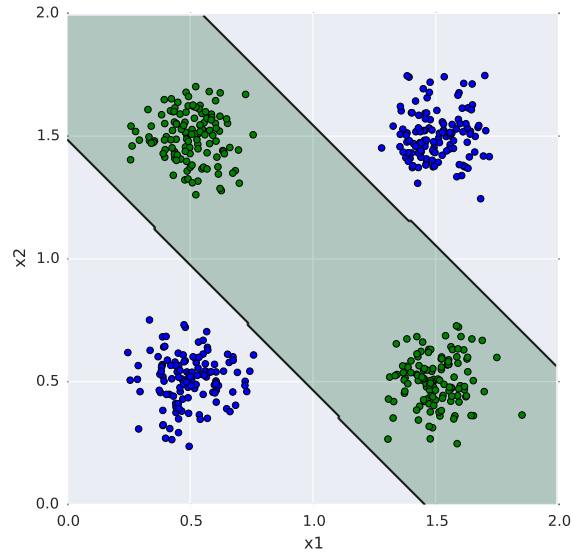


Figure 9: Demonstration of the decision boundaries for a *multi-layer* perceptron.

of multilayer perceptrons was recognized early on, it became popular opinion that these models could not be used to solve complex problems. In addition to theoretical skepticism, those who continued to pursue neural network research were also faced with an array of practical challenges. Neural networks were prone to over-fitting, extremely slow to train, and difficult to use with more than a few layers. Independently, these issues might merely have been viewed as common research obstacles. However, coupled with the failed promises of early progress, such difficulties would malign the pursuit of neural network research for some time.

1.2 Scientific Milestones (1980–2010)

In the face of widespread pessimism toward neural computing, some researchers would persevere through this “AI Winter”. Over the course of thirty some years, these diligent efforts would result in crucial breakthroughs that, in com-

bination with the steady churn of technological progress, would fundamentally change the landscape of neural network research. While no one facet can truly be credited with reviving the field, each would play an integral role in helping the research community again warm up to neural networks.

1.2.1 Efficient Error Backpropagation

Via the perceptron algorithm, a machine effectively “learns” to classify data by finding parameters that optimize a given objective function. This learning strategy must be modified in the multilayer case, as it is not possible to directly update the parameters through the heaviside, or unit step, function. Early researchers noted that if the discontinuous activation function is replaced with a logistic, or sigmoid, the composite multilayer network is *differentiable*. Thus, as will be discussed in more detail in Section 2.2.2, it is possible to use this gradient information to optimize the parameters of a network given an objective fitness measure, effectively *back-propagating* error through the network (G. E. Hinton, 1986).

This approach was not without its deficiencies, however. First and foremost, the error term was typically computed over the entire dataset, which often proved computationally expensive for sufficiently large collections. Additionally, early efforts found this particular approach often resulted in poor answers, and struggled to update parameters of networks with many layers. In time, a *stochastic* variant of this algorithm was shown to be far more efficient, and even circumvented some of these issues (Y. A. LeCun, Bottou, Orr, & Müller, 1998). Whereas conventional optimization occurs over an entire collection of datapoints, this efficient version randomly subsamples the training set and computes the error term over fewer datapoints. Though the error es-

timate is much noisier, the strategy is still effective, significantly faster, and may even be less susceptible to bad parameters as a result.

1.2.2 Convolutional Neural Networks

Though equal parts remarkable and encouraging, the universal approximation theorem has significant practical limitations. First, the theorem only holds for a sufficiently, and perhaps infinitely, wide network. Additionally, it says nothing about how one might parameterize such a function in order to achieve some desired behavior. Thus, despite such overwhelming promise, early neural networks often resulted in poor approximations of the functions being modeled, because the flexibility of the model vastly surpassed the amount of data available to train it. This flexibility was apparent in computer vision tasks, as multilayer perceptrons struggled to cope with simple homographic deformations, such as translation, scaling, or rotation.

Natural vision systems, on the other hand, are quite robust to these kinds of variation, and, as before, researchers looked to biology for inspiration in modeling these processes. One particularly insightful study was that of Hubel and Wiesel in 1959, who experimented on the neural behavior of the feline visual cortex (Hubel & Wiesel, 1959). By presenting visual stimuli while probing the brain with electrodes, they discovered two different kinds of neurons —simple cells and complex cells— organized hierarchically. Experimental results indicated that the former is responsible for local feature extraction, while the latter combines the inputs of simple cells to absorb small amounts of variation in the observed stimuli.

These ideas were first incorporated in the Neocognitron of (Fukushima, 1988), and later realized successfully as a convolutional neural network for

shape recognition and handwritten digit classification (LeCun, Bottou, Bengio, & Haffner, 1998), offering three key design features. *Local receptive fields*, where compute features from a small neighborhood of pixels, exploiting the observation that nearby pixel values tend to be highly correlated. Smaller receptive fields require fewer parameters, thus reducing the overall size of the model and the amount of data needed to train it. Applying local receptive fields as a convolution results in effectively sharing the weights over all positions in an image. Thus *weight sharing* reduces the number of parameters even further, while allowing the network to identify features regardless of position in an image. Finally, *max-pooling* reduces small neighborhoods to the largest value within it, introducing not only shift but a small amount of scale and rotation invariance. Taken together, these advantageous attributes directly resolved many of the issues plaguing multilayer perceptrons, and proved to be the most successful instance of neural computing for nearly two decades.

1.2.3 Proliferation of Digital Data

Given the ubiquity of digital information in the modern era, it is easy to forget that neural networks were first developed in a considerably different day and age. The very existence of digital audio and images, for example, did not become commonplace until the end of the 20th century. Even then, these signals were costly to create, cumbersome to work with, and generally difficult to share. Furthermore, once obtained, the process of annotating this information for use in supervised learning requires a great deal of time and effort. Thus, in the vein of speech recognition or computer vision for example, machine perception research was forced to work with small datasets as a result. Given the versatility of neural networks to model complex data, it was typically

trivial to over-fit these small training, while failing to generalize to new data. Neural networks developed the reputation of being “data hungry”, requiring a large amount of training data in order to do anything useful in practice.

While this was, and in some cases still is, a valid consideration for neural networks, the problem of data scarcity is far less of an issue now. For many well worn tasks, researchers have had ample time to curate massive labeled datasets. In the late 1980s, LeCun et al oversaw the development of a hand-written digit dataset, comprised of 60k 28x28 pixel images (Y. LeCun, Cortes, & Burges, 1998); for comparison, the ImageNet dataset consists of millions of tagged, high resolution images (Deng et al., 2009). Similar efforts have been undertaken in the fields of speech recognition (Fisher, Doddington, & Goudie-Marshall, 1986), face recognition (Huang, Ramesh, Berg, & Learned-Miller, 2007), or natural language processing (Lewis, Yang, Rose, & Li, 2004), to name only a few. Additionally, given the rise of the Internet, it became possible to leverage a variety of information sources to obtain labeled data, such as user-generated content, via Last.fm* or Twitter†, weak feedback signals, e.g. Pandora Radio‡, or even as a means of distributing the annotation effort (Von Ahn, Blum, Hopper, & Langford, 2003). Combined, the range of information available for large-scale supervised learning grew considerably, diminishing the issues posed by data-hungry algorithms.

As a by-product of the global transition to the digital realm though, an even larger portion of *unlabeled* data was at the disposal of machine learning researchers. Coupling the easy availability of digital information with the idea

*<http://www.last.fm/>

†<http://www.twitter.com/>

‡<http://www.pandora.com/>

that “the human brain isn’t *really* supervised”, much effort was invested in the space of *unsupervised* learning algorithms. Finally, in 2006, one such method, referred as *contrastive divergence*, was able to successfully exploit unlabeled data to improve the performance of a neural network (G. E. Hinton, Osindero, & Teh, 2006). Using a cascade of of undirected models, known as a Restricted Boltzmann machine (RBM), a neural network was trained in a greedy, layer-wise manner to reproduce realistic data. After this process of learning how real data behaves, the model could be “fine-tuned” with a smaller amount of labeled data to realize impressive, state of the art performance. Referred to as “pre-training”, learning on unsupervised data allowed the model to discover parameters closer to a good final solution. Though later discoveries would demonstrate pre-training to be unnecessary under certain conditions (Zeiler et al., 2013), this breakthrough was perhaps the first to forcefully recapture the attention of the machine learning community.

1.2.4 Advances in Computing Technology

While enough cannot be said about the scientific contributions of neural network researchers over the last 40 years, it is critical to appreciate how computing technology has evolved over that time span. In the days of Rosenblatt and Minsky, computers consisted of transistors visible to the naked eye, filled entire rooms, and cost a small fortune. More recently, personal computers of the 1980s had kilobytes of memory, central processing units (CPUs) operated in the range of tens of megahertz, and were still far too expensive for all but the elite or dedicated. Computation was still quite slow as a result, and thus the process of training neural networks took an impressive amount of time. To combat these difficulties, researchers often attempted to cut corners by using

smaller models, smaller datasets, and training with fewer iterations. Somewhat unsurprisingly, the common experience with neural networks was quite unfavorable.

Eventually, technology would catch up to the theory. Hardware became increasingly smaller, memory grew to the size of gigabytes, processors accelerated for a time seemingly without bound, and the cost of computers dropped to the point of near ubiquity. On the heels of these developments, other hardware and tools were adapted to facilitate neural network research, such as parallel computing with graphics processing units (GPUs) and accessible software tools, e.g. Theano (Bergstra et al., 2010) or Torch (Collobert, Kavukcuoglu, & Farabet, 2011). Combined, significantly better technology directly enabled research at unprecedented scale, accelerating progress and relaxing computational constraints imposed by technical limitations.

1.3 Modern Renaissance (post-2010)

Following the many key advances named above, deep learning quickly accelerated into the academic, industrial, and public lexicon. It did not take long to convince some research communities of the newfound promise of neural networks. In hardly any time at all, deep neural networks surpassed the state of the art in automatic speech recognition, systems tuned over the course of several decades (G. Hinton et al., 2012). Similarly compelling results were obtained in computer vision (Krizhevsky et al., 2012) and natural language processing (Sutskever, Martens, & Hinton, 2011). Acknowledging the usefulness of such high-performing systems, companies like Google, Facebook, Microsoft, IBM, and Baidu began investing in industrial strength deep learning teams and infrastructure (Dean et al., 2012).

Complementing the migration of ideas and individuals from academia to industry, the idea of “deep learning” and thinking machines has again struck a chord with both the press and general public. Google’s research efforts in 2012, for example, yielded a deep network that automatically learned the concepts of “cats” and “human faces”, trained on millions of still video frames from YouTube (Q. Le et al., 2012). Understandably, the story was picked up by *The New York Times* and *WIRED*, among countless others, drawing widespread attention. Coupled with charismatic interviews from many of the field’s preeminent leaders, neural networks have been thrust back into the limelight.

2 Core Concepts

Reflecting on the historical lineage of neural networks, it becomes apparent that deep learning is not one single thing, but rather a collection of ideas and approaches toward neural computing. Thus, to help define the scope and limits of this work, this section introduces the fundamental components that contribute to a modern definition of the topic. Here, “deep learning” is defined as an approach to designing complex information processing systems, exhibiting two key traits; one, discussed in Subsection 2.1, the system architecture can be expressed as a composite nonlinear function, composed of many simpler operations; and two, presented in Subsection 2.2, the parameters of this function can be “learned” by combining an objective function with numerical optimization methods. Finally, modern neural network research has produced a useful array of “tricks of the trade”, detailed in Subection 2.3, which often help squeeze every last bit of performance from a model.

2.1 Modular Architectures

It is a hallmark of deep neural networks that system architectures are constructed from only a handful of unique processing units, repeated and arranged to form complex structures. This modular approach to system design allows the researcher to focus on each operation at different levels of abstraction; from a few basic building blocks, it is straightforward to create elaborate architectures (Szegedy et al., 2014).

In the broadest of terms, a neural network transforms an input X into an output Z via a composite nonlinear function $\mathcal{F}(\cdot|\Theta)$, given a parameter set Θ . This is traditionally, but not necessarily, achieved as a sequential cascade of L self-contained operations, $f_l(\cdot|\theta_l)$, referred to as *layers*, *nodes*, or *sub-functions*, the order of which is given by l :

$$Z = \mathcal{F}(X|\Theta) = f_L(\dots f_2(f_1(X|\theta_1)|\theta_2))\dots|\theta_L) \quad (2)$$

In this formulation, $\mathcal{F} = \{f_1, f_2, \dots f_L\}$ is the set of layers, $\Theta = \{\theta_1, \theta_2, \dots \theta_L\}$ is the corresponding set of parameters, the output of one layer is passed as the input to the next, as $X_{l+1} = Z_l$. Generally speaking, the overall *depth* of a network is denoted by L ; more accurately, however, processing depth as a means to representational power is determined by the number of nonlinear operations in a network. The *width* of a network, on the other hand, is determined by the dimensionality of the intermediary representations between sub-functions, and often varies within the network.

It is worth noting that this sequential cascade of operations is only common practice and not a hard and fast rule. Some networks make use of interesting types of connectivity between layers, referred to as “skip-layer” connections;

others adopt an explicit graphical interpretation, encouraging the description of the network in terms of nodes and edges. Therefore the design of a network architecture must address several interrelated questions: what is the function being modeled? what domain knowledge can be leveraged to inform this design? and how do these relate to the data being processed, or the problem being addressed? The considerable flexibility afforded by this modular design approach is both one of the greatest advantages, and criticisms, of deep neural networks.

While many mathematical operations can be incorporated into a neural network, provided they are approximately differentiable, there are a handful of common operations and conventions worth discussing.

2.1.1 Affine Transform

The fundamental building block of the classic neural network is the *affine transform*:

$$Z_l = f_l(X_l|\theta_l) = h(W X_l + b), \theta_l = [W, b] \quad (3)$$

Here, the input to the l^{th} layer, X_l , is flattened to a column vector of length N , projected against a weight matrix, W , of shape (M, N) , added to a vector bias term, b , of length M , and passed through a transfer function, $h(\cdot)$. As addressed shortly, the transfer function is generally some pointwise nonlinearity; the linear matrix or dot product is then a special case of Eq. 4. Alternatively, this operation is also known as a, *dense*, *fully-connected*, or *multiperceptron* layer, to distinguish its topology from other kinds of connectivity. This common sub-function descends directly from the perceptron, and is the core op-

eration of the multi-layer perceptron. Additionally, the affine transformation is a crucial processing block, because, as it will soon be discussed, many other sub-functions can be interpreted as a special instance of a matrix-product.

2.1.2 Local Affine Transformation

Though a recent development, the *local* affine transformation is a cousin of the general affine transformation, operating only on sparsely connected neighborhoods of an input. Based on observations of visual systems, local receptive fields exploit correlations found in spatial pixel neighborhoods to reduce the complexity of features to be learned (Q. V. Le et al., 2010). Notably, local affine transformations can be realized as a full matrix product where most weights are zero, expressed as the following:

$$Z_l = f_l(X_l|\theta_l) = h\left(\sum_{m=0}^M X_l[m+n]W_k[m] + b_k\right), n < N - M + 1, k < K, \theta_l = [W, b] \quad (4)$$

Here, M adjacent features are projected against K different combinations of weights per shift, n , and this locally dense connectivity is translated across all $N - M + 1$, possible locations of a given input; the dimensionality of the output, Z_l , is $K * (N - M + 1)$. This formulation is particularly interesting, as it illustrates the inherent difficulty posed by the universal approximation theorem. Technically, a local receptive field can be implemented as a significantly larger fully-connected matrix product; learning this behavior directly, however, can be quite difficult. First off, this particular sparse connectivity would need to be learned redundantly in all locations of an input. Additionally, such true sparsity is challenging to learn, and it is unlikely such behavior would

be so well localized. Finally, the prospect of learning this particular topology from a fully-connected matrix demands a prohibitive number of parameters, throttling both computation and the learning process.

Furthermore, this spatially correlated topology is unique to visual signals, and such connectivity assumptions may not map to other domains. Natural sound, for example, is harmonic in nature, and it is reasonable to assume that octave relationships may have a stronger connection than neighboring frequencies. Some research has argued that different types of connectivity could be determined from data (Coates & Ng, 2012), but this has yet to be extensively explored in the deep learning literature.

2.1.3 Convolutions

Extending the topology of local receptive fields, convolutions further simplify the learning process by *sharing* parameters across all locations in an input representation. Also known as weight tying, a convolution is generically expressed by the following:

$$f_l(X_l|\theta_l) = h(X_l \circledast W + b), \quad \theta_l = [W, b] \quad (5)$$

In deep learning literature, there are typically two kinds of convolution operations indicated by the \circledast . The first, and perhaps original, interpretation is referred to as a “2D” convolution:

$$\hat{Z}[m, a, b]_l = \sum_{i=0}^N \sum_{j=-\infty}^{\infty} \sum_{k=-\infty}^{\infty} X_l[i, x, y] W[m, a - j, b - k] \quad (6)$$

Here the valid convolution is computed by convolving a 3D input tensor, X_l , consisting of N *feature maps* or channels, with a collection of M 2D weight

kernels, W , followed by the addition of a vector bias term b . In this formulation, X_l has shape (N, d_0, d_1) where (d_0, d_1) is the shape of each map, W has shape (M, m_0, m_1) , where (m_0, m_1) define the size of the kernel, and the output, Z_l , has shape $(M, d_0 - m_0 + 1, d_1 - m_1 + 1)$. Note that in this 2D formulation, the activation of a kernel is summed equally across the separate feature maps, and thus a single kernel is shared across channels and spatial translations.

Alternatively, the other common variant of the \otimes operator is the 3D convolution, written as follows:

$$\hat{Z}_l[m, a, b] = \sum_{i=0}^N \sum_{j=-\infty}^{\infty} \sum_{k=-\infty}^{\infty} X[i, x, y] W[m, i, a - j, b - k] \quad (7)$$

The input, X , is again a 3D tensor of feature maps, but now the weight tensor used in the convolution, W , is 4D, with shape (M, N, m_0, m_1) . Whereas before, in the 2D case, each kernel translated across feature maps, the 3D case aligns a kernel with the number of feature maps, N . Thus, in the instance that $N = 1$, both operations are equivalent. Due to the local connectivity across feature maps, 3D convolutions require N times more parameters than their 2D cousin, but far fewer than an equivalent full matrix product. Using kernels that are sensitive to characteristics across different simultaneous feature maps allows for correlations to be discovered across the same representation.

2.1.4 Nonlinearities

A seemingly simple piece of the deep learning toolkit, pointwise nonlinearities are crucial to the construction of complex networks. In the absence of nonlinear behavior, a cascade of linear systems is itself another linear system.

Nonlinear operations, however, change the overall behavior of the composite function, and are the source of representational power in deep networks. Conventionally referred to as *transfer* or *activation* functions, these operations are typically applied to the output of other linear functions, e.g. after an affine transformation.

The two basic nonlinear functions in deep networks are the logistic, or sigmoid, and the related hyperbolic tangent:

$$\begin{aligned} \text{logistic}(x) &= \frac{1}{1 + e^{-x}} \\ \text{tanh}(x) &= \frac{1 - e^{-2x}}{1 + e^{-2x}} \end{aligned}$$

Shown in Figure ??, both functions have inflection points at zero, saturate toward infinity in both directions, and are everywhere differentiable. While the logistic initially came into practice as a smooth approximation to the unit step function, the hyperbolic tangent has gained favor for being centered about the origin. The saturating behavior of these functions is significant, because training a poorly initialized network may be painfully slow, if even possible, due to small gradients at the limits.

More recently, the *Rectified Linear Unit* (ReLU), or halfwave rectification, has seen a considerable up-tick in usage in deep networks:

$$\text{relu}(x) = \max(x, 0) = x * (x > 0)$$

Notably, it was demonstrated in (Zeiler et al., 2013) that, with sufficient data, rectified linear units could achieve state of the art performance without unsupervised pretraining. Though the theory is still catching up to the practice, there are two widely held rationalizations of this behavior. Most importantly, the function does not saturate in the positive region, and thus gradients propagate freely through an arbitrarily deep network. Additionally, the hard threshold results in naturally sparse representations, an advantageous property for classification systems (Bengio & LeCun, 2007). Noting that no information flows in the negative mode, others have extended this idea into the “leaky ReLU”, but these have seen far less use in practice as they undermine the benefits of sparsity (Maas, Hannun, & Ng, 2013).

2.1.5 Pooling

In a broad sense, *pooling* operations compute local statistics over representations. This can be understood as a form of summarization, which trades accuracy of locality for various forms of invariance. Original drawing inspiration from the Hubel and Weisel experiments, the most common pooling operation is *max-pooling*, which mimics the behavior of complex cells in the visual cortex by passing only the largest value within a local neighborhood. This offers, depending on the size of the neighborhood, invariance to scale, translation, and rotation in visual tasks, and contributes significantly to the

success of convolutional networks. the parameter p_0, p_1 is a two-element tuple that defines the neighborhood of the max operator along these dimensions:

$$\hat{Z}[x, y] = \max(\mathbf{1}_{p=0}^P \mathbf{1}_{q=0}^Q X[x + p, y + q]) \quad (8)$$

Other forms of pooling exist, though less common in practice. Similar in nature, *L2-pooling* computes the magnitude vector of a neighborhood in Euclidean space (Sermanet, Chintala, & LeCun, 2012). This can be interpreted as a softer form of max-pooling, where all inputs contribute to the output, albeit dominated by the maxima. Finally, other normalized statistics like the *mean*, *min*, or *standard deviation* have found use in temporal pooling (Hamel, Lemieux, Bengio, & Eck, 2011). It is a particular advantage of these pooling functions that they can be applied globally to variable length inputs as a means of producing equivalently-shaped summary statistics, a common challenge faced in processing music signals, e.g. songs, of different durations.

2.1.6 Classification Functions

While the system components named previously are sufficient for modeling continuous variables, classification systems require a decision function to select a discrete class as the “winner”. Perhaps the simplest decision rule identifies the position of a global optima in a representation, i.e. the *argmax* or *argmin*, which return the index of the largest or smallest value, respectively. These two functions are often used in two semantically opposite conditions: the former is used in systems that estimate similarity or likelihood, where larger values are better; and inversely, the latter is used in systems that estimate dissimilarity or distance, where smaller values are better.

One common approach to obtaining a likelihood estimate from a neural network is to flatten the output of the final, often linear, layer and pass it through the *softmax* function, defined as follows:

$$\sigma(Z|\beta) = \frac{\exp(\beta Z)}{\sum_{k=1}^K \exp(\beta Z_k)} \quad (9)$$

Here, Z is the output of the final layer, f_L , K is the dimensionality of the classifier, equal to the number of classes in the problem, and β is a hyperparameter that controls the dynamic range sensitivity of the function. For $\beta \rightarrow 0_+$, all classes trend to a uniform probability, while the inverse is true for $\beta \rightarrow \text{inf.}$ Geometrically, the softmax operation can be understood as continuous mapping of coordinate points to the surface of a K -dimensional simplex, and thus its outputs have unit magnitude in L_1 space.

The other common approach to classification in neural networks is to identify the nearest target, template, or centroid to the output, given some notion of distance. In this case, not only is the network itself a variable, but also the representations used for computing distance. Thus, it may be advantageous to incorporate these target representations into the network itself and take the winning class to be the one with the lowest energy, via the *radial basis* function:

$$f(x|\theta) = \sum (x - \mathbf{w})^2, \theta = [w] \quad (10)$$

Functionally equivalent to template matching or nearest centroid classification in Euclidean space, the advantage of incorporating the radial basis function *in* a network is two-fold: first, parameters can be optimized to a set of chosen targets, which may have semantic importance, as in (LeCun et al., 1998);

and second, these class targets can be learned from the data. Care must be exercised when attempting to learn these bases though, as a trivial solution exists at the point where all distances are zero. Without additional constraints on the learning problem, a network may learn to drive the input to the radial basis function to zero, thus making all distances arbitrarily small.

2.2 Automatic Learning

Recall that the overarching goal here is to model the relationship between a collection of input data and desired outputs. The building blocks described previously are used to construct powerful, complex functions, but this is only half the design problem; it is then necessary to find the parameters that enable this general function to perform some specific behavior. Analogously to objected oriented programming, a network's equation family defines its *class*, whereas the parameters define its *instance*.

Importantly, assuming the desired outputs are known, it is possible to measure how well a given function approximates this relationship. Now consider that there exists a hypothetical space of all possible solutions and, in it, at least *one* function that optimally satisfies this measure. In this interpretation, the process of finding such a function can be thought of as a *search*, and thus the ideas of “learning” and “searching” can be used interchangeably. Unfortunately, the space of all possible solutions is effectively infinite and cannot be explored exhaustively, automatically or otherwise. However, conceptualizing functions in terms of classes and instances provides an elegant way of making this search significantly more manageable: a function's class, which can be purposefully designed, greatly reduces the space of all possible instances to a much smaller subspace that *can* be explored.

With this in mind, the distinction between function classes and instances is a practical one to make. By heavily restricting the space of possible solutions, automatically searching this smaller subset becomes feasible. Furthermore, differentiable* fitness measures allow for the application of gradient-based optimization methods to find good parameters, freeing the researcher to focus less on the specific parameters of a system and more on the abstract design of the underlying model. Learning is advantageous because it simplifies the search for good parameters, accelerates research, yields flexible systems that can adapt to new data, and facilitates the discovery of new solutions not previously considered. While it might seem like this violates the “No Free Lunch” theorem, training a network shifts the focus on system design from a micro to macro level, comprised of two components: defining a suitable objective function, and numerically optimizing the parameters over a collection of training data.

2.2.1 Loss Functions

It is worth noting that there are various, roughly equivalent interpretations of neural network training and optimization. While largely inspired by (Y. LeCun, Chopra, Hadsell, Ranzato, & Huang, 2006), the perspective presented here takes a far more constrained approach to *inference*, whereby prediction is achieved by the forward propagation of an input through a model. All additional operations necessary to measure the quality of a model can be considered supplementary *meta-function* that sits on top of the desired model during training only.

*Technically speaking, some functions, like rectified linear units, are not everywhere-differentiable, but will work in practice.

In the most general sense, this evaluative component is known as a *loss* function, L , quantifying the degree of fitness between a model, \mathcal{F} , its parameters, Θ , and a collection of N labeled data, $\mathcal{D} = \{(X_i, Y_i)\}$, where X_i is the i^{th} observation and Y_i its corresponding desired output:

$$\mathcal{L}(D|\Theta) = \frac{1}{M} \sum_{m=0}^{1 \leq M \leq N} L(Y_m|X_m, \{\cdot, \Theta\}) \quad (11)$$

Here, the scalar loss, \mathcal{L} , is computed by averaging the per-sample loss over some number of observations, M , referred to as the batch size. Thus, having designed or constrained the network, $\{\cdot\}$, the optimal parameters, $\hat{\Theta}$, are those that minimize the loss over the data:

$$\hat{\Theta} = \min_{\Theta \in} \mathcal{L}(D|\Theta) \quad (12)$$

Given this common formulation, the choice of loss function for a particular problem becomes another modular design decision. While there is a good deal of flexibility in the composition of a loss function, there are a few standard measures of note. Perhaps the most common loss function is the *mean squared error*, which measures the squared Euclidean distance between the desired output and the estimated output produced by the network:

$$L_{mse}(Y|X, \{\cdot, \Theta\}) = \|Y - \{(X|\Theta)\}\|_2^2 \quad (13)$$

A versatile function, the squared error loss is a common choice for regression problems where domain knowledge does not immediately offer a better intuition for computing distance between representations. Generally, this loss function is only suitable if the error can be reasonably modeled as Gaussian.

Additionally, the squared error loss is functionally equivalent to minimizing the correct output of a radial basis function when a finite number of target values are used as a fixed basis. The choice to include this operation internal to a model, rather than incorporating it into the loss function, is determined by the intended mode of operation of the model, i.e. for regression or classification.

The *margin* is another common component in loss function design, finding an intersection with support vector machines (Bishop, 2006). Mathematically similar to the rectified linear unit described previously, the margin's role in loss function design is to incur no penalty when its input drops below a given value m :

$$L_{margin}(x|m) = \max(0, x - m) \quad (14)$$

The margin is often used in conjunction with operations that measure relative quality, i.e. distance. For example, a margin could be used to penalize a model when the distance between inputs of the same class is outside a given radius; conversely, the opposite behavior is achieved by negating the argument.

The remaining loss function of interest here is the *negative log-likelihood* loss, a common criterion for classification problems. For networks that produce probability mass functions over a set of K classes, e.g. via the softmax operation, the parameters can be optimized by minimizing the correct class' negative log-likelihood:

$$\mathcal{L}_{nll} = -\log(P(\mathcal{F}(X^k|\Theta) = Y^k)) \quad (15)$$

Expressed in this manner, X^k and Y^k are the input data and predicted output corresponding to a class label, k . Optimizing this measure maximizes the class

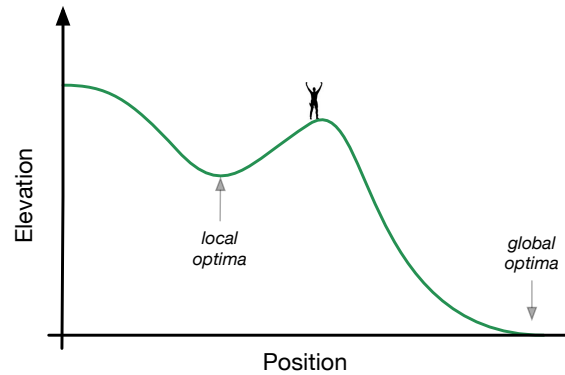


Figure 10: Hill-climbing analogy of gradient descent.

conditional probabilities over a collection of data. Importantly, due to the unit magnitude constraint offered by the softmax operation, pushing down on the correct answer has the effect of pulling *up* on the values of all other classes.

Finally, it is worth noting that it is often the case that a loss function is used as a proxy for some other measure of interest, e.g. accuracy, which may be difficult to optimize directly. Some work has explored the use of other objective functions that may be more appropriately aligned with the given task, such as minimum classification error (MCE) (Juang, Hou, & Lee, 1997), but alternative functions often entail their own unique challenges.

2.2.2 Numerical Optimization

Once the quality of a model’s configuration can be expressed as a single scalar quantity, it becomes immediately possible to automate the search of this space. Given infinite time and computational power, this could at least theoretically be achieved by the application of an exhaustive, brute force search. Practically speaking, however, such naive strategies are unlikely to uncover any solutions of value due to the immensity of the search space.

Gradient methods, on the other hand, exploit “local” improvements as a means of proceeding toward better answers. A simple analogy helps illustrate both the intuition, and some deficiencies of the approach. Consider the task of climbing down a hill blindfolded, illustrated in Figure 10. At any given time, one might poke about in every direction in an effort to find the steepest decline, and then, take a single step in the most immediately promising direction. Eventually, save for some technical exceptions, this strategy will lead to a condition in which a locally optimal condition has been reached, i.e. any step increases elevation. But is this the bottom of the mountain, or has our climber gotten stuck in a valley? Thus it is an inherent difficulty of such an algorithm that it will lead to *an* answer, but not necessarily the globally optimal one. Additionally, given its greedy nature, the best step *now* might not be the best step overall; this is apparent in the diagram, where the best initial step is in the opposite direction, to the left, of the global optima, to the right. This further illustrates the importance of initialization, and the effect that an unfortunate starting point can have on the optimization process.

Expressed more formally, the gradient descent update rule over a set of parameters, Θ , is defined as its difference with the gradient of the scalar loss \mathcal{L} over a set of datapoints, $\hat{D} \in D$, with respect to these parameters, weighted by a learning rate η :

$$\Theta_{n+1} \leftarrow \Theta_n - \eta * \frac{\nabla \mathcal{L}(\hat{D} \mid \Theta_n)}{\nabla \Theta_n} \quad (16)$$

It is apparent from this formulation that a model’s ability to learn is a direct result of the estimate of the scalar loss, which depends not only on the model, its parameters, and the choice of loss function, but *also* the data over which it

is measured. While early applications of gradient descent for training neural networks would compute parameter updates using the entire collection of datapoints available for training, deep learning typically leverages batch methods to approximate the overall loss.

Though the work presented here will focus exclusively on stochastic gradient descent as the numerical optimization method of choice, it is worth noting that there are alternative methods and extensions one could employ. In addition to known issues regarding poor local minima, there are also degenerate cases in which a first-order method like this may be very slow to converge. In practice however, gradient descent is often sufficient with some slight modifications, such as the use of slight Nesterov momentum (Sutskever, Martens, Dahl, & Hinton, 2013). Alternatively, *Quasi-Newton* methods compute approximations to higher order partial derivatives, which may help the optimization process recover from problematic regions in the loss surface and aid in convergence (Liu & Nocedal, 1989). These higher order methods often entail a higher computational cost, however, and it is not uncommon to employ them, or their approximations, in tandem with stochastic gradient descent (Kavukcuoglu et al., 2010).

2.3 Tricks of the Trade

Armed with versatile complex functions and a means to automatically learn parameters, deep learning seems perfectly posed to tackle a vast array of information processing problems. For some time however, however, only a handful of researchers were able to get positive results, thus earning it its reputation as a “dark art.” As with many things, the devil is truly in the details, and over

time the various tricks and less principled practices necessary for success have been consolidated into a set of common strategies (Bengio, 2012).

2.3.1 Penalizing Bad Behavior

The goal of architecting a deep network is adjust the representational power of the model to a given problem, incorporating known constraints where possible. More often than not, however, a the inherent complexity of a particular model vastly exceeds the latent complexity of the data at hand. As a result, the learning process will result in parameter configurations that perform well on the training data, but fail to generalize to unseen data. Conceptually, this can be understood as the model having too many options when achieving its goal state, and happening to pick one that is bad for reasons that do not factor into its objective function. It can be said that the learning problem is under-determined, and thus additional penalties can be added to the overall loss. Interestingly enough, this penalty-based approach is common practice in other greedy optimization systems, like capitalistic markets, where taxes and fines serve to discourage certain outcomes, e.g. dumping industrial by-products in a river.

While it is conceivable that any number of undesirable system behaviors could be quantified by some scalar measure, there are two general approaches common across deep learning. The first, referred to as weight decay, ridge regression, or L_2 -regularization, is classic across machine learning, and finds use in a variety of learning algorithms:

$$\mathcal{L}_{decay} = \sum_i \lambda ||\Theta_i||_2^2 \tag{17}$$

Here, the vector magnitude of the i^{th} parameter, Θ_i , is weighted by a hyperparameter, λ_i , and summed together as a scalar penalty. Conventionally, the same weight is applied to all parameters, although differently shaped parameters will bias the cumulative term. During optimization, this has the effect of driving parameters toward, but not exactly, zero, providing an intuitive interpretation consistent with Occam’s razor, i.e. prefer simpler solutions. This often prevents the network from overfitting the training data exactly, and resulting in solutions that generalize better.

Whereas weight decay is applied to the parameters of network, *sparsity* penalties are applied over intermediary representations in a network, taking the form of a cumulative weighted vector magnitude in L_1 space:

$$\mathcal{L}_{sparsity} = \sum_i \lambda ||Z_i||_1 \quad (18)$$

Ideally, it would be more preferable to optimize the L_0 loss. This this problem is generally *np*-hard, however, and the L_1 loss, its convex envelope, serves as a differentiable approximation. While it is not guaranteed that the solutions of each may coincide, optimizing the L_1 generally yields good results in practice. As a result, this penalty will prefer sparse representations, which help *disentangle* factors of variation in the data (Bengio, 2009).

2.3.2 Parameter Initialization

As shown previously in the discussion of gradient descent, poor initial conditions can delay, or in some cases even prevent, an optimization algorithm from finding good solutions. Attempts to train deep networks with error back-propagation can compound this issue, as an error signal can become

insignificantly small after several layers under certain conditions. While this is somewhat sensitive to the architectural decisions made, such as the choice of non-linearities used, all networks are sensitive to parameter initialization.

In general, sampling coefficients from appropriately tuned random distributions leads to reasonably good results. There is little consensus on the advantages of a uniform versus normal distribution for initialization, but the primary factor to control in either case is the dynamic range or scale. As a rule of thumb, this can be automated somewhat by using the *fan-in*, or dimensionality of the input, to keep the activations of within the operating region of saturating transfer functions, i.e. sigmoid or hyperbolic tangent. Rectified linear units, on the other hand, are far less sensitive to the choice of initialization, as half of its operating region is non-saturating, contributing to their rise in popularity. As a result, sufficiently small, centered normal distributions, $\mathcal{N}(\mu = 0, \sigma \approx 0.01)$, work well in practice.

Alternatively, unsupervised *pre-training* methods, credited with effectively jump-starting the renaissance of deep learning, can be used to initialize networks in a data-driven manner. Though there are different perspectives on how this can be achieved, the core concept is the same: using a large amount of unlabeled data, train a deep network to reconstruct realistic observations in a greedy, layer-wise manner. Then, once all layers have been initialized, conventional supervised training can be applied. In theory, this data-driven process works by getting the parameters of the network closer to a good solution. As mentioned previously, the first successful realization of this idea used restricted Boltzmann machines (RBMs) (G. E. Hinton et al., 2006), leveraging Gibbs sampling and contrastive divergence to produce realistic samples from the model. The deterministic variant is the autoencoder, which use data aug-

mentation, sparsity, or weight tying to train a pair of deep, invertible functions (Bengio, Courville, & Vincent, 2013). Most recently, unsupervised pre-training has fallen out of favor somewhat, as comparable results can be obtained with rectified linear units and randomized weights, provided there is sufficient labeled data. That said, this strategy may still prove useful for problems in which it is difficult to obtain a large amount of labeled information, such as personalized systems.

2.3.3 Dropout

Another recent addition to canon of deep learning practicum is the training strategy known as *dropout* (G. E. Hinton, Srivastava, Krizhevsky, Sutskever, & Salakhutdinov, 2012). As the name suggests, some percentage of parameters in a network are randomly “dropped” during training in estimating the loss and computing an update step. The details of how exactly this is done vary slightly from sub-function to sub-function, but a description in terms of an affine transformation is sufficiently illustrative:

$$Z = f(Z|p) = \frac{1}{(1-p)} Z \mathcal{B}(1, p)^M$$

Here, the activations of an affine transformation, Z , a column vector of length M , are masked by a binomial distribution, \mathcal{B} , with probability p . To offset the effect of smaller magnitudes resulting from fewer units being active during training, these outputs are scaled by one minus the probability parameter, such that, in expectation over \mathcal{B} , the use of dropout matches the original function. Applied in this manner, this prevents rows, or bases, in the matrix

projection from contributing to the final output, inhibiting the co-adaptation of parameters. Thus dropout is effective because the parameters are unable to depend on one another for effectiveness, and tend to learn independently good features.

Additionally, dropout has an interesting relationship with model averaging, or *bagging*. Masking parameters has the effect of selecting a subset of parameters from the given model, and therefore one of many complementary models is updated at each training iteration. This reduces the theoretical bound on generalization error, and has been proven to greatly improve results for models prone to overfitting.

2.3.4 Data Augmentation

A common strategy among deep learning practitioners is to leverage domain knowledge wherever possible. While this has an obvious connection to the architectural design and choice of loss function, another oft exploited, though less documented, opportunity is through the application of data augmentation. The main idea here is that a collection of labeled data can be manipulated in varying degrees of realism. This exploits the common scenario that modeling the synthesis process is typically easier than the analysis process. While the most effective distortions are rather domain specific, common deformations include operations such as translation, scaling, additive noise, nonlinear distortion. In this space of music, this could range from signal-level attributes, like perceptual codecs or production effects, to musically inspired augmentations, such as pitch shifting or time-stretching.

2.3.5 Normalization

Finally, enough cannot be said about the importance of proper data normalization. The simplest form of data normalization is maximum scaling, such that all datapoints are bounded on the same input region. Another form of dynamic range control is achieved by normalizing inputs to have unit magnitude in some L_p -space, e.g. Euclidean. Alternatively, coefficient-wise *standardization* is strongly advocated, e.g. subtract the mean and divide by standard deviation. This can be extended via principal components analysis (PCA), which offers the added benefit of dimensionality reduction, or PCA-whitening (alternatively, ZCA), which “whitens” the data by scaling coefficients by their eigenvalues. Lastly, *local contrast normalization* (LCN) has proven to be a particularly useful approach to data normalization (Sermanet, Kavukcuoglu, Chintala, & LeCun, 2013). Finding inspiration from biological processes, LCN performs a form of automatic gain control over local neighborhoods, and can lead to surprisingly discriminative features even in the absence of training (Kavukcuoglu et al., 2010).

In addition to constraining the input space, it is also common to constrain the *parameters* within a network, often taking two forms. In some cases, like (G. E. Hinton et al., 2012), parameters are simply constrained inside the volume of the given hypersphere, such that any time parameters are updated to values that result in a magnitude larger than 1, they are rescaled. This constraint gives parameters freedom to adapt to the nuances of the data without growing arbitrarily large to offset the contributions of small weights. Bounding parameters shares a loose connection to weight decay, in parameters are prevented from growing arbitrarily large, without the need for additional

penalties or hyperparameters. Alternatively, there are also successful instances of weights being constrained to the surface of the L_2 hypersphere, a common approach in various forms of sparse coding.

3 Summary

Building upon formal logic and biological analogy, neural networks were devised in the 1960s as an approach to information processing. However, after initial promise, they were largely met with skepticism, indifference, or worse through the remainder of the century. The few who persevered made various discoveries that, coupled with steady advances in computing technology, would eventually return neural computation to the fore: mini-batch stochastic gradient descent made optimization computationally efficient and less susceptible to poor local minima, and encouraged further exploration of numerical optimization methods; convolutional networks reduced model complexity and over-fitting through scale and translation invariance; and lastly, larger labeled datasets reduced overfitting, and abundant unlabeled data could be used to better initialize networks in an unsupervised manner.

Deep learning is therefore based on two principles: first, that complex problems can be decomposed into a series of simpler subproblems; and second, what exactly these simpler subproblems are or should be can be discovered from a corpus of data.

CHAPTER IV

TIMBRE SIMILARITY

Timbre has proven to be a difficult attribute to define in acoustic perception, and there is little consensus as result in its underpinnings or the efforts to model it computationally. Psychoacoustics has long sought to better understand the space of timbre using subjective pairwise ratings between acoustic stimuli, but this information is costly to obtain and the generalizability of conclusions ultimately dependent on the palette of sounds considered. This chapter explores an objective, data-driven approach to the development of relative timbre spaces as a scalable alternative to this line of research. Here, instrument taxonomies are used to as a proxy for timbre similarity, and a deep convolutional network is used to project time-frequency representations of audio into a low-dimensional, semantically organized space. The quality of the resulting embeddings is demonstrated through a series of experiments, indicating that this approach shows significant promise for organizing large collections of audio samples by timbre.

1 Context

Despite its common usage in the various forms of music for centuries, a satisfactory definition of *timbre* remains elusive to this day; in fact, the one adopted by the American National Standards Institute embodies this challenge, arriv-

ing at a concept through the exclusion of others (“Psychoacoustic terminology S3:20”, 1973):

Timbre is that attribute of auditory sensation in terms of which a subject can judge that two sounds similarly presented and having the same loudness and pitch are dissimilar.

As evidenced by this definition, the very notion of “timbre” is still an open research topic in psychoacoustics. This reality is captured quite succinctly by Phillippe Manoury, who offered the following insight (Manoury, 1991), as translated by (Donnadieu, 2007):

One of the most striking paradoxes concerning timbre is that when we knew less about it, it didn’t pose much of a problem.

There are many advantages to developing a deeper understanding of timbre, from both an artistic and scientific perspective. Of particular interest to this work, however, the absence of a constructive definition —timbre is a result of X, Y, and Z— makes it difficult to directly build computational systems to characterize and compare timbres. Thus, before proceeding, it is valuable to review what is known of timbre, and prior efforts to transfer this knowledge into engineering systems.

1.1 Psychoacoustics

The perception of timbre falls under the umbrella of *psychoacoustics*, a topic of study that sits at the boundary between acoustics and psychology. Some of the earliest research in psychoacoustics was pioneered by von Helmholtz in his inquiries into the sensations of pitch and loudness (Bregman, 1994). Inquiries

specific to timbre would not come until much later, due to two difficulties in experimental design. One, whereas pitch and loudness can be ranked on a one dimensional scale, it is unclear from personal introspection what the salient dimensions of timbre might be. Instead, it is often necessary to use metaphors as a means of comparing and relating the perception of timbre, such as describing a sound as “bright” or “muddy”. Additionally, researchers were limited by the kinds of stimuli they could create and use in perceptual experimentation, and thus were constrained in the space of possible parameters to explore.

With the advent of computers and continued scientific advances through the 20th century, these issues could be addressed directly, and several researchers set out to identify the existence of fundamental dimensions. This work, performed by Plomp (Plomp, 1976) and Grey (Grey, 1977), among others, adopted a similar experimental design. Human subjects are presented pairs of sound stimuli and asked to rate the similarity between the two. Having collected an exhaustive set of pairwise ratings from a number of participants, multi-dimensional scaling is then used to project the stimuli into a low-dimensional space such that the reported relationships between these datapoints are minimally distorted (Grey, 1977); an example space is shown in Figure 11. Using this similarity model, the researcher then considers a wide array of time-frequency signal statistics, or *features*, in order to identify those that best correlate with the different dimensions. This approach has produced a useful, albeit large, set of features on which computational models have been constructed. Among the earliest were those of log-attack time, spectral centroid, and spectral spread, and were later corroborated by other researchers,

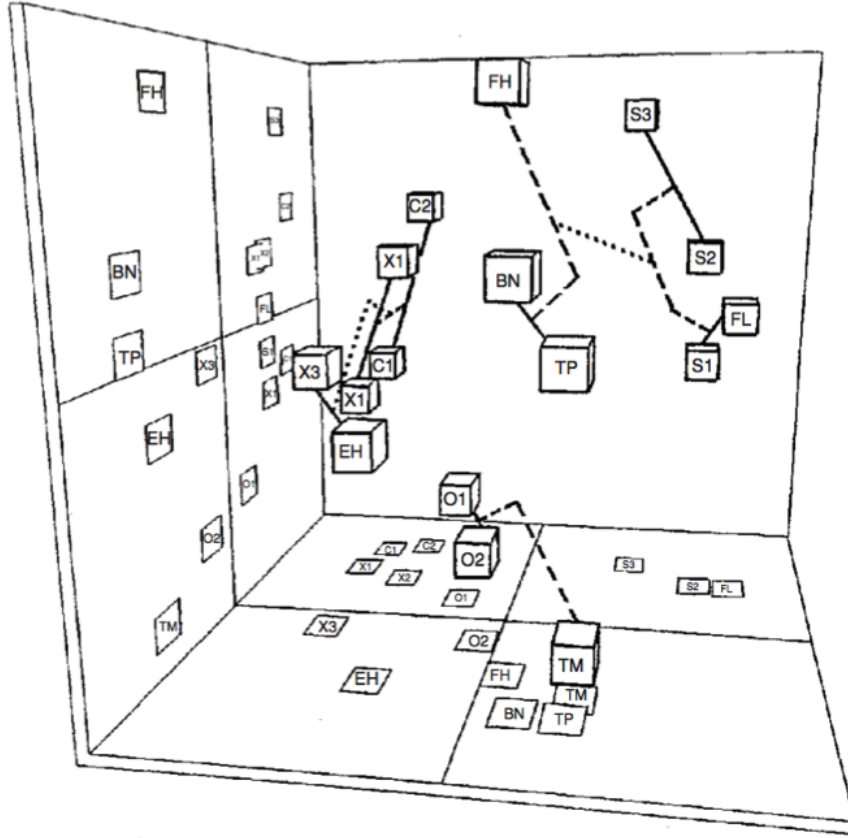


Figure 11: The resulting MDS model developed in the work of Grey.

as in the work of (McAdams, Winsberg, Donnadieu, De Soete, & Krimphoff, 1995).

More recently, however, some have begun to recognize a few shortcomings of this approach to timbre research (Glennon, 2014). First, a timbre space derived from the multidimensional scaling of pairwise ratings is limited to the sonic palette used to produce it, and the inclusion of additional stimuli is likely to rearrange how the space is organized. For instance, the MDS model for a collection of orchestral instruments will be quite different with and without considering electronic synthesizers. This also has significant implications on the granularity of sounds considered. In (Iverson & Krumhansl, 1993), the

attack and sustained regions of a sound were considered separately, resulting in slightly different MDS models. This is not to say that a space derived from relative comparisons is necessarily deficient, however, simply that it must be adapted in the presence of new or different information. Second, the process of finding well-correlated features to explain the resulting MDS model is difficult and time consuming. A researcher must repeat the involved process of feature exploration for every model obtained through a different combination of stimuli. Furthermore, as noted by Caclin et al., “Given the multiplicity of acoustical parameters that could be proposed to explain perceptual dimensions, one can never be sure that the selected parameters do not merely covary with the true underlying parameters.” (Caclin, McAdams, Smith, & Winsberg, 2005). In other words, correlation does not imply causation, and features identified by inspection entail some degree of uncertainty. Finally, the process of collecting subjective pairwise ratings is especially costly, because the number of possible comparisons increases quadratically with number of unique stimuli considered. This places a practical constraint on the generality of a timbre space, as it quickly becomes impossible for subjects to exhaustively rate all combinations. In the absence of complete information, statistical methods, such as imputation, are necessary to interpolate a sparse set of responses.

1.2 Computational Modeling of Timbre

Most previous approaches to computationally modeling timbre instantaneously can be grouped into one of two categories: signal statistics and basis projections. The first follows from the perceptual research described above, whereby specific features are designed to encode some high level semantic concept, e.g. log-attack time or spectral brightness. Initially these corresponded to the fea-

tures named by in the work of Grey or Krumhansl, but have expanded over time to include a wide array of creative and clever measures. The interested reader is directed to (Essid, Richard, & David, 2006) for a comprehensive space of possible features.

From an often complementary perspective, other music researchers have utilised transform-based approaches to project signals into representations with various desirable properties. One of the earliest and most common approaches is the use of Mel-frequency Cepstral Coefficients (MFCCs) for timbre-oriented tasks. Originally designed for speech coding in the 1980s (Davis & Mermelstein, 1980), the first significant contribution in MIR to call attention to MFCCs as useful music features was that of Logan in 2000 (Logan, 2000). MFCCs have, at least in practice, become nearly synonymous with timbre-centric MIR, now being used in a wide array of systems for instrument classification (Herrera-Boyer, Peeters, & Dubnov, 2003), tagging (Barrington, Yazdani, Turnbull, & Lanckriet, 2008), genre prediction (Tzanetakis & Cook, 2002), mood estimation (Schmidt & Kim, 2010) or structural analysis (Paulus, Müller, & Klapuri, 2010), to name only a few works. As described in detail in Chapter ??, the general process of computing MFCCs proceeds as follows: an input audio signal is divided into overlapping, short-time *frames*, on the order of tens to hundreds of milliseconds; a filterbank, perceptually scaled in frequency, is then applied to each short-time frame and log-compressed; finally, a discrete cosine transform (DCT) is applied to these frequency coefficients, characterizing the shape of the spectrum (or the spectrum of the spectrum, referred to as the *cepstrum*). Only the first dozen or so coefficients are used, based on the principle that they compactly describe the spectral contour, though this is more convention than rule. Some have even gone so far

as to literally *equate* MFCCs and timbre, concluding that specific coefficients are responsible for various perceptual dimensions (Terasawa, Slaney, & Berger, 2005).

Similar in principle, though less widely adopted, is to instead *learn* the set of bases against which a time-frequency representation is projected. One such instance is observed in (Jehan, 2005), which preserves the first 12 coefficients of a trained PCA decomposition. In this scenario, the projection into the PCA subspace serves to decorrelate the principal axes of the data in the input space, much like the Discrete Cosine Transform. The primary difference here, however, is that the bases are learned from a sample of observations, rather than defined analytically.

1.3 Motivation

While many computational approaches have proven useful for various classification or recognition tasks, none directly result in a notion of timbre similarity, a useful concept with a variety of applications. One notable instance is the difficulty faced in the search and navigation of large sound sample libraries. Queries are predominantly forced to take the form of text, as in the Freesound* archive shown in Figure 12, which is problematic for at least two reasons. On one hand, it can be challenging to describe a specific query semantically, and often metaphors and figurative language are used to relate the experience of a sound; a distorted guitar might be referred to as ‘crunchy’, or a trumpet as ‘bright.’ Conversely, this kind of descriptive language is far from standardized and varies in meaning from one individual to the next. Furthermore, such

*<http://www.freesound.org/>

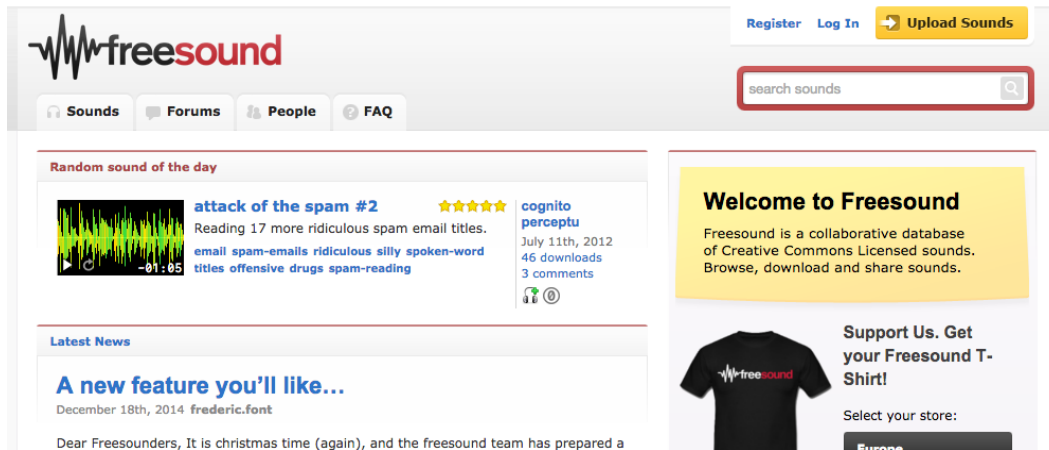


Figure 12: Screenshot of the Freesound homepage. Immediately visible are both the semantic descriptors ascribed to a particular sound (left), and the primary search mechanism, a text field (right).

descriptions are not always associated with every sound in a collection, and typically only at the granularity of the entire recording. As a result, the task of navigating a sound library is often reduced to that of an exhaustive, brute force search.

The development of a robust timbre space would not only make it possible to search for sounds with sounds, bypassing the linguistic intermediary, but also facilitate the ranking of potentially relevant results by providing a notion of distance. This concept of a metric timbre space is also particularly attractive in the realm of user interfaces and visualization. Euclidean spaces are easily relatable by physical analogy, and visualization allows for acoustic information to be understood in an intuitive manner. The ability to explore familiar ideas from an unfamiliar perspective holds considerable merit for artistic exploration and new approaches to composition.

1.4 Limitations

It is valuable to note that despite the difficulty inherent to defining timbre, all computational research must adopt some working concept of it, implicitly or otherwise. Generally two facets to timbre; sound quality and sound source. They are related but not exactly equal. The work presented here operates on the assumption that the perception of timbre is tightly coupled with the experience of discriminating between unique sound sources. This is not intended to be a true equivalence with timbre, but a functional approximation that allows the research to proceed. Compromise and simplification; You'll pay out somewhere. Human judgments of similarity absolve you assumptions about the relationship between source and quality, but costly to curate at scale. A data-driven approach, on the other hand, offers the opposite scenario; it is relatively simple to collect this data, but requires that assumptions be made regarding the quality being related.

Ensemble versus solo. Here focus entirely on the latter.

2 Learning Timbre Similarity

From the previous review of psychoacoustics research and efforts to computationally explain timbre, there is an important series of observations to consider. Classic timbre features are discovered through an involved process of designing a number of signal-level statistics and identifying which correlate with the dimensions of a model. Importantly, those statistics that best explain a timbre space are only valid in the context of the sound sources considered, and thus the process should be repeated for different sonic palettes. Additionally, the subjective ratings necessary to conduct this kind of research are costly to ob-

tain. Therefore, taking cues from the discussion of Chapter II, these practical challenges encourages the use of feature learning to automate the development of timbre similarity models.

Having discussed the value and applications of computational timbre similarity space, it is worthwhile to outline the goals for such a system. First and foremost, one would learn, rather than design, signal-level features relevant to achieve the given task and circumvent the issues identified previously. This idea is based on the combination of an inability to clearly define the sensory phenomenon, while affording the flexibility to change the space of timbres considered. Additionally, sound should be represented in an intuitive manner, such that distance between points is semantically meaningful. In other words, signals from the same source should be near-neighbors, whereas sounds from different sources should be far apart. Finally, the ideal similarity space is perceptually *smooth*, meaning that a point that interpolates the path between two others should be a blend of the two, e.g. a tenor saxophone might fall between a clarinet and a French horn.

These objectives share conceptual overlap with dimensionality reduction methods and instrument classification systems, on which this work builds. In lieu of precise information regarding the relationship between two given sounds, music instrument classes are used as a proxy for timbre similarity. The approach presented here consists of four components, as diagrammed in Figure 13, and discussed in the following subsections. First, all audio is transformed into a time-frequency representation (Subsection 2.1). The main component of the system is a deep convolutional network, which maps tiles of these time-frequency coefficients into a low-dimensional space (Subsection 2.2). A pairwise training harness is made by copying this network, and parameters

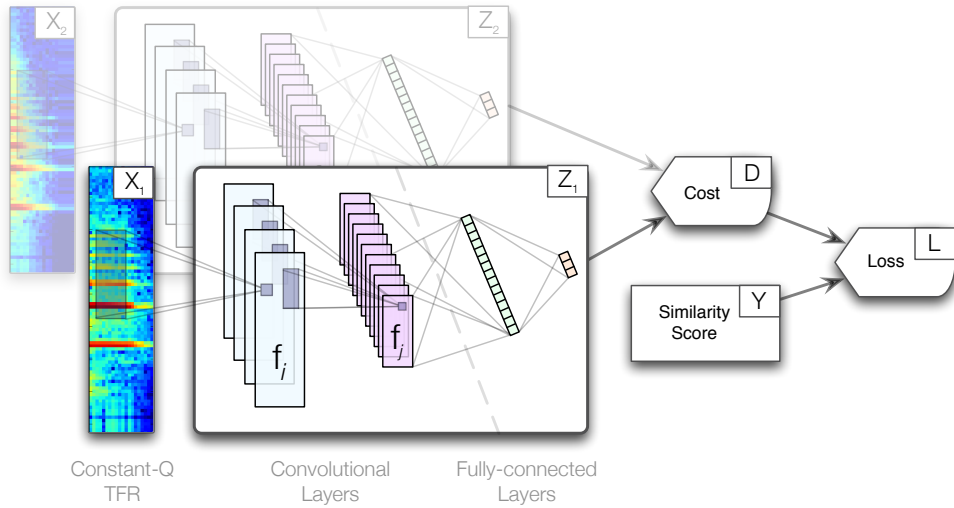


Figure 13: Diagram of the proposed system: a flexible neural network is trained in a pairwise manner to minimize the distance between similar inputs, and the inverse of dissimilar ones.

are learned by minimizing the distance between observations of the same sound source and maximizing the distance otherwise (Subsection 2.3). At test time, the pairwise harness is discarded, and the resulting network is used to project inputs to the learned embedding space.

2.1 Time-Frequency Representation

Time-domain audio signals are first processed by a Constant-Q transform (CQT). The most important benefit of this particular filterbank is that the CQT is logarithmic in frequency. While this serves as a reasonable approximation of the human auditory system, it has the practical benefit of linearizing convolutions in pitch as well as time. It is generally agreed upon that timbre perception is, at least to some degree, invariant to pitch, and this allows the network to behave similarly.

The constant-Q filterbank is parameterized as follows: all input audio is first downsampled to 16kHz; bins are spaced at 24 per octave, or quarter-tone resolution, and span eight octaves, from 27.5Hz to 7040Hz; analysis is performed at a framerate of 20Hz uniformly across all frequency bins. Logarithmic compression is applied to the frequency coefficients with an offset of one, i.e. $\log(X + 1.0)$.

2.2 Deep Convolutional Networks for Timbre Embedding

Noting that the details of deep learning and convolutional networks are discussed at length previously, only those decisions unique to this task are addressed here; for clarity regarding the mathematical or conceptual definitions of these terms, refer to Chapter III.

A five-layer neural network is designed to project time-frequency inputs into a low-dimensional embedding. The first three layers make use of 3D-convolutions, to take advantage of translation invariance, reduce the overall parameter space, and act as a constraint on the learning problem. Max-pooling is applied in time and frequency, to further accelerate computation by reducing the size of feature maps, and allowing a small degree of scale invariance in both directions. The final two layers are fully-connected affine transformations, the latter of which yields the embedding space. The first four hidden layers use a hyperbolic tangent as the activation function, while the visible output layer is linear, i.e. it has no activation function in the conventional sense.

Hyperbolic tangents are chosen as the activation function for the hidden layers purely as a function of numerical stability. It was empirically observed that randomly initialized networks designed with rectified linear units instead were near impossible to train; perhaps due to the relative nature of the learning

problem, i.e. the network must discover an equilibrium for the training data, the parameters were routinely pulled into a space where all activations would go to zero, collapsing the network. Conversely, hyperbolic tangents, which saturate and are everywhere-differentiable, did not suffer the same fate. It is possible that the use of activation functions that provide an error signal everywhere, such as sigmoids or “leaky” rectified linear units, or better parameter initialization might avoid this behavior, but neither are explored here.

It was observed in the course of previous research, that the use of a saturating nonlinearity at the output of the embedding function can lead to problematic behavior (Humphrey et al., 2011). As will be discussed in more detail shortly, bounded outputs makes the choice of hyperparameters crucial in order to prevent the network from pushing datapoints against the limits of its space, and thus the output layer is chosen here to be linear. The absence of boundaries allows the network to find an appropriate scale factor for the embedding. This is similar in principle to the practice of linear “bottleneck” layers in other embedding systems (Yu & Seltzer, 2011).

The input to the network is a 2D tile of log-CQT coefficients with shape $(20, 192)$, corresponding to time and frequency respectively. The frequency channels of the CQT span eight octaves, from 27.5 to 7040 Hz, with quarter-tone resolution. The first convolutional layer uses 20 filters with shape $(1, 5, 13)$ and max-pooling with shape $(2, 2)$. The max-pooling in time introduces a small degree of temporal scale invariance, while the same operation in frequency serves to reduce quartertone to semitone resolution. The second convolutional layer uses 40 filters with shape $(20, 5, 11)$ and max-pooling with shape $(2, 2)$, and the third convolutional layer uses 80 filters with shape $(1, 1, 9)$ and max-pooling with shape $(2, 2)$; in both instances, max-pooling is used to further

reduce dimensionality while introducing more scale invariance. The fourth layer is fully-connected and has 256 output coefficients, while the final layer is also fully connected and has 3 output coefficients.

2.3 Pairwise Training

In the absence of this subjective pairwise instrument ratings, instrument taxonomies are used as a proxy for timbre similarity. This approach to defining timbre “neighborhoods” is used to extend the work of (Hadsell, Chopra, & LeCun, 2006) to address this challenge of learning a timbre similarity space. Referred to by the authors as “dimensionality reduction by learning an invariant mapping” (DrLIM), a deep network was trained in a pairwise manner to minimize the distance between “similar” data points in a learned, nonlinear embedding space, and vice versa. Similarity was determined in an unsupervised manner by linking the k -nearest neighbors in the input space. Though left as future work, the authors propose that other information, such as class relationships, might be leveraged to learn different embeddings. This is an important consideration for the problem of timbre, because fundamental frequency and amplitude are likely to dominate the graph of nearest neighbors defined in the input space alone.

The intuition behind DrLIM is both simple and satisfying: datapoints that are deemed “similar” should be close together, while those that are “dissimilar” should be far apart. Though the precise distance metric is a flexible design decision, it is used here in the Euclidean sense. A collection of similar and dissimilar relationships can be understood by analogy to a physical system of attractive and repulsive forces, where learning proceeds by finding a

balance between them; and furthermore, this analogy illustrates the need for contrasting forces to achieve equilibrium.

At its core, DrLIM is ultimately a pairwise training strategy. First, a parameterized, differentiable function, $\mathcal{F}(X|\Theta)$, e.g. a neural network, is designed for a given problem; in the case of dimensionality reduction, the output will be much smaller than the input, and typically either 2 or 3 for the purposes of visualization. During training, the function \mathcal{F} is copied and parameters, Θ , *shared* between both > Two inputs, X_1 and X_2 , are transformed by their respective functions, \mathcal{F}_1 and \mathcal{F}_2 , to produce the outputs, Z_1 and Z_2 . A metric, e.g. Euclidean, is chosen to compute a distance, D between these outputs. Finally, a similarity score, Y , representing the relationship between X_1 and X_2 , is passed to a contrastive loss function, which penalizes similar and dissimilar pairs differently. Generalizing the original DrLIM approach, different margin terms are applied in the two conditions. For similar pairs, the loss will be small when the distance is small, or zero within the margin m_s ; for dissimilar pairs, the loss will be small when the distance is large, or zero outside a dissimilar margin, m_d . This formal definition is summarized symbolically by the following:

$$Z_1 = \mathcal{F}_1(X_1|\Theta), Z_2 = \mathcal{F}_2(X_2|\Theta)$$

$$D = ||Z_1 - Z_2||_2$$

$$\mathcal{L}_s = \max(0, D^2 - m_s)$$

$$\mathcal{L}_d = \max(0, m_d - D)^2$$

$$\mathcal{L} = Y * \mathcal{L}_s + (1 - Y) * \mathcal{L}_d$$

Note that similarity is given by $Y = 1$, for consistency with boolean logic. As a result, the first term of the loss function is only non-zero for similar pairs, and the inverse is true for the second term.

Returning to the previous discussion regarding the dynamic range of the output layer, it should now be clear that the choice of margin only influences the learned embedding relative to a scale factor when the output is unbounded. The two loss terms are mirrored parabolas, and changing the margin, or horizontal offset, only serves to shift the vertical line about which they reflect. The curvature, and thus the gradient, of the loss function is left unchanged.

Whereas the differential margin controls the spread of all points in space, the similar margin will control the spread of a similarity neighborhood. In the original formulation, where implicitly $m_{sim} = 0$, the loss is lowest when all inputs are mapped to *exactly* the same point; for the purposes of similarity, a more diffuse distribution of points is desirable. It is worth noting the slight parallel to linear discriminant analysis, a statistical method that seeks to jointly minimize intraclass variance and maximize interclass variance. Given

the relative nature of this trade-off, it is sufficient to pick a single ratio between the margins, eliminating the need to vary both hyperparameters separately.

In practice, training proceeded via minibatch stochastic gradient descent with a constant learning rate, set at 0.02 for 25k iterations, or until a batch returned a total loss of zero. Batches consisted of 100 comparisons, drawn such that a datapoint was paired with both a positive and negative example.

3 Methodology

To assess the viability of data-driven nonlinear semantic embeddings for timbre similarity, and thus address the goals outlined at the outset of Section 2, two experiments are used to quantify different performance criteria. First, the local structure and class boundaries of the learned embeddings are explored with a classification task. Second, global organization of the space is measured by a ranked retrieval task. Additionally, in lieu of a subjective evaluation of perceptual “smoothness” of the resulting timbre space, the learned embeddings are investigated through confusion analysis and visualization. In each instance, the approach presented here is compared to a conceptually similar, albeit admittedly simpler, system.

Finally, the formulation described in the previous section presents two system variables, thus giving rise to two additional considerations:

1. What is the effect of using different margin ratios?
2. How does the sonic palette considered impact the learned embedding?

3.1 Data

The data source used herein is drawn from the Vienna Symphonic Library (VSL), a truly massive collection of studio-grade orchestral instrument samples recorded over a variety of performance techniques*. In aggregate, the VSL contains over 400k sound recordings from more than 40 different instruments, both pitched and percussive. Sorting instrument classes by sample count yields 27 instruments with at least 5k samples; three of these instruments, however, are not reasonably distinct from other sources, e.g. “flute-1” and “flute-2”, and discarded rather than risk introducing conflicting information. This decision yields the set of instruments contained in Table 1 for experimentation.

The distribution of sound files for these instruments, grouped by class, is given in Figure 14. As discussed previously, it is an inherent difficulty of pairwise similarity models that the resulting relationships are limited by the number of unique classes considered. Fortunately, there is no added cost to considering a wider palette of sound sources here because the label information is objective. Therefore, building upon previous work (Humphrey et al., 2011), three configuration subsets are repeated from the pilot study as well as a fourth consisting of all 24 classes, given in Table 2.

For each instrument class, 5k samples are drawn, without replacement, to build a uniformly distributed collection. This step simplifies the process of data sampling during stochastic training of the network, which may be sensitive to class imbalances. The collection of instrument samples is stratified into five equal partitions for cross validation, used at a ratio of 3-1-1 for training,

*<https://vsl.co.at/en>

Table 1

Instruments considered and their corresponding codes.

Instrument	Code	Instrument	Code
French Horn	ho	Tuba	tu
Violin	vi	Cimbasso	ci
Bb Clarinet	klb	Piccolo	pt
Tenor Trombone	tp	Oboe	ob
C Trumpet	trc	Bass Clarinet	bkl
Bass Trombone	bp	Wagner Tuba	wt
Acoustic Concert Guitar	akg	Contra Bassoon	kfa
Bassoon	fa	English Horn	eh
Cello	vc	Bass	kb
Bass Trumpet	bt	Soprano Saxophone	sxs
Distorted Guitar	eg	Tenor Saxophone	sxt
Flute	fl	Alto Flute	afl

validation, and testing, respectively. The partitions are circularly rotated such that each is used as the test set once, i.e. (1, 2, 3)-4-5, (2, 3, 4)-5-1, and so on.

3.2 Margin Ratios

Though the pairwise training strategy described in Section 2.3 consists of two margin hyperparameters, it is ultimately the ratio between the two that governs how the space will be shaped. In isolation, the exact choice of dissimilar term’s margin, m_d , is inconsequential and determines the radius of the bounding sphere. Going forward, this value is fixed to $\sqrt{12}$, corresponding to the radius of the sphere that intersects the coordinate (2, 2, 2). Moving the similar term’s margin, m_s , relative to this value will lead to different embeddings, and

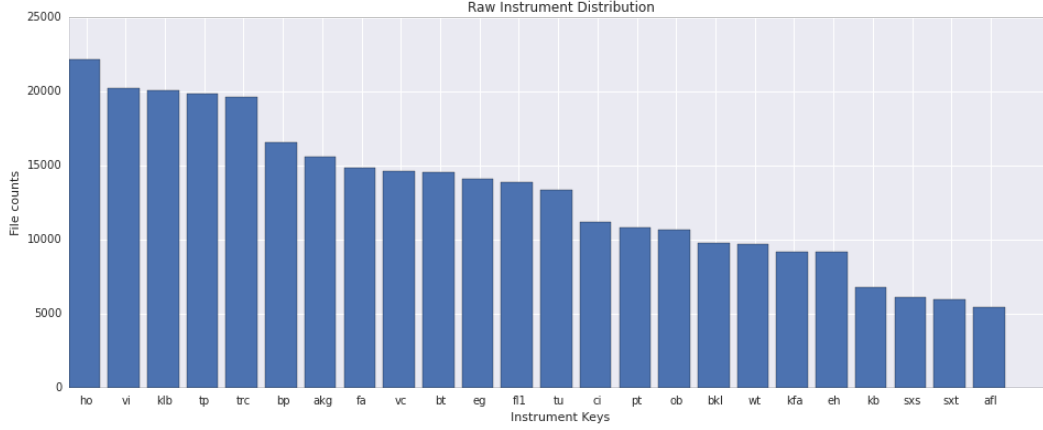


Figure 14: Distribution of instrument samples in the Vienna Symphonic Library.

Table 2

Instrument set configurations.

Key	Instrument Codes
c5	tu, ob, klb, vc, fl
c8	trc, ho, ob, eh, klb, sxt, vi, vc
c12	c8 + {tp, tu, fa, fl}
c24	c12 + {bp, akg, bt, eg, ci, pt, bkl, wt, kfa, kb, sxs, afl}

three ratios of $r_m = \frac{m_s}{m_d}$ are considered here: 0, $\frac{1}{4}$, and $\frac{1}{2}$. The corresponding loss functions are shown in Figure 15.

3.3 Comparison Algorithm

For the purposes of comparison, a similarly motivated system is constructed using the combination of principal components analysis (PCA) and linear discriminant analysis (LDA). Previous work explored the application of PCA alone and locally linear embedding as alternative approaches to dimensionality reduction. Both are unsupervised methods, and do not make for the most fair comparison against a supervised neural network. LDA, however,

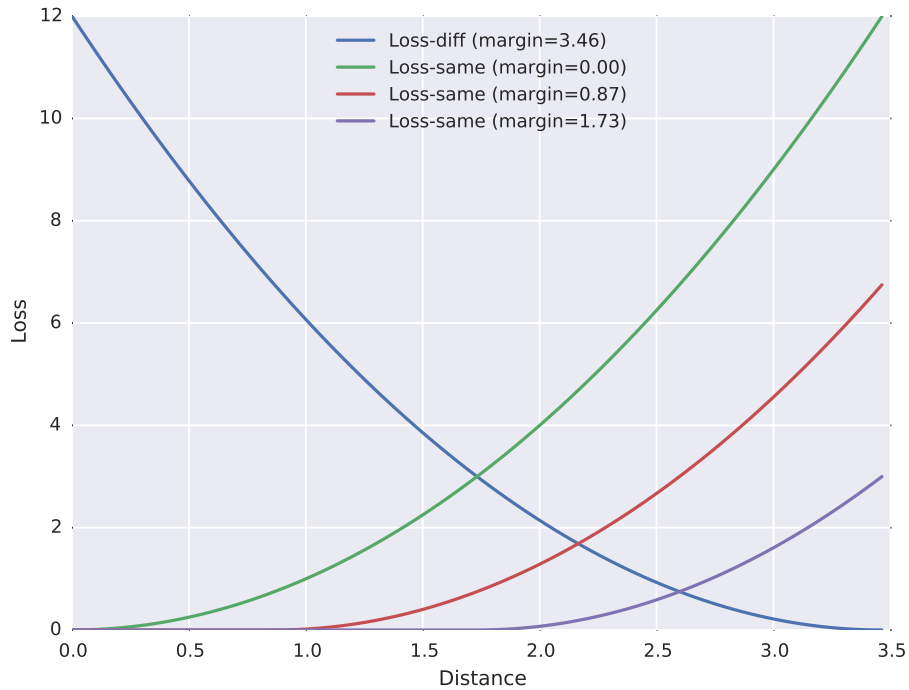


Figure 15: Loss contours for different margin ratios.

is a supervised approach to dimensionality reduction, and shares at least a conceptual parallel to the proposed system, as mentioned briefly in Section 2.3.

It is important to note though that LDA can exhibit odd behavior in high dimensional spaces, and projecting into a PCA subspace first can help alleviate these issues (Ji & Ye, 2008). This subspace projection is further motivated by computational efficiency concerns, where the input dimensionality is prohibitive to training. Additionally, the cascade of PCA followed by LDA mimics a two-layer neural network, and is interchangeable with the framework described here. Using the same input dimensions, 20×192 , a whitened PCA transform is fit to a large sample of the training set. The principal 256

components are preserved, based on an empirical exploration of the explained variance as well as a midway point in the dimensionality reduction of the system, i.e. the number of coefficients decreases near-equally between the PCA and LDA stages. After applying the PCA transform to the training sample, an LDA transform is fit to the same data and its corresponding instrument classes, yielding a 3-dimensional embedding.

3.4 Experimental Results

As an initial quantitative inquiry, trained models are tested on a classification task using the k-Nearest Neighbors classifier in scikit-learn*. First, networks are trained across the 4 instrument configurations, 3 margin ratios, and 5 folds, and all data are projected into the resulting embedding space. From here, a collection of points are sampled from each partition —50k, 10k, 25k— for training, validation, and test, respectively. The training set is used to fit the classifier, while the validation set is used to identify an optimal setting for the parameter k , corresponding to the number of neighbors considered in the decision rule. Classification accuracy is then computed across all three sets, and tallied across folds to produce averages and standard deviations across the various conditions; these results are given in Tables 3-5.

A few conclusions are immediately obvious from these results. Most striking is the performance discrepancy between the NLSE and the PCA-LDA models. Previous work demonstrated a significant margin between the unsupervised dimensionality reduction methods, and this result shows that the difference is indeed a function of complexity, not just the supervised learning

*<http://scikit-learn.org/>

Table 3

kNN classification results over the training set.

config	c5	c8	c12	c24
NLSE, $r_m = 0.0$	93.81 ± 0.53	89.97 ± 0.40	86.70 ± 0.39	74.17 ± 0.79
NLSE, $r_m = 0.25$	94.21 ± 0.18	90.25 ± 0.54	87.17 ± 0.35	73.91 ± 0.61
NLSE, $r_m = 0.5$	93.04 ± 0.29	89.16 ± 0.27	86.08 ± 0.26	71.59 ± 0.88
PCA-LDA	64.44 ± 0.47	56.58 ± 0.60	47.22 ± 0.45	35.81 ± 0.28

Table 4

kNN classification results over the validation set.

config	c5	c8	c12	c24
NLSE, $r_m = 0.0$	92.37 ± 0.64	87.94 ± 0.45	84.52 ± 0.97	70.86 ± 1.51
NLSE, $r_m = 0.25$	93.75 ± 0.53	88.62 ± 0.58	85.65 ± 0.26	71.46 ± 0.63
NLSE, $r_m = 0.5$	91.75 ± 0.67	87.78 ± 0.85	83.78 ± 0.53	66.37 ± 1.69
PCA-LDA	59.97 ± 0.96	52.49 ± 2.68	39.25 ± 2.01	24.32 ± 0.97

Table 5

k-Neighbors classification results over the testing set.

config	c5	c8	c12	c24
NLSE, $r_m = 0.0$	92.49 ± 0.41	88.26 ± 0.74	84.35 ± 0.41	70.67 ± 0.55
NLSE, $r_m = 0.25$	92.97 ± 0.41	88.67 ± 0.79	85.16 ± 0.24	70.28 ± 0.86
NLSE, $r_m = 0.5$	91.96 ± 0.35	87.83 ± 0.25	84.04 ± 0.57	66.91 ± 0.57
PCA-LDA	59.91 ± 0.77	50.24 ± 1.52	39.32 ± 0.86	24.77 ± 0.51

process. To a lesser extent, all models show some degree of over-fitting, but the effect is more severe for the PCA-LDA model than any NLSE. It is interesting to note that a non-zero similarity margin, m_s , leads to slightly better classification results than the centered loss function. One explanation for such behavior is that introducing a small region of zero-loss within a class may allow

the network to emphasize dissimilar relationships more as training proceeds. It would appear too much freedom, on the other hand, leads to fuzzy boundaries between classes and begins to compromise local structure.

The outcome of the classification experiment can also be used to inform how smooth or intuitive this space might be. To do so, confusion matrices are shown for the c12 configuration for the best NLSE, with a margin ratio of 0.25, and the PCA-LDA model, in Tables 6 and 7, respectively.

Though more confusions are to be expected in the PCA-LDA model, given the classification accuracy, it is important to note that these errors are distributed somewhat uniformly across classes, rather than correlated with instrument similarity. This higher noise-floor indicates that the instruments’ distribution exhibit a good deal of overlap in space. Some logical confusions seem unavoidable, such as French horn (ho) and trombone (tp), or flute (fl) and oboe (ob), occurring in both models. The former makes sense given common instrument families, i.e. brass, while the latter likely arises from the upper range of the instruments, which has fewer harmonics.

Other instrument relationships also appear to confound some element of pitch height in similarity, particularly for the PCA-LDA model. This is observed in the confusions between tuba, bassoon, and French horn. In the NLSE model, tuba is confused with French horn more often than bassoon; for the PCA-LDA model, however, the inverse is true. Intuitively, the two brass instruments should share the higher confusion rate, and thus pitch is being used by the LDA model as a feature with which to distinguish between classes. The convolutional model, on the other hand, is forced to embrace a considerable amount of pitch invariance, and is prevented from making the same error.

Table 6

Confusion Matrix for c12; NLSE with a margin ratio of 0.25.

Reference	Estimated											
	eh	fa	fl	ho	klb	ob	sxt	tp	trc	tu	vc	vi
eh	85.52	1.10	0.46	3.05	1.76	3.05	0.44	0.23	2.53	0.30	0.51	1.64
fa	1.74	85.82	0.05	3.93	0.26	0.19	0.63	0.81	0.51	3.64	2.48	0.51
fl	0.80	0.10	85.20	0.90	2.19	6.00	1.67	0.14	2.33	0.07	0.33	1.17
ho	0.76	1.88	0.07	82.52	0.26	0.29	0.33	6.50	1.18	2.73	0.88	1.26
klb	1.23	0.40	2.46	1.16	86.57	3.02	1.80	0.11	1.01	0.25	1.37	1.05
ob	2.90	0.06	3.09	1.12	2.56	81.22	0.44	0.17	5.29	0.04	0.04	1.39
sxt	0.24	0.38	1.01	0.51	1.24	0.84	86.34	0.14	0.48	0.65	4.97	2.78
tp	0.39	0.87	0.10	11.87	0.03	0.49	0.20	80.96	2.38	2.73	0.95	0.59
trc	1.14	0.11	1.77	3.84	0.56	4.13	0.59	1.74	83.45	0.08	0.09	2.47
tu	0.04	1.55	0.04	5.32	0.04	0.01	0.57	2.18	0.09	86.44	2.82	0.57
vc	0.27	0.53	0.24	1.51	0.79	0.49	2.68	0.27	0.63	2.32	89.74	1.49
vi	0.49	0.23	0.61	2.44	0.46	1.20	2.46	0.48	2.05	0.41	2.06	87.32

Table 7

Confusion Matrix for c12; PCA-LDA.

Reference	Estimated											
	eh	fa	fl	ho	klb	ob	sxt	tp	trc	tu	vc	vi
eh	46.10	3.01	11.27	6.79	2.61	10.92	0.21	9.69	9.85	0.79	1.20	1.04
fa	5.12	46.84	0.37	14.42	0.31	0.27	1.88	7.55	0.58	17.15	2.47	0.57
fl	13.33	1.62	20.82	6.93	4.01	17.74	1.79	4.26	16.32	1.63	2.10	1.85
ho	5.45	19.49	2.11	43.53	1.77	0.29	0.94	15.47	1.38	7.51	1.53	4.70
klb	10.80	4.88	11.69	10.62	9.04	9.61	4.84	5.47	11.99	6.51	7.96	5.07
ob	15.45	0.40	17.80	1.67	3.09	31.93	0.56	2.42	19.92	0.69	0.81	2.14
sxt	0.48	2.77	2.28	3.62	2.33	0.97	47.47	1.40	3.45	9.48	14.48	15.77
tp	15.98	9.58	6.45	19.43	2.02	5.74	0.52	18.93	9.42	4.33	1.15	2.07
trc	10.72	0.20	14.14	3.64	3.50	19.71	0.77	5.67	36.01	0.27	1.12	4.97
tu	0.04	12.39	0.06	7.13	0.77	0.03	4.35	3.68	0.03	62.74	7.67	0.26
vc	0.39	1.65	2.87	2.66	2.09	2.36	18.99	1.16	4.29	10.02	44.95	12.05
vi	0.93	1.78	2.63	5.04	1.62	3.22	13.43	1.44	7.86	1.43	4.49	56.47

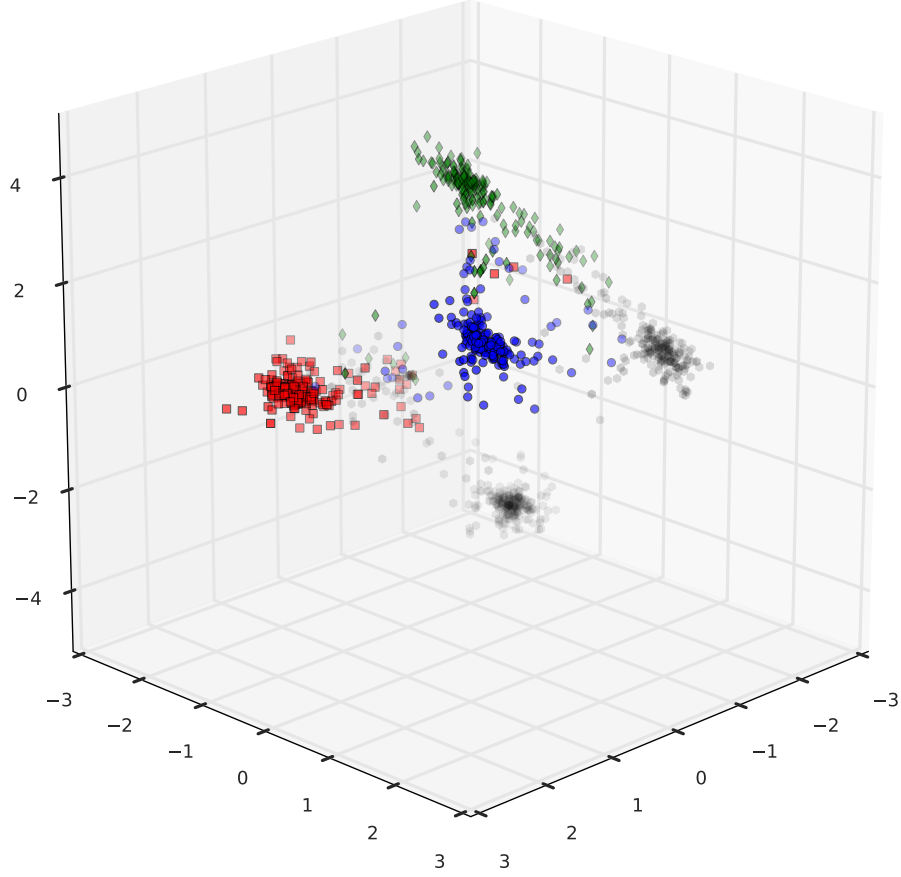


Figure 16: Embeddings of clarinet (blue circles), oboe (green diamonds), and cello (red squares) observations across models trained with the “c5” instrument configurations.

To help illustrate the semantic organization of the learned embedding, 3D scatter plots are given in Figures 16–19 following observations of the three instruments common to all configurations —clarinet, oboe, cello— across the different embeddings for $\frac{m_s}{m_d} = 0.25$. Other instruments are displayed as semi-transparent black to help highlight the three instruments of interest, while giving an impression of the overall space. There are several takeaways of note

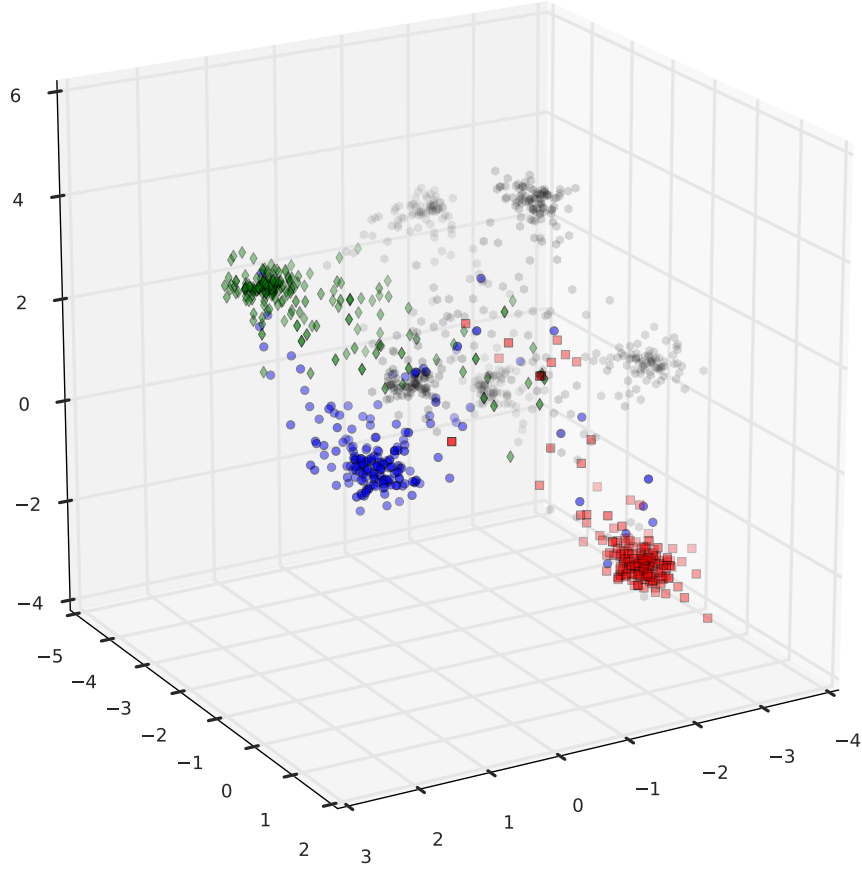


Figure 17: Embeddings of clarinet (blue circles), oboe (green diamonds), and cello (red squares) observations across models trained with the “c8” (top) and “c8” (bottom) instrument configurations.

revealed through visualization. As to be expected, the various sonic palettes used to learn the embedding yield different organizations of points in space. That said, the relationship between the three sources is relatively consistent, as the cluster of clarinet always sits between oboe and cello. Somewhat undesirably, the NLSEs do not achieve very good diffusion in space. Though there are some interpolating points, clusters are clearly visible in at least the first

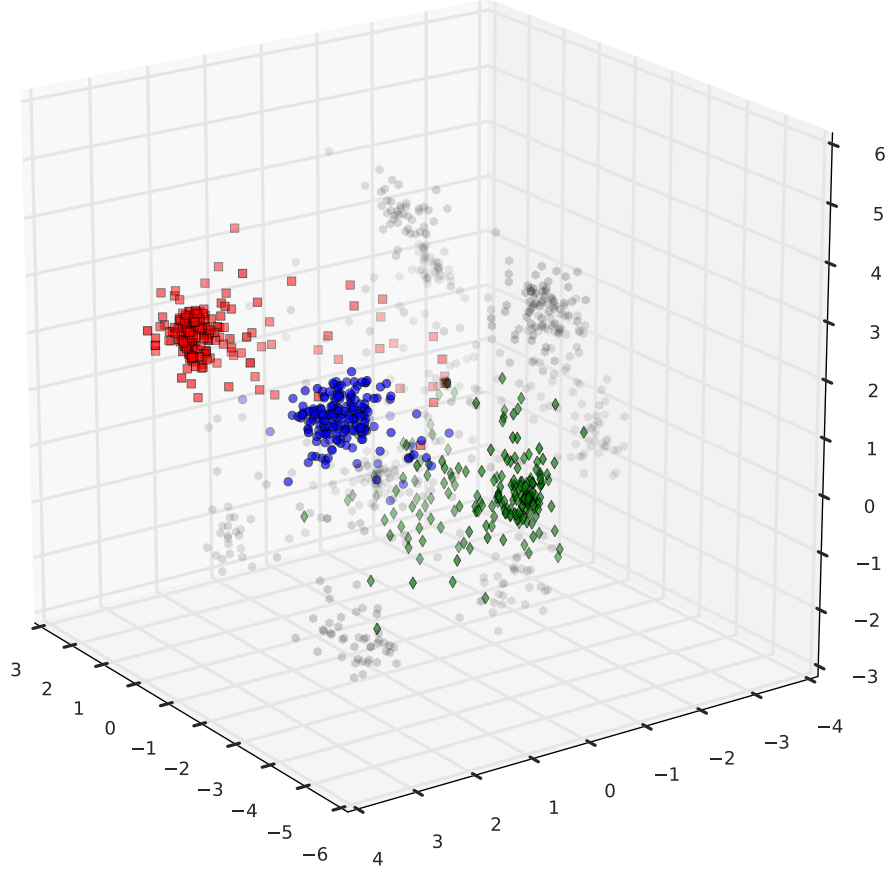


Figure 18: Embeddings of clarinet (blue circles), oboe (green diamonds), and cello (red squares) observations across models trained with the “c12” instrument configurations.

three instrument configurations. In the “c24” condition, good cluster definition begins to break down slightly, although the three instruments of interest remain clearly clustered. Compared to results obtained previously, the use of a linear output layer has indeed eliminated undesirable boundary behavior, producing embeddings that comfortably occupy a volume roughly centered about the origin. An unbounded output has the potential to drift infinitely in some

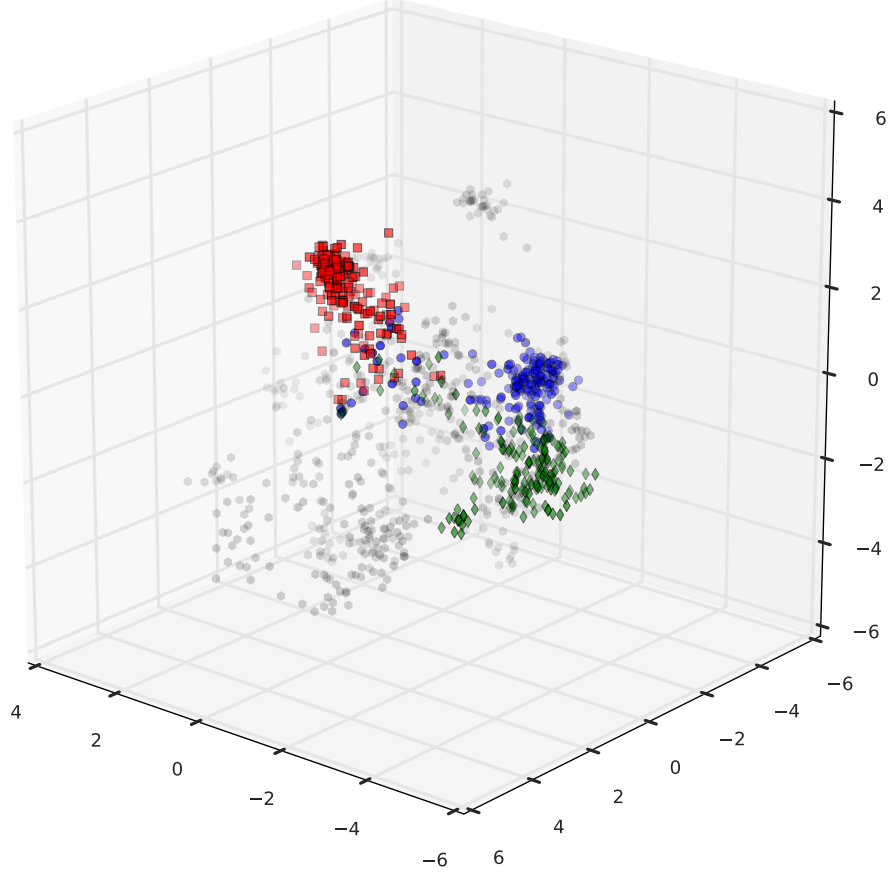


Figure 19: Embeddings of clarinet (blue circles), oboe (green diamonds), and cello (red squares) observations across models trained with the “c24” instrument configurations.

direction, which might happen in the presence of biased or noisy data, and so it is encouraging that the NLSEs remain centered.

To further test the semantic organization of the learned embeddings, the outputs are used as features for a ranked retrieval task, using Euclidean distance as a scoring function. Whereas kNN classification quantifies the local organization of points in space, measuring precision as a function of recall

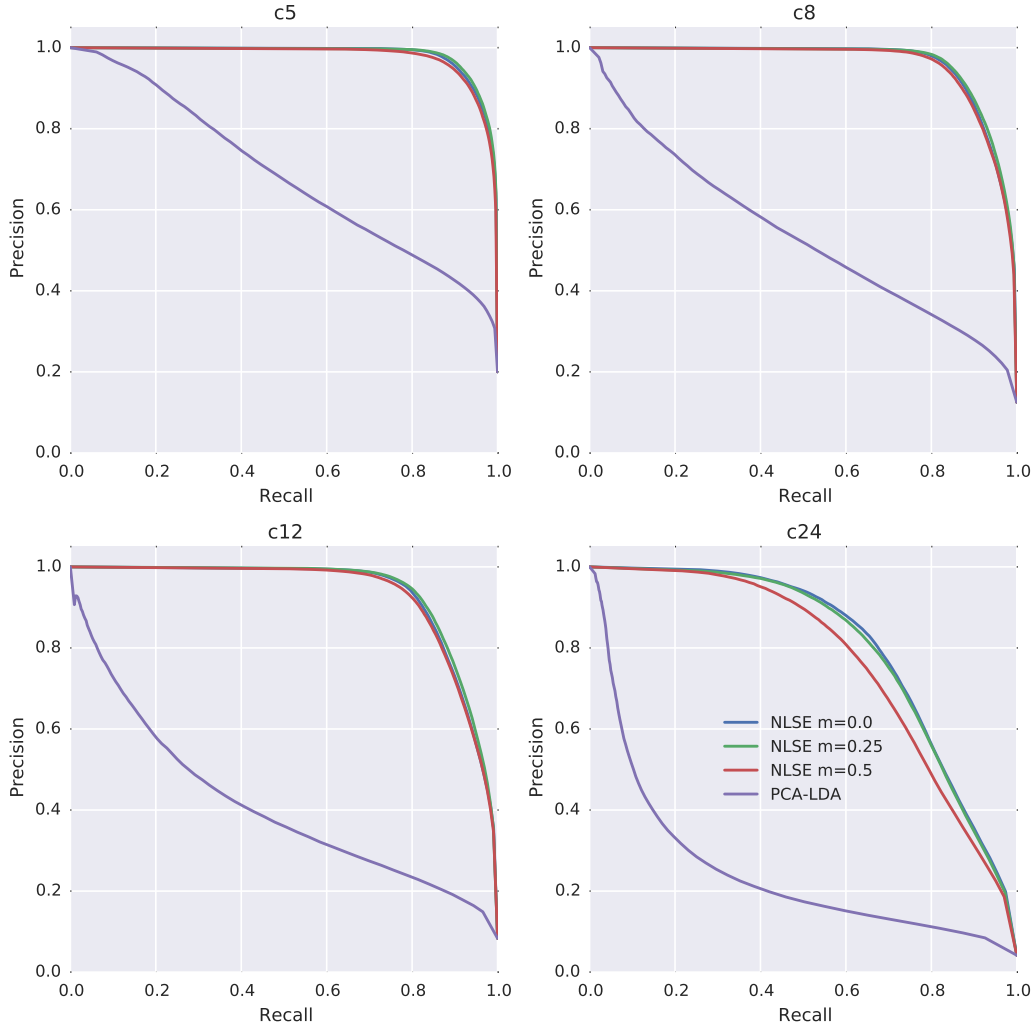


Figure 20: Recall-Precision curves over the four instrument configurations.

helps characterize how the data are organized globally. This method of analysis provides insight into how useful the embedding would be as a distance-based instrument sample search engine. Recall-precision (RP) curves are computed using the scikit-learn toolkit and averaged over the five folds; the resulting curves for the four configurations are shown in Figure 20. An RP-curve illustrates how precision varies as more relevant items are returned; thus, in the

ideal condition, an RP-curve approaches the upper-right corner of the plot, versus the lower-left in the worst-case scenario.

Given the classification accuracy and confusion matrices above, the performance gap between the NLSE and PCA-LDA models is unsurprising. Still, it is interesting to consider what the shape of these recall-precision curves indicates. The two characteristics to observe are the concavity of the contour and the “knee” at which it breaks downward. In all instrument configurations, with the slight exception of “c5”, the NLSE models and the PCA-LDA model exhibit opposite second-derivatives. This behavior can be understood as the acceleration with which precision changes as a function of recall. For the NLSEs, precision degrades slowly until reaching a crossover point, referred to here as the knee, where precision drops off rapidly. The PCA-LDA model does the opposite, where precision drops quickly close to a query point, and slows as recall increases. Therefore, as encouraged by the visualizations, the NLSEs contain better separated class clusters than the PCA-LDA embeddings. Furthermore, the knee of a recall-precision curves belies an interesting region in the document space, indicating that the edge of a cluster has been reached. This is a useful observation for determining early-stopping criteria in the display of ranked results, as well as identifying boundary regions in the embedding that may present interesting opportunities for sonic exploration.

Looking to differences between NLSEs, there are two variables to consider: instrument configuration and margin ratio. For the first three instrument configurations, precision is roughly 100% out past a recall of 0.6; this can be understood as the top 60% of results for a given query will correctly match that instrument. In the final condition though, “c24”, this break occurs just past a recall of 0.2, at which point other instrument samples would start

to appear in the list of results. This is still quite good from the perspective of precision at the top of the ranked list, but it illustrates the point at which the embedding space is beginning to get crowded. A higher dimensional embedding might alleviate this at the expense of visualization, but would still be suitable for a retrieval system. With respect to the margin ratio, the first three instrument conditions are again roughly equivalent, whereas “c24” begins to show more contrast between the embeddings. In all conditions $\frac{m_s}{m_d} = 0.5$ yields the lowest performing NLSE, and is most pronounced in the final condition. This is consistent with expectation, because the larger value of m_s de-emphasizes the tightness of instrument clusters, causing more overlap between the different classes.

4 Conclusions

In this chapter, an approach to building a computational model of timbre similarity was presented, which achieved three goals. First, the system is able to automatically learn relevant signal level features, circumventing the challenge posed by the lack of a constructive definition of timbre. Second, the resulting timbre space is semantically well-organized, as demonstrated by classification and ranked retrieval measures, and intuitive, based on a Euclidean notion of distance in a dimensionality that can be easily visualized. Lastly, the space is quantitatively smooth, such that what confusions exist correspond to instrument families or other like sounds. This was made possible by leveraging objective information about a collection of sound sources, eliminating the need for costly subjective ratings. Together, this approach to learning a timbre

space shows promise for visualization and user-facing applications, such as the search and navigation of large sound libraries.

That said, there is considerable future work to be considered. All evaluation performed here is quantitative, and arguably disconnected from all subjective experience. User studies would serve to further investigate the ideas of perceptual smoothness and if or how is obtained by the learned space. Efforts to analyze the learned features may further help elucidate the latent factors that contribute to the percept of timbre. Additionally, though this approach is able to make use of objective instrument taxonomies, any similarity space obtained through pairwise comparisons is always limited by the range of inputs considered. Therefore, in order to obtain a more general timbre space, a much wider set of sound sources would need to be considered. Conversely, the intended use case of such a system may provide a constrained palette with which to operate, e.g. instrument sounds for recording engineers and environmental sounds for acoustic ecologists. Finally, there are at least two other ways the sound source information could be used to train a system in a supervised manner. One, it may be advantageous to obtain subjective pairwise ratings not between all possible sounds, but rather groups or classes of sounds. These pairwise ratings could be used to train a system with soft, continuous-valued similarity ratings, rather than the binary comparison scores used here. Two, rather than defining an entire class to be similar, a hybrid approach to similarity based on distance-based neighborhoods in the input space constrained to a single class may also lead to interesting embeddings. It is unlikely such an embedding would exhibit spherical clusters as was produced here, but points are likely to be more uniformly distributed, or diffuse, in space.

CHAPTER V

AUTOMATIC CHORD ESTIMATION

The focus of this study now turns to automatic chord estimation (ACE), one of the oldest subtopics in the field of content-based MIR. Complementing the previous chapter, ACE presents a more challenging, well-worn problem with a long research lineage and myriad applications. In addition to the standard objective of advancing the state of the art in a given application domain, this chapter also aims to use deep learning to explore challenges inherent to complex music intelligence tasks. Adopting a similar approach as before, deep convolutional neural networks are used to estimate the likelihoods of different chord classes from time-frequency representations of audio. A thorough investigation serves to not only offer insight into the application of deep learning methods to future problems in content-based MIR, but also culminate in a better understanding of the chord estimation task itself.

1 Context

In this section, the prerequisite knowledge relevant to a thorough treatment of automatic chord estimation systems is addressed. First, a basic review of Western tonal theory introduces core music concepts, and thus provides a language with which to discuss the problem. This is followed by a brief outline of the syntax used here to compactly notate chords. The problem domain is then jointly motivated by potential use-cases and the observed needs of modern

musicians. Finally, known limitations are detailed to identify assumptions and subsequently contextualize any conclusions drawn from this work.

1.1 Musical Foundations

To understand the chord estimation task is to understand the concepts on which it is built. Therefore, given the significant emphasis on signal processing and machine learning in this work, it is therefore valuable to provide a basic review of chords and harmony for the benefit of the technical reader unfamiliar with music theory.

The most atomic unit in the acoustic realm of chords is a *tone*, defined here as a pitched sound object. Pitch is defined as the perceptual phenomena whereby a sound stimulus can be matched to the frequency of a pure sinusoid, known as the fundamental frequency (Krumhansl, 1979). As a result, sounds can be naturally ordered by fundamental frequency on a scale from low to high. An *octave* is defined as the doubling of a quantity, and exhibits a special relationship in human perception by which two tones, differing in fundamental frequency by an integer power of 2, are perceived as being similar; this phenomena is referred to as octave equivalence.

A *note* is the musical counterpart of a tone, and the two are related by way of a tuning system. While there are a multiplicity of tuning systems, this work will exclusively consider 12-Tone Equal Temperament, or 12-TET, named for the 12 discrete, equally spaced tones within an octave. The relationship between notes and tones in 12-TET is defined by the following:

$$f_{pitch} = f_{tuning} * 2^{\exp(\frac{n_{index} - 48}{N})} \quad (19)$$

where N is the number of notes per octave, n_{index} is an integer note index, f_{tuning} is a reference tuning frequency, in Hertz, and f_{pitch} is the fundamental frequency, in Hertz, of the corresponding tone. Standard contemporary tuning equates $A4 = 440Hz$, although it should be noted that this is more convention than rule and is subject to change. In 12-TET, the unique *pitch classes* within an octave are named, given by the ordered set, \mathcal{P} :

$$\mathcal{P} = \{C, C\sharp/D\flat, D, D\sharp/E\flat, E, F, F\sharp/G\flat, G, G\sharp/A\flat, A, A\sharp/B\flat, B\} \quad (20)$$

Here, sharps, \sharp , and flats, \flat , are symbols used to indicate the raising or lowering of a note by one *semitone*, respectively. Due to this particular tuning system, some pitch classes can be spelled with either a sharp or a flat, e.g. $A\sharp = B\flat$, a property known as *enharmonic equivalence*. An absolute note name, consisting of a pitch class, p , and an octave, o , is given by the following as a function of absolute note index, such that $n_{index} = 0 \rightarrow C0$, $n_{index} = 12 \rightarrow C1$, etc.:

$$p = \mathcal{P}[\text{mod}(n_{index}, 12)]$$

$$o = \lfloor n_{index}/12 \rfloor$$

Most real music combines different notes, and thus it is useful to define an *interval* as the relative semitone distance between two notes. Intervals are critical in the understanding of harmony because they are generally perceived as similar regardless of absolute pitch height. For example, the interval from

C5 to G5 sounds the same as the interval from F3 to C4, both being seven semitones.

From here, three interdependent harmonic concepts are built through the simultaneous and sequential combination of intervals: scales, chords, and keys. It is crucial to recognize that each is an emergent quality of intervallic relationships, and efforts to define one purely in terms of another are somewhat circular. That said, an ordered set of intervals is known as a scale, of which the *diatonic* is the most widely used in common practice music. It consists of seven intervals, given by the following:

$$\{+2, +2, +1, +2, +2, +2, +1\}$$

Rotating this sequence circularly results in different *modes*, of which two are common in contemporary popular music: *major*, with a rotation of 0; and *minor**, with a rotation to the right of 3. Each scale's *degrees* are expressed by the following, shown here as a semitone distance from a starting note:

$$\text{major} = \{0, 2, 4, 5, 7, 9, 11, 12\}$$

$$\text{minor} = \{0, 2, 3, 5, 7, 8, 10, 12\}$$

* More accurately, this is the natural minor scale, but the distinction is not terribly important here.

The pitch class on which the scale starts is referred to the *tonic*, and lends its name to the scale. The sequences in (1.1) can be used to recover a scale by circularly indexing the set of pitch classes given in (20). To illustrate, a C Major and A minor scale are given by the following:

$$\mathcal{C}_{major} = \{C, D, E, F, G, A, B\}$$

$$\mathcal{A}_{minor} = \{A, B, C, D, E, F, G\}$$

It should be noted that these two scales, despite different modes and tonics, are composed of identical notes. These scales are known as each other's relative major and minor, respectively, and share a strong perceptual affinity.

Proceeding from scales, a *chord* is classically conceived as a simultaneous grouping of notes. One of the most important chord types in Western tonal theory is the *triad*, on which many other theoretical constructs are built. Comprised of three notes, a triad is built by taking the third and fifth scale degrees from a chosen starting point, referred to as the *root*. For example, this expansion is given for the major scale in Table 8.

For a given chord, the *root* is defined as the home note on which the intervals are stacked, and the *quality* determined by the relationship of the intervals to the root. An interval of 4 semitones is known as a *major third* and 3 semitones a *minor third*, sharing a commonality with the third scale degree of the major and minor scales, respectively. Therefore, the qualities of the first six chords in the table are named for their relationship with the corresponding

Table 8

Roman numeral, quality, semitones, and adjacent intervals of triads in the Major scale.

Roman Numeral	Quality	Semitones	Intervals
I	Major	{0, 4, 7}	{+4, +3}
ii	minor	{2, 5, 9}	{+3, +4}
iii	minor	{4, 7, 7}	{+3, +4}
IV	Major	{5, 9, 12}	{+4, +3}
V	Major	{7, 11, 2}	{+4, +3}
vi	minor	{9, 0, 4}	{+3, +4}
vii°	diminished	{11, 2, 5}	{+3, +3}
I	Major	{0, 4, 7}	{+4, +3}

major and minor scales; the vii° chord, however, is “diminished” because it contains two minor thirds.

Sharing much common ground with scales and chords, a *key* identifies a point of harmonic stability in reference to a tonic and its corresponding triad. Accordingly, the naming of a “key” takes a pitch class and either a major or minor quality, e.g. E-major or C♯-minor. Key is integral to the discussion here, because its impression or expectation establishes a harmonic framework with which one can parse music, facilitating the understanding of notes and intervals as they relate to scales and chords. For a more detailed review, the curious reader is directed to (Laitz & Bartlette, 2009).

1.2 What is a “chord”?

While much of this theory can be detailed specifically, real music is by no means so well behaved. As a result, a more practical definition of a “chord” is

open to some interpretation, and may take multiple meanings. For example, (McVicar, 2013) collects three possible definitions, restated here:

1. *Everyone agrees that chord is used for a group of musical tones.*
2. *Two or more notes sounding simultaneously are known as a chord.*
3. *Three or more pitches sounded simultaneously or functioning as if sounded simultaneously.*

Additionally, (Harte, 2010) expands the scope of (2) in order to describe all tonal music, “allow[ing] a chord to comprise zero or more notes.” The various flavors of definitions begs an obvious question: what makes the concept of a chord so hard to pin down?

Much of this difficulty stems from the fact that the relative importance of the individual notes in a collection may change in different contexts. In practice, a chord is named based on three, potentially subjective, criteria: its root, its contributing intervals, and how these two relate to the perceived key. Therefore, as will be shown shortly, a chord can easily take a different name if any of these decisions are changed or re-weighted.

Having briefly reviewed Western tonal theory, a deeper understanding of the variation inherent to defining a chord can be obtained by exploring a few simple examples. The one invariant property shared by all definitions named previously is the idea that a pitch collection may be understood as a single harmonic object. The time span over which this phenomena may unfold, however, is flexible. To illustrate the point, consider the three bars notated in Figure 21, where a major chord is written as a true simultaneity, an arpeggiation, and as an series of non-overlapping quarter notes, respectively.

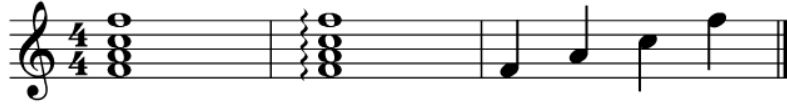


Figure 21: A stable F major chord played out over three time scales, as a true simultaneity, an arpeggiation, and four non-overlapping quarter notes.



Figure 22: A stable C major chord is embellished by passing non-chord tones.

In this instance, the degree of overlap in time is expanded until it no longer exists, and yet the collection of notes continues to function as a coherent harmonic object.

On the other hand, as shown in Figure 22, the simultaneous sounding of different notes does not necessarily give rise to the perception of different chords. Here, a major triad is sustained under the first several degrees of its scale. While three notes in the upper voice are contained in the C-major triad, the others –the D, F, and A– are referred to as “nonchord” tones. These extra notes are explained away in the overall harmonic scene, as they fall on metrically weak beats, are comparatively short in duration, and quickly move to notes that *are* in the chord. These embellishments do not contribute to the harmonic center of the phrase, and the bar can still be understood as a stable C major chord.

A last example, shown in Figure 23, illustrates the level of complexity and decision making that may arise in the process of describing music in the lan-

The image shows a handwritten harmonic analysis of a piano piece, identified as '161. Dance' by Schubert. The analysis is written on a four-staff system, with the original musical notation in the top two staves and the harmonic analysis in the bottom two staves. The key signature is three flats (B-flat, E-flat, A-flat), and the time signature is 3/4. The piece is marked 'p' (piano) and 'f' (forte). The analysis includes various chord symbols: I (tonic), (i) (first inversion tonic), (iii) (third inversion tonic), V^7 (dominant seventh), (vi) (sixth degree), and IAC (first inversion augmented triad). The analysis also includes a 'repeated phrases' bracket and an 'anticipating period' bracket. The piece is marked 'Schubert' and '161. Dance'.

Figure 23: A sample harmonic analysis of a piano piece, performed as a music theory exercise.

guage of chords. Referred to as *harmonic analysis*, this exercise is performed in an effort to understand the harmonic content in a theoretical manner, such that chords are notated alongside the original score. There are many observations one might draw from this example, but a few are of particular interest here. First, even in an instance such as this, where an individual is able to operate directly on the symbolic representation, it is common to find alternate reasonable interpretations of the same musical content. As notated in measure 2 for example, the combination of the A and F can be understood both as an

embellishment on the *I*, or as an implied *iii* chord. When performed, however, the musician can influence how one might perceive this simultaneity through the use of expressive timing or dynamics. Additionally, this instance focuses on a harmonically simple excerpt for solo piano. The introduction of other voices will only serve to complicate the resulting musical surface, especially where timbre is considered. Lastly, this kind of theoretical analysis is developed in, and largely tailored to, the tradition of Western tonal music. While contemporary popular music is certainly influenced by this tradition, it by no means adheres to the same rules and conventions.

1.3 Chord Syntax

It is a pragmatic but necessary prerequisite step to define a standard syntax for compactly notating chords. Much of the pioneering work in this space was performed by Harte (Harte, Sandler, Abdallah, & Gómez, 2005), and many of these conventions are utilized here. Going forward, chords expressed in this scheme are stylized with a fixed-width font, e.g. `A:min`.

Firstly, Harte’s general chord notation is described by the following four-part symbolic description:

$$\text{root} : \text{quality} (\text{intervals}) / \text{bass} \quad (21)$$

Every chord name begins with a `root` in the form of a pitch class, optionally modified by zero or more sharps or flats, or one of two reserved characters: `N` for the “null” no-chord condition, or `X` for the special case in which the musical content cannot be described harmonically.

The root is potentially followed by a `quality` shorthand, separated by a

Table 9

Chord quality names and corresponding relative semitones.

Name	Shorthand	Semitones
Major	maj	{0, 4, 7}
Minor	min	{0, 3, 7}
Major 7	maj7	{0, 4, 7, 11}
Minor 7	min7	{0, 3, 7, 10}
Dominant 7	7	{0, 4, 7, 10}
Major 6	maj6	{0, 4, 7, 9}
Minor 6	min6	{0, 3, 7, 9}
Diminished	dim	{0, 3, 6}
Augmented	aug	{0, 4, 8}
Suspended 2 nd	sus2	{0, 2, 7}
Suspended 4 th	sus4	{0, 5, 7}
Fully-diminished 7	dim7	{0, 3, 6, 9}
Half-diminished 7	hdim7	{0, 3, 6, 10}

colon and implying a particular set of note intervals. Though there are a large number of possible chord qualities, this is often limited to a particular subset. Those considered in this work are indicated in Table 9.

The third field provides a set of **intervals**, wrapped by parentheses. In practice, there are two reasons for representing information intervallically. One such instance is, through a combination of additional degrees and asterisks, the modification of a quality shorthand in order to represent a non-standard, but related, chord. An example of this might be the chord name **A:min(*b3, b7)**, indicating that the minor third (*C*) is absent and a minor 7 (*G*) has

been added. The other instance occurs when the intervals are certain but the quality is ambiguous, such as $\mathbb{C}:(1, 5)$.

The final field of this chord syntax is the **bass** interval, which indicates the scale degree of the lowest contributing pitch. Typically this is also the root of the chord, and is implied as such in the absence of an explicit bass interval. However, it is necessary to state that the scale degrees of the chord —given by the quality shorthand and the interval set— can be further augmented by the inclusion of a bass interval. For example, the chords $\mathbb{C}:\text{maj}/\text{b}7$ and $\mathbb{C}:7$ would be understood as containing the same pitch classes, but are spelled differently.

1.4 Motivation

Even from the earliest efforts in content-based MIR, automatic music transcription has stood as one of the Holy Grails of the field. Time would prove this to be an exceptionally difficult problem, and fracture this common cause into a variety of smaller, and hopefully more manageable, subtopics. Automatic chord estimation materialized as one such task, now receiving healthy attention for more than a decade, and is established as a benchmark challenge at the annual MIREX event*.

Given the prerequisite skill necessary to produce chord transcriptions manually from recorded audio, there is considerable motivation to develop automated systems capable of reliably performing this task. As evidenced by large online communities surrounding websites like e-chords[†] or Ultimate Guitar[‡], countless individuals invest considerable time and effort in the cura-

*http://www.music-ir.org/mirex/wiki/MIREX_HOME

[†]<http://www.e-chords.com>

[‡]<http://www.ultimate-guitar.com>

tion and consumption of popular music transcriptions. Often this is driven by desire to learn and perform music for which symbolic notation does not exist. Conversely, automatic chord estimation systems would be particularly useful in the areas of composition, recording, and production. Furthermore, the curation of this content would enable large-scale musicological analysis of contemporary music.

In addition to the concerns of individual users, computational systems capable of reliable chord estimation are directly useful inside the domain of content-based MIR. Chords can serve as a robust mid-level representation with which to build systems and extract higher level musical knowledge, and have been used for cover song retrieval (Bello, 2007) and genre recognition (Anglade, Ramirez, & Dixon, 2009). Such systems would also facilitate data collection for other tasks, aiding in visualization and other facets of music transcription.

From a more philosophical perspective, the identification of chords is also intriguing as an intelligent musical behavior, being a high level cognitive process that is often open to multiple interpretations between knowledgeable experts. Experiential bias of the annotator may manifest in the subjective decisions made by an observer, where a pianist may arrive at a different harmonic analysis than that of a guitarist due to how a collection of notes might be produced. Finally, the knowledge and skill of the one recognizing chords in music will affect the resulting interpretations. Beginners will likely prefer simple or more common descriptions, whereas experts will be more aware of complex nuances and have better command over a larger harmonic vocabulary.

1.5 Limitations

It should be acknowledged that this inquiry is subject to the limitations of tonal theory and chords as a language with which to describe a piece of music. Primarily, this work is interested in the tradition of tonal Western music, with a particular focus on popular contemporary music from the last century, in 12-TET. Within this large body of music content, the use of harmony and chords has steadily evolved over time. Classically, music theorists have long sought to characterize musical works via analysis and reduction as a means to understanding, typically operating from a symbolic representations, i.e. a score. As a result, the language of “chords” developed as an expressive, yet compact, language with which one might describe a piece of music harmonically. However, Western “pop music”, infused with elements of folk, blues, jazz, rock and countless other influences, is not required to adhere to or consider the rules of traditional tonal theory (Tagg, 1982). Thus efforts to understand the former in the language of the latter is ultimately limited by the validity in doing so.

Even in the constrained space of Western tonal music set forth here, not all musical works will be well-described by the language of harmonic analysis, and thus chords may be a clumsy language with which to describe such music. An alternative approach to analysis, such as voice leading, might make more sense in this instance; in others, such as “math rock”, a lack of clearly structured harmonic content may arguably render the goal of harmonic analysis irrelevant (Cateforis, 2002). As genre is itself an amorphous and ill-defined concept, the degree to which a piece of music might be understood harmonically will vary, both absolutely and internally.

2 Previous Research in Automatic Chord Estimation

Building upon the conceptual foundations addressed previously, automatic chord estimation research can be described in three parts. First, an effort is made to formally define the goals of computational systems. The research lineage is then surveyed, identifying commonalities between this work and highlighting the state of the art. Approaches to evaluation methodology are discussed last, including a review of data used to benchmark the research presented here.

2.1 Problem Formulation

Following the motivations outlined in 1.4, the goal of an automatic chord estimation (ACE) system is –or, at least, has been– to produce “good” time-aligned sequence of chords from a given music signal. As discussed in 1.3, it is a particular nuance of chord notation that the space of valid spellings is effectively infinite. To constrain the complexity of the task at hand, ACE systems are traditionally designed to estimate chords from a finite *vocabulary*, defined *a priori*. This simplification reduces the chord estimation to a classification problem, where all observations are assigned to one of K chord *classes*.

Historically, the choice of chord vocabulary has been anything but standard, influenced primarily by the data available to a researcher. Supervised machine learning approaches, for example, can be sensitive to the amount of labeled data available for training, in which case it might be advantageous to only consider sufficiently represented chord classes. Furthermore, not all researchers have access to the same data, introducing another degree of vari-

ability. As a result, it can be challenging, if not impossible, to compare the performance of systems designed for different chord vocabularies.

To address this challenge, one common strategy employed by the research community is that of Major-Minor chord resolution. Based on the common 24 Major and minor keys, this formulation proceeds by mapping all chords in a collection to either a Major or Minor chord with the same root (McVicar, 2013). While this results in some musically reasonable chord mappings, e.g. `maj7` \rightarrow `maj`, others are more difficult to justify, e.g. `dim7` \rightarrow `min` or `aug` \rightarrow `maj`.

Having framed chord estimation as a classification problem, there are two critical assumptions to note going forward. First, the classification paradigm operates on the notion that the relationship between an observation and its corresponding class is stable. Chord estimation research has classically leveraged expert musicians in the spirit of achieving objectivity, but this is ultimately an approximation to some unknown degree. Second, flat classification problems—those in which different classes are conceptually independent—are built on the assumption of mutually exclusive relationships. In other words, assignment to one class precludes the valid assignment to any other classes considered. For example, “cat” and “dog” are mutually exclusive classes of “animal”, but “cat” and “mammal” are not. Returning to chords, `C:dim7` and `C:maj` are clearly mutually exclusive classes, but it is difficult to say the same of `C:maj7` and `C:maj`, as the former *contains* the latter.

2.2 Computational Approaches

Considering the space of ACE research, it is observed by (Cho, 2014) all approaches to the task adopt the same basic architecture, diagrammed in Fig-

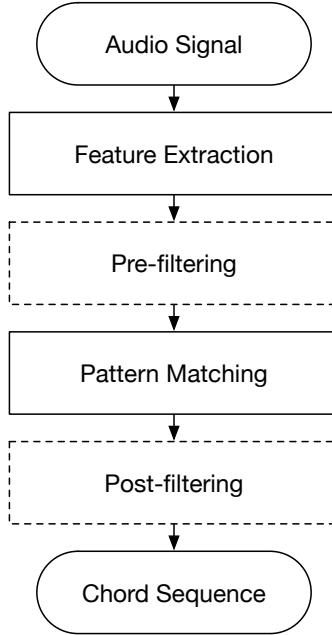


Figure 24: Block-diagram of the common building blocks in modern automatic chord estimation systems.

Figure 24. First, harmonic features, referred to as pitch class profiles (PCP) or *chroma*, are extracted from short-time observations of the audio signal. Initially proposed for use in chord estimation systems by Fujishima (Fujishima, 1999), chroma features attempt to measure the amount of energy in the signal corresponding to the 12 pitch classes named in Eq. 20. These features may then be processed by any number of means, referred to in the literature as *pre-filtering*. Importantly, this is done prior to the next stage of *pattern matching*, which is performed on the final feature representation to measure how similar the observed signal is to a set of chord names. The process of pattern matching, a relatively local operation, is mapped over a much longer signal, e.g. a full recording, yielding a time-varying estimate of the various

chord types the model can represent. Finally, *post-filtering* is applied to the output of the pattern matching stage, resulting in a sequence of chord names over time.

Though the implementation details have continued to evolve over the last decade, the brunt of chord estimation research has concentrated not on the fundamental system per se, but rather the tuning of its components. In particular, much time and energy has been invested in developing not just better features, but specifically better *chroma* features (Müller & Ewert, 2010). Complementing chroma features, others have explored the use of multi-band chroma to model bass frequencies separately (Mauch & Dixon, 2010b) or a Tonnetz representation in an effort to better encode harmonic relationships between chords (Lee & Slaney, 2007). Acknowledging the challenges inherent to designing good features, Pachet et al pioneered work in automatic feature optimization (Cabral & Pachet, 2006), and more recently deep learning methods have been employed to learn robust Tonnetz features (Humphrey, Cho, & Bello, 2012). Early methods focused on local smoothing, such as low-pass or median filtering as a form of pre-filtering (Cho et al., 2010), but more recently some methods have attempted to leverage the repeated nature of music to yield more stable estimates of the harmonic composition at a given point in time (Cho & Bello, 2011). Various classification strategies have been investigated such as binary templates (Oudre, Grenier, & Févotte, 2009), Dirichlet distribution models (Burgoyne & Saul, 2005), or Support Vector Machines (SVMs) (Weller, Ellis, & Jebara, 2009), but Gaussian Mixture Models (GMM) are by and large the most common feature modeling approach (Cho, 2014). The choice of post-filtering methods has been shown to significantly impact system performance, and much research has focused on properly tuning Hid-

den Markov Models (HMMs) (Cho et al., 2010), first introduced by (Sheh & Ellis, 2003). Recently, in an effort to continue to advance the state of the art, researchers have begun exploring more complex post-filtering methods such as Dynamic Bayesian Networks (DBNs) (Mauch & Dixon, 2010a), Conditional Random Fields (Sumi et al., 2012), and Variable-Length Markov Models (Chordia et al., 2011).

It is worth noting that in this lineage, the systems that do make use of data-driven learning typically only do so in disjoint stages. More often than not, machine learning is only performed at the pattern matching stage, where increasingly powerful models are fit to hand-crafted features. A few works do attempt to learn features, such as (Mauch & Dixon, 2010a; Humphrey, Cho, & Bello, 2012), but the different stages are optimized independently. Though it is standard practice to train a GMM/HMM jointly, some have observed that learning the parameters of the HMM, i.e. the transition probabilities, yields no significant benefit over a uniform probabilities with a strong self-transition affinity (Cho, 2014). One notable work that attempts to jointly optimize multiple stages is that of (Cho, Kim, & Bello, 2012), which optimizes the GMM to a minimum frame classification error, rather than a conventional maximum likelihood formulation.

2.3 Evaluation Methodology

In order to objectively measure the quality of a proposed ACE system, it is necessary to address two related components: the collection of ground-truth data, and the manner in which estimations are compared to this reference data.

The first major effort to curate ground truth chord transcriptions was

led by Harte in the mid-2000s, referred to as the Isophonics dataset*, where a small team of researchers transcribed the entire discography of The Beatles. Containing chord annotations for 180 tracks, this was a landmark dataset in the field of MIR and shaped years of ACE research. Importantly, this transcription effort leveraged professional transcriptions of the music under consideration, and was verified for accuracy multiple times. However, despite this rigor, the data is drawn from a single artist and very well known to the research community; some have argued that ACE research has begun to manually overfit this collection.

To combat these issues, two datasets were released following the 2011 Conference of the International Society of Music Information Retrieval (ISMIR), one of over 700 tracks led by J. Ashley Burgoyne (Burgoyne, Wild, & Fujinaga, 2011) and another of 295 tracks led by Tae Min Cho, from the Music and Audio Research Lab at NYU†; here, the former is referred to as “Billboard” and the latter as “MARL-Chords”, corresponding to their related projects. In parallel, an additional, comparatively small, dataset was released, containing 20 tracks by the band Queen, provided by Matthias Mauch as an extension to the Isophonics set. In all four cases, the chord transcriptions are provided as “ground truth”, on the premise that the data corresponds to the expert perspective. To help prevent errors and resolve judgment calls, these additional annotation efforts employed a review process, where the transcriptions of one or more annotators were verified by a different individual.

Leveraging this ground truth data, it is possible to quantitatively assess

*<http://isophonics.net/content/reference-annotations>

†<https://github.com/tmc323/Chord-Annotations>

the outputs of a computational system. Expressed formally, the conventional approach to scoring an ACE system is a weighted measure of chord-symbol recall, R_W , between a reference, \mathcal{R} , and estimation, \mathcal{E} , chord sequence as a *continuous* integral over time, summed over a collection of N pairs:

$$R_W = \frac{1}{S} \sum_{n=0}^{N-1} \int_{t=0}^{T_n} C(\mathcal{R}_n(t), \mathcal{E}_n(t)) dt \quad (22)$$

Here, C is a chord *comparison* function, bounded on $[0, 1]$, t is time, n the index of the track in a collection, T_n the duration of the n^{th} track. S corresponds to the cumulative amount of time, or *support*, on which C is defined, computed by a similar integral:

$$S = \sum_{n=0}^{N-1} \int_{t=0}^{T_n} (\mathcal{R}_n(t), \mathcal{E}_n(t) \in \mathfrak{R}) dt \quad (23)$$

Defining the normalization term S separately is useful when comparing chord names, as it relaxes the assumption that the comparison function is defined for all possible chords. Furthermore, setting the comparison function as a free variable allows for flexible evaluation of a system’s outputs, and thus all emphasis can be placed on the choice of comparison function, C . In practice, this measure has been referred to as *Weighted Chord Symbol Recall* (WCSR) (Harte, 2010), *Relative Correct Overlap* (TCO) (McVicar, 2013), or *Framewise Recognition Rate* (Cho, 2014), but it is, most generally, a recall measure.

As discussed, most ACE research typically proceeds by mapping all chords into a smaller chord vocabulary, and using an enharmonic equivalence comparison function at evaluation, e.g. $\text{C\#}:\text{maj} == \text{Db}:\text{maj}$. Recently, this approach was generalized by the effort behind the open source evaluation tool-

Table 10

Chord comparison functions and examples in `mir_eval`.

Name	Equal	Inequal	Ignored
Root	G#:aug, Ab:min	C:maj/5, G:maj	—
Thirds	A:maj, A:aug	C:maj7, C:min	—
Triads	D:dim, D:hdim7	D:maj, D:aug	—
Sevenths	B:9, B:7	B:maj7, B:7	sus2, dim
Tetrads	F:min7, F:min(b7)	F:dim7, F:hdim7	—
majmin	E:maj, E:maj7	E:maj, E:sus2	sus2, dim
MIREX	C:maj6, A:min7	C:maj, A:min	

box, `mir_eval` (Raffel et al., 2014), introducing a suite of chord comparison functions. The seven rules considered here are summarized in Table 10.

The meaning of most rules may be clear from the table, but it is useful to describe each individually. The “root” comparison only considers the enharmonic root of a chord spelling. Comparison at “thirds” is based on the minor third scale degree, and is equivalent to the conventional mapping of all chords to their closest major-minor equivalent. In other words, a chord with a minor-third is minor, e.g. `dim7` \rightarrow `min`, and *all* other chords map to major, e.g. `sus2` \rightarrow `maj`. The “triads” rule considers the first seven semitones of a chord spelling, encompassing the space of major, minor, augmented, and diminished chords. The “sevenths” rule is limited to major-minor chords and their tetrad extensions, i.e. major, minor, and dominants; chords outside this set are considered “out-of-gamut” and ignored. The “tetrads” comparison extends this to all chords contained within an octave, e.g. six chords and half-diminished sevenths. The “Major-minor” comparison is limited to major

and minor chords alone; like “sevenths”, other chords are ignored from evaluation. Unlike the other rules, “MIREX” compares chords at the pitch class level, and defines equivalence if three or four notes intersect. Comparing the pitch class composition of a chord allows for a slightly relaxed evaluation, allowing for misidentified roots and related chords. Finally, rules that ignore certain chords only do so when they occur in a reference annotation. In other words, an estimation is not held accountable for chords deemed to be out of gamut, but predicting such chords is still counted as an error.

Complementing these rules, it was recently proposed by Cho in (Cho, 2014) that, when working with larger chord vocabularies, special attention should be paid to performance across all chord qualities. The motivation for additional measures stems from the reality that chord classes are not uniformly distributed, and a model that ignores infrequent chords will not be well characterized by global statistics. Instead, Cho proposes a chord quality recall measure, R_Q whereby all chord comparisons are rotated to their equivalents in C , and averaged without normalizing by occurrence.

$$R_Q = \sum_{q=0}^{Q-1} \frac{1}{W_q} \sum_{n=0}^{N-1} \int_{t=0}^{T_n} C(\mathcal{R}_n(t), \mathcal{E}_n(t)|q) \partial t \quad (24)$$

Referred to originally as *Average Chord Quality Accuracy* (ACQA), this metric weights the contributions of the individual chord qualities equally, regardless of distribution effects. Notably, as the overall chord distribution becomes more uniform, this measure will converge to Eq. (22). However, given the significant imbalance of chord classes, large swings in any overall weighted recall statistic may result in small differences of the quality-wise recall, and vice versa. It

should also be noted that the only comparison function on which quality-wise recall is well defined is strict equivalence.

3 Pilot Study

Here, a preliminary study conducted by the author in 2012, and presented at the International Conference of Machine Learning and Applications (ICMLA 2012), is revisited and expanded upon to frame subsequent work (Humphrey & Bello, 2012). Approaching ACE from the perspective of classifying music audio among the standard 24 Major-Minor classes, in addition to a no-chord estimator, a deep convolutional network is explored as a means to realize a full chord estimation system. Doing so not only addresses the questions of relevance or quality toward chroma as a representation, but error analysis of an end-to-end data-driven approach can be used to gain insight into the data itself. This observation gives rise to two related questions: one, how does performance change as a function of model complexity, and two, in what instances can the model *not* overfit the training data?

3.1 Experimental Setup

Audio signals are downsampled to 7040Hz and transformed to a constant-Q time-frequency representation. This transform consists of 36 bins per octave, resulting in 252 filters spanning 27.5–1760Hz, and is applied at a framerate of 40Hz. The high time-resolution of the constant-Q spectra is further reduced to a framerate of 4Hz by mean-filtering each frequency coefficient with a 15-point window and decimating in time by a factor of 10. As discussed previously, a constant-Q filterbank front-end provides the dual benefits of a

reduced input dimensionality, compared to the raw audio signal, and produces a time-frequency representation that is linear in pitch, allowing for convolutions to learn pitch-invariant features.

The input to the network is defined as a 20-frame time-frequency *patch*, corresponding to 5 seconds. A long input duration is chosen in an effort to learn context, thereby reducing the need for post-filtering. Local-contrast normalization is applied to the constant-Q representation, serving as a form of automatic gain control, and somewhat similar in principle to log-whitening used previously in chord estimation (Cho et al., 2010). As an experimental variable, data are augmented by applying random circular shifts along the frequency axis during training within an octave range. The linearity of pitch in a constant-Q representation affords the ability to “transpose” an observation as if it were a chord of a different root by shifting the pitch tile and changing the label accordingly. Every data point in the training set then contributes to each chord class of the same quality (Major or minor), having the effect of inflating the dataset by a factor of 12. The two conditions —before and after augmentation— are referred to henceforth as “As-Is” and “Transposed”.

A five-layer 3D convolutional network is used as the general model, consisting of three convolutional layers and two fully-connected layers. Six different model complexities are explored by considering two high-level variables, given in Table 11. The width of each layer, as the number of kernels or units, increases over a small (S), medium (M), and large (L) configuration. Two different kernel shapes are considered, referred to as 1 and 2. Note that only the first convolutional layer makes use of pooling, and only in the frequency dimension, by a factor of three in an effort to learn slight tuning invariance.

Table 11

Model Configurations - Larger models proceed down the rows, as small (S), medium (M), and large (L); two different kernel shapes, 1 and 2, are given across columns.

	1	2
S	K:(1, 4, 6, 25), P:(1, 3)	K:(1, 4, 5, 25), P:(1, 3)
	K:(4, 6, 6, 27)	K:(4, 6, 5, 13)
	K:(6, 8, 6, 27)	K:(6, 8, 5, 13)
	W:(480, 50)	W:(2560, 50)
	W:(50, 25)	W:(50, 25)
M	K:(1, 6, 6, 25), P:(1, 3)	K:(1, 6, 5, 25), P:(1, 3)
	K:(6, 9, 6, 27)	K:(6, 9, 5, 13)
	K:(9, 12, 6, 27)	K:(9, 12, 5, 13)
	W:(720, 125)	W:(3840, 125)
	W:(125, 25)	W:(125, 25)
L	K:(1, 16, 6, 25), P:(1, 3)	K:(1, 16, 5, 25), P:(1, 3)
	K:(16, 20, 6, 27)	K:(16, 20, 5, 13)
	K:(20, 24, 6, 27)	K:(20, 24, 5, 13)
	W:(1440, 200)	W:(7680, 200)
	W:(200, 25)	W:(200, 25)

The output of the final layer is passed through a softmax operator, producing an output that behaves as a likelihood function over the chord classes.

As this work predates access to the Billboard and Queen datasets, only the MARL-Chords and Beatles collections are considered, totaling 475 tracks. All chords are resolved to their nearest major-minor equivalent, as discussed in Section 2.1, based on the third scale degree: `min` if the quality should contain a flat third, otherwise `maj`. The collection of 475 tracks are stratified into five folds, with the data being split into training, validation, and test sets at a ratio of 3–1–1, respectively. The algorithm by which the data are stratified

Table 12

Overall recall for two models, with transposition and LCN.

	L-1			S-1		
Fold	Train	Valid	Test	Train	Valid	Test
1	83.2	77.6	77.8	79.6	76.9	76.8
2	83.6	78.2	76.9	80.5	77.0	76.8
3	82.0	78.1	78.3	80.0	77.2	78.2
4	83.6	78.6	76.8	80.2	78.0	75.8
5	81.7	76.5	77.7	79.5	75.9	76.8
Total	82.81	77.80	77.48	79.97	77.00	76.87

is non-trivial, but somewhat irrelevant to the discussion here; the curious reader is referred to the original publication for more detail. Model parameters are learned by minimizing the Negative Log-Likelihood (NLL) loss over the training set. This is achieved via mini-batch stochastic gradient descent with a fixed learning rate and batch size, and early stopping is performed as a function of classification error over the validation set. Training batches are assembled by a forced uniform sampling over the data, such that each class occurs with equal probability.

3.2 Quantitative Results

Following the discussion of evaluation in 2.3, the only comparison function used here is “thirds”, and all statistics correspond to a weighted recall measure.

As an initial benchmark, it is necessary to consider performance variance over different test sets. The outer model configurations in the first column of Table 11 (Arch:L-1 and Arch:S-1) were selected for five-fold evaluation, influenced by run-time considerations. Overall recall is given in Table 12, and

offers two important insights. One, deep network chord estimation performs competitively with the state of the art at the major-minor task. Previously published numbers on the same dataset fall in the upper 70% range (Cho et al., 2010), and it is encouraging that this initial inquiry roughly matches state of the art performance. Noting the variation in performance falls within a 2% margin across folds, a leave-one-out (LoO) strategy is used for experimentation across configurations, with and without data transposition.

The overall recall results are given in Table 13. Perhaps the most obvious trend is the drop in recall on the training set between data conditions. Transposing the training data also improves generalization, as well as reducing the extent to which the network can overfit the training data. Transposing the input pitch spectra should have a negligible effect on the parameters of the convolutional layers, and this is confirmed by the results. All models in the second column, e.g. X-2, have smaller kernels, which leads to a much larger weight matrix in the first fully connected layer, and worse generalization in the non-transposed condition. It is reasonable to conclude that over-fitting mostly occurs in the final layers of the network, which do not take advantage of weight tying. Transposing the data results in an effect similar to that of weight tying, but because the sharing is not explicit the model must learn to encode this redundant information with more training data.

3.3 Qualitative Analysis

Having obtained promising quantitative results, the larger research objectives can now be addressed. As indicated by Table 13, transposing the data during training slightly improves generalization, but does more to limit the degree to which the models can overfit the training data. These two behaviors are not

Table 13

Performance as a function of model complexity, over a single fold.

Arch	As-Is			Transposed		
	Train	Valid	Test	Train	Valid	Test
S-1	84.7	74.9	75.6	79.5	75.9	76.8
M-1	85.5	75.0	75.5	80.6	75.6	77.0
L-1	92.0	75.2	75.5	81.7	76.5	77.7
S-2	87.0	73.1	74.5	78.4	75.5	76.2
M-2	91.2	73.9	74.0	79.4	75.4	76.6
L-2	91.7	73.6	73.8	81.6	76.3	77.4

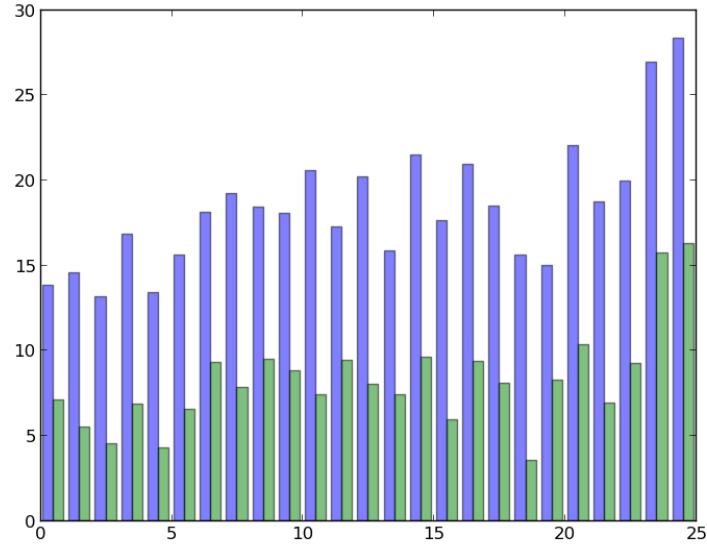


Figure 25: Accuracy differential between training and test as a function of chord class, ordered along the x-axis from most to least common in the dataset for As-Is (blue) and Transposed (green) conditions.

necessarily equivalent, and therefore whatever these networks learn as a result of data augmentation is preventing it from overfitting a considerable portion of the training set.

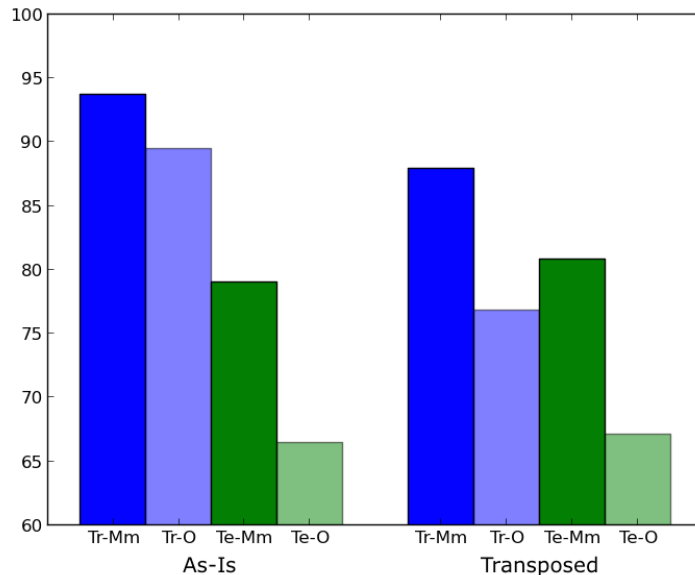


Figure 26: Effects of transposition on classification accuracy as a function explicitly labeled Major-Minor chords (dark bars), versus other chord types (lighter bars) that have been resolved to their nearest Major-Minor equivalent, for training (blue) and test (green) in As-Is (left) and Transposed (right) conditions.

One potential cause of over-fitting is due to an under-representation of some chord classes in the dataset. If this were the case, the most frequent classes should be unaffected by data augmentation, while less common classes would exhibit drastic swings in performance. Focusing here on Arch:L-1, Figure 25 shows the change in accuracy between data conditions for both training and test sets as a function of chord class, sorted by most to least common in the dataset. This plot indicates that, while transposing data during training reduces over-fitting, it does so uniformly across chord classes, on the order of about 10%. Therefore, all chord classes benefit equally from data augmentation, which is characteristic of intra-class variance more so than inadequate data for less common classes.

If this is indeed the case, there are likely two main sources of intra-class

variance: the practice of resolving all chord classes to Major-Minor, or error in the ground truth transcriptions. As a means to assess the former, Figure 26 plots the accuracy for chords that strictly labeled root-position Major-minor (Mm) versus all other (O) chords that are mapped into these classes in the train (Tr) and test (Te) conditions, with and without transposition. This is a far more informative figure, resulting in a few valuable insights. First, there is a moderate drop in performance over the training set for strictly Major-minor chords when data are transposed ($\approx -5\%$), but this causes a noticeable increase in generalization for strictly Major-minor chords in test set ($\approx +3\%$). Other chords, however, experience a significant decrease in performance within the training set ($\approx -11\%$) with transposition, but register a negligible improvement in the test set ($> 1\%$). One interpretation of this behavior is there is too much conceptual variation in the space of Other chords to meaningfully generalize to unseen data that is also not strictly Major-minor. This definition by exclusion gives rise to a class subset that is less populated than its strict counterpart, but will inherently contain a wider range of musical content. Though a sufficiently complex model may be able to overfit these datapoints in the absence of transposition, counting each observation toward every pitch class distributes the added variance across all classes evenly. This causes the model to ignore uncommon modes in the class distribution as noise, while reinforcing the strict Major-minor model in the process.

In addition to the effects of vocabulary resolution, there is also the consideration as to where estimation errors reside in the data. Due to the naturally repetitive nature of music, it is expected that problematic chords will often come from the same track. More importantly, it is because of this strong internal structure that these chords are likely problematic for similar reasons, and

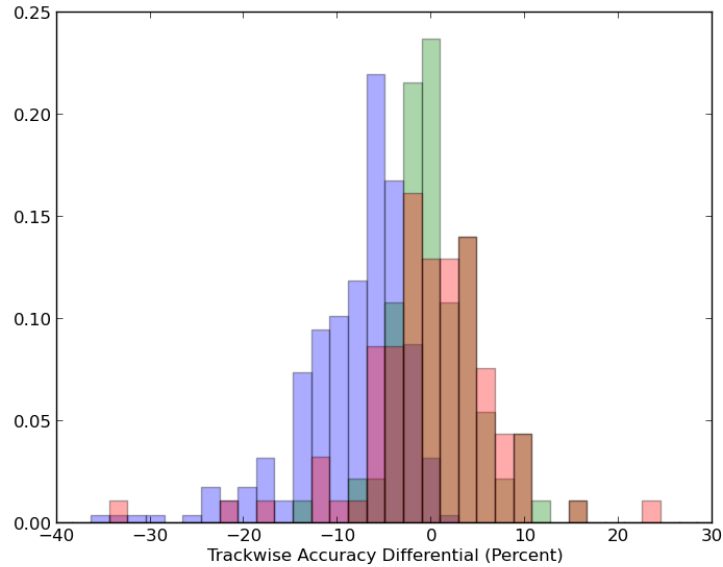


Figure 27: Histograms of track-wise recall differential between As-Is and Transposed data conditions, for training (blue), validation (red) and test (green) datasets.

in a manner that might not reveal itself when viewed independently. Therefore, tracks in the training set that exhibit significantly different performance between data conditions may help answer another question: why might transposition prevent the model from overfitting certain chords? To investigate this behavior, track-wise histograms of recall differential are computed with and without data transposition for the training, validation, and test splits, shown in Figure 27. Interestingly, performance over most tracks is unaffected or only slightly changed by the transposed data condition, as evidenced by the near-zero mode of the distributions. Some tracks in the training set, however, yield considerably worse results when the data is transposed. While this is consistent with intuition, , it indicates that error analysis is sufficiently motivated at

the track, rather than instance, level, and may offer insight into future areas of improvement.

One such problem track is “With or Without You” by U2. Here, the ground truth transcription consists primarily of four chords: D:maj, D:maj/5, D:maj6/6, and D:maj(4)/4. When resolved to the Major-minor vocabulary, the transcription is reduced entirely to D:maj. In the As-Is data condition, the model is able to call nearly the entire track D:maj; when training data are transposed, however, the model is unable to reproduce the ground truth transcription and instead tracks the bass motion, producing D:maj, A:maj, B:min, and G:maj, a very common harmonic progression in popular music. As far as quantitative evaluation is concerned, this second estimation exhibits a high degree of mismatch with the reference transcription, but is qualitatively reasonable and arguably far more useful to a musician. Importantly, this illustrates that the process of mapping chords to a reduced vocabulary can cause objective measures to deviate considerably from subjective experience, and thus confounding evaluation.

However, perhaps even more critically, the reliability of the reference annotation is somewhat dubious. Returning to the original song, one finds reasonably ambiguous harmonic content, consisting of a vocal melody, the moving bass line mentioned previously, a string pad sustaining a high-pitched D, and a moving guitar riff. Therefore, as a point of comparison, an Internet search yields six guitar chord transcriptions from the website Ultimate Guitar*. These alternative interpretations are consolidated in Table 14, alongside the reference, noting both the average and number of ratings, as well as the

*http://tabs.ultimate-guitar.com/u/u2/with_or_without_you_crd.htm, accessed 19 April 2015.

number of views the tab has received. Though the view count is not directly indicative of a transcription’s accuracy, it does provide a weak signal indicating that a large number of users did *not* rate it negatively. In considering this particular example, there are a handful of takeaways to note. First, all but the sixth of the user-generated chord transcriptions are equivalent by the conventional major-minor mapping rules, which is, interestingly enough, the same one produced by the model presented here. Second, this rather large community of musicians shows, at least for this song, a strong preference for root position chords. While it is difficult to determine why an annotator might choose one interpretation than another, it would appear general, root-position chords are preferred to nuanced chord spellings, e.g. `G:maj` over `D:maj(4)/4`. Finally, this raises questions surrounding the practice of using such precise chord labels for annotation. If nothing else, the flexibility afforded by this particular chord syntax allows annotators to effectively “build” their own chords through non-standard intervals or various bass intervals, amplifying the role subjectivity can play in transcription. This is not only problematic from a practical standpoint—are various annotators using this syntax consistently?—but atypical chord spellings are most likely to appear when the music content being described is especially ambiguous.

3.4 Conclusions

Following this initial inquiry, there are a few important conclusions to draw that should influence subsequent work. First and foremost, the common practice of major-minor chord resolution is responsible for a significant amount of error, both in training and test. While this approach simplifies the problem being addressed, it appears to introduce uninformative variation to classes

Table 14

Various real chord transcriptions for “With or Without You” by U2, comparing the reference annotation with six interpretations from a popular guitar tablature website; a raised asterisk indicates the transcription is given relative to a capo, and transposed to the actual key here.

Ver.	Chord Sequence				Score	Ratings	Views
Ref.	D:maj	D:maj/5	D:maj6/6	D:maj(4)/4	—	—	—
1	D:maj	A:maj	B:min	G:maj	4/5	193	1,985,878
2	D:5	A:sus4	B:min7	G:maj	5/5	11	184,611
3*	D:maj	A:maj	B:min	G:maj	4/5	23	188,152
4*	D:maj	A:maj	B:min	G:maj7	4/5	14	84,825
5*	D:maj	A:maj	B:min	G:maj	5/5	248	338,222
6	D:5	A:5	D:5/B	G:5	5/5	5	16,208

during training, and thus noise in the resulting evaluation. Therefore, for this reason alone, future work should consider larger vocabulary chord estimation, so that each chord class can be modeled explicitly. Additionally, an investigation into sources of error revealed that the performance for some tracks changes drastically between data conditions. Further exploration encouraged the notion that chord annotations with modified intervals or bass information may amplify the subjectivity of a transcription, and thus introduce noise in the reference chord annotations. Ignoring over-specified chord names would serve as an approach to data cleaning, maximizing confidence in the ground truth data and resulting in more stable evaluation.

4 Large Vocabulary Chord Estimation

Combining observations resulting from the previous study with other recent trends in ACE research, the focus now turns to the task of large vocabulary

ACE. There is a small body of research pertaining to vocabularies beyond the major-minor formulation, exploring different mixtures of chord classes and inversions (Mauch & Dixon, 2010b; Ni, McVicar, Santos-Rodriguez, & De Bie, 2012). However, for the same reasons discussed in 2.1, comparing new endeavors to these efforts is problematic due to differences in the vocabularies considered and data used. Perhaps the most advanced large-vocabulary ACE systems to date is the recent work of Cho (Cho, 2014). The design of this system is largely consistent with the previous overview of ACE research. A multiband chroma representation is computed from beat-synchronous audio analysis, producing four parallel chroma features. Each is modeled by a separate Gaussian Mixture Model, yielding four separate observation likelihoods as a function of time. These four posteriors are then decoded jointly, using a k-stream HMM, resulting in a time-aligned chord sequence. In addition to being one of the highest performing systems at the recent iteration of MIREX, a software implementation was obtained, thereby enabling direct comparisons with this work. It also considers the largest vocabulary, and presents an even greater challenge to the application of deep learning to ACE.

4.1 Data Considerations

Following the previous work of Cho (Cho, 2014), thirteen chord qualities, given in Table 9, in all twelve pitch classes and one no-chord class are considered here, for a total of 157 chord classes. Having all four datasets at hand, these collections are merged into the largest collection of chord transcriptions used to date, totaling 1235 tracks. Given that the collections were curated in isolation of each other, it is a necessary first step to identify and remove duplicates to avoid data contamination during cross validation. To these ends, each record-

ing is checked against the EchoNest Analyze API* and associated with its track and song identifiers, corresponding to the recording and work, respectively. Though multiple track IDs will map to the same song ID, uniqueness is defined at the level of a song to ensure duplicates are removed. This identifies 18 redundant songs, and all but one is dropped for each collision from the total collection, resulting in a final count of 1217 unique tracks.

Based on conclusions of the pilot study, the decision is made to ignore all chord labels that do not strictly match one of the considered qualities, i.e. chords that specify interval modifications, e.g. `A:min(*b3)`, or non-root inversions, e.g. `C:maj/5`. The motivation for doing so is two-fold. First, the increased number of chord qualities makes it difficult to map certain chords into one class or another, such as `D:sus4(b7)`, which sits halfway between a `D:sus4` and a `D:7`. Second, cleaning the reference data on this criteria can only improve annotation consistency; considering that the cumulative data is compiled from multiple sources and several dozen annotators over the course of a decade, it is quite unlikely that such nuanced conventions were used identically by all subjects involved. This ignored subset comprises only a small percentage of the overall data, and helps filter out suspicious chord spellings, such as `D:maj(1)/#1` or `A:maj(2,*3)/2`.

Unsurprisingly, as the data is collected from real music, the distribution of absolute chord classes is extremely imbalanced. In fact, some chord qualities do not occur in every root, and stratifying the data for training, validation, and testing only exacerbates the issue. Much previous work, including the previous discussion, has demonstrated that chord names can be rotated

*<http://developer.echonest.com/docs/v4>

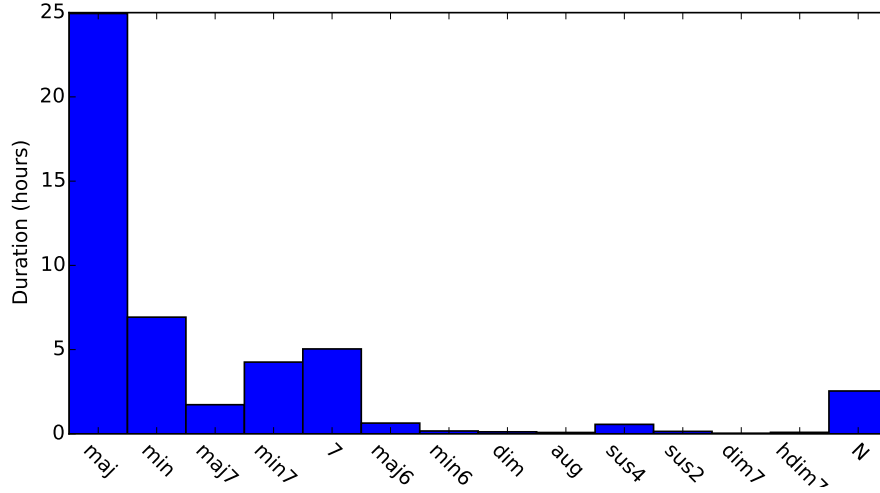


Figure 28: Histogram of chord qualities in the merged data collection.

to distribute instances across qualities, rather than absolute classes, motivating root-invariant analysis. A root-invariant histogram of the chord qualities contained in the merged dataset, given in Figure 28, clearly shows there is severe relative and absolute class imbalance. To the former, a stark division exists between the majority classes (`maj`, `min`, `maj7`, `min7`, `7`, and `N`), and the minority classes (`maj6`, `min6`, `dim`, `aug`, `sus4`, `sus2`, `dim7`, `hdim7`). The ratio, for example, between the most and least common qualities, `maj` and `dim7` respectively, is nearly three orders of magnitude (≈ 700). Arguably, the more challenging imbalance is an overall lack of data for some minority classes. Over all roots, the total duration of `dim7` is on the order of hundreds of seconds. Considering the repetitive structure of music, it is reasonable to assume that these few instances also occur in the same small number of tracks, limiting the variability of this data.

4.2 Experimental Setup

This work proceeds directly from the previous study, and takes a similar approach in many facets. There are a handful of important distinctions to make between the two, however, and these differences are detailed here.

4.2.1 Input Representation

A comparable constant-Q transform is applied to the audio at a framerate of 20Hz, without subsequent low-pass filtering or decimation, and time-frequency patches are formed from 20 frames, corresponding to 1 second. Whereas the previous study aimed to learn context directly, there is the inherent concern that a low input framerate will reduce the amount of data available for training beyond what is required by the model. In lieu of learning this context, standard post-filtering will be applied in the form of a uniform-transition HMM with a tunable self-transition penalty, consistent with (Cho, 2014); this is introduced in greater detail shortly, in ??.

Additionally, local contrast normalization is included as a standard pre-processing stage, with minor modifications. In previous work, a threshold is placed on the scaling term given by the average standard deviation over the entire input. While this may be suitable in the field of computer vision, where spatial dimensions are equivalent in 2-space, audio data behave differently in time and frequency. Namely, complex sounds often exhibit an overtone series, and energy becomes more densely concentrated in higher frequencies. This can have the undesirable effect of inadequately gaining regions that are more sparse. To correct for this behavior, the scaling coefficient is adjusted

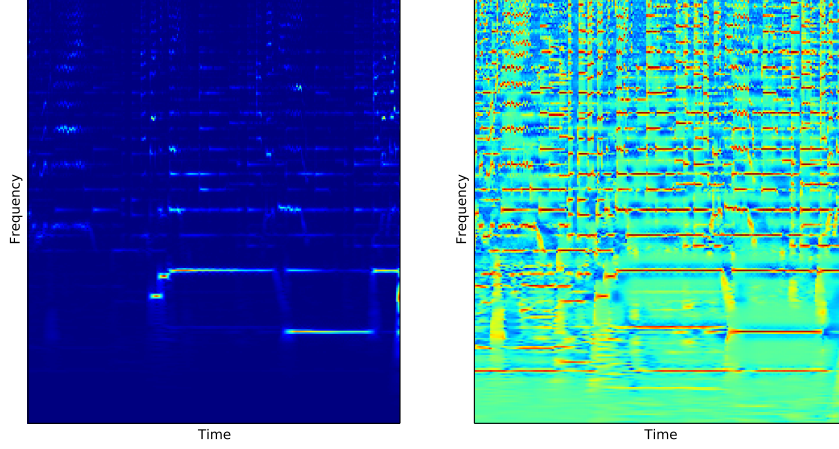


Figure 29: The visible effects of octave-dependent LCN, before (left) and after (right).

such that the frequency range considered is a piecewise function of frequency height, defined as follows:

$$\begin{aligned}
 X_{filt} &= (X \circledast W) \\
 V &= X_{filt} - X \\
 S_k &= V \circledast W_k \\
 S_k &= \max(S_k, \mu_{S_k}) \\
 Y &= \sum_{k=0}^{K-1} g_k * \frac{V}{S_k}
 \end{aligned}$$

To illustrate the benefit of this piecewise combination, an example track is given in Figure 29, both with and without the octave-dependent modification.

4.2.2 Designing a Root-Invariant Classifier

One of the key findings from the previous study of deep networks for ACE, consistent with previous research, is the importance of enforcing or encouraging root-invariance in the model. With GMMs, this is typically achieved by rotating all data, i.e. chroma features, to the same root and fitting a model for each quality. Then, when applying the model, likelihoods are estimated for each root by circularly rotating chroma through all twelve positions to recover the full space of chord classes. In the pilot study on chord estimation, this concept was mimicked by rotating the data in the input domain. While this data augmentation helped produce better results, it is a somewhat inelegant approach to realize pitch-invariance. During training, the model must learn the same representation for all 12 pitch classes in order to represent each quality in every root. Not only does this require more parameters to capture this redundant information, but it will likely require more training iterations for the model to do so.

Alternatively, pitch invariance can be built directly into the model by tying the weights of the classifier for different chord qualities across the twelve roots. This is realized here by defining a four layer network composed entirely of convolution operations, diagrammed in Figure 30. As convolutional networks amply demonstrated, weight tying is an effective way to achieve translation invariance with fewer parameters and less training data. This is achieved as follows: the penultimate representation is designed to yield a matrix with shape $(12 \times N)$, corresponding to the number of pitch classes and the chosen output dimensionality of the second to last layer, respectively; the chord classifier is then applied by taking the inner product with a weight ma-

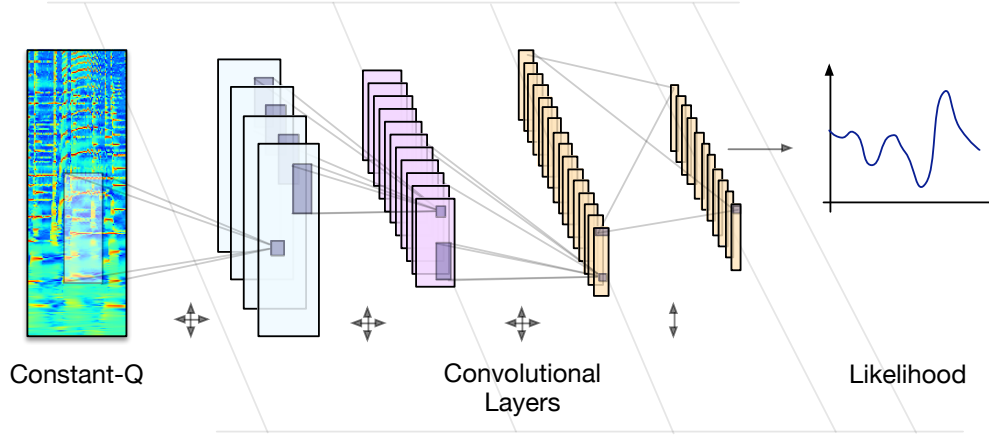


Figure 30: A Fully Convolutional Chord Estimation Architecture.

trix with shape $(N \times 13)$, corresponding to the dimensionality of the previous layer and the number of chord qualities, respectively. For ease and efficiency this is implemented as a convolution, but the result is equivalent. This produces a (12×13) matrix, which is flattened to represent the 13 qualities in all possible roots.

The no-chord class is not captured by this operation, however, and a separate fully connected layer is applied in parallel to the flattened penultimate representation to estimate this class independently. This one-dimensional no-chord estimator is then concatenated with the flattened chord estimator, and this combined representation of 157 classes is normalized by the softmax operation to yield the probability mass function over all classes.

4.2.3 Convolutional Dropout

Here, the principles of dropout, discussed in Chapter III, are extended to 3D convolutions. In the weight-matrix case, training with dropout effectively

ignores activations of an transformed output, setting them to zero. Considering each output coefficient as a measure of activation for a given “feature”, the act of dropout can be interpreted as sub-sampling the feature extractors learned by the model.

By extension, the same principle could be applied to convolutional layers. In the absence of any known prior effort to do so, this is achieved here by dropping out a full 3D kernel. Expressed formally, the i^{th} kernel, W_i , can be masked with probability p , resulting in the possibly empty feature map, Z_i :

$$Z_i = \text{binomial}(p) * h(X \otimes W_i + b_i) / (1.0 - p) \quad (25)$$

where the output is also scaled by the complement of the probability. Here, it is expected that each kernel learns a collection of feature extractors that, on average, work well together. In the language of co-adaptation, the tensor can be seen as a “team” of feature detectors, and as such correlations are broken up at this mid, as opposed to global, level.

There are two small implementation details worth noting. First, the same parameters in the model are dropped out over the entire batch, and not separately for each datapoint in the batch. In the model averaging interpretation of dropout, this is analogous to updating one possible model at each update step, and offers interesting parallels to coordinate block descent. Additionally, the original proposal of dropout suggests that the activations of all outputs be halved when using the full model at test time. This is somewhat cumbersome in practice, and, as indicated in Eq. (25), scale *up* the parameters during training by the complement of the dropout ratio, allowing models in test to be agnostic of this process.

Table 15

Parameter shapes in the three model complexities considered.

layer	L	XL	XXL	pooling
0	(16, 1, 5, 13)	(20, 1, 5, 13)	(24, 1, 5, 13)	(2, 3)
1	(32, 16, 5, 37)	(40, 20, 5, 37)	(48, 24, 5, 37)	(2, 1)
2	(64, 32, 1, 33)	(80, 40, 1, 33)	(96, 48, 1, 33)	(2, 1)
3.a	(13, 64, 1, 1)	(13, 80, 1, 1)	(13, 96, 1, 1)	–
3.b	(768, 1)	(960, 1)	(1152, 1)	–

4.2.4 Architectural Considerations

Given the theoretical relationship between dropout and model averaging, it is reasonable to expect that larger dropout ratios will necessitate larger architectures. Therefore, a variety of model complexities are explored by changing the number of kernels in each layer; the primary weight shapes of the networks are given in Table 15:

The first four layers are 3D-convolutions, and thus the weights are 4 dimensional, corresponding to the number of kernels, the number of input feature maps, filter size in the time dimension, and filter size in the frequency dimension, respectively. The first three of these layers make use of max-pooling in time by a factor of 2, but only the first also performs max-pooling in frequency, by a factor of 3. As before, downsampling in frequency is performed in the spirit of learning slight tuning invariance, consistent with 12-TET. The final layer, 3.b, corresponds to the fully-connected no-chord regressor, and the shape of this weight matrix is given as the input and output dimensionality, respectively.

4.2.5 Correcting Class Imbalance with Scaled Likelihood Estimation

Having defined the output of the model as a probability function, it is straightforward to again optimize the parameters to the negative log-likelihood over the dataset. However, there are two difficulties that prohibit the uniform presentation of classes, as in previous work. First, due to the increased number of classes, mini-batches would either need to consist of many datapoints, increasing the processing time of each batch, or the learning rate would need to be smaller or diminishing over time to prevent oscillation, increasing the number of iterations necessary to converge. To the latter point, it is easy to imagine scenarios where gradient descent wobbles back and forth between updates, as each batch pulls the parameters in a slightly different direction. The other challenge this raises is a result of the considerable class imbalance, where uniform presentation would both spend too much time on minority classes and inhibit the speed at which the model could learn the full extent of the majority classes.

The solution to this problem is found in Bayes' theorem, where the network is understood as yielding a class posterior probability, $P(Y|x)$, for the chord class, Y , given the observation, x :

$$P(x|Y) = \frac{P(Y|x)P(x)}{P(Y)} \quad (26)$$

The observation likelihood, $P(x|Y)$, is then a function of the posterior, the class prior, $P(Y)$, and probability of the observation itself, $P(x)$. This final quantity is independent and thus can be ignored:

$$P(x|Y) \propto \frac{P(Y|x)}{P(Y)} \quad (27)$$

While the deep network is trained to produce the class posterior, the class prior can be measured empirically over the training set, and divided out after the fact. Referred to as *scaled likelihood estimation*, this strategy has proven effective at reducing the effects of class imbalances in the functionally related domain of automatic speech recognition (Dahl, Yu, Deng, & Acero, 2012). Notably, scaled likelihood estimation is particularly attractive here because it scales well with the number of estimated classes.

As a final comment, a possible pitfall when applying likelihood scaling is a matter of numerical stability arising from the least represented classes, or classes that might not even occur in the training set. Whereas this can be mitigated with pseudo-counting—setting a heuristically determined, non-zero lower bound—the class prior here is computed by counting each observation toward the same quality in all roots. Though this prior could be seen as a coefficient vector to optimize in a more direct way, this approach works reasonably well in practice.

4.2.6 Training and Early Stopping

In contrast to the previous study, which converged to a stable result in a few thousand iterations, it takes considerably more effort to train models for this task, on the order of hundreds of thousands of iterations. Given the lengthy run time, parameters are saved every 1k iterations during training, and frame the problem of early stopping as a brute-force search over a finite set of parameter configurations. Due to the application of Viterbi and the coupling with the self transition penalty, validation is non-trivial and computationally expensive. Therefore, exhaustive validation is performed every 10k iterations, starting at 5k. The best model and self-transition penalty are chosen by finding the

Table 16

Weighted recall across metrics over the training data.

Model		triads	root	MIREX	tetrads	sevenths	thirds	majmin
Cho, 2014		0.8053	0.8529	0.8205	0.6763	0.6823	0.8261	0.8109
L	0.0	0.9087	0.9249	0.9140	0.8600	0.8601	0.9177	0.9097
	0.125	0.8706	0.9018	0.8785	0.7798	0.7792	0.8895	0.8718
	0.25	0.8382	0.8811	0.8490	0.7266	0.7277	0.8637	0.8405
	0.5	0.7916	0.8491	0.8079	0.6577	0.6641	0.8262	0.7970
XL	0.0	0.9236	0.9353	0.9277	0.8903	0.8906	0.9300	0.9244
	0.125	0.8899	0.9145	0.8962	0.8217	0.8214	0.9049	0.8908
	0.25	0.8504	0.8888	0.8610	0.7374	0.7385	0.8734	0.8527
	0.5	0.7972	0.8541	0.8118	0.6632	0.6683	0.8329	0.8014
XXL	0.0	0.9462	0.9528	0.9487	0.9297	0.9300	0.9498	0.9466
	0.125	0.8994	0.9209	0.9048	0.8386	0.8374	0.9133	0.8997
	0.25	0.8701	0.9041	0.8777	0.7833	0.7828	0.8921	0.8710
	0.5	0.8043	0.8573	0.8184	0.6783	0.6820	0.8374	0.8080

configuration with the highest harmonic mean over all evaluation metrics given in Section 2.3.

4.3 Experimental Results

Following from the setup defined above, the three model complexities —X, XL, and XXL— are trained with four dropout values, $p_{dropout} \in \{0.0, 0.125, 0.25, 0.5\}$ across five folds of the data. Note that when $p_{dropout} = 0.0$, this is equivalent to training a model without dropout. Additionally, the system presented in (Cho, 2014), referred to here simply as “Cho”, is trained on identical partitions of the data and evaluated alongside the deep networks. Weighted recall for the various comparison functions, averaged across folds, is given in Tables 16 and 17 for the training and test splits, respectively.

There are several observations that may be drawn from these two tables. First, in the absence of dropout, the deep network models considered here

Table 17

Weighted recall across metrics over the test (holdout) data.

Model		triads	root	MIREX	tetrads	sevenths	thirds	majmin
Cho, 2014		0.7970	0.8475	0.8147	0.6592	0.6704	0.8197	0.8057
L	0.0	0.7939	0.8442	0.8102	0.6583	0.6725	0.8135	0.8041
	0.125	0.7951	0.8465	0.8109	0.6516	0.6616	0.8203	0.8028
	0.25	0.7882	0.8445	0.8039	0.6509	0.6592	0.8175	0.7950
	0.5	0.7762	0.8372	0.7936	0.6358	0.6442	0.8115	0.7832
XL	0.0	0.7939	0.8432	0.8098	0.6589	0.6736	0.8122	0.8042
	0.125	0.7995	0.8493	0.8145	0.6673	0.6788	0.8227	0.8077
	0.25	0.7950	0.8479	0.8114	0.6493	0.6580	0.8215	0.8023
	0.5	0.7773	0.8401	0.7940	0.6351	0.6430	0.8147	0.7836
XXL	0.0	0.7969	0.8463	0.8130	0.6583	0.6741	0.8136	0.8080
	0.125	0.7993	0.8477	0.8140	0.6633	0.6745	0.8215	0.8075
	0.25	0.7947	0.8497	0.8092	0.6592	0.6686	0.8241	0.8020
	0.5	0.7768	0.8369	0.7941	0.6392	0.6468	0.8121	0.7830

are able to overfit the training data. This is an important finding in so far as making sure that the fully-convolutional architecture is not overly constrained, and indicates that the XXL model is a reasonable upper bound on complexity. The effect of dropout on performance over the training set is significant, as it reduces overfitting consistent with increased values.

Shifting focus to performance on the test set, it is obvious that these differences in training set performance have little impact on generalization, and all models appear to be roughly equivalent. A small amount of dropout – 0.125 or 0.25– has a slight positive effect on generalization; too much dropout, on the other hand, seems to have a negative effect on performance. There are two possible explanations for this behavior: one, a high degree of convolutional dropout is more destabilizing than in the fully-connected setting; and two, these models were not finished learning, and stopped prematurely.

Overall, the best deep networks appear to be essentially equivalent to

Table 18

Quality-wise recall statistics for train and test partitions, averaged over folds.

train	0.0	0.125	0.25	0.5
L	0.8858	0.8263	0.7403	0.5838
XL	0.9049	0.8569	0.7652	0.6147
XXL	0.9421	0.8838	0.8232	0.6459
test	0.0	0.125	0.25	0.5
L	0.4306	0.5029	0.5240	0.5135
XL	0.4174	0.4887	0.5281	0.5253
XXL	0.3935	0.4825	0.5127	0.5257

the state of the art comparison system, referred to henceforth as “Cho”; XL-0.125 just barely eclipses Cho in every metric but “MIREX”, while XL-0.25 is right on its heels. The different metrics indicate that confusions at the strict level are predominantly musically related, i.e. descending in order from root, thirds, triads, sevenths, tetrads. Interestingly, the performance gap between the “root” and “triads” scores is quite small, $\approx 5\%$, while the gap between “root” and “tetrads” is nearly 20%, for all models considered. One way to interpret this result is that these systems are quite robust in the estimation of three-note chords, but struggle to match the way in which reference annotators use sevenths.

The results for quality-wise recall are given in Table 18. While dropout is again able to considerably reduce over-fitting in the training set, it appears to have a more profound effect here towards generalization. Whereas before a 0.5 dropout ratio seemed to result in the “worst” deep networks, here it leads to the best generalization across all chord qualities. Furthermore, the best

Table 19

Individual chord quality accuracies for the XL-model over test data, averaged across all folds.

	support (min)	0.0	0.125	0.25	0.5	Cho
C:maj	397.4887	0.7669	0.7390	0.6776	0.6645	0.7196
C:min	105.7641	0.5868	0.6105	0.6085	0.6001	0.6467
C:7	68.1321	0.4315	0.5183	0.5783	0.5362	0.5959
C:min7	63.9526	0.4840	0.5263	0.5954	0.5593	0.5381
N	41.6994	0.7408	0.7679	0.7875	0.7772	0.5877
C:maj7	23.3095	0.5802	0.6780	0.7268	0.7410	0.6587
C:sus4	8.3140	0.2380	0.3369	0.3811	0.4231	0.3894
C:maj6	7.6729	0.1929	0.2908	0.3847	0.3540	0.3028
C:sus2	2.4250	0.1921	0.3216	0.3698	0.3995	0.1993
C:dim	1.8756	0.4167	0.4105	0.4140	0.3955	0.5150
C:min6	1.5716	0.2552	0.3870	0.4505	0.5076	0.3129
C:aug	1.2705	0.3730	0.5078	0.5346	0.5521	0.3752
C:hdim7	1.1506	0.3840	0.5688	0.6659	0.6140	0.4593
C:dim7	0.5650	0.2012	0.1790	0.2186	0.2296	0.0643
total	–	0.4174	0.4887	0.5281	0.5253	0.4546

performing models, according to weighted recall, are not the best performing models in this table. Thus these results allude to the notion that overall weighted recall may have an inverse relationship with quality-wise recall.

To further assess this claim, the individual chord quality accuracies are broken out by class for XL-0.125, and compared alongside Cho, given in Table 19. Immediately obvious is the influence of sharp distribution effects in generalization. Performance for the majority chord qualities, in the upper half of the table, is noticeably higher than the minority classes, in the lower half. The one exception is that of dominant 7 chords, which seems relatively low, especially compared to Cho; this is likely a result of V vs V7 confusions, but the annotations do not easily provide this functional information to validate

the hypothesis*. XXL-0.25 yields near identical weighted recall statistics to Cho, but achieves a significant increase in quality-wise recall, R_Q , 0.5127 to 0.4546 ($\delta 0.0581$).

Notably, the only chord quality to decrease in accuracy with dropout is major, indicating that, with the addition of dropout, the decision boundary between major and its related classes, e.g. major 7 and dominant 7, shift. However, because of the significant imbalance in the support of each quality, given here in minutes, a small drop in accuracy for major yields a considerable drop in weighted recall overall. To illustrate, between the 0.125 and 0.25 dropout ratios, major accuracy drops 6%, while dominant 7 and major 7 accuracy increase 6% and 5%, respectively. In terms of overall time though, this comes at the expense of 24 minutes of major now being classified “incorrectly”, compared to a combined 5 minutes of dominant and major 7’s now being “correct”. Therefore, it seems there is a trade-off between these two measures, and thus the representational power of the model does not really change. Rather, the decision boundaries between overlapping classes must prefer one over the other, and thus model selection may ultimately be a function of use-case. For example, is it better to have a model that predicts simpler chords, i.e. major, most of the time? Or to have a model that makes use of a wider vocabulary of chords? Lastly, it is necessary to recognize that, while quality-wise recall provides a glimpse into more nuanced system behavior, the severe class imbalance results in a rather volatile metric.

As a final investigation of algorithm performance with this particular collection of data, having the estimations of two very different computational

*The Billboard dataset *does* provide tonic information, and thus this relationship could be recovered from the data; however, it is left here as a fertile area for future work

Table 20

Weighted recall scores for the two algorithms scored against each other, and the better match of either algorithm against the reference.

	triads	root	MIREX	tetrads	sevenths	thirds	majmin
XL-0.25 vs Cho	0.7835	0.8406	0.8044	0.6769	0.7072	0.8148	0.8095
Cho vs XL-0.25	0.7835	0.8406	0.8044	0.6770	0.6982	0.8148	0.8035
Ref. vs max(XL-0.25 Cho)	0.8243	0.8705	0.8402	0.7037	0.7157	0.8452	0.8331

systems affords the opportunity to explore where they do or do not agree. First, the two algorithms are scored against each other by using one as the reference and the other as the estimation. Then, both algorithms are evaluated against the human reference, keeping the maximum of the two scores for each track. This second condition is similar in concept to model averaging, and serves to further highlight differences between estimations. The results of each given in Table 20.

It is interesting to consider that, despite nearly equivalent performance on the metrics reported above, the algorithms match each other about as well as either matches the reference annotations. If the systems made same mistakes, there would be better agreement between their respective outputs. Since this isn't the case, it is safe to conclude that the errors made by one system are sufficiently different than those of the other. This is a particularly valuable discovery, as these systems offer two sufficiently distinct, automated perspectives and can be leveraged for further analysis. Additionally, combining the systems' estimations shows that there are some instances where one model outperforms the other, encouraging the exploration of ensemble methods in future work.

Expanding on this analysis, it is of considerable interest to compare scores between algorithms versus the best match with the reference on a track-

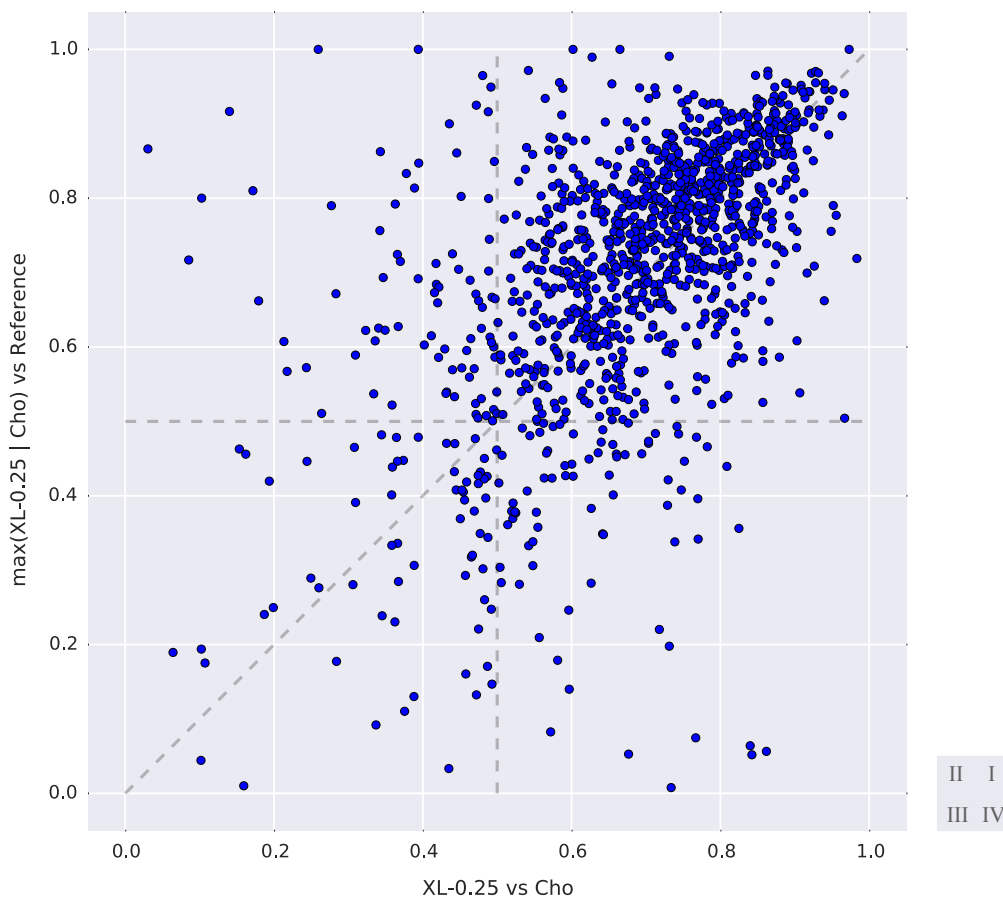


Figure 31: Track-wise agreement between algorithms versus the best match between either algorithm and the ground truth data.

wise basis; this is given in Figure 31 for the “tetrads” metric. Here, each track is represented as a point in coordinate space, with algorithmic agreement along the x -axis and best agreement with the ground truth annotations along the y -axis. For clarity, this scatter plot can be understood in the following way: the line $x = y$ corresponds to an equal level of agreement between all three chord transcriptions; bisecting the graph horizontally and vertically yields four quadrants, enumerated I-IV in a counterclockwise manner, starting from the

upper right. Tracks that fall in each quadrant correspond to a different kind of behavior. Points in Quadrant I indicate that both estimations and the reference have a high level of agreement ($x > 0.5$, $y > 0.5$). Quadrant II contains tracks where the algorithms disagree significantly ($x < 0.5$), but one estimation matches the reference well ($y > 0.5$). Tracks in Quadrant III correspond to the condition that no transcription agrees with another, ($x < 0.5$, $y < 0.5$), and are particularly curious. Finally, Quadrant IV contains tracks where the algorithms estimate the same chords ($x > 0.5$), but the reference disagrees with them both ($y < 0.5$).

With this in mind, three-part annotations can be examined for a point in each quadrant, consisting of a reference (Ref), an estimation from the best deep neural network (XL-0.25), and an estimation from the baseline system (Cho). In all following examples, chords are shown as color bars changing over time, from left to right; as a convention, the pitch class of the chord’s root is mapped to color hue, and the darkness is a function of chord quality, e.g. all **E:*** chords are a shade of green. No-chords are always black, and chords that do not fit into one of the 157 chord classes are shown in gray.

A track from Quadrant I is given in Figure 32. As to be expected, the reference and both estimated chord sequences look quite similar. The most notable discrepancy between the human-provided transcription and the two from the automatic methods is at the end of the sequence. While the reference annotation claims that the transcription ends on a **G:maj**, both algorithms estimate the final chord as a **C:maj**. This occurs because the song is in the key of G major, and thus the human chooses to end the song on the tonic chord. However, the song —“Against the Wind” by Bob Seger— is recorded with a fade-out, and the last audible part of the recording is in fact the **C:maj**, when

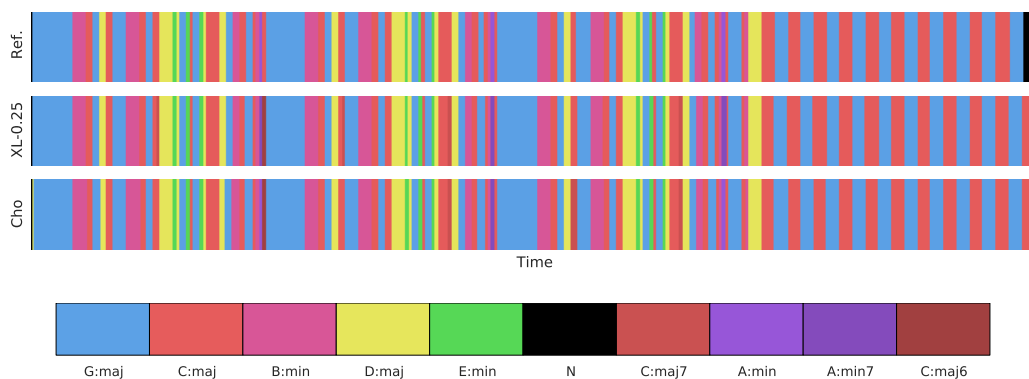


Figure 32: Reference and estimated chord sequences for a track in Quadrant I, where both algorithms agree with the reference.

playback volume is adjusted accordingly. Therefore, this is an instance of the automatic systems being more precise than the human annotator.

Next, a track from Quadrant II —“Smoking Gun” by Robert Cray— is considered in Figure 33. While the baseline system, Cho, agrees strongly with the reference that the predominant chord is an $E:min7$, the deep network, XL-0.25, prefers $E:min$, and produces a poor “tetrads” score as a result. This confusion is an understandable one, and highlights an interesting issue in automatic chord estimation evaluation. Depending on the instance, it could be argued that some chords are not fundamentally different classes, and thus “confusions” in the traditional machine learning sense, but rather a subset of a more specified chord, e.g. $E:min7 \supset E:min$. Traditional 1-of-k classifier schemes fail to encode this nuance, whereas a hierarchical representation would better capture this relationship.

The track drawn from Quadrant III —“Nowhere to Run” by Martha Reeves and the Vandellas— is shown in Figure 34, and especially interesting for three reasons. First, most of the considerable disagreement between the reference and both estimated chord sequences can be explained by a tuning

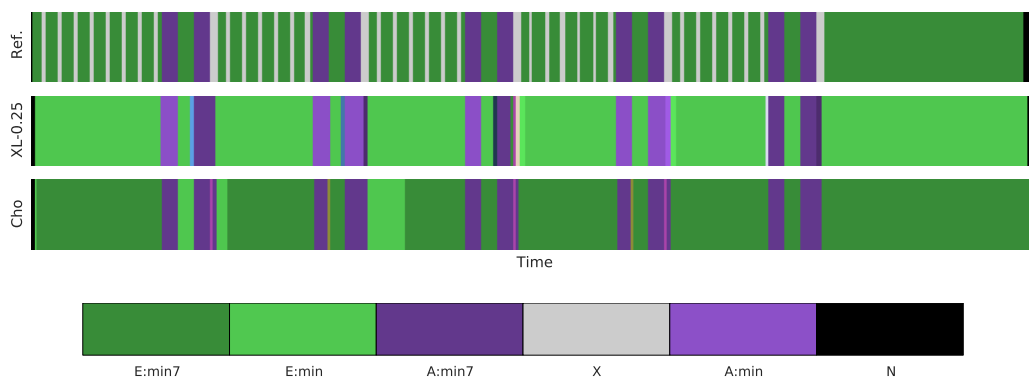


Figure 33: Reference and estimated chord sequences for a track in Quadrant II, the condition where algorithms disagree sharply, but one agrees strongly with the reference.

issue. The automatic systems predict the tonic as **G**, while the human reference is based on **Ab**. Matching a pure tone to this recording places the tonic around 400 Hz; for reference to absolute pitch, the notes **G3** and **Ab3** correspond to roughly 391 Hz and 415 Hz, respectively. Therefore, the human annotator and automatic systems disagree on how this non-standard tuning should be quantized. Second, the two automatic systems again differ on whether to label the chord a **ma**j or 7 chord; this time, however, the reference annotation prefers the triad. Lastly, there are three instrumental “breaks” in the piece where the backing band drops out to solo voice and drumset. While the reference annotation marks the first occurrence in the song with an **X** chord label, the other two are not marked similarly despite sounding nearly identical. The deep network model labels all three of these instances as “no-chord,” shown as black regions in the middle track. This raises interesting questions regarding, among general annotator guidance, aspects of temporal continuity in a transcription. How literal should an annotator be when marking a silence as “no-chord”? Is there a duration at which a gap becomes a pause? And, of significant

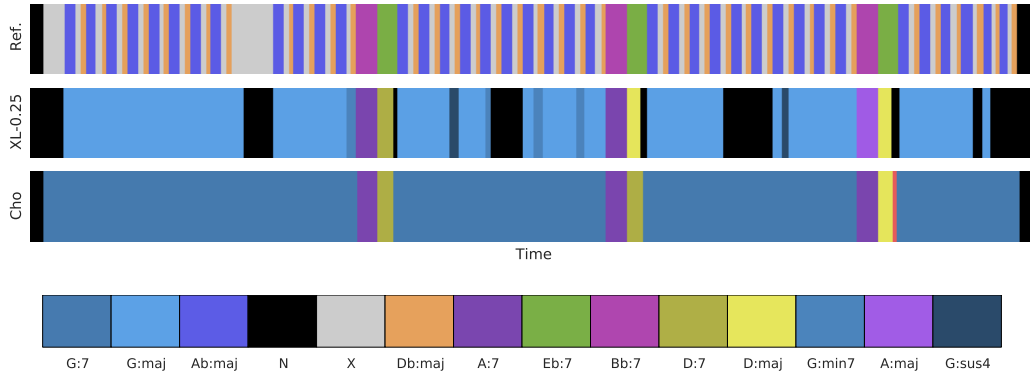


Figure 34: Reference and estimated chord sequences for a track in Quadrant III, the condition where neither algorithm agrees with the reference, nor each other.

importance when merging various datasets, are different annotators applying the same rules and decision criteria? Taken together, this example serves to illustrate how fragile the notion of “ground truth” can be, due to practical issues of calibration, annotator consistency, and the instructions given during the annotation process.

As a final example from this analysis of two-part estimations, a track is considered from Quadrant IV, shown in Figure 35, corresponding to “Some Like It Hot” by The Power Station. Evidenced by the large regions of estimated no-chord, both automatic systems struggle on this particular track. Listening through, there are likely two contributing factors. First, the song is very percussive, and heavily compressed wideband noise probably disrupts the harmonic estimates made by the models. Second, the song makes very little use of true pitch simultaneities, and much of the harmony in this song is implied. The song is also sparsely populated from an instrumental perspective, resulting in erroneous estimations. While both systems appear to fall victim to this kind of content, the example speaks to the limitations of the task as

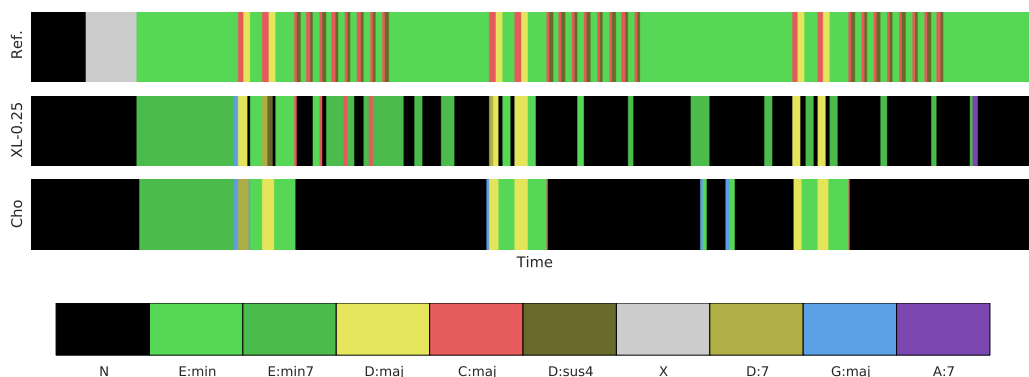


Figure 35: Reference and estimated chord sequences for a track in Quadrant IV, the condition where both algorithms agree with each other, but neither agrees with the reference.

it is currently defined, as well as the importance of ensuring that the content included in a dataset is relevant to the task at hand.

4.4 Rock Corpus Analysis

Much can and has been said about the consistency, and thus quality, of the reference annotations used for development and evaluation of chord estimation systems. The majority of human-provided chord annotations are often singular, either being performed by one person or as the result of a review process to resolve disagreements. The idea of examining, rather than resolving, annotator disagreements is an interesting one, because there are two reasons why such discrepancies might occur. The first is simply a matter of human error, resulting from typographical errors and other similar oversights. The second, and far more interesting cause, is that there is indeed some room for interpretation in the musical content, leading to different acceptable annotations. Most chord annotation curation efforts have made an explicit effort to resolve

all discrepancies to a canonical transcription, however, and it is not possible to explore any such instances in the data used so far.

Fortunately, the *Rock Corpus* dataset, first introduced in (De Clercq & Temperley, 2011), is a set of 200 popular rock tracks with time-aligned chord and melody transcriptions performed by two expert musicians: one, a pianist, and the other, a guitarist. This insight into musical expertise adds an interesting dimension to the inquiry when attempting to understand alternate interpretations by the annotators. This collection of chord transcriptions has seen little use in the ACE literature, as its initial release lacked timing data for the transcriptions, and the chord names are provided in a Roman Numeral syntax. A subsequent release fixed the former issue, however, in addition to doubling the size of the collection. The latter issue is more a matter of convenience, as key information is provided with the transcriptions and this format can be translated to absolute chord names, consistent with the syntax in Section 1.3. This dataset provides a previously uncaptialized opportunity to explore the behavior of ACE systems as a function of multiple reference transcriptions.

As an initial step, the two annotators, referred to here as DT and TdC, are each used as a reference and estimation perspective in order to quantify the agreement between them. The results, given in Table 21, indicate a high, but imperfect level of consistency between the two human perspectives. Following earlier trends, this is also a function of the chord spelling complexity, whereby equivalence at the “root” is much higher than for “tetrads”. Additionally, it is worth noting the asymmetry in chord comparisons. Thus, depending on the perspective used as the reference, a estimation may match better or worse with it. Finally, it is curious that the “MIREX” score is not perfect, a measure

Table 21

Weighted recall scores for the two references against each other, each as the reference against a deep network, and either against the deep network.

	triads	root	MIREX	tetrads	sevenths	thirds	majmin
DT vs TdC	0.8986	0.9329	0.9180	0.8355	0.8380	0.9042	0.9008
TdC vs DT	0.9117	0.9465	0.9168	0.8477	0.8537	0.9174	0.9176
DT vs XL-0.25	0.7051	0.7816	0.7180	0.5625	0.5653	0.7314	0.7084
TdC vs XL-0.25	0.7182	0.7939	0.7314	0.5786	0.5822	0.7444	0.7228
max(DT TdC) vs XL-0.25	0.7306	0.8010	0.7431	0.5998	0.6032	0.7569	0.7348

that focuses on pitch composition rather than contextual spelling. One would assume that the difficulty in naming a chord is more a function of the latter than the former, but this proves not to be the case.

Continuing, the deep network used in the previous analysis, XL-0.125, is again considered here. Adopting a similar methodology as before, the estimations of the deep network are compared against both references separately, as well as against the references together and taking the better match. Overall performance is generally worse than that observed over the holdout data in the previous investigation. This is predominantly caused by a mismatch in the chord distributions between the datasets, as the RockCorpus only contains half the chords known to the automatic system. Scores at the comparison of the “root” are much closer than that of the “tetrads” level, for example. Comparing the algorithmic estimations against the intersection of the human references results in a small performance boost across all scores, indicating that the machine’s “errors” are musically plausible, and might be validated by an additional perspective.

Keeping with previous methodology, annotator agreement is compared to the better match between the estimation and either reference on a track-wise basis; this is given in Figure 36 for the “tetrads” metric. Similar to

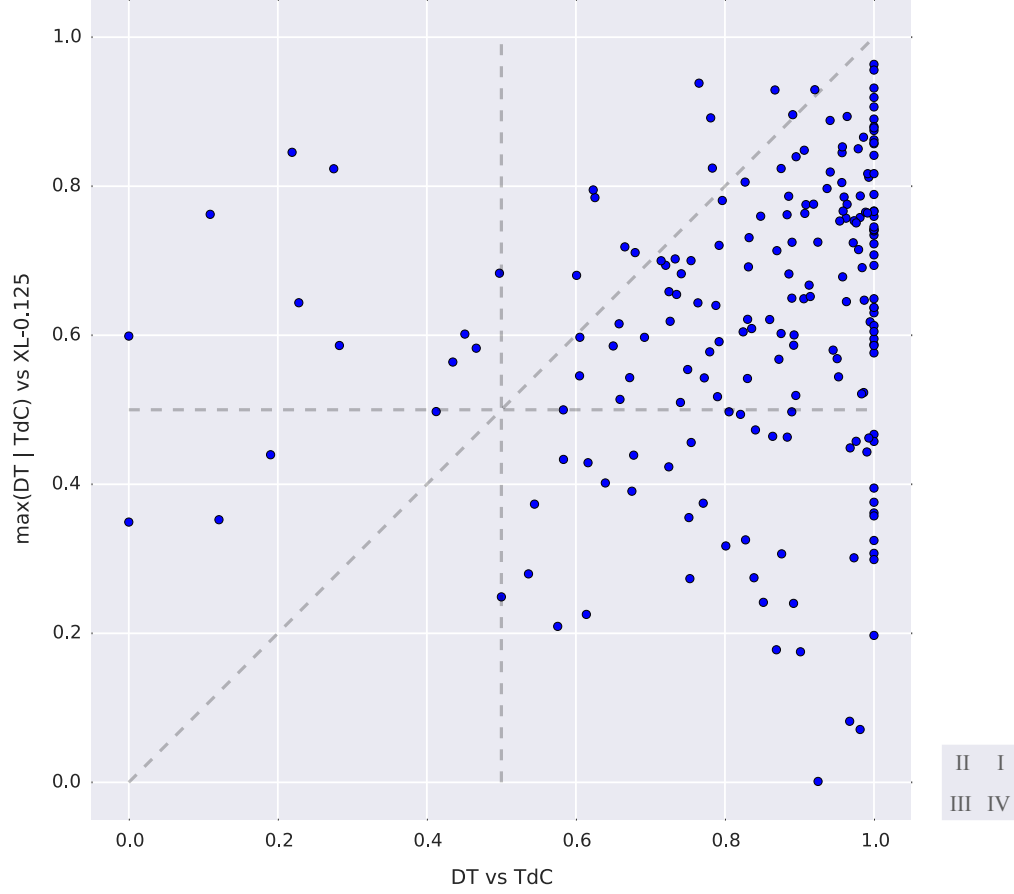


Figure 36: Track-wise agreement between annotators versus the best match between either annotator and the best performing deep network.

Figure 31, the former is shown along the x -axis, and the latter along the y -axis. This scatter plot can be understood similarly to before, with a few slight differences. As before, Quadrant I contains tracks where all transcriptions agree estimation ($x > 0.5, y > 0.5$), and Quadrant III, where all transcriptions disagree ($x < 0.5, y < 0.5$). Quadrant II now contains tracks where the *annotators* disagree significantly ($x < 0.5$), but one annotator matches the

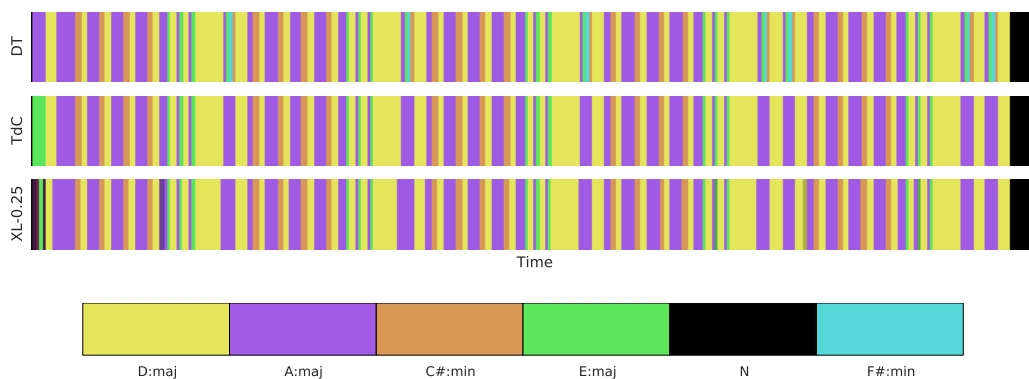


Figure 37: Reference and estimated chord sequences for a track in Quadrant I, where the algorithm agrees with both annotators.

reference well ($y > 0.5$), and Quadrant IV contains tracks where the annotators agree ($x > 0.5$), but the estimation disagrees with them both ($y < 0.5$).

Figure 37 shows the three-part transcription for “The Weight” by The Band, a track in Quadrant I. Again, the three chord sequences score well against each other, and offer a high degree of visual similarity. Notably though, the algorithmic estimation independently agrees more with the interpretation of TdC than DT. For example, there are slight discrepancies in root movement at the end of the sequence, where DT expands the A:maj of TdC and XL-0.25 into A:maj-F#:min-C#:min, or relatively in Roman numeral form, I-vi-iii. This motion occurs multiple times in the song, and each annotator’s interpretation is internally consistent.

Being that the human annotators tend to agree more with each other than the two automatic systems previously considered, fewer tracks fall in the left half of the scatter plot in Figure 36 than Figure 31. One of these from Quadrant II, “All Apologies” by Nirvana, is shown in Figure 38. Here, the human annotators have disagreed on the harmonic spelling of the verse, with DT and TdC reporting C#:maj and C#:7, respectively. On closer inspection,

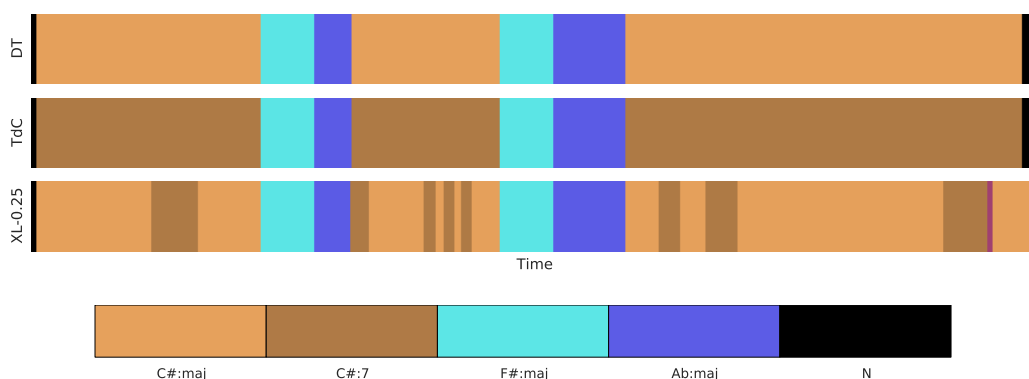


Figure 38: Reference and estimated chord sequences for a track in Quadrant II, the condition where the annotators disagree sharply, but one agrees strongly with the algorithm.

it would appear that both annotators are in some sense correct; the majority of the verse is arguably **C#:maj**, but a cello sustains the flat-7th of this key intermittently. These regions that this occurs are clearly captured in the XL-0.25 annotation, corresponding to its **C#:7** predictions. This proves to be an interesting discrepancy, because one annotator (DT) is using long-term structural information about the song to apply a single chord to the entire verse.

Another of these select few, this time from Quadrant III, is “Papa’s Got a Brand New Bag” by James Brown, shown in Figure 39. In this instance, all perspectives involved disagree substantially, not only at the level of sevenths but also the quality of the third. Exploring why this might be the case, one finds the song consists of unison riffs and syncopated rhythms with liberal use of rests. In the absence of sustained simultaneities, most of the harmony in the song is implied, creating even more room for subjectivity on behalf of the annotators. The automatic system, alternatively, flips back and forth between the various perspectives of the two reference annotations. Again, the chord

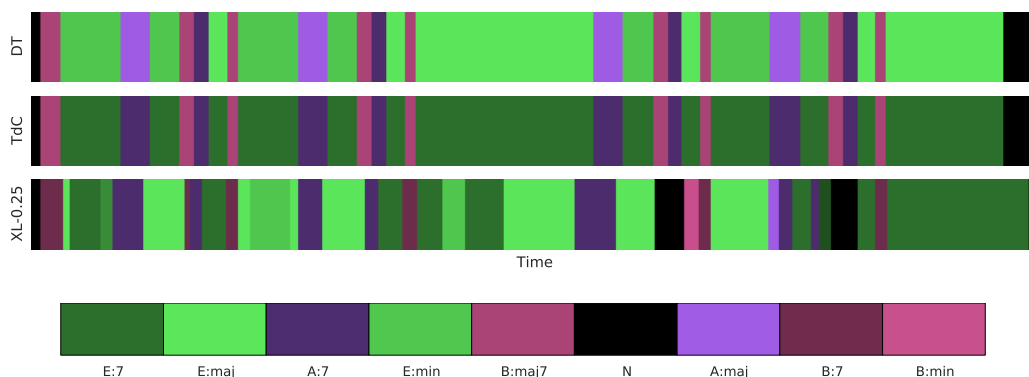


Figure 39: Reference and estimated chord sequences for a track in Quadrant III, the condition where neither annotator agrees with the algorithm, nor each other.

estimation machine has little choice but to be literal in its interpretation of solo vocal phrases, and labels such regions “no-chord” in the middle of the song. This kind of repeat behavior calls attention to what is fundamentally an issue of problem formulation. Chord *transcription* is a more abstract and ultimately different task than chord *recognition*, taking into consideration high-level concepts long term musical structure, repetition, segmentation or key, but conventional methodology conflates these two to some unknown degree.

Finally, a track from Quadrant IV —“Every Breath You Take” by The Police— is given in Figure 40. Similar to the track considered from the same region before, the estimation is consistently a half-step flat from the references. This pattern is indicative of a similar tuning discrepancy as before, and tuning a pure sinusoid to the track finds the tonic at approximately 426Hz, putting the song just over a quartertone flat. Though functionally equivalent to the previous example of tuning being an issue, this instance is interesting because two annotators independently arrived at the same decision. One reason this might have occurred is that, as a rock song, it makes far more sense to play

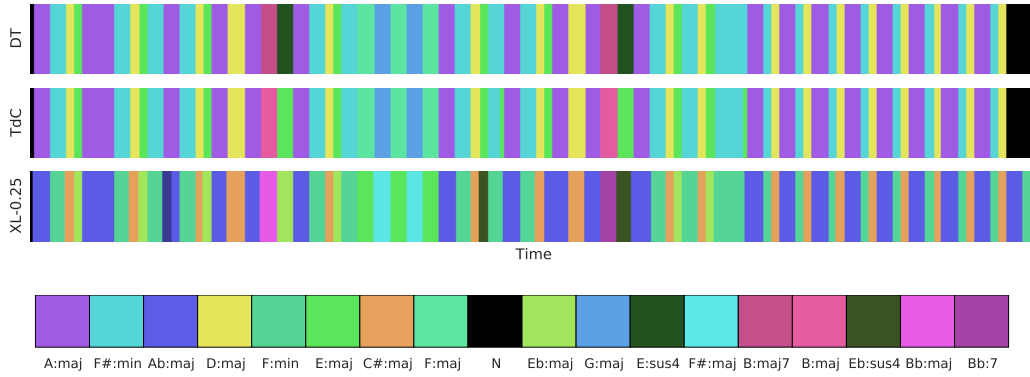


Figure 40: Reference and estimated chord sequences for a track in Quadrant IV, the condition where both annotators agree with each other, but neither agrees with the algorithm.

in $A:maj$ on guitar than $Ab:maj$. The chord shapes involved are much easier to form in $A:maj$, and therefore more likely. Additionally, a guitarist would probably need to change tunings to play the $Eb:maj$ in the right voicing. Taken together, it is noteworthy that such extrinsic knowledge can, and perhaps should, play a role in the process of automatic chord estimation.

4.5 Conclusions & Future Work

Based on conventional methodology, the proposed deep learning approach leads to results on par with the state of the art, just eclipsing the baseline in the various metrics considered. Though fully convolutional models can be complex enough to over-fit the dataset, a small amount of dropout during training helps reduce this over-fitting, while leading to slightly better generalization. Dropout also raises quality-wise recall in minority classes significantly. Given the significant imbalance of chord classes in the dataset, however, this is a very unstable measure of system performance. Small shifts in the quality-wise recall of a majority class can result in large performance swings, and

vice versa. Thus, the criteria for what makes for a “good” system may be motivated by use case, to determine which of these metrics correspond to the desired behavior.

Additionally, the suite of chord comparison functions demonstrate that most errors between reference and estimation are hierarchical, and increase with specificity of the chord, e.g. from root to triads to tetrads. One-of-K classifiers struggle to encode these relationships between certain chords, e.g. `A:maj` and `A:maj7`. By analogy, this is like building a classifier that discriminates between, among other classes, animals and cats, respectively; all cats are animals, but not all animals are cats. This is problematic in a flat classifier, because it is attempting to linearly separate a pocket of density contained within another. To address this issue, a structured output and loss function would be better suited to model these relationships. Directly encoding the knowledge that `maj7` chords are a subset of `maj`, will make it easier for a machine to learn these boundaries. One such hierarchy for the chords considered in this work is given in Figure 41. Here, the degree of specificity increases as the decision tree is traversed to a leaf node, and could be achieved with more structured output, such as a hierarchical softmax or conditional estimator.

In comparing the estimations of the two computational systems, deeper insight is gained into the ground truth data used for development, and the kinds of behavior, both the good and bad, that such approaches are prone to exhibit. First, it is observed that they make rather different predictions and their responses could be combined, motivating the exploration of ensemble methods. Looking at performance at the track-level, alternatively, helps identify problematic or noisy data in a collection. This is similar in spirit

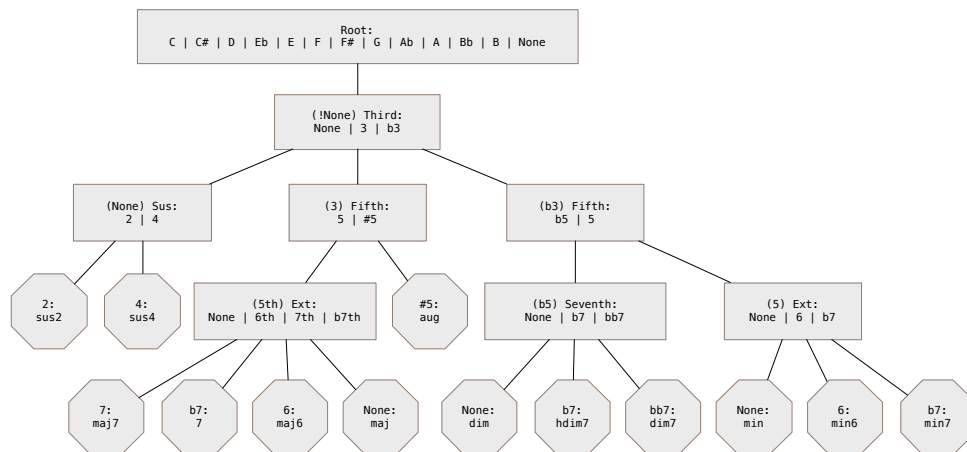


Figure 41: A possible chord hierarchy for structured prediction of classes. Decisions blocks are rectangular, with the result of the previous node shown in parentheses, the semantic meaning of the node is given before a colon, and the set of valid responses is pipe-separated. Stopping conditions are given as octagons.

to a growing body of work in MIR (Zapata, Holzapfel, Davies, Oliveira, & Gouyon, 2012), but it also provides some unique insight into the ACE task itself. Computational systems tend to be precise in ways humans are not, such as continuing to predict chords during a fade-out, or reporting “no-chord” during a musical break. The issue of hierarchical class relationships manifests in both models, but instances of algorithm disagreement point to music content that falls near a boundary between classes. Such knowledge could be used to single out and review the reference chord annotation more closely. These systems can also be sensitive to inexact tuning, and half-step deviations can be a large source of error. Additionally, implied harmony or temporally sparse

patterns can be especially problematic, resulting in an over-estimation of the “no-chord” class.

Conversely, leveraging multiple annotations for a given track provides a deeper understanding of the errors a computational system might make. As seen here, the assessment of a system will change depending on which interpretation is taken as a reference, and a computational system may agree with another human’s perspective, consistent with the work of McVicar (Ni, McVicar, Santos-Rodriguez, & De Bie, 2013). Most important is the recognition that human annotators do not agree all the time, and that describing some music content in the language of chords is inherently subjective. In such cases, there is no “ground truth” to speak of, and multiple chord labels may be acceptable. This is a critical observation, and one that cuts strongly against the long-standing methodology of curating ground truth references in ACE. Stated previously, the sole purpose of an objective measure is that it serves as a reasonable proxy for subjective experience. If the goal of ACE is to produce a chord annotation on par with that of a qualified human—a musical Turing test of sorts—then the reference annotation at hand may be only one of many valid perspectives. As a result, evaluating against a singular perspective is leading to results that may be inconsistent with subjective experience.

Instead, embracing multiple perspectives, rather than attempting to resolve them into a canonical target, would allow for more stable evaluation of computational models. One such way this could be achieved is by obtaining multiple chord annotations and creating a time-aligned bag of words reference. For example, knowing that 97 of 100 annotators labeled a chord as **A:7** is very different from the scenario that 52 did, while the other 48 reported **A:maj**. Curating a dataset with so many perspectives is unlikely to happen, but per-

haps multiple computational systems could be used to find boundary chords that *do* warrant multiple perspectives. Ultimately, this study demonstrates that future chord datasets should strive to capture the degree of subjectivity in an annotation, thus enabling objective measures better correspond with subjective experience.

5 Summary

In this chapter, the application of deep learning to ACE has been thoroughly explored. By standard evaluation practices, competitive performance is demonstrated on both a major-minor and large-vocabulary formulation of the task. Importantly, much effort is invested in understanding both the behavior of the computational systems discussed, as well as the reference data used to evaluate performance. Perhaps the most important finding is that the current state of the art may have truly hit a glass ceiling, due to the conventional practice of building and testing against “ground truth” datasets for an all too often subjective task. This challenge is further compounded by approaches to prediction and evaluation, which attempt to perform flat classification of a hierarchically structured chord taxonomy. Thus, while there is almost certainly room for improvement, the exploration here indicates that the vast majority of error in modern chord recognition systems is a result of invalid assumptions baked into the very task.

Notably, four issues with current chord estimation methodology have been identified in this work. One, it seems necessary that computational models embrace structured outputs; one-of-K class encoding schemes introduce unnecessary complexity between what are naturally hierarchical relationships.

Two, the community fundamentally needs to better distinguish between the two tasks at hand, being chord recognition —I am playing this literal chord on guitar— and chord transcription —finding the best chord label to describe this harmonically homogeneous region of music— and how this is articulated to reference annotation authors. Three, chord transcription requires more explicit segmentation, rather than letting such boundaries between regions of harmonic stability result implicitly from post-filtering algorithms. Lastly, the often subjective nature of chord labeling needs to be acknowledged in the process of curating reference data, and the human labeling task should combine multiple perspectives rather than attempt to yield a canonical “expert” reference.

CHAPTER VI

FROM MUSIC AUDIO TO GUITAR TABLATURE

In the previous chapter, it was demonstrated that state of the art ACE systems perform at a relatively high level, often producing reasonable, if not exact, chord estimations. Deploying these systems in the wild for real users, however, presents two practical difficulties: one, performing a given chord sequence requires that the musician knows how to play each chord on their instrument of choice; and two, the performance of classification-minded chord estimation systems does not degrade gracefully, especially for those lacking a good music theory background. Recognizing that guitarists account for a large majority of the music community, both challenges are addressed here by designing a deep convolutional network to model the physical constraints of a guitar’s fretboard, directly producing *human-readable* representations of music audio, i.e. tablature.

Approaching chord estimation through the lens of guitar tablature offers a variety of critical benefits. Guitar chord shapes impose an explicit hierarchy among notes in a chord family, such that related chords are forced to be near-neighbors in the output space. This constrains the learning problem in musically meaningful ways, and enables the model to produce outputs beyond the lexicon of chords used for training. The human-readable nature of the system’s output is also valuable from a user perspective, being immediately useful with minimal prerequisite knowledge. Furthermore, a softer prediction

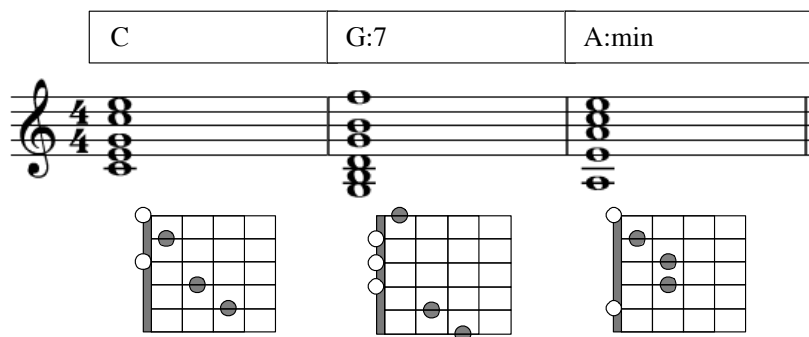


Figure 42: A chord sequence (top), traditional staff notation (middle), and guitar tablature (bottom) of the same musical information, in decreasing levels of abstraction.

surface results in more informative errors, thus allowing for more graceful degradation of performance. Finally, with an eye toward future work, the estimation of tablature makes it far easier for large online guitar communities to validate and, as needed, correct system outputs, regardless of skill level.

Therefore, this chapter extends a novel approach to bootstrapping the task of automatic chord estimation to develop an end-to-end system capable of representing polyphonic music audio as guitar tablature. To encourage playability, a finite vocabulary of chord shape templates are defined, and the network is trained by minimizing the distance between its output and the best template for a given observation. Experimental results show that the model achieves the goal of faithfully mapping audio to a fretboard representation, as well as further advancing the art in some quantitative evaluation metrics.

1 Context

To date, the majority of research in automatic chord estimation is based on the two-fold premise that (a) it is fundamentally a classification problem, and (b) the ideal output is a time-aligned sequence of singular chord names. That

said, it is worthwhile to reconsider how the development of such systems is motivated by the goal of helping the ambitious musician learn to play a particular song. Notably, today guitarists comprise one of the largest groups of musicians attempting to do just that. Over the last century, the guitar, in all of its forms, has drastically risen in popularity and prevalence, both in professional and amateur settings. Given the low start-up cost, portability, favorable learning curve, and —courtesy of musicians like Jimi Hendrix or *The Beatles*— an undisputed “cool factor” in Western popular culture, it is unsurprising that guitars dwarf music instrument sales in the United States. Based on the 2014 annual report of the National Association of Music Merchants (NAMM), a whopping 2.47M guitars were sold in 2013 in the United States, accounting for a retail value of \$1.07 *billion* USD (of Music Merchants, 2014); as a point of comparison, all wind instruments *combined* —the next largest instrument category— totaled just over half that figure, at \$521M USD.

The fact that guitarists comprise such a large portion of the musician community is important, as it affects how they might prefer to interact with a chord estimation system. While most instruments make use of traditional staff notation, fretted instruments, like lute or guitar, have a long history of using *tablature* to notate music. Illustrated in Figure 42, tablature requires minimal musical knowledge to interpret, and thus offers the advantage that is easier to read, particularly for beginners. Whereas staff notation explicitly encodes pitch information, leaving the performer to translate notes to a given instrument, tablature explicitly encodes performance instructions for a given instrument. Though it can be difficult to accurately depict rhythmic information with tablature, this is seldom an obstacle for guitarists. Chords-centric

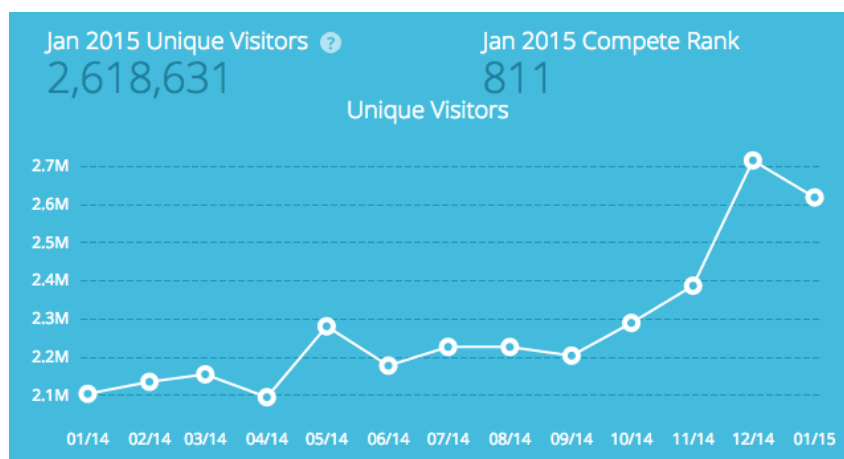


Figure 43: Visitor statistics for the tab website *Ultimate Guitar*, as of January 2015.

guitar parts typically place less emphasis on rhythm, and changes are usually aligned with lyrics or metrical position.

From the earliest days of personal computing, contemporary guitarists have embraced technology en masse for the creation and dissemination of “tabs.” Initial bandwidth and memory limitations, however, prevented the curation of high resolution images of sheet music, and symbolic representations, like MIDI, required specialized programs to render the music visually. With small file sizes and compatibility with common text editors, ASCII “tabs” made it comparatively trivial to create, share, and store guitar music. Thus, combining easy readability and a sufficient level of musical detail with technological constraints of the time period, guitar tablature spiked in popularity towards the end of the 20th century. As evidenced by heavily trafficked, user-curated websites like Ultimate Guitar*, modern online guitar communities continue to

* <http://www.ultimate-guitar.com/>

place a high demand on tablature. Shown in Figure ??, this website alone sees, on average, over 2M unique visitors* per month in the United States.

Taken together, these observations motivate an obvious conclusion. Guitarists comprise a significant portion of the global music community, and are actively creating and using tablature as a means of learning music. An automatic chord estimation system would be extremely valuable to this demographic, but such a system should be sensitive to the common preference for tablature. Therefore, this chapter is an effort to steer automatic chord estimation toward a specific application, in order to address a potential use case for a very real demographic.

2 Proposed System

Building on previous efforts in automatic chord estimation, the best performing configuration presented in Chapter 5, XL-0.125, is extended here for guitar chord estimation. As such, many design decisions remain consistent with the previous description, such as the constant-Q representation, dataset, or evaluation methodology, and are omitted from the discussion. There are a few important differences, however, which are addressed below. First, the architecture is modified slightly to produce fretboard-like outputs. A strategy is then discussed for incorporating guitar chord shapes into the model. The modified objective and decision functions are presented last, detailing how the model is optimized and use to estimate discrete chord classes.

Finally, it is worth mentioning that, despite the desire to map notes to a guitar fretboard, this approach is much closer in principle to chord estimation

*Based on Compete.com analytics data, accessed on 15 March, 2015.

than automatic transcription methods. Thus, while some previous work embraces this position in the realm of transcribing guitar recordings (Barbancho, Klapuri, Tardón, & Barbancho, 2012) or arranging music for guitar (Hori, Kameoka, & Sagayama, 2013), the only previous work in estimating guitar chords as tablature directly from polyphonic recordings is performed by the author, in (Humphrey & Bello, 2014).

2.1 Designing a Fretboard Model

So far, this study has shown deep trainable networks to be a powerful approach to solving complex machine learning problems. Another particular advantage of deep learning is that it can be remarkably *flexible*, whereby functionally new systems can be developed quickly by making different architectural decisions*. This high-level design strategy is exploited here by modifying the convolutional neural network used in the previous chapter to produce an output representation that behaves like the fretboard of a guitar.

To understand the proposed model, it is helpful to first review the physical system of interest. The modern guitar consists of six parallel strings, conventionally tuned to E2, A2, D2, G3, B3, and E4, and thus can simultaneously sound between zero and six notes. A guitar is also fretted, such that the different pitches produced by a string are quantized in ascending order as a function of fret, resulting from shortening the length of the vibrating string in discrete quantities. Continuous pitch may be achieved by various means, such as bending the strings, but such embellishments are beyond the scope of consideration here. Thus, it can be said that each string only takes a finite

*Provided its “quality” can be expressed as a differentiable objective function, that is.

number of states: off (X), open, (0), or a number corresponding to the fret at which the string is held down. Most real guitars have upwards of 20 frets, but, as a simplification, all chords will be voiced in the first seven frets; therefore, in this model, each string will take one of nine mutually exclusive states.

Framed as such, the strings of a guitar can modeled as six correlated, but ultimately independent, probability mass functions. This is achieved by passing the output of an affine projection through a softmax function, as described in Chapter ??, yielding a non-negative representation with unity magnitude. Starting with the first three layers of the “XL” model defined previously, six independent softmax layers are used to model each string independently, and concatenated to form a 2-dimensional heatmap of the fretboard:

$$Z_i = f_i(X_{l-1}|\theta_i) = \sigma(W_i \bullet X_{l-1} + b_i), i \in [0 : 6), \theta = [W_i, b_i s]$$

The activation of the i^{th} string, Z_i , is computed by projecting the output of the penultimate layer, X_{L-1} , of an L -layer network against the weights, W_i , and added to a bias term, b_i . This linear transformation is normalized by the softmax function, σ , and repeated for each of the six strings. The overall model is diagrammed in Figure 44.

2.2 Guitar Chord Templates

Having designed a convolutional network to estimate the active states of a fretboard, it is necessary to devise a mapping from chord transcriptions to fingerings on a guitar. As the annotations available were curated for generic chord estimation, they do not offer insight into how a given chord might best

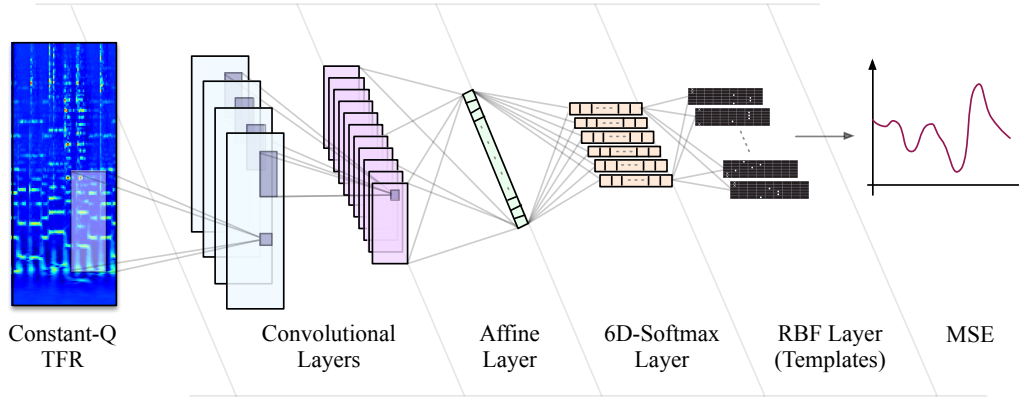


Figure 44: Full diagram of the proposed network during training.

be voiced on a guitar. Therefore, using the same vocabulary of 157 chords as before, a canonical chord shape “template” is chosen for each. This is done in such a way so as to prefer voicings where all quality variations over a given root are maximally similar, i.e. chords of the same root are near-neighbors in fretboard space. To illustrate, tablature representations for all templates are given in Figure ??.

It is worthwhile at this point to acknowledge the natural multiplicity of chord voicings on the guitar. In addition to the normal variation that may occur in stacking the notes of a chord, e.g. “open” or “closed” position, there are other factors that influence the actual pitches that are played. First, the same note can often be played in multiple positions on the fretboard. For example, E3 can be played on the 12th fret of the first string, the 7th of the second, or the 2nd of the third. Additionally, some standard chord voicings cannot be formed on the guitar, comfortably or otherwise. For this reason it is quite common to play the 3rd scale degree of a chord above the octave, as the 10th, though there are a few exceptions resulting from open fingerings. Context may also influence how a chord is played, such as the instance in

which it is easier to move from one chord shape to the next. Rather than attempt to address all of these issues now, the canonical template approach is chosen as a practical means to simplify overall system design. The choice of one template over another likely has nontrivial implications for the behavior of the model, but this is left as a variable to be explored in the future.

2.3 Decision Functions

While estimating frets is sufficient for human interpretation, it is necessary to design two related decision functions in order to allow the machine to operate automatically. First, the templates defined above must be incorporated into an objective function, such that the machine can learn to optimize this measure over the training data. Finding inspiration in (LeCun et al., 1998), a Radial Basis Function (RBF) layer is added to the network, given as follows:

$$\mathcal{E}(Z|W_T) = \sum (Z_{out} - W_T[k])^2 \quad (28)$$

where Z is the output of the fretboard model, W_T is a tensor of chord shape templates with shape $(K, 6, 9)$, such that K is the number of chord templates, and k the index of the reference class. Note that these templates will impose the proper organization on the output of the model, and thus remain fixed during the learning process. Since these weights are constant, minimizing this function does not require a contrastive penalty or margin term to prevent it from collapsing, i.e. making all the squared distances zero.

Additionally, for the purposes of Viterbi post-filtering and fairly comparing with previous results, the energy surface must be inverted into a likelihood function. This is achieved by negating the energy function, E , and normalizing as a Gibbs distribution:

$$\mathcal{L}(Y_k|X, \Theta) = \frac{\exp(-\beta E(Y_k|X, \Theta))}{\sum_i^K \exp(-\beta E(Y_k|X, \Theta))} \quad (29)$$

For the experiments detailed below, $\beta = 1$. It is conceivable that the choice of hyperparameter may impact system behavior when coupled with the Viterbi algorithm, but this value was empirically observed to give good results and not explored further.

3 Experimental Method

The focus of this chapter now shifts toward the experimental method adopted to investigate the behavior of this basic approach. Herein, the training strategy, corresponding variables, and subsequent quantitative results are addressed in turn.

3.1 Training Strategy

Though it is able to incorporate musical knowledge into the architectural design, the model proposed here is unable achieve root-invariant weight sharing as in the previous chapter. This is due to root-dependent chord shapes resulting from the nonuniform arrangement of chords on the neck of the guitar. It is important to consider the effect this has on system performance, as well as other means of achieving “root invariance”, and thus three different training strategies are employed here.

As a baseline condition, the model is trained with the natural distribution of the data (“as-is”). Note that it is reasonable to expect these models to be deficient, as there may be chord class mismatch between training and test conditions, i.e. chord classes in the test partition do not occur in the training

set. To address the imbalanced learning problem, the second training condition scales the loss of each training observation by a class-dependent weight (“scaled”). These weights are determined by computing the root-invariant prior over the training partition, taking its inverse, and standardizing the coefficients to unit mean and standard deviation. The third and final training condition couples loss scaling with data augmentation, such that during training each datapoint is circularly shifted in frequency on the interval $[-12, 12]$ (“augmented”). This allows the variance of each chord quality to be evenly distributed across classes, and helping prevent any missing class coverage in the training set.

Identical partitions of the dataset used in the previous chapter are employed here; the dataset is split 68:12:20, into training, validation, and test, respectively, and the partitions are rotated so that all data is used as a holdout set once. All models are trained with mini-batch stochastic gradient descent at a batch size of 100, learning rate of 0.02, and dropout ratio of 0.125. Training proceeded for several hundred iterations, ultimately bounded by a ceiling of 24 hours, and parameters saved every 10k iterations. Model selection was performed as a brute force search over both the parameter checkpoints and self-transition penalty for Viterbi, from -20 to -40 in steps of 5. The best model was chosen by the maximum of the harmonic mean of the chord evaluation metrics outlined previously.

3.2 Quantitative Evaluation

In the absence of a thorough user study, the proposed approach is evaluated against the chord estimation task posed in Chapter 5.4. Applying the methodology outlined above, the three training conditions were run to completion and

Table 22

Weighted recall scores over the test set for two previous models, and the three conditions considered here.

	triads	root	MIREX	tetrads	sevenths	thirds	majmin
Cho	0.7970	0.8475	0.8147	0.6592	0.6704	0.8197	0.8057
Root-invariant (Chapter 5)	0.7995	0.8493	0.8145	0.6673	0.6788	0.8227	0.8077
As-Is	0.8234	0.8705	0.8352	0.6855	0.7084	0.8376	0.8394
Scaled	0.8156	0.8644	0.8283	0.6791	0.6994	0.8308	0.8295
Augmented	0.8294	0.8715	0.8420	0.6989	0.7167	0.8440	0.8412

used to predict into the space of 157 chord classes. Given the intersection with the previous chapter, these three systems are compared to the system presented in (Cho, 2014), referred to as “Cho”, and the model related to those explored here, “XL-0.125”, referred to now as the “root-invariant” condition.

Overall performance is measured as weighted recall across metrics, as per the previous chapter, and given in Table 22. These results indicate that, over all the data, the three fretboard models perform far better than either of the two previous ones. Additionally, the combination of weight scaling and pitch shifting leads to the best overall performance. As the second, weight scaled condition fares slightly worse than training the models with the data as-is, it is reasonable to assume that applying pitch shifting only during training likely would have resulted in higher overall scores.

However, quality-wise recall, given in Table 23, offers a more nuanced depiction of the effect of these strategies. Going from the “as-is” to “scaled” conditions, the slight reduction in micro-recall is traded for an increase in averaged quality-wise recall. This result is intuitively satisfying, as the introduction of class-dependent weights into the training process should help raise the preference for the long tail chords. Though this can help attenuate the model’s strong preference for majority chord classes, it does nothing to help

Table 23

Quality-wise recall across conditions.

quality	Root-invariant	As-is	Scaled	Augmented	Support (min)
maj	0.7390	0.8572	0.8413	0.8417	397.4887
min	0.6105	0.6516	0.6312	0.6645	105.7641
7	0.5183	0.2928	0.3001	0.3367	68.1321
min7	0.5263	0.4556	0.4670	0.5077	63.9526
N	0.7679	0.6670	0.6712	0.6942	41.6994
maj7	0.6780	0.4143	0.4614	0.5525	23.3095
maj6	0.2908	0.0259	0.0682	0.1061	7.6729
sus4	0.3369	0.0252	0.0952	0.1747	8.3140
sus2	0.3216	0.0098	0.0146	0.2216	2.4250
aug	0.5078	0.0093	0.1431	0.3365	1.2705
dim	0.4105	0.2898	0.4030	0.3803	1.8756
min6	0.3870	0.0367	0.1611	0.3011	1.5716
hdim7	0.5688	0.0000	0.0610	0.3913	1.1506
dim7	0.1790	0.0040	0.0453	0.0391	0.5650
average	0.4887	0.2671	0.3117	0.3963	

address the overall deficiency of minority classes. Therefore, when loss scaling is combined with data augmentation, the increase in performance is profound; nearly all statistics, at both the micro and quality-wise macro level, improve, some significantly.

Additionally, it can be seen that the higher overall scores in the guitar model are a result of over-predicting majority, and in particular “major”, chords Compared to the root-invariant model of the previous chapter, the guitar models are over 10% better at predicting major chords alone. Somewhat surprisingly, the root-invariant model is nearly 20% better at dominant seven chords than any new model trained here. This can be understood as an artifact of the intersection in fretboard space between major and dominant seven chords, coupled with the significant bias toward major chords. Finally, as to be expected, the imbalanced distribution of chord qualities prevents the lower

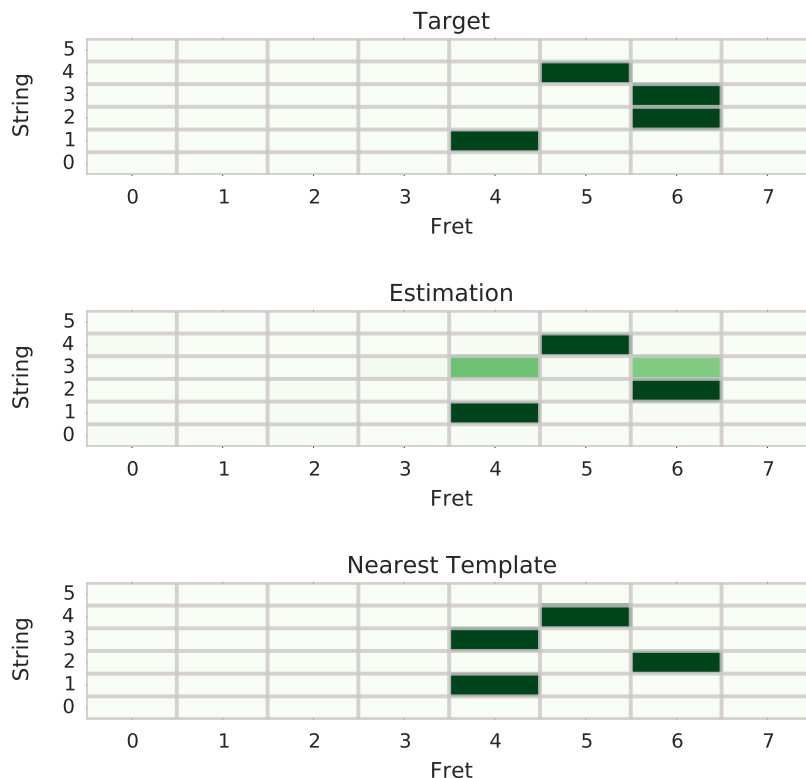


Figure 45: Understanding misclassification as quantization error, given a target (top), estimation (middle), and nearest template (bottom).

quality-wise scores from being reflected in the overall, micro-recall statistics. This behavior alludes to the previous discussion regarding the apparent trade-off between global performance and quality-wise accuracy.

Another important behavior to consider here is the idea that the direct estimation of a fretboard representation affords some benefit over simply projecting the predicted labels of a standard chord classifier *back* onto guitar chord templates. Typical chord estimation systems, and especially those that rely heavily on Viterbi, like (Cho, 2014), produce a sequence of discrete chord labels, and are thus effectively “classifiers”. Comparatively, fretboard estimation can be seen as regression, given the continuous output surface, but can be

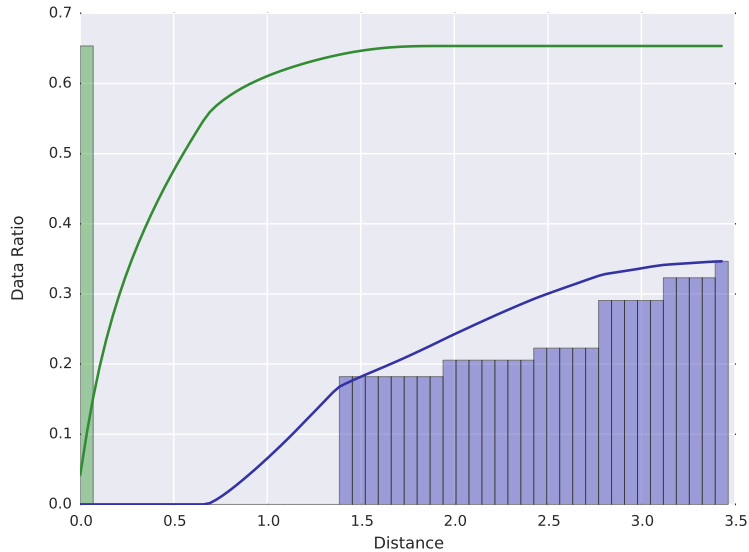


Figure 46: Cumulative distribution functions of distance are shown for correct (green) and incorrect (blue) classification, in the discrete (classification) and continuous (regression) conditions.

used for classification by identifying the closest known chord template, referred to as *vector quantization*. Thus, illustrated in Figure 45, misclassification can be understood as a type of quantization *error*; here a prediction close to, but on the wrong side of, the decision boundary is assigned to a different chord shape. However, since the representation is human readable, it is trivial to correct the error from a `C#:min7` to `C#:min`.

In the space of fretboard estimation, this can be quantified by comparing the distances between predictions both before and after classification, partitioned on whether or not the datapoint is correctly assigned. First, the distances between the continuous-valued fretboard and the corresponding target template are computed and tallied as a histogram. Next, the estimated fretboard representations are assigned to the *nearest* template, and distance is computed to the target; these distances, are also aggregated into a histogram,

but take only a finite number of values. Both probability density functions are integrated into continuous distribution functions to illustrate what ratio of the data lives within the radius of a bounding volume, shown in Figure 46. Here, it can be seen that classification clearly widens the gap between correct and incorrect answers. Alternatively, in the regression scenario, nearly 20% of these errors are closer to their correct class, accounting for well over 80% of the data. Importantly, the improved proximity of these errors mean that they should be easier to correct by potential users.

4 Discussion

This chapter has proposed a method of extending a previous automatic chord estimation by introducing the physical constraints of a guitar. Not only does this yield a higher performing system, based on overall metrics, but the outputs are directly human readable making the system attractive from a user experience standpoint. While the system seems to struggle more than previous efforts across all chord qualities, this is arguably an instance in which performing better over *all* the data is preferable. The sample sizes of rare chords in the dataset are at times too small to draw meaningful conclusions about performance at the class level. Further complicating matters, there is also some unknown degree of subjectivity in the reference annotations, used for both training and evaluation. Pragmatically speaking though, consistently tracking root motion, e.g. being in the right place on the neck, is probably sufficient for most guitarists to deem such a system useful. The vast majority of music can be simplified as “power chords”, consisting of the root and the fifth, and often more detailed chords can be realized by modifying this basic

shape. Therefore, while many methodological challenges inherent to chord estimation persist in this study, predicting chord shapes as tablature softens the degree to which they impact usability.

Setting aside the limitations of quantitative evaluation in chord estimation, the proposed system warrants a subjective, user experience study in the future. Many of the gains identified here can only be qualitatively assessing how such a system achieves its goals, such as the degree to which performance *does* in fact degrade gracefully. Countless instances can be identified where system “errors” can be reasonably interpreted as the chord name provided, but only a study with real users will indicate how useful this is.

Perhaps most importantly though, the next logical research step is to develop an interactive user interface and deploy the system at scale. In addition to distributed instruction, deploying an “autotabbing” system provides a means to collect and clean reference annotations. Approaching this task from the perspective of human-readable representations holds considerable promise in the space of data curation, as editing a chord transcription is generally a far simpler task than creating one from scratch. This reality is only amplified for annotators who lack formal ear training. Importantly though, this reduces the prerequisite skill level, the amount of time needed to complete the task, or both, required in the data authoring task. Therefore, this system creates the potential to include a larger number of musicians across a wide array of skills in the data collection process. More annotators not only opens the door for more annotations, but multiple perspectives of the *same* music content. This will be critical to the role subjectivity plays in chord estimation research, now and in the near future.

CHAPTER VII

WORKFLOWS FOR REPRODUCIBLE RESEARCH

In recent years, the philosophy of open source software has begun taking root in scientific research, particularly in the field of computer science. There are several reasons why open research is beneficial to the greater body of human knowledge, but three are of particular value here. First, sharing code and data allows others to reproduce previous results, a fundamental tenet of the scientific method. Open source implementations are invaluable for sufficiently complex systems. It may be near impossible to describe every minute detail in a publication necessary for someone else to replicate the obtained results; for some works, in fact, the only artifact that can do this unambiguously is the source code itself. Second, and in a related vein, open research makes it both easier and faster to build upon and extend the previous work of others. Even in the instance a researcher is able to recreate a published system, the time and effort necessary to get to this point is significant and arguably unnecessary. Granted, while there is an educational component inherent to the re-implementation of previous work, the situation is akin to long division: it is certainly valuable to learn *how* to divide by hand, but no one shuns the use of a calculator on a day to day basis. Lastly, it is a good and responsible act to contribute tools and software back to the larger research community. All research stands on the shoulders of previous efforts, from improving on a recently published algorithm to the decades-old linear algebra routines doing

all its number crunching. The reality is that no one individual has ever truly solved anything on their own, and sharing the fruits of one’s research endeavors serves the common good.

With these motivations in mind, this chapter details the several open source contributions made in the course of this work, culminating in a single code repository used to produce the results contained herein. These software tools consist of the following: Section 1 describes `jams`, a JSON annotated music specification and python API for rich music descriptions; Section 2 introduces `biggie`, an HDF5 interface for interacting with notoriously big data; Section 3 details `optimus`, a user-friendly library for building and serializing arbitrary acyclic processing graphs; Section 4 discusses relevant contributions to `mir_eval`, a transparent framework for evaluating MIR systems; and finally, Section 5 outlines `dl4mir`, the framework used here to complete this research.

1 `jams`

Music annotations—the collection of observations made by one or more agents about an acoustic music signal—are an integral component of content-based music informatics methodology, and are necessary for designing, evaluating, and comparing computational systems. For clarity, the scope of an annotation is constrained to time scales at or below the level of a complete song, such as semantic descriptors (tags) or time-aligned chords labels. Traditionally, the community has relied on plain text and custom conventions to serialize this data to a file for the purposes of storage and dissemination, collectively referred to as “lab-files.” Despite a lack of formal standards, lab-files have been, and

continue to be, the preferred file format for a variety of music description tasks, such as beat or onset estimation, chord estimation, or segmentation.

Meanwhile, the interests and requirements of the community are continually evolving, thus testing the practical limitations of lab-files. Reflecting on this tradition, there are three unfolding research trends that are demanding more of a given annotation format:

- **Comprehensive annotation data:** Rich annotations, like the Billboard dataset (Burgoyne et al., 2011), require new, content-specific conventions, increasing the complexity of the software necessary to decode it and the burden on the researcher to use it; such annotations can be so complex, in fact, it becomes necessary to document how to understand and parse the format (De Haas & Burgoyne, 2012).
- **Multiple annotations for a given task:** The experience of music can be highly subjective, at which point the notion of “ground truth” becomes tenuous. Recent work in automatic chord estimation, both here and in (Ni et al., 2013), has shown that multiple reference annotations should be embraced, as they can provide important insight into both system evaluation and the problem formulation itself.
- **Multiple concepts for a given signal:** Although systems are classically developed to describe observations in a single namespace, e.g. chords, there is ongoing discussion toward integrating information across various musical concepts (Vincent, Raczynski, Ono, Sagayama, et al., 2010). This has already yielded measurable benefits for the joint estimation of chords and downbeats (Papadopoulos & Peeters, 2011) or chords

and segments (Mauch, Noland, & Dixon, 2009), where leveraging multiple information sources for the same input signal can lead to improved performance.

It has long been acknowledged that lab-files cannot be used to these ends, and various formats and technologies have been previously proposed to alleviate these issues, such as RDF (Cannam, Landone, Sandler, & Bello, 2006), HDF5 (Bertin-Mahieux et al., 2011), or XML (McKay, Fiebrink, McEnnis, Li, & Fujinaga, 2005). However, none of these formats have been widely embraced by the community. Considering these options, the weak adoption of alternative formats is likely due to the combination of multiple factors. For example, new tools can be difficult, if not impossible, to integrate into a research workflow because of compatibility issues with a preferred development platform or programming environment. Additionally, it is a common criticism that the syntax or data model of these alternative formats is non-obvious, verbose, or otherwise confusing. This is especially problematic when researchers must handle format conversions. Therefore, a JSON Annotated Music Specification (JAMS) was developed to address these various needs.

1.1 Core Design Principles

In order to craft an annotation format that might serve the community into the foreseeable future, it is worthwhile to consolidate the lessons learned from both the relative success of lab-files and the challenges faced by alternative formats into a set of principles that might guide the design of a new format. With this in mind, it is argued that usability, and thus the likelihood of adoption, is a function of three criteria: simplicity, structure, and sustainability.

1.1.1 Simplicity

The value of simplicity is demonstrated by lab-files in two specific ways. First, the contents are represented in a format that is intuitive, such that the document model clearly matches the data structure and is human-readable, i.e. uses a lightweight syntax. This is a particular criticism of RDF and XML, which can be verbose compared to plain text. Second, lab-files are conceptually easy to incorporate into research workflows. The choice of an alternative file format can be a significant hurdle if it is not widely supported, as is the case with RDF, or the data model of the document does not match the data model of the programming language, as with XML.

1.1.2 Structure

It is important to recognize that lab-files developed as a way to serialize tabular data (i.e. arrays) in a language-independent manner. Though lab-files excel at this particular use case, they lack the structure required to encode complex data such as hierarchies or mix different data types, such as scalars, strings, multidimensional arrays, etc. This is a known limitation, and the community has devised a variety of ad hoc strategies to cope with it: folder trees and naming conventions, such as “{X}/{Y}/{Z}.lab”, where X, Y, and Z correspond to “artist”, “album”, and “title”, respectively*; parsing rules, such as “lines beginning with ‘#’ are to be ignored as comments”; auxiliary websites or articles, decoupled from the annotations themselves, to provide critical information such as syntax, conventions, or methodology. Alternative representations are able to manage more complex data via standardized markup and

*<http://www.isophonics.net/content/reference-annotations>

named entities, such as fields in the case of RDF or JSON, or IDs, attributes and tags for XML.

1.1.3 Sustainability

Recently in MIR, a more concerted effort has been made toward sustainable research methods, which we see positively impacting annotations in two ways. First, there is considerable value to encoding methodology and metadata directly in an annotation, as doing so makes it easier to both support and maintain the annotation while also enabling direct analyses of this additional information. Additionally, it is unnecessary for the MIR community to develop every tool and utility ourselves; we should instead leverage well-supported technologies from larger communities when possible.

1.2 The JAMS Schema

A JAMS object is hierarchically structured to capture all relevant information in a logically organized manner. The primary record is an *annotation*, of which a JAMS object may contain zero or more. Annotations are comprised of *observations*, which maintain a set of properties: time, duration, value, confidence, namespace. Observations have two variants, to better handle *sparse* or *dense* data, such as onsets or melody, respectively. The semantic context of an observation is specified by its *namespace*, providing information about how the data in the observation should be understood. Thus a namespace allows for easy filtering and interpretation of the data in an annotation for different music description tasks.

An annotation also maintains a *metadata* object. Annotation metadata allows for rich descriptions of *who* and *how* a particular record was produced.

Currently, metadata has properties such as *corpus*, *annotator*, *validation*, *curator*, to name a few fields. Not only does this information make it easier to develop and evaluate systems with an eye to subjectivity, but it enables deeper meta-analyses of the annotations themselves. This could be achieved by considering the observations made by annotators with different musical backgrounds or degrees of formal training, for example.

In addition to an array of annotations, JAMS objects also maintain a top-level file metadata object. While annotation metadata sponges information about observer, file metadata tracks global information about the corresponding audio signal, with properties like *title*, *artist*, *duration*, or *identifiers*. As there is currently no standard convention for uniquely specifying audio recordings in a global manner, file metadata exists to help link the JAMS object with the appropriate signal.

Finally, *sandboxes* exist in both the top-level and annotation objects to facilitate the growth and extensibility of the format. These are unconstrained objects that can be used as needed for anything not covered by the formal schema. This is done in the hope that sandboxes might identify information that could be incorporated into future versions.

1.3 Python API

To facilitate the use and integration of JAMS in software projects, a Python library is publicly available and under active development*. This application programming interface (API) provides a simple interface for reading, writing, and manipulating the information in a JAMS object. Many common datasets

*<http://github.com/marl/jams>

are also provided in this format to further encourage adoption. Complementing the creation and use of human annotations, JAMS makes it easier to operate on this information, such as augmenting audio and annotations in parallel*.

2 biggie

Common practice in machine learning frameworks, like scikit-learn[†], often proceeds by massaging the training data into something like a large table, i.e. rows are individual datapoints, and columns correspond to different features, coefficients, etc. Attempts to map this paradigm to time-varying data, such as audio, can be problematic and cumbersome in practice. It is typically advantageous to learn on several consecutive frames of features, referred to as tiles, windows, or shingles, as was the case in all work presented here. These tiles could be sharded from longer feature sequences into some number of discrete, equivalently shaped observations, but this is generally undesirable. Doing so limits the flexibility of the researcher considering different tile sizes, requiring that the data be sharded again. More difficult, this practice increases the footprint of the dataset linearly with the size of each observation. Lastly, it is helpful to keep the entire feature sequence in tact, as it facilitates the application of any additional time-dependent processing, e.g. Viterbi decoding.

To achieve these ends, **biggie**[‡] is the high-dimensional, signal-level equivalent to JAMS, built on two basic objects. An *entity* is a struct-like object designed to keep various numerical representations together, regardless of samplerate. These objects are then freely written to and read from a *stash*, a

*<http://github.com/bmcfee/muda>

†<http://scikit-learn.org/>

‡<http://github.com/ejhumphrey/biggie>

persistent, i.e. on-disk, key-value store. Here, each entity is given a unique key, making it easy to align a dictionary of signals with a dictionary of annotations. Leveraging the `h5py`^{*} library under the hood and based on HDF5[†], biggie scales to arbitrarily large datasets; however, though it shares a common interface with in-memory dictionaries, the footprint of a stash scales gracefully with available computational resources by lazily loading data into memory. Biggie also allows random access and data caching, greatly facilitating stochastic sampling for on-line learning[‡]. Perhaps most practically though, biggie serves as *the* data interchange format between processing stages in the larger framework, eliminating the need for filename parsing or content-specific naming conventions; operations are written to consume and return stashes with data under the same keys. While biggie helps address similar pain points as other libraries, like `pandas`[§], the HDF5 back-end is crucial for serializing a large collection numerical tensors in a simple, Pythonic manner.

3 optimus

Deep learning tools have matured rapidly in the last half decade, with powerful choices across several programming languages. Developed under the direction of Yoshua Bengio at the University of Montreal, Theano[¶], is the leading Python library for deep learning. Boasting a large and growing user base, Theano offers all the pieces necessary for deep learning research, including

^{*}<http://www.h5py.org/>

[†]<http://www.hdfgroup.org/HDF5/>

[‡]<https://github.com/bmcfree/pescador>

[§]<http://pandas.pydata.org/>

[¶]<http://deeplearning.net/software/theano/>

symbolic differentiation, optimized C-implementations, and seamless integration with GPUs. Though extremely powerful, useful, and expressive, there are two facets of the library that proved troublesome in the course of this work. Serialization is can be tricky, especially for networks under active development. The common approach to saving objects in Python, known as “pickling”, is sensitive to modifications to the code, and thus something pickled previously may not be recoverable in the future. Additionally, designing neural networks in Theano can be rather time-consuming, and is not the most user-friendly experience. This is especially true when constructing non-standard architectures, such as guitar fretboard models or pairwise training strategies.

Optimus* is therefore an effort to address both of these problems in a maximally versatile and intuitive manner. To simplify the creation, training, and application of neural networks, common building blocks are provided as natively serializable objects that can be wired together in a graphical manner. Arbitrary acyclic, i.e. no loops, directed processing graphs can be architected from a large collection of *nodes*, including inputs, functions, like “Affine” or “Conv3D”, and outputs. While a large space of loss functions can be realized from these basic building blocks, a handful of standard *losses* are provided, simplifying design further. The topology between these parts is given by a routing table, and passed off to a *graph*, which connects the dots and returns a callable function. Furthermore, given the flexibility to define topologies in a processing graph, it is simple to explore a variety of modifications, such as layer-wise hyperparameters, tapping various intermediary representations as outputs, or “cloning” a processing node to create a deep copy of its parameters.

*<http://github.com/ejhumphrey/optimus>

In addition to the user-facing advantages, robust serialization is achieved by expressing processing graph definitions as JSON and archives of multidimensional arrays. This offers the additional benefit that it would be straightforward to port optimus models, as graph definitions and parameter bundles, across languages in the future. Finally, as parameter assignments are expressed through named nodes and fields, it is trivial to not only save parameters but easily initialize them by arbitrary means, such as pre-training or using supervised learning on hand-crafted functions.

4 mir_eval

As addressed at the outset of this work, the conventional formulation of many music description tasks attempts to model the behavior of an expert human listener. Framing the problem as a mapping between inputs and outputs allows for objective quantitative evaluation, thus providing common dimensions on which to compare algorithms and systems against each other. The particular dimensions on which an algorithm is evaluated is almost by definition specific to the application, and different metrics have evolved over time to provide musically meaningful assessment. Many such scoring functions are nontrivial to implement, however, and small details can give rise to variability in resulting metrics. The Music Information Retrieval Evaluation eXchange (MIREX), the community’s annual algorithm evaluation event, has helped provide common ground on which to compare different systems. This implementation is seldom used outside MIREX due to a variety of practical difficulties, however, and instead, researchers often resort to reimplementing the same evaluation metrics. Unfortunately these personal implementations are not standardized, and may

differ in important details, or even contain bugs, confounding comparisons. Therefore, the `mir_eval` project aims to address these challenges for a variety of well-established tasks (Raffel et al., 2014).

Though larger in scope, the contributions to the chord estimation evaluation are particularly relevant to this work. Despite being one of the oldest MIREX tasks, evaluation methodology and metrics for automatic chord estimation is an ongoing topic of discussion, due to issues with vocabularies, comparison semantics, and other lexicographical challenges unique to the task (Pauwels & Peeters, 2013). One source of difficulty stems from an inherent subjectivity in “spelling” a chord name and the level of detail a human observer can provide in a reference annotation (Ni et al., 2013). As a result, a consensus has yet to be reached regarding the single best approach to comparing two sequences of chord labels, and instead are often compared over a set of rules, i.e Root, Major-Minor, and Sevenths, with or without inversions.

To efficiently compare chords, a given chord label is first separated into its constituent parts, based on the syntax of (Harte, 2010). For example, the chord label `G:maj(6)/5` is mapped to three pieces of information: the root (“G”), the root-invariant active semitones as determined by the quality shorthand (“maj”) and scale degrees (“6”), and the bass interval (“5”). Based on this representation, an estimated chord label can be compared with a reference by defining a comparison function between these invariant representations. Any encoded chords that are deemed to be “out of gamut” return a negative score to be easily filtered. Track-wise scores are computed by weighting each comparison by the duration of its interval, over all intervals in an audio file. This is achieved by forming the union of the boundaries in each sequence, sampling the labels, and summing the time intervals of the “correct” ranges.

The cumulative score, referred to as *weighted chord symbol recall*, is tallied over a set audio files by discrete summation, where the importance of each score is weighted by the duration of each annotation (Mauch, 2010).

5 dl4mir

Leveraging these various software components, an aggregate framework representing all work presented herein is also made available, referred to as `dl4mir*`. At a high level, this resource contains the additional functionality necessary to reproduce this work. For convenience, the majority of processes can be executed via a series of shell scripts to perform feature extraction, training, and evaluation. All reported results and figures provided in this document are survived in IPython notebooks for both reference and repeatability. Furthermore, dependencies have been minimized to make this framework sufficiently independent. In addition to the immediate goal of reproducing experimental results, this is done in the hope that it may facilitate future extensions of this work. Serialized versions of important network models are also included to make comparisons against similar work easier, or to simply build them into other systems. Finally, the source code is provided so that it will serve as another example of how one might architect a flexible deep learning framework.

6 Summary

This chapter has covered the various libraries and programming tools developed in the course of this work: `jams`, a JSON format and Python API for music annotation; `biggie`, an HDF5 interface for managing large amounts

*<http://github.com/ejhumphrey/dl4mir>

of numerical data; `optimus`, a versatile yet intuitive approach to building, training, and saving deep networks; and `mir_eval`, an open framework for evaluation. Not only are many of these components independently useful to deep learning workflows, but the software necessary to repeat the research reported here is made publicly available online, in the form of `dl4mir`. Following the spirit of reproducible research, these efforts aspire to make it easier to repeat, compare against, and extend the work presented, ultimately serving the greater music informatics community.

CHAPTER VIII

CONCLUSION

This thesis has explored the application of deep learning methods to the general domain of automatic music description, focusing on timbre similarity and automatic chord estimation. Encouraging performance is obtained in both areas, advancing the state of art in the latter, while providing deeper insight to the tasks at hand. These observations encourage the slight reformulation of chord estimation as a representation learning, rather than a classification, problem, resulting in a high performing system with myriad practical benefits. This chapter summarizes the main contributions of this work, and offers perspectives for the future, including an assessment of outstanding challenges and the potential impact of continued research in this domain.

1 Summary

Automatic music description is at the heart of content-based music informatics research. This is necessary for problems where manual annotation does not scale, such as acoustic similarity, as well as problems where most people lack the musical expertise to perform the task well, such as transcription. While this topic is independently valuable, it would seem that progress is decelerating, and thus any efforts to correct this course must first determine why. In Chapter II, common practice in automatic music description was revisited, leading to the identification of three deficiencies worth addressing: hand-crafted feature

design is not sustainable, shallow architectures are fundamentally limited, and short-time analysis alone fails to capture long-term musical structure. Deep architectures and feature learning were shown to hold promise in music analysis tasks, evidenced both conceptually and by its growing success, motivating the exploration of deep learning in automatic music description.

At this point, it was necessary to consider what is “deep learning”, and why is this an option now? In Chapter III, the history of the field was first re-examined, showing that after an over-hyped introduction, neural networks languished through the latter part of the 20th century. This period of skepticism and disinterest gave technology time to catch up to the theory, and after a series of significant research contributions, deep learning made a triumphant return to the fore of computer science, toppling longstanding benchmarks seemingly overnight. While this has brought about a second wave of hype and interest, it also encouraging the curation of a more established theory of deep networks. As reviewed, the modern practice of deep learning consists of a handful of modular processing blocks, strung together in differentiable functions and numerically optimized to an objective function via gradient-based methods, complemented by a growing suite of practical tricks of the trade.

Having reviewed the modern core of deep learning, this work shifted focus in Chapter IV to explore these methods directly. As a first inquiry, a deep convolutional network was applied to the task of timbre similarity, achieving three goals: the model is able to learn relevant signal-level features that give rise to source identification; the resulting output space is organized in a semantically meaningful way; and the smoothness of the space is indicated by error analysis. The approach presented here also offers novel extensions to previous efforts in pairwise training, achieving extremely robust representa-

tions despite a considerable reduction in dimensionality. And perhaps most importantly, these results are obtained without the need for costly subjective pairwise ratings of content.

Whereas timbre similarity served as a relatively constrained problem, Chapter V sought to test the limits of deep learning as applied to automatic chord estimation, a well-established music informatics challenge. Competitive performance is achieved with a deep convolutional neural network, evaluated in both a conventional and large-vocabulary variant of the task. Somewhat more interestingly, rigorous error analysis reveals that efforts in automatic chord estimation are converging to a glass ceiling, due in large part to the objective formulation of an often subjective experience. The problems caused by the tenuous nature of “ground truth” annotations are exacerbated by efforts to treat chord estimation as a flat, rather than hierarchical, classification task. Therefore, the single most critical advancement facing the topic of automatic chord estimation is a re-evaluation of the task the community is attempting to solve and the data used to do so.

Despite these difficulties, the chord estimation data is leveraged to ask a slightly different question: can a model be built to automatically predict chords as guitar tablature? Therefore, in Chapter VI, again using a deep convolutional architecture, global performance statistics are improved over the general chord estimation system, while offering significant practical benefits. In addition to being a high-performing system, the fretboard estimations are immediately human-readable and thus attractive from a user experience perspective. Such a representation is also advantageous from a data collection, correction, and validation standpoints, significantly reducing the degree of prerequisite skill necessary to contribute annotations.

Finally, the various open source software artifacts developed in the course of this research are introduced and detailed in Chapter VII: **jams**, the structured music annotation format designed for multiplicity of both annotator perspective and task namespace; **biggie**, an approach to managing large collections of numerical data for training stochastic learning algorithms; **optimus**, a user-friendly library for describing and serializing trainable processing graphs; **mir_eval**, a suite of evaluation tools for benchmarking music description algorithms; and finally **dl4mir**, a common framework for the systems and results presented here.

2 Perspectives on Future Work

Based on the work performed herein and observations made in the process, this section offers several perspectives on deep learning, music informatics, and the intersection between them, in the spirit of helping guide future work.

2.1 Architectural Design in Deep Learning

Among the most common questions currently facing the application of deep learning to any problem is that of architectural design. “How many layers,” one might ask, “or how many kernels are necessary for this to work? Should I use convolutions, or matrix products, or something else altogether? And furthermore, if and when it does actually work, what on earth is it doing?” Admittedly, the combination of numerical optimization and extremely versatile functions often results in systems with opaque mid-level representations, earning deep networks the reputation as “black boxes” and the study of them a “dark art”. However, while these enigmatic functions might cause under-

standable confusion, the design of deep architectures is not necessarily devoid of intuition.

But where to start? The simplest way one might begin to explore deep learning for some problem of choice is to build some previously published algorithm and use gradient descent to fine-tune the hand crafted weights. There are countless MIR systems could be reformulated in the deep learning framework, such as onset detection, instrument identification, or pitch estimation. Most importantly, doing so eliminates the need to compare minor implementation details, like specific hyperparameters or window coefficients; just learn it and let it all come out in the wash. Additionally, introducing the process of learning to classic music informatics systems makes it easier to *combine* multiple systems to reap the benefits of model averaging. The key takeaway here is the notion that good architectures already exist for some problems, and that better performance can be obtained by using numerical optimization to expand the space of parameters considered.

Critically, these lessons and practices transcend known problems to new ones. As demonstrated with the fretboard architecture of Chapter VI, systems can be quickly constructed by appropriately constraining the behavior. Since a guitar has six strings, and each can only be active in one place, it makes sense to model each as a probability surface. That said, there is much more could be done here. Perhaps transition constraints could be imposed, such that the model would prefer common chord shapes, or positional constraints, whereby nearby frets are preferred to large jumps. In this manner, end-to-end systems can be designed and developed at a high level, and numerical optimization can be leveraged to work out the fine details. Furthermore, while learning can discover useful features that were previously overlooked or not

considered, this advantage is amplified for new challenges and applications that do not offer much guiding intuition. For tasks like artist identification or automatic mixing, it is difficult to comprehend, much less articulate, exactly what signal attributes are informative to the task and how an implementation might robustly capture this information.

Thus, deep learning, as an approach to system design, transforms the challenge from “how do I *implement* the desired behavior?” to “how do I *achieve* the desired behavior?” The nature of this advantage is illustrated by the relationship between programming in a high-level language, like Python, and a low-level one, like assembly. Technically speaking, both can be used to the same ends, but high-level languages allow for faster development of complex systems by abstracting away the minute details, like memory management or the laborious task of moving data around registers. In both cases, precise control is traded for power, leaving the developer to focus on the task at hand with greater speed and efficiency. Note, however, that abstraction doesn’t eliminate the need for sound architecture, only the need to worry about certain facets. The fundamental design challenge is the same, but operating at a higher level of abstraction allows the deep learning researcher to build bigger systems faster.

Therefore, it is worthwhile to note that music informatics researchers are quite proficient at leveraging domain knowledge, engineering acumen, and a bit of intuition to architect signal processing systems; how many principal components should one keep of a feature representation? what is a suitable window size for a Fourier transform? how many subbands should a given filterbank have? The same intuition can and should be used to design deep networks, as discussed in the learning of chroma, the design of a tempo es-

timization system, or constructing instrument-specific models. Ultimately, the process of designing a deep architecture is as arbitrary or intentional as one makes it; it's only guesswork if you're guessing.

2.2 Practical Advice for Fellow Practitioners

While the previous discussion hopefully serves to address some of the mystery inherent to deep learning, it certainly entails the disclaimer of “your mileage may vary.” The following are a handful of guidelines accumulated in the course of research; far more suggestion than direction, they have served well in practice.

1. **Data is fundamental:** The data-driven researcher will live and die by the quality of data available. It is widely held that lots of weakly labeled data will often trump a small amount of strongly labeled data. The cousin of this sentiment is once you have enough data, everything will come out in the wash. Take care to note that though this may hold for training, with caveats, the inverse is true for evaluation. Furthermore, beware obscure biases in data. Greedy optimization will happily yield bad solutions because of oversights in the curation of training data. This is particularly true of regression problems. It is possible to compensate for biased data in classification via uniform class presentation or likelihood scaling, but this can be far less obvious for continuous valued outputs.
2. **Design what you know, learn what you don't:** As mentioned, neural networks offer the theoretical promise of the universal approximation theorem, but realizing such general behavior is far from trivial. It is

therefore crucial to leverage domain knowledge where possible. This will typically take two forms: one, simplify the learning problem by removing degrees of freedom known to be irrelevant to the problem; two, constrain the learning problem to encourage musically plausible solutions. If loudness doesn't impact one's ability to recognize chords, for example, the data should probably be normalized. Music informatics researchers have a diverse background on which to draw, and this knowledge can be incorporated into the model or training strategy. Notably, curriculum learning will likely become a much larger topic in the near future, and much can be incorporated from music education and pedagogy in this process.

3. **Over-fitting is your friend:** Long heralded as the boon of deep networks, over-fitting is arguably a *good* sign, and far more desirable behavior than the inverse. Simply put, if a deep network is unable to over-fit the training data, something is likely wrong. This is often due to one or more of the following three issues; one, it is indicative of a problem with the data, e.g. the observations and targets are uncorrelated or, worse, conflicting; two, the chosen model lacks the representational power to fit the training data; or three, and most problematic, the learning problem is poorly posed and optimization is getting stuck in a bad local minima. The methods for dealing with such issues are not so well codified, but consist of data cleaning, exploring “bigger” models, unsupervised pre-training, changing the loss function, etc.; that said, the efficacy of such approaches will vary case by case.

4. **Get good mileage out of greedy optimization:** Gradient descent

and other such greedy methods are certainly prone to bad local minima, but it is not impossible to take active measures to discourage unfortunate solutions. Additionally, it may be easier to define the kinds of things a model *shouldn't* do than the things it should. For example, a fretboard prediction network could incorporate a penalty whereby “unplayable” chord shapes incur significant loss to help keep outputs in a realistic space.

5. **The gap between “real” and synthetic music is closing:** As more modern music transitions to a digital environment, the difference in quality between a real sound recording and one synthesized for research purposes is converging to zero. Generally speaking, samplers, synthesizers, and other music production software are underutilized in data-driven music research. These high quality tools can also be used for data augmentation to make algorithms robust to irrelevant deformations, such as perceptual codecs, background noise, tempo modulation, or pitch shifting. By generating an unprecedented amount of realistic training data, can we functionally solve tasks such as onset detection, pitch detection, or tempo estimation?

2.3 Limitations, Served with a Side of Realism

As some of its forebears recognize and advocate, especially those who persevered through the first AI winter, it is crucial to maintain reasonable expectations for what can be achieved with deep learning. Shown in Figure 47, it is interesting to consider that, to date, the path of deep learning has roughly followed Gartner’s hype cycle for emerging technologies: after a most promising

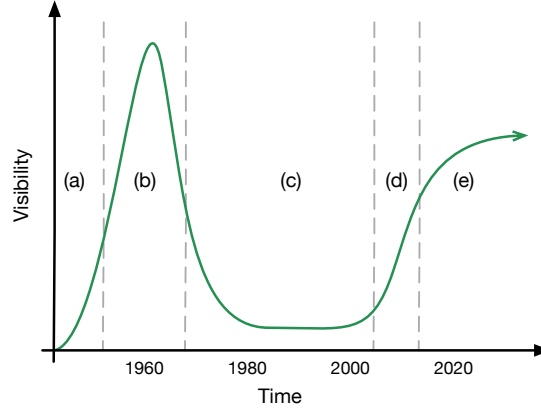


Figure 47: Gartner Hype cycle, applied to the trajectory of neural networks, consisting of five phases: (a) innovation, (b) peak of inflated expectations, (c) trough of disillusionment, (d) slope of enlightenment, and (e) plateau of productivity.

start followed by a period of disinterest, neural networks have sharply returned to prominence.

It goes almost without saying that excitement and interest in deep learning is spiking across computer science, in academia and industry alike. Research groups are forming based primarily on deep learning, it is being used to win increasing number of data science competitions, and the topic has become common fodder for popular science articles and interviews. With all the success and attention, it is easy to get carried away in thinking that deep learning is the proverbial “magic bullet”, that it might topple all problems in due time.

The reality, however, is far more modest. Deep learning *is* indeed several important things. It is a powerful approach to non-linear system design. Deep networks make fantastic models of physical phenomena, and could have profound use in the fields of acoustics, immersive audio, or amplifier modeling. It is extremely versatile, and can be easily adapted in application-specific ways that other powerful machines, such as SVMs, cannot. And, given significant

gains in compute power, the combination of architectural flexibility and numerical optimization makes it an arguably efficient research strategy, at least more so than graduate students manually tuning parameters.

That said, there are a few important things deep learning is not. It is by no means the best answer to every problem. Deep learning is, in its current forms, still relatively data hungry and often computationally expensive. Even in the case of unsupervised pre-training, sufficient volumes of realistic data may not be trivial to obtain, as in the case of accelerometers or gyroscopes. To the latter point, any performance gains obtained with a deep network could be effectively negated by disproportionate increase in processing time. Both of these limitations are like to become less important over time, but remain relevant today. More conceptually, albeit somewhat contentiously, nor can the modern flavors of deep learning be called “intelligent”; echoing the ghost of Searle, a system may certainly *behave* intelligently without truly *being* so. Despite the imagery evoked by metaphors and figurative language that pervade the field, deep learning has as much in common with humanity as a player piano, and learning is merely a means of programming in a data-driven manner. This is not to say that deep learning can or will not lead to such breakthroughs, but care should be taken when differentiating metaphor from reality.

What should one make of deep learning then? Suffice it to say that deep learning is just another tool—a powerful tool, but a tool nonetheless—to be included in the toolbelt of the information science practitioner. Similar to the trajectories of other, now standard, methods, such as principal components analysis, Gaussian mixture models, or support vector machines, deep learning is settling into the final stage of its hype cycle, the point at which it becomes

a means to solve a problem. *Is deep learning some magic bullet?* Of course not. *Is it intelligent?* Hardly. But is it useful? Can it accelerate the process of system design and implementation? Can it allow the clever researcher to quickly vet ideas and develop complex, robust systems? The answer to all of these is *yes*.

2.4 On the Apparent Rise of Glass Ceilings in Music Informatics

One of the motivating factors of this work was to understand and potentially address the increasing prevalence of diminishing returns in various music description tasks, like chord estimation, genre classification, or mood prediction. The main hypothesis resulting from an initial survey was the idea that common approaches to system design were inadequate, and another approach to system design, i.e. deep learning, might afford significant performance gains. However, one of the most significant outcomes of this work is in some sense the most unexpected: subtle deficiencies in methodology may be contributing as much or more to unsatisfactory results than the algorithms or approaches used to achieve them.

This finding reflects a small but growing trend in music informatics of critically assessing how the scientific method is applied to machine listening tasks, with meta-analyses of genre recognition (Sturm, 2014a), rhythmic similarity (Esparza, Bello, & Humphrey, 2014), and music segmentation (Nieto, 2015), to name a few. Looking to the intersection of these areas, the research methodology of automatic music description consists of five basic steps:

1. Define the problem.
2. Collect data.

3. Build a solution.
4. Evaluate the model.
5. Iterate.

With this method as a common underpinning, the evolution of content-based music informatics unfolds logically. Though a young field, the majority of current research has converged to a handful of established tasks, as evidenced by those conducted at MIREX. Labeled music data is notorious difficult to amass in large quantities, but resources have grown steadily for those well-worn problems. In cases where it has not, researchers are faced with one of two options: curate the data themselves, or adapt an existing dataset to their problem. Having developed a solution, it is necessary to benchmark a proposed algorithm against previous efforts. However, to make such comparisons fairly, it is typically necessary to compute the same metrics over the same data, as the systems themselves are seldom made public. Thus, most research efforts today focus almost exclusively on (3) and (5), adopting or otherwise accepting the other three.

It is critical to note, though, that these other methodological stages — problem formulation, data curation, and evaluation— have developed naturally over time at the community level, based not on globally optimal design but rather a combination of evolving understanding, inertia, and convenience. At this point, it serves to return to an initial question posed by this work: why *are* music description systems yielding diminishing returns? The findings of this work, particularly in the space of automatic chord estimation, corroborate the growing trend that perhaps the biggest problem facing content-based music informatics is one of methodology.

With this in mind, reconsider the case of automatic chord estimation. What is the scope of the problem being addressed? “Can an agent provide acceptable chord transcriptions?” is a very different question from “Can an agent reproduce *these* chord transcriptions?” Analysis of the Rock Corpus transcriptions showed that comparing the outputs of two expert musicians can achieve a “yes” and “no” respectively. Does the data reflect these requirements? Chord annotations consist of single perspectives from several anonymized annotators. It is doubtful that all annotators are using the chord labels the same way. How do we know when the problem is solved? Does weighted chord symbol recall with different comparison rules correspond to subjective experience? Not all substitutions are equal, as the distance in harmonic function between a I:7 and a I is quite different from a V:7 and a V, for example.

Understandably, it is easy to lose sight of the fact that research is not just the process of iterative system development, but the entire arc of the scientific method. At this point in the trajectory of music informatics, it is conceivable that several well-worn tasks could use a reassessment of what constitutes methodological best practices. This is hardly a novel realization, but one that warrants greater awareness within the music informatics community. It is necessary, but ultimately insufficient, to tirelessly pursue better solutions; we must be as diligent in our pursuit of better problems.

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