An Interpretable Machine Learning Framework for Modeling High-Resolution Spectroscopic Data*

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ABSTRACT

Comparison of échelle spectra to synthetic models has become a computational statistics challenge, with over ten thousand individual spectral lines affecting a typical cool star échelle spectrum. Telluric artifacts, imperfect line lists, inexact continuum placement, and inflexible models frustrate the scientific promise of these information-rich datasets. Here we debut an interpretable machine-learning framework blasé that addresses these and other challenges. The semi-empirical approach can be viewed as "transfer learning"-first pre-training models on noise-free precomputed synthetic spectral models, then learning the corrections to line depths and widths from whole-spectrum fitting to an observed spectrum. The auto-differentiable model employs back-propagation, the fundamental algorithm empowering modern Deep Learning and Neural Networks. Here, however, the 40,000+ parameters symbolize physically interpretable line profile properties such as amplitude, width, location, and shape, plus radial velocity and rotational broadening. This hybrid data-/model- driven framework allows joint modeling of stellar and telluric lines simultaneously, a potentially transformative step forwards for mitigating the deleterious telluric contamination in the near-infrared. The blasé approach acts as both a deconvolution tool and semi-empirical model. The general purpose scaffolding may be extensible to many scientific applications, including Precision Radial Velocities, Doppler Imaging, chemical abundances for Galactic archaeology, line veiling, magnetic fields, and remote sensing. Its sparse-matrix architecture and GPU-acceleration make blasé fast. The open-source PyTorch-based code blase includes tutorials, Application Programming Interface (API) documentation, and more. We show how the tool fits into the existing Python spectroscopy ecosystem, demonstrate a range of astrophysical applications, and discuss limitations and future extensions.

Keywords: High resolution spectroscopy (2096), Stellar spectral lines (1630), Astronomy data modeling(1859), GPU Computing (1969), Calibration (2179), Radial Velocity (1332), Maximum likelihood estimation (1901), Deconvolution (1910), Atomic spectroscopy (2099), Stellar photospheres (1237)

1. INTRODUCTION

1.1. Spectral fitting past and present

Tens of thousands or more individual spectral lines give rise to a sea of undulations that imbue each stellar spectrum with its characteristic appearance. The identification and understanding of these lines have defined a large category of astrophysics over the last century. The field grew from by-eye catalogs of stellar templates (Cannon & Pickering 1901) to quantifying the role of atomic ionization balance (Payne 1925), to modern synthetic forward models including millions or billions of lines (e.g. Husser et al. 2013; Marley et al. 2021; van

den Bekerom & Pannier 2021). As technology has improved, our data and models have become more voluminous, more precise, and more complicated. The mere act of comparing models to observed spectra can now resemble a computational statistics challenge as much as a scientific one. Here we introduce a new machine-learning-based framework blasé aimed at solving computational, statistical, and scientific challenges associated with data-model comparisons for modern astronomical spectroscopy.

The metaphorical holy grail of astronomical spectroscopy would be a function that takes in an observed stellar spectrum and reports back the position, amplitude, width, and shape of all of its spectral lines, automatically, accurately, and precisely. The function would go further. It would report back the systemic radial

^{*} Open source code at https://github.com/gully/blase

velocity (RV) and rotational broadening $(v \sin i)$ and fundamental stellar properties, including T_{eff} , $\log q$, and [Fe/H]. Finally, the function would—in a feat of artificial intelligence—provide what it believes to be the interpretable generating function that produced this data in the first place, so that we may gain insights on future examples of this or other stars. Solving this problem is hard, for at least four reasons. First, the spectral lines may overlap, and so the assignment of one line may be partially degenerate with the assignment of some other adjacent line. Second, extremely wide line wings blend into the continuum, such that the placement of the continuum level may become ill-defined. Third, the extent of line blending and realized line shape depends strongly on the spectral resolution of the spectrograph, the rotational broadening of the star, and possibly the instrumental configuration at the time of observation. Finally, telluric absorption lines commingle with the astronomical spectral lines of interest, censoring some spectral regions entirely, or partially confounding other lines with chance alignments.

Addressing these and other challenges forms the backbone of *spectral calibration*, an increasingly valuable specialty as the deficits in our models become intolerable with greater data quantity and quality. Luckily, many scientific applications in astrophysics do not need the technically demanding noise-free template, nor catalog-of-all-spectral-lines. A few lines suffice. For those applications, human inspection of isolated lines and semi-automated equivalent width determination have been—and will remain—adequate.

But many new and important questions in the fields of stars and exoplanets aspire to reach the margins of what the entire dataset can inform. In particular, data from high-grasp échelle spectrographs possess simultaneously high spectral resolving power and high bandwidth, yielding tens of thousands or possibly millions of independent spectral resolution elements for each star, substar, or exoplanet. Scientific applications that seek to gain signal by "stacking" spectral lines or cross-correlating with templates can hypothetically gain huge boosts in the accessible signal-to-noise ratio compared to a single or few lines. Most manual and semi-automated methods cannot take advantage of the entire spectral bandwidth, or rely on exact knowledge of the underlying templates and may fail to achieve the hypothetical promise of these high-bandwidth spectrographs (Hood et al. 2020).

For example, exoplanet cross-correlation spectroscopy (Birkby et al. 2013) hinges on accurate molecular spectral templates to detect and characterize the atmospheres of exoplanets. Imperfections in these templates

can mute the perceived signal strength of these atmospheric features (Hoeijmakers et al. 2015).

In extreme precision radial velocity (EPRV) applications, cross-correlation methods work (Dumusque 2018a), but have many limitations (Zhao et al. 2022). Among the many such limitations, one pernicious noise floor stands out as enigmatic: telluric mitigation. Many practitioners today simply mask these telluric regions, yet micro-tellurics still inject variance into the spectrum that cannot be easily accounted for with existing methods. Instead, a robust accounting of telluric absorption at the cm/s level may require joint modeling of the star and the Earth's atmospheric absorption before convolution with an instrumental kernel. This telluric joint modeling capability does not yet exist at a precision that can meet these strenuous demands.

In the case of Doppler imaging, an accurate underlying spectral template is needed to detect longitudinally symmetric structures (Vogt & Penrod 1983; Luger et al. 2021) such as polar spots (Roettenbacher et al. 2016) or zonal bands (Crossfield et al. 2014; Apai et al. 2021). There exists a nearly circular reasoning: we need to know the extent of line profile perturbations to reveal the underlying spectral template, but we need the underlying spectral template to estimate the line profile perturbations. The approaches introduced here offer a path forward on these long-standing friction points.

1.2. Automatic differentiation technology

Existing open-source frameworks have overcome some of these challenges, or have been purpose-built for specialized applications. These frameworks include ROBOSPECT (Waters & Hollek 2013), specmatch (Petigura 2015), specmatch-emp (Yee et al. 2017), wobble (Bedell et al. 2019), starfish (Czekala et al. 2015), sick (Casey 2016), psoap (Czekala et al. 2017), FAL (Cargile et al. in prep), CHIMERA (Line et al. 2015), the Cannon (Ho et al. 2017), MOOG (Sneden et al. 2012), MOOGStokes (Deen 2013), MINESweeper (Cargile et al. 2020), and recently ExoJAX (Kawahara et al. 2022). The designs of these frameworks necessarily have to make a choice in the bias-variance tradeoff: is the tool more data-driven or more model-driven? The statistical tradeoff can be viewed as a concession in physical self-consistency for model flexibility: more or fewer parameters; more accurate or more precise.

A key new enabling technology breaks these classical tradeoffs in data-model comparisons for astronomical spectroscopy. Automatic differentiation ("autodiff" or "autograd", Gunes Baydin et al. 2015; Maclaurin 2016) and its affiliated backpropagation algorithm (Kelley 1960; Linnainmaa 1976; Rumelhart et al. 1986;

Dreyfus 1990) has revolutionized machine learning and neural network architecture design, and is increasingly applied in astrophysical data analysis contexts, e.g. kernel phase coronography with poppy (Pope et al. 2021), and exoplanet orbit fitting with exoplanet (Foreman-Mackey et al. 2021a). Of the spectroscopy frameworks mentioned above, the TensorFlow-based (Abadi et al. 2015) wobble and the JAX-based (Bradbury et al. 2018) ExoJAX employ autodiff technology. wobble treats each pixel as a tunable control point, producing $\sim 10^5$ parameters for a modern stellar spectrum. The ExoJAX framework has only ~dozens of tunable parameters that describe the fundamental physical properties controlling a brown dwarf atmosphere. These two autodiff-aware frameworks span the extreme ends of non-parametric and parametric modeling for spectroscopy.

In this paper, we show that autodiff-aware semiempirical models offer an appealing middle ground: informed by self-consistent models but refined with data. This sweet spot in the bias-variance tradoff can be thought of as a hybrid data-and-model driven approach. The algorithm presented here focuses on modeling the spectra of stars and brown dwarfs. Existing stellar models (e.g. Husser et al. 2013) and substellar models (e.g. Marley et al. 2021) laboriously solve for a self consistent thermal structure in the atmosphere given the copious opacity sources that themselves depend on temperture and pressure. Here we build upon that hard work by cloning pre-existing synthetic stellar or substellar models (Section 2), and optionally by cloning models of Earth's atmospheric "telluric" absorption (Section 3). We introduce the interpretable forward-model design and its PyTorch-based (Paszke et al. 2019) implementation, blasé. In Section 5 we describe how to adapt both stellar and telluric cloned models simultaneously, using a transfer-learning step. We obtain semi-empirical models by comparing to real-world échelle data in Section 6. Finally, we discuss perspectives on how to think of blasé (Section 7) and chronicle many conceivable extensions for unlocking new science (Section 8).

2. METHODOLOGY I: CLONING STELLAR SPECTRA

2.1. Overall Architecture and Design Choices

We start with a high resolution pre-computed synthetic stellar or substellar model spectrum, $S_{\rm abs}(\lambda)$ at its native resolution sampling and with its original absolute physical flux units. The procedure that follows is largely agnostic to the exact details of how this spectrum was made, or what physics or chemistry it may represent. For the purposes of this paper, we will showcase examples from two well-known families of precom-

puted synthetic astronomical spectra: PHOENIX (Husser et al. 2013) for stellar spectra ($T_{\rm eff} \in [2300, 10000]~K$) and Sonora (Marley et al. 2021) for substellar spectra ($T_{\rm eff} \in [200, 2300]~K$). The algorithms in the framework may also work for precomputed synthetic spectra of reflected light exoplanets, supernovae, galaxies, or even further afield such as laboratory physical chemistry, plasma physics, materials science, or remote sensing.

We place mild demands on the precomputed spectra. They should have sporadic regions of discernable continuum devoid of lines, and the continuum shape should vary smoothly in wavelength. The spectral lines or pseudo-lines should be resolved (and not sub-sampled). We suspect most stellar spectra meet these criteria, except for the coolest M dwarfs and brown dwarfs. The method can handle spectra without discernable continua, with some fine-adjustments to the preprocessing steps. We truncate the red and blue limits of the precomputed synthetic spectrum to match a highbandwidth echelle spectrograph, extended with a buffer at the edges of size $\pm \Delta \lambda_{\text{buffer}}$, chosen to account for plausible radial velocity and rotational broadening of real stars. A generous buffer of $v \sin i < 500 \text{ km s}^{-1}$ and $|RV| < 500 \text{ km s}^{-1} \text{ yields a typical buffer of about}$ 30 Å.

The choice of limiting the bandwidth to a region of interest around a single echelle spectrograph bandwidth stems from computational constraints. In principle, there is no fundamental limit to the bandwidth one could clone with the method presented here, up to and including the entire precomputed synthetic spectral model bandwidth. We adopt the exact native wavelength sampling with no smoothing or interpolation, yielding a wavelength vector λ_S with length N_S equal to the number of pixels within the extents of our region of interest including the buffers.

At this stage, we have the choice of whether to work in linear or log scale flux units. Adopting the log of the flux would ensure that the cloned model possesses only positive flux values, a desirable trait of any physical spectral model. We have implemented both modes in blasé, allowing users to choose their preference. We only narrate the linear flux unit description in the main text of this document for the sake of clarity, and since most practitioners may tend to think of flux in terms of linear flux units. The data-model comparison step will always take place in linear flux units, so the only operational difference is the behavior for deep and saturated lines. Appendix A lists the equations adjusted to log flux units.

2.2. Initialization

We initialize the cloned model with a series of preprocessing steps. We divide the entire spectrum by a black body $B(\lambda_S)$ of the same effective temperature T_{eff} as the model template. The resulting signal usually still has smooth wiggles around the continuum. An optional continuum flattening step ensures that subsequent spectral line finding steps get applied uniformly. This high-pass filtering step should be set to capture the genuine spectral shape, without over-fitting broad line wings such as those in deep Hydrogen and sodium lines. Any high-pass filter will work, a Gaussian Process approach would be ideal. Instead we apply a simple and familiar heuristic: fit a polynomial $P(\lambda_S)$ to a few continuum peaks and divide out the trend.

The result should be a flattened "continuumnormalized" spectrum familiar to practitioners in highresolution spectroscopy, with the continuum level close to unity. It is this spectrum that will serve as the centerpiece of subsequent training steps. We, therefore, drop any subscript and simply refer to this flattened spectrum as S:

$$S = S_{\rm abs}/B/P \tag{1}$$

where the division indicates element-wise division of these arrays or "vectors" of flux values.

We emphasize that a recreation of the unvarnished input spectrum—if desired—can be obtained by multiplying the continuum-flattened signal by the "perturbed black body", $B(\lambda_S) \odot P(\lambda_S)$, that symbolizes the black body modulated by continuum opacity or broad-band radiative transfer effects. This smooth spectrum may be useful for applications that need to keep track of broad-band flux, such as low-resolution spectra, or regions with molecular band heads. The "perturbed black body" continuum model contains $n_{\text{poly}} + 1$ fixed-butpossibly tunable lookup parameters, plus the fixed input $T_{\rm eff}$. For most practitioners these terms serve as nuisance parameters and are perfunctorily discarded.

Next, we identify the spectral lines. We apply a local-minimum-finding algorithm (Virtanen et al. 2020) on the spectrum by defining a prominence threshold $P_{\rm rom} \in (0.005, 0.02)$. This threshold dictates the number of lines that will be modeled: a lower prominence finds more, weaker lines, and a larger prominence finds fewer, deeper lines. The prominence algorithm successfully finds lines that reside on top of broad line wings, or unresolved band heads provided that the individual lines exceed the prominence threshold in their local region. The number of lines N_{lines} depends on the bandwidth, prominence, and the intrinsic properties of the

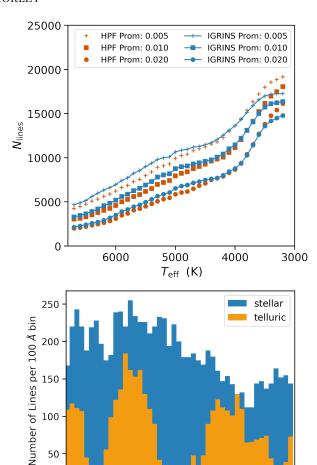


Figure 1. Scaling of prominence and density of spectral lines. Top: Number of lines versus effective temperature for PHOENIX models truncated to IGRINS (blue, connected points) and HPF (red, free-standing points) bandwidths, for different prominence thresholds of 0.02, 0.01, and 0.005. Bottom: Number of lines per 100 Å wavelength bin for stellar (blue, upper envelope of steps) and telluric (orange, lower envelope of steps), illustrated for a $T_{\text{eff}} = 4700 \text{ K}$, $\log g = 4.5$ PHOENIX model and a T = 290K, relative humidity of 40% TelFit model.

10000

 λ (\mathring{A})

11000

12000

50

0

9000

input spectrum, principally effective temperature and metallicity.

For this paper, we illustrate examples for two échelle spectrographs with particularly large spectral grasp: the Habitable Zone Planet Finder (HPF, Mahadevan et al. 2014) on the Hobby-Eberly Telescope at McDonald Observatory in Fort Davis, Texas; and the Immersion Grating Infrared Spectrograph (IGRINS, Park et al. 2014) currently on the Gemini South Telescope on Cerro Pachón in Chile. The R = 55,000 HPF has a native bandwidth of 8079 - 12785 Å, which we expand

to 8049 - 12815 Å including the edge buffers. IGRINS has two cameras for H and K band, with the combined spectrum spanning 14267 - 25217 Å including the edge buffers and the region in-between the two cameras, all at a resolving power of (R = 45,000). The spectrograph acquisition, reduction, and post-processing steps yield data $D(\lambda_D)$, where λ_D is the wavelength vector at the instrumental resolution and sampling of each instrument, generally much coarser than the resolution and sampling of the precomputed synthetic spectra. The data wavelength vector may also contain gaps between échelle orders, whereas the precomputed wavelength coordinates are usually contiguous. HPF may have up to $2048 \times 28 = 57,344$ pixels, and IGRINS has typically about 75,000 pixels, after common trimming of noisy edge pixels and unusable telluric regions. Meanwhile, the HPF-truncated model spectra have $N_s = 335,849$ native resolution samples, comparable to the IGRINStruncated model spectra, $N_s = 330,052$.

Figure 1 shows how the number of detected lines $N_{\rm lines}$ scales with effective temperature and prominence threshold $P_{\rm rom}$ for the PHOENIX grid, truncated to the bandwidths-plus-buffers for HPF and IGRINS. We see between about 2,000 and 20,000 lines depending on the $T_{\rm eff}$ and $P_{\rm rom}$. HPF and IGRINS have a comparable number of lines, and halving the prominence increases the number of lines by about 20-30% in these ranges. The number of lines monotonically increases towards cooler effective temperatures.

So far we have only one piece of information about the spectral lines: their location. Next, we derive coarse properties about each detected peak: its amplitude and width, again using the prominence algorithms implemented in scipy (Virtanen et al. 2020).

There does not exist a general-purpose, single-shot algorithm for obtaining the lineshape in the presence of overlapping spectral lines: where do the wings of one line begin and the wings of another adjacent line end? We, therefore, do not attempt to determine anything about the lineshape at this stage and instead assume that the lines resemble a Voigt profile, with a guess width about equally split between Lorentzian and Gaussian.

2.3. The blasé Stellar Clone Model

We have now arrived at the blasé clone model $S_{\text{clone}}(\lambda_S)$ for a flattened synthetic spectrum S: it is the cumulative product of transmission through the sea of all overlapping spectral lines:

$$\mathsf{S}_{\text{clone}} = \prod_{j=1}^{N_{\text{lines}}} 1 - a_j \mathsf{V}_j \tag{2}$$

where V_j is the Voigt profile $V(\lambda_S - \lambda_{c,j}, \sigma_j, \gamma_j)$ with Gaussian standard deviation σ , Lorentzian half-width γ , at line center position λ_c , for the j^{th} spectral line. The amplitude a is always expected to be positive for absorption lines.

The Voigt profile $V(\lambda, \sigma_j, \gamma_j)$ can be computed in exact closed-form using the Voigt-Hjerting function (Hjerting 1938) as the real part of the complex Fadeeva function (e.g. Zaghloul & Ali 2011). Evaluation of the Fadeeva function can be computationally costly, and so approximate forms may be desirable. Here we adopt the pseudo-Voigt approximation (Ida et al. 2000).

2.4. Goodness of fit metric

The model evaluated with its coarse initial values would have terrible performance: it would only vaguely resemble the synthetic spectral model, with up to $\pm 50\%$ undulations from the inexact assignment of widths, lineshapes, and amplitudes. Instead, we tune the parameters of the model, starting from these coarse initial values. This model has between $N_{\text{lines}} \times 1$ and $N_{\text{lines}} \times 4$ free parameters, depending on how many of the 4 physical line properties you wish to tune. We default to fitting three parameters per line, where the center wavelength is held fixed and the amplitude, width, and lineshape are allowed to vary. We minimize a scalar "goodness-of-fit" metric, aka loss scalar \mathcal{L} , chosen as the mean squared error (MSE), which is proportional to χ^2 , the sum of the squares of the residual vector $R \equiv S - S_{\rm clone}$ but has no notion of per-pixel noise since the precomputed synthetic spectrum has no uncertainty:

$$\mathcal{L} = \sum_{i}^{N_S} (S_i - S_{\text{clone},i})^2 = \mathsf{R}^{\mathsf{T}} \cdot \mathsf{R}$$
 (3)

As seen in Figure 1, the number of lines can exceed 7,000, meaning the clone model has over 21,000 free parameters. Fitting more than about 300 parameters is difficult with conventional optimizers that rely on minmizing a single scalar loss: the search space becomes too large. Here we employ a variant of Stochastic Gradient Descent (SGD), a transformative optimization technique that can scale to a virtually unlimited number of parameters (Ruder 2016). This technique computes the derivative of the loss scalar with respect to each of the parameters, the so-called Jacobian: $\vec{\nabla} \mathcal{L} = (\frac{\partial \mathcal{L}}{\partial a_j}, \frac{\partial \mathcal{L}}{\partial \sigma_j}, \cdots, \frac{\partial \mathcal{L}}{\partial \gamma_j})$. The Jacobian indicates how the MSE would decrease with a change in the parameter-of-interest, or put simply "which way and by how much" you have to change each individual line property to get a better fit.

The optimizer updates the a_j, σ_j, γ_j parameters by a small fraction of the Jacobian—called the learning rate

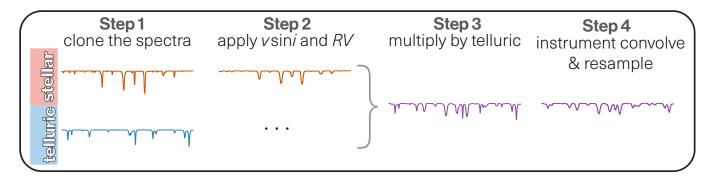


Figure 2. Visual flowchart of the blasé forward model. Step 0 (not shown) is to choose a precomputed synthetic stellar spectrum—and, optionally, a precomputed synthetic telluric spectrum—with physical properties close to the target and observing conditions. Both the stellar and telluric spectra get cloned (Step 1). The stellar model is warped to its extrinsic properties (Step 2), and then the stellar and telluric models get multiplied together (Step 3). Next, this joint model is convolved with an instrumental kernel and resampled to the wavelength coordinates of the data spectrum (Step 4). It is this forward model that gets directly compared to the observed spectrum (not shown).

(LR)—towards the direction that would improve the fit, for all parameters *simultaneously*. The Jacobian is calculated behind the scenes with automatic differentiation implemented as the so-called backpropagation algorithm or simply "backprop" (Gunes Baydin et al. 2015). We choose the PyTorch framework that computes these Jacobians efficiently for all of the mathematical primitives in our blasé implementation (Paszke et al. 2019).

It is this ability to automatically compute Jacobians that sets PyTorch (and JAX and TensorFlow) apart. These frameworks give exact Jacobians, instantaneously, for free (or cheap). Without exact Jacobians a conventional optimization step only obtains one piece of information: how much the overall loss changed. With exact Jacobians, we obtain $N_{\rm params}$ pieces of information for each evaluation of the forward model. The power of gradient descent becomes transformative as the number of parameters grows into the tens of thousands, as is the case for $blas\acute{e}$.

2.5. GPU and Autodiff specific considerations

Forward modeling with tens of thousands of physics-informed parameters may seem like such a significant paradigm shift that it can feel too good to be true. In this section we introduce the non-negligible architectural design tradeoffs that arise when adopting an autodiff framework—such as PyTorch—for physics based forward modeling.

First, we make a few tweaks to the implementation for numerical purposes. We enforce that all Gaussian and Lorenzian widths are positive by tuning the natural log of the widths, then exponentiating them before inclusion in Equation 2. Based on the minus sign in Equation 2, an amplitude a_j could hypothetically take on either positive values (flux loss, absorption) or negative values (flux gain, emission). For now we focus on

photospheric absorption lines—as opposed to, say, chromospheric emission lines—and therefore enforce all the amplitudes to be positive by tuning $\ln a_j$ and then exponentiating in the same strategy as above. Emission lines—if desired—could be included with a mere sign flip to Equation 2. Lines that can manifest either in absorption or emission could hypothetically relax the natural log pre-processing step for isolated lines. We narate only the absorption scenario moving forward.

The autodiff machinery has a convenient way to set which parameters are held fixed and which are iteratively fine-tuned. One simply disables the autodiff flag for the fixed parameters: we set the requires_grad=True property for any PyTorch tensor that we want to vary. This design allows us to easily explore whether, say, allowing the λ_c parameter to vary significantly improves the fit.

The computational bottleneck occurs at the evaluation of Equation 2, which can be viewed as having an $N_{\text{lines}} \times N_S$ matrix \bar{F} assembled by stacking each Voigt absorption profile $V_j(\lambda_s)$ on top of each other:

$$\begin{pmatrix} 1 - a_1 \mathsf{V}_1(\boldsymbol{\lambda}_s) \\ 1 - a_2 \mathsf{V}_2(\boldsymbol{\lambda}_s) \\ \vdots \\ 1 - a_{N_{\text{lines}}} \mathsf{V}_{N_{\text{lines}}}(\boldsymbol{\lambda}_s) \end{pmatrix}$$
(4)

An element of this matrix, F_{ji} , will have the flux value for a given j^{th} line at a given i^{th} wavelength coordinate. Equation 2 performs a type of matrix contraction, turning an $N_{\text{lines}} \times N_S$ matrix into a length N_S row vector. The number of Floating Point Operations (FLOPS) scales with the number of entries in this matrix. So we face a tradeoff of wanting to make the matrix large for accuracy and small for computational expedience.

We can rewrite Equation 2 as a sum by taking the log of both sides and dropping in this \bar{F} matrix:

$$\ln \mathsf{S}_{\mathrm{clone}} = \sum_{j=1}^{N_{lines}} \ln F_{ji} = \mathbf{1} \cdot \ln \bar{\mathbf{F}}$$
 (5)

where **1** is a $1 \times N_{\text{lines}}$ row vector of all-ones. We reëmphasize that—in its current form—each spectral line has to be painstakingly evaluated across the entire spectral bandwidth. Efficient GPU algorithms exist for voluminous matrix manipulations such as this one, so this voluminous computation will proceed as quickly as possible on modern machines. In particular, the proprietary CUDA architecture for NVIDIA® GPUs contains Tensor cores with specialized matrix math. The chief bottleneck occurs when the storage of the $ar{F}$ matrix exceeds the available RAM of a GPU or CPU: the computation will fail with an "Out of Memory" exception. Modern NVIDIA GPUs have 8–40 GB of RAM, which translates roughly to a few thousand spectral lines across $\sim 300,000$ pixels. It is generally not possible to construct Equation 5 in its entirety in one-fell-swoop, even on a GPU. A remedy is needed.

2.6. Sparsity

The $\ln \bar{F}$ matrix is sparsely populated: most of the entries far from the line center are vanishingly close to zero. Here we take advantage of that mostly empty matrix using the mathematics of sparse matrices (Saad 2003).

We retain a relatively small number of pixels $N_{\rm cut}$ adjacent to the line center. Setting this wing cut produces a speedup by a factor of $\frac{N_S}{N_{\rm cut}}$, which can exceed $100\times$ for wide bandwidth spectra. The choice of $N_{\rm cut}$ is nuanced. It should be set large enough that truncation effects are not seen for the broadest lines. But even more, $N_{\rm cut}$ has to be future-proofed for Dopplershifting. Extreme Doppler shifts could hypothetically send line cores entirely outside the extents of N_{cut} if set too low. We therefore typically set wingcuts comparable to the buffer size $2\Delta\lambda_{\text{buffer}}$, even though most weak lines only perceptibly affect < 1 Å. We coerce all wing cuts to be the same number of pixels, typically 6000 pixels, $\sim 30-60$ Å for PHOENIX, with the middle pixel being at the line center position, and about 3000 pixels to the red and blue side of the line. We populate a new approximate sparse matrix $\ln \hat{F}$ with only these 6000 pixels per line and assume zeros everywhere else.

The remapping of the sparse matrix can be pictured as having shifted all lines to the center of this new matrix $\hat{\mathbf{F}}$, visualized pictorially in Equation 6. The algorithmic machinery keeps track of each (i, j, F_{ji}) trio of coordinates and flux values.

$$\bar{F} = \begin{pmatrix} \dots & & & \dots & & \\ \dots & & & & \dots & \\ \vdots & & & & \dots & \\ \vdots & & & & \dots & \\ \bar{F} \mapsto \hat{F} = \begin{pmatrix} & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & &$$

Sparse matrix methods generally support an operation known as *coalescing*, which sums values with repeated indices. Each pixel may get computed about ~ 100 times in this sparse implementation, which is about $70\times$ faster than each pixel getting computed $N_{\rm lines} \sim 7000$ times in the dense approach. Efficient algorithms for assembling and coalescing sparse matrices exist in PyTorch. Some GPUs now support additional hardware-acceleration of sparse matrices, providing even greater speedups.

2.7. Optimization and training

We use the Adam optimizer (Kingma & Ba 2014) with a typical learning rate $LR \in (0.005, 0.1)$ and all the defaults for PyTorch v1.11. We defined the number of training epochs $N_{epoch} = 100-10,000$ depending on the application. The user can optionally monitor a live view of the training progress with Tensorboard (Abadi et al. 2015) to gain an intuition for the training efficiency.

Figure 3 shows a portion of a PHOENIX spectrum cloned with blasé. The 1000 epochs of training took 56 seconds on an NVIDIA® RTX2070 GPU with PyTorch v1.11, CUDA v11.1, and Intel® Core™ i7-9750H CPUs at 2.60GHz, with all tensors as FP64. The same computation on a 2020 M1 Macbook Air took 1^h25^m with PyTorch v1.9, $90\times$ slower than the GPU counterpart.

We store the model parameters to disk and refer to the entire collection of parameters as a *pre-trained model*. More specifically this fine-tuned model represents an evaluable and interpretable clone of the original static pixel-by-pixel flux values.

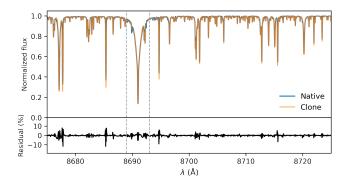


Figure 3. PHOENIX spectrum cloned with blasé. This $T_{\rm eff}=4700~{\rm K},~{\rm log}~g=4.5$ solar metallicity model has 9,028 individual cloned spectral lines, each with 3 tuned parameters. The pictured 50 Å chunk contains 121 spectral lines and represents about 1% of the entire spectral bandwidth that was cloned. Some flaws can be seen near the cores of deep lines, or wings of broad lines.

3. METHODOLOGY II: CLONING TELLURIC SPECTRA

Ground-based near-IR échelle spectra possess thousands of depressions attributable to molecular line absorption in Earth's atmosphere. These telluric lines hamper the unbiased interpretation of échelle spectra, so some treatment plan is needed. Often the regions of known, deep tellurics are simply discarded. In other cases, the lines are modeled with first principles lineby-line radiative transfer (e.g. TelFit, Gullikson et al. 2014; Clough et al. 2005) or through data-driven means (e.g. wobble, Bedell et al. 2019). The most demanding EPRV applications require a precision characterization of telluric lines that the astronomical community has not yet been able to achieve, and that may rival even the abilities of Earth Science practitioners. A hybrid data-/model- driven approach was among the chief recommendations of the Telluric Hack Week Workshop¹ aimed at improving mitigation of the atmosphere's deleterious effects (David W. Hogg, priv. comm.). The blasé framework achieves a key milestone by introducing a hybrid approach to tellurics.

3.1. The blasé Telluric Clone Model

We start with a precomputed synthetic telluric model, T with associated wavelength coordinates λ_T . We employ a TelFit model, though any precomputed synthetic telluric model will work, such as MOLECFIT (Smette et al. 2015). The TelFit model does not contain any continuum sources of opacity, so we can skip the flattening procedure described in Equation 1. We

orchestrate the same initialization and line finding as in the stellar models and obtain a coarse clone.

The number of pixels in the telluric model can be chosen at the time of running TelFit. Here we choose a spectral resolution $R \sim 10^6$, adequate for resolving narrow telluric lines, and yielding about 2 million pixels across the entire HPF bandwidth. This pixel sampling is about $6\times$ finer than the native PHOENIX pixel sampling. The number of telluric lines depends on the atmospheric properties, in particular, the local surface temperature T_{\oplus} and relative humidity RH. For a surface temperature of $T=290\mathrm{K}$ (62° F), relative humidity of 40%, and typical conditions for McDonald Observatory, we anticipate 3615 telluric lines across the entire HPF bandwidth, distributed as shown in the bottom panel of Figure 1.

One guiding principle departs from the stellar case: telluric lines do not require future-proofing for large radial velocity shifts, hypothetically allowing us to reduce the number of pixels needed for a wingcut. Small radial velocity shifts are possible due to bulk motions in the Earth's atmosphere, but those bulk motions should be much smaller than the speeds of stars towards and away from Earth. So we can hypothetically tolerate a much smaller wing cut for telluric lines. In practice, telluric lines can be saturated, and accurately cloning the resulting broad telluric wings still benefits from $N_{\rm cut} \sim 6000$ pixels, comparable to the stellar scenario.

We optimize the sparse telluric clone, achieving comparable computational speed as the PHOENIX cloning task. We are left with $T_{\rm clone}(\lambda_T)$, the tunable telluric clone model evaluated at its original native coordinates. Figure 4 shows a before-and-after view of the blasé cloning procedure for a TelFit model zoomed in on a J-band region with dense Telluric lines. We see that the blasé fine-tuning appears nearly indistinguishable from the native resolution TelFit precomputed model.

4. CLONING PERFORMANCE

We compute the residual $R(\lambda_S)$ of native-minuscloned PHOENIX model, illustrated in the bottom panel of Figure 3. We see an RMS residual of 1.2%/pixel at native resolution. The telluric clone shows a comparable level of performance. These residuals tend to pile up in local symmetrically balanced clusters that get cancelled out once convolved with coarser instrumental line profiles, so their overall effect at instrumental resolution is typically negligible: the clones are almost perfect. We identify three main categories of cloning flaws that may not be negligible depending on the science application.

The first—and expected—source of large residuals is simply missing line opacity due to our finite prominence

¹ https://speakerdeck.com/dwhgg/telluric-line-hack-week-wrap-up

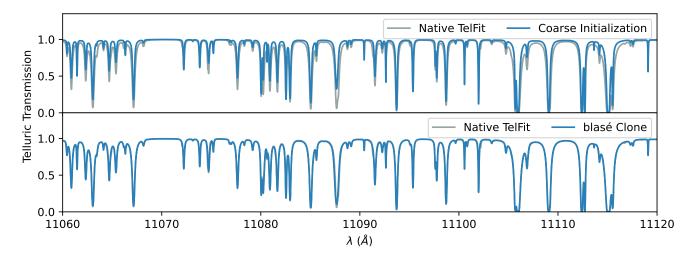


Figure 4. Pre-computed TelFit model cloned with blase. **Before:** The top panel shows a forward model evaluated with coarsely initialized line-by-line properties, based on simple threshold-based peak-finding and crude estimation of the shape of the line. **After:** The bottom panel shows the *blasé* fine-tuned clone, with nearly indentical spectral structure.

threshold. Lines with prominence less than $P_{\rm rom}$ yield residual notches with strengths comparable to $P_{\rm rom}$. Including smaller prominence lines by lowering $P_{\rm rom}$ produces smaller residuals, at the tradeoff of computing more lines and yielding higher computational cost. But at some point, turning down $P_{\rm rom}$ yields diminishing returns, as other imperfections provide a noise floor. We have experimentally determined this noise floor to occur near $P_{\rm rom}=0.01$.

Second, another anticipated flaw occurs in the line cores of relatively narrow lines, where the pseudo-Voigt profile becomes a poor approximation of the exact Voigt profile. The cloned model tends to overestimate the flux at the core and underestimate the flux along the slopes of the lines.

Finally, and most perniciously, a large category of residuals appear near the wings of the deepest and broadest lines—such as Hydrogen and neutral alkali metal lines. The true lines exhibit advanced lineshapes, such as non-Lorentzian line wings that are not captured with the overly simplistic Voigt line profile. Figure 5 highlights super-Lorentzian line wings around a line at 8691 Å. Narrow lines devolve into missing line wing opacity, the favored tradeoff when the continuum estimate's poor performance outweighs the pain of a narrow-but-tolerably-small spike. This flaw can be seen where a line initialized at 8692.5 Å and another pair of lines at 8690.0 all melt into line wings.

Cloning telluric lines suffers from one additional problem. Telluric lines can be extremely deep, exhibiting almost vanishing transmission with saturated line cores common in-between the atmospheric windows that de-

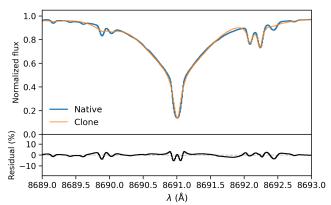


Figure 5. Zoom in of the region between the vertical gray bars in Figure 3. The cloned model has 7 spectral lines describing the 400 pixels in this 4 Å chunk. The native PHOENIX pixel sampling can be see as the boxy steps in both the native and cloned model.

fine the $I,\ J,\ H,$ and K bands. The blasé method can cope with these saturated lines, but often it treats nearby-and-blended saturated lines as one single line. This glomming together of lines has little practical effect since few astronomical practitioners can make use of such profoundly saturated data.

5. SEMI-EMPIRICAL MODELS WITH TRANSFER LEARNING TECHNIQUES

The cloned models already represents a useful intermediate product: the distillation of $N_s=335,849\times 2$ pixel flux values and their wavelength coordinates into a more compact quartet of properties for a list of <10,000 spectral lines, a dimensionality reduction of $18\times$ for the

cost of 1.2%/pixel in accuracy. So the process so far can be myopically viewed as a physics-informed compression algorithm. But the cloned model serves as a mere stepping stone in our principal quest: the comparison of models to real data.

5.1. Augmenting the stellar clone with radial velocity and rotational broadening

Real stars possess two key extrinsic properties. Rotational broadening $v\sin i$ and radial velocity RV depend on the observer's viewing location. We follow Czekala et al. (2015) by emphasizing the qualifier extrinsic, to distiguish between stellar intrinsic properties, such as $T_{\rm eff}$, $\log g$, and [Fe/H]. Intrinsic properties appear the same from any viewing location—at least for stars with isotropic surfaces—while extrinsic properties do not. The distinction is important because the extrinsic terms act as simple convolutions and translations to the cloned spectrum, and can be treated after the cloning procedure. We therefore define an augmented model, which we designate the "extrinsic" model, $S_{\rm ext}$:

$$S_{\text{ext}}(\lambda_Z) = S_{\text{clone}}(\lambda_Z - \frac{RV}{c}\lambda_c) * \zeta\left(\frac{v}{v\sin i}\right)$$
 (7)

where ζ is the convolution kernel for rigid body rotation (e.g. Kawahara et al. 2022), \boldsymbol{v} is the spectral axis represented as relative velocity coordinates, and * denotes the convolution operator.

Most autodiff-aware convolve operators act in pixel-space, approximating kernels as numerically sampled functions. There exists a special exponential spectral sample spacing that allows the convolution operators to work out of the box for rotational broadening. The design of blasé permits the stellar and telluric models to be reëvaluated at any wavelength coordinate vector, provided that it adequately samples the underlying lines. We, therefore, change the sampling from the native stellar and telluric wavelength coordinate grids, λ_S and λ_T , to this special exponentially sampled wavelength grid, denoted with the subscript Z:

$$\lambda_Z = \lambda_0 \exp\left\{\frac{\boldsymbol{v} - v_0}{c}\right\} \tag{8}$$

where $v - v_0$ is the velocity vector going from zero to the velocity associated with the largest wavelength, with *linear* spaced velocity samples. We choose a sampling in velocity space of 0.5 km/s, which corresponds to about $10\times$ finer than the instrumental resolving power of HPF, and delivers a minimum and maximum wavelength spacing of 0.013 and 0.024 Å/pixel respectively for the HPF bandwidth.

Operationally, the radial velocity shift RV gets applied to the line center positions rather than scaling the entire wavelength grid point coordinates, λ_Z . This choice yields a convenience: it cleanly makes the RV autodiff-aware, meaning that an infinitesimal change to the RV value can be sensed through backpropagation by affecting only the line center positions.

The $S_{\rm ext}$ spectrum is shown in Step 2 of Figure 2. There is currently no equivalent post-processing of the telluric spectrum T. As mentioned previously, we assume motions of the Earth's atmosphere are much less than the desired stellar radial velocity precisions. However, demanding Extreme Precision Radial Velocity (EPRV) exoplanet searches may need to consider minuscule systematic RV shifts and broadening of the telluric templates, arising from the turbulent and bulk motions of the Earth's atmosphere. A $T_{\rm ext}$ could hypothetically be implemented in $blas\acute{e}$ to achieve these strenuous precision demands.

5.2. Joint Stellar and Telluric Model

Figure 2 shows a visual guide to all the steps in blase. We have arrived at what may be the most intriguing-and-yet-simple of these steps: we simply multiply the rotationally-broadened-and-RV-shifted stellar model by the telluric transmission:

$$\mathsf{M}_{\mathrm{joint}} = \mathsf{S}_{\mathrm{ext}}(\boldsymbol{\lambda}_Z) \odot \mathsf{T}_{\mathrm{clone}}(\boldsymbol{\lambda}_Z) \tag{9}$$

It is only at this stage that we may apply the instrumental broadening kernel. The instrumental resolving power, R, acts as a convolution with a Gaussian line profile of width $\sigma = \frac{c}{2.355R}$. Real astronomical instruments usually have wavelength-dependent resolving power, which complicates the implementation for high-grasp spectra. The extent to which this effect matters will depend on the science application. For now, blasé simply assumes a fixed resolving power.

Notice that the order of Steps 3 and 4 in Figure 2 cannot be swapped. Mathematically speaking, multiplication and convolution do not commute. While the distinction may seem negligible, it matters at the level of precisions sought in EPRV applications (Suvrath Mahadevan, Arpita Roy, Sharon Xuesong Wang priv. comm.). Water vapor lines in our own atmosphere can "beat" with water vapor in the spectrum of, say, an M-dwarf atmosphere. The systematic telluric miscancelation would imbue a Moiré pattern of residuals that is most acute for sources with sharp lines, namely low projected rotational broadening $(v \sin i \sim \frac{c}{2R})$. The approach in blasé may therefore unlock a level of telluric calibration that has evaded previous efforts.

As noted, typical data-pixel sampling λ_D is much coarser than the model pixel sampling λ_Z . We therefore resample the model to the data spectrum in the following way. We evaluate the joint model $M_{\rm joint}$ at all of the super-resolution wavelength coordinates, and then compute the mean value of those pixels within the bounds of each coarse data pixel. The resampling procedure is autodiff-aware: the same clusters of high resolution coordinates map to the same data pixel coordinates, no matter what the RV is. The RV only dictates what flux values are realized within those pixel bounds.

The final forward model for $blas\acute{e}$ is designated simply as M without subscripts to emphasize that we have achieved the desired goal of a plausible end-to-end physics-informed yet highly flexible forward model for each datum in the 1D observed spectrum:

$$\mathsf{M}(\boldsymbol{\lambda}_D) = \text{resample} \left[\mathsf{M}_{\text{joint}}(\boldsymbol{\lambda}_Z) * g(R) \right]$$
 (10)

where g is the Gaussian instrumental convolution kernel and the resample[] operation indicates the average of model pixels that fall within each data pixel's red and blue boundaries.

5.3. Regularization

Equation 10 has $\sim 21,000$ tunable parameters from the star, $\sim 9,000$ tunable parameters from the Earth's atmosphere, plus $v\sin i$ and RV. That adds up to about 30,002 model parameters. The resolving power may also be treated as tunable if it is not known or varies slightly with e.g. seeing, slit-or-fiber illumination, or instrumental configuration: 30,003.

It may appear desirable to simply optimize all of these parameters in a *laissez-faire* manner, allowing them all to take on whatever value the data dictates. Such a stratagem would overfit the data, resulting in unphysically perverse lineshapes that do not reflect the air of reality we aspire to impose on our synthetic spectral models. Lines would haphazardly fit noise-spikes, and conspire together to warp spectral shapes in unexpected ways. This overfit model may suit some rare purposes. But most of the time, we prefer to strike a better balance in the bias-variance tradeoff.

We apply some amount of regularization, a restriction on the allowed values the model parameters can take on. Fortunately, we have a firm theoretical basis to justify this regularization. We believe our precomputed synthetic spectral models are quasi-statically correct: the predicted spectra resemble the unobserved "True spectrum" with lines in the correct place, but just with the wrong area under the curve. This statement may stem from the fact that it is easier to predict the mere exis-

tence of some energy transition of atoms and molecules than it is to predict their transition rates, abundances, temperature and pressure effects, and all the other line strength effects that flow down to how much light a line ultimately absorbs in a stellar atmosphere.

The degree of regularization will control the extent of overfitting or underfitting. The most extreme regularization—the antithesis of the laissez-faire scenario—would yield a model too rigid to respond to the data at all, yielding a model entirely unchanged from the cloned PHOENIX and TelFit models, the extreme end of underfitting. So regularization constitutes the only hyperparameters worthy of tuning in blasé. The choice of how to set the regularization is problem-specific. We default to the following choice. We fix all line parameters except for amplitude, which receives an L1 loss—namely we penalize the absolute value of departures from an amplitude's starting place:

$$\mathcal{L}_{reg} \equiv \sum_{j=1}^{N_{\text{lines}}} \left| \frac{\ln a_j - \ln \hat{a}_j}{\Lambda} \right| \tag{11}$$

where the hat notation demarcates the amplitudes obtained from cloning, *i.e.* the "initial, theory-inspired amplitude". We assign $\Lambda \sim 5$. The total, overall loss then becomes:

$$\mathcal{L}_{tot} = \mathcal{L}_{MSE} + \mathcal{L}_{reg} \tag{12}$$

This weak degree of regularization has the effect of permitting refinement of only the most conspicuous data-model mismatches; the weakest lines do not bother to move from their initial state because doing so would penalize the regularization without enough reduction of the overall loss. The extrinsic $v\sin i$ and RV have no regularization, but in practice, they need to be initialized close to their plausible values.

6. RESULTS: COMPARISON TO DATA

6.1. WASP 69 with HPF

The planet-host star WASP 69 makes a great benchmark because it has a low $v \sin i = 2.2 \pm 0.4$ km/s (Casasayas-Barris et al. 2017), making its lines sharp and easy to perceive. The K5 dwarf has an effective temperature of about 4700 K, $\log g = 4.535 \pm 0.023$, and slightly super-solar metallicity (Anderson et al. 2014).

Figure 6 shows a portion of an HPF spectrum of WASP 69 centered on a region devoid of telluric lines. This figure highlights a baseline case with no line-by-line fitting, simply conventional template matching: comparing HPF data of WASP 69 to the closest PHOENIX

template, convolved and resampled to the HPF resolution and sampling. Conspicuous residuals of $\pm 10\%$ appear throughout the spectrum, with lines in the correct place, but with the amplitudes systematically biased. These lines residuals arise from bona-fide imperfections in the PHOENIX spectrum, with a minor contribution from the coarse sampling of the PHOENIX grid.

Figure 7 shows the same data spectrum compared to a pixel-level model trained using the *blasé* technique described in Section 5. The model fit appears much better, with typical residuals approaching the photon noise of the data themselves. The model is not perfect however, especially around line cores. These residuals stem from a combination of causes. First, our finite regularization restricts line amplitudes and widths from straying too far from their values. This computational tug-ofwar makes line cores land just-short of the values the otherwise would have obtained in the absence of regularization. Second, the spectral resolution kernel may get biased from the domineering model-misspecification of broad lines, setting up a slightly sub-par performance for all the other lines.

The inferred, semi-empirical high-resolution model for WASP 69 is therefore the transfer-learned model \hat{S}_{clone} , unadorned with the extrinsic and instrumental properties. This semi-empirical model is shown in Figure 8. The departures from the PHOENIX model appear dramatic at this native resolution. The transfer-learned semi-empirical template exhibits both deeper and shallower lines than the native PHOENIX model.

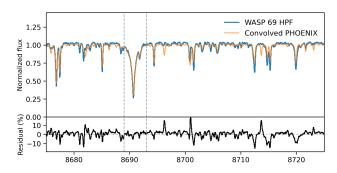


Figure 6. WASP 69 observed with HPF, compared to a $T_{\rm eff}=4700~{\rm K}, \log g=4.5~{\rm solar}$ metallicity PHOENIX model warped to $v\sin i=2.2~{\rm km/s},~RV=-9.6~{\rm km/s},$ and HPF resolving power.

The low $v \sin i$ of WASP 69 means that the line profile broadening arises principally from the finite instrumental resolution. The interplay of telluric lines and stellar lines within an instrumental resolution element is exactly one of the challenges blasé was designed to solve, as discussed in Section 3. In Figure 9 we show a multi-panel dissection of the HPF spectrum in a wave-

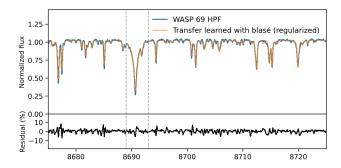


Figure 7. Semi-empirical model of WASP 69 transfer-learned with blasé, employing a regularization prior on the learned amplitudes.

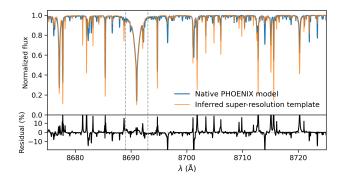


Figure 8. The native-resolution semi-empirical model transfer-learned from WASP 69 HPF data. The revised model can be viewed as a super-resolution deconvolution of the HPF spectrum.

length region in which conspicuous stellar and telluric lines coexist. The blasé end-to-end model exhibits excellent agreement with the data. The trustworthiness of the line properties inferred in locations where stellar and telluric lines exactly overlap remains an open research question: a tie-breaker is needed to distinguish these overlapping line inferrences. We discuss conceivable tie-brakers later in the paper.

6.2. IGRINS spectrum of a T6 Ultracool Dwarf

The T6 ultracool dwarf 2MASS J08173001-6155158 was recently observed with IGRINS, revealing a rich spectroscopic atlas of molecules in its $T_{\rm eff}=1060\pm50$ cloud-free atmosphere (Tannock et al. 2022). The sea of molecular lines are so rich as to blur entirely the notion of isolated lines. Many lines should instead be considered pseudo lines. The boundaries of lines for 2MASS J08173001-6155158 become even more amorphous in the presence of its moderately high $v\sin i=22.5\pm0.5$ km/s rotational broadening (Tannock et al. 2022). This T6 IGRINS spectrum therefore represents an extreme end of a test-case, in which the underlying notions of $blas\acute{e}$ go outside the comparatively safe as-

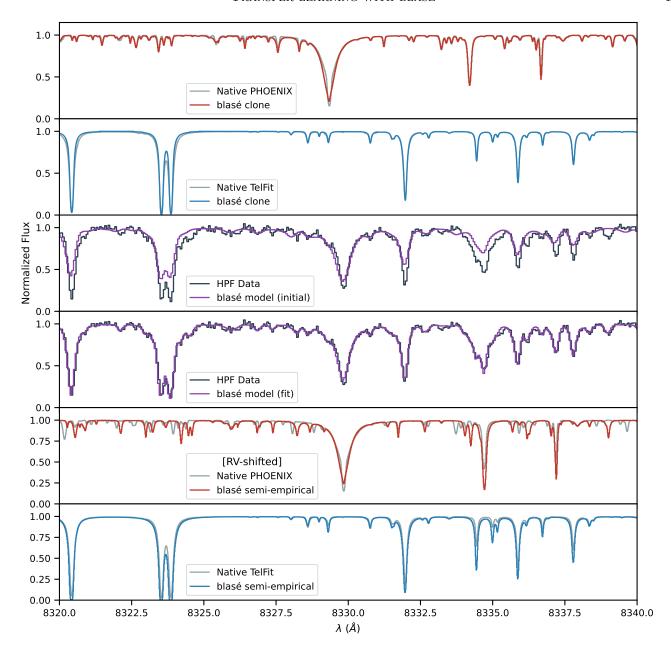


Figure 9. End-to-end training on a portion of HPF spectrum of WASP-69. The top two panels show the cloning in from Step 1 of the visual guide. The middle two panels depict the before-and-after of the instrument-convolved and resampled joint model illustrated in Step 4 of the visual guide. The final two panels show the underlying semi-empirical models learned in the process.

sumptions in stellar spectra. The application of blasé to $2MASS\ J08173001-6155158$ can be viewed as conducting a deconvolution step simultaneously paired with line-by-line inference.

We initialize blasé with the nearest Sonora template with $T_{\rm eff}=1100$ K, $\log g=5.0$, and solar metallicity (Tannock et al. 2022). Figure 10 shows the clone of this native spectrum near the peak of the H-band. The cloning performance appears adequate to capture most of the molecular pseudo-lines, with a few notable shortcomings. The high signal-to-noise ratio of this

IGRINS spectrum lays bare the departures from the broadened Sonora-Bobcat model, albeit with surprisingly close agreement given the challenge of substellar atmosphere modeling. The line-by-line departures from Sonora exceed any minor flaws from the cloning process, indicating genuine opacity differences between the data and Sonora template. Our technique automatically detects and reports the underlying structure of data-model mismatches, a key milestone for the assembly of refined opacity tables in this cool dwarf temperature regime.

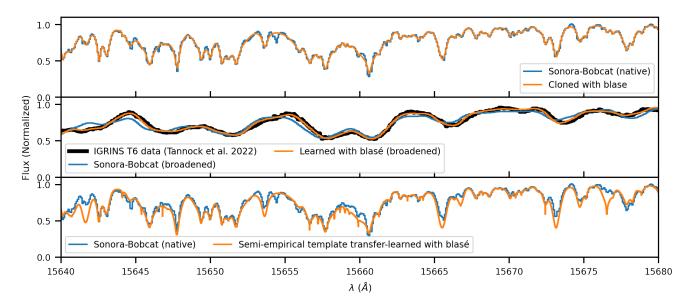


Figure 10. Line-by-line dissection of the 2MASS J08173001-6155158 IGRINS spectrum. The *top* panel compares the native Sonora-Bobcat spectrum to the blasé clone. We match the RV shift of the observed spectrum (thick black line) and convolve the initial and learned Sonora template to the large $v \sin i$. The bottom panel shows the locations and extents of line mismatches at the native resolution of Sonora.

7. DISCUSSION

7.1. How to interpret line-by-line properties

Up to this point we have presented the mechanics of blasé, and shown that it applies to a wide range of applications including stars, substars, and Earth's atmosphere. The interpretation of the approach and its line-by-line outputs may carry different meaning to different practitioners. Here we caricature some of these perspectives:

"Reverse-engineering the line lists" - Equation 2 (and its exponential twin Equation A2) could have been populated with initial line-by-line properties from atomic and molecular line lists, instead of the coarse detectionthreshold approach we presented in Section 2.2. So our cloning procedure can be viewed as reverse-engineering the quantum mechanical properties housed in line lists such as HITRAN, HITEMP, EXOMOL, VALD, or countless other primary source documents. This interpretation is only partly accurate. The line lists generally store the temperature and pressure scaling terms that get included in 1-D thermal structure calculations with distinct Voigt coefficient terms computed for each layer in the stellar or substellar or Earth atmosphere. The cloned properties therefore represent the flux-weighted average line property, having undergone an integral through the radiative transfer in the atmosphere.

Our deliberate choice not to initialize from the atomic and molecular lines means we do not have to compute an expensive multi-layer atmosphere at each inference step. The precomputed models have already taken care of that step, and so our cloning procedure encodes all the quantum mechanics and stellar atmospheres knowledge written down at the time of the computing of PHOENIX, Sonora, or LBLRTM.

"Equivalent Width machine"— The area under the curve of each jth blasé line represents its (neighborweighted) Equivalent Width (EW). That means that blasé can automatically and quickly distill a massive high-bandwidth spectrum into a catalog of line positions and EWs. Even more, the catalog would have some resilience to rotational broadening, line blending, and other friction points that hamper such an industrialscale approach with existing tooling. Current applications of Equivalent Widths rely on either isolated lines or—for multi-epoch applications—the assumption of non-variable neighboring lines. Blasé does not have these limitations. The interpretations of these EWs may have to be treated with care if intercompared with EWs obtained in the conventional way. This approach may suit stellar abundance applications, such as Galactic archaeology and near-field cosmology.

"Fancy deconvolution"— The blasé approach can be viewed as a fancy deconvolution procedure, in which the convolution kernels get inferred alongside the imperfections of the model template. This perspective may suit practitioners who wish to obtain a robust rotational broadening estimate in the presence of imperfect templates, and vice versa: robust templates in the presence of imperfect rotational broadening.

"A surrogate model, emulator, or discrepancy modeling"— The machine learning community is increasingly incorporating physics-based emulation into machine learning models. Blasé can be viewed as such a technique, often labeled as "surrogate models".

"A good enough substrate" - The imperfections in precomputed stellar and substellar models have hampered their use in high-fidelity applications, where model flaws overwhelm the sigal-to-noise in the data. The adaptability of blasé can be viewed as increasing the number of applications that could conceivably benefit from the rich-albeit-imperfect information encoded in these model grid spectra. In the next section we illustrate how these adaptable models can serve as a subtrate for applications in which the imperfections in precomputed models would have otherwise precluded their use.

8. UNLOCKING NEW SCIENCE

The scaffolding of blasé is designed in a way to promote extensibility, so many scientific questions can be written down in the flexible language presented in this paper. In some cases the flexibility, precision, speed, and ease-of-use may unlock new approaches to long-standing astrophysical questions and practical challenges. Here we enumerate some planned or conceivable extensions. We break up the themes into two categories: scientific extensions (this Section) and technical innovations (Section C).

8.1. Extreme Precision Radial Velocity

Already blasé is equipped to fit every single line with its own systemic RV, by tuning the line center position $\lambda_{\rm c}$ at each training epoch. There exists both empirical evidence and some theoretical motivation that RVjitter varies from line-to-line (Dumusque 2018b). Importantly, the extent of this line-by-line RV jitter could be predicted in part by the depth of line formation (Cretignier et al. 2021; Al Moulla et al. 2022). A future extreme-precision-RV (EPRV) version of blase could leverage this information. As an example, the depth-offormation for all (or a subset) of lines could be obtained and associated with each spectral line. A regularization scaling term could be introduced to allow the line positions to vary, but only in proportion to their depth of formation.

8.2. Line-by-line fundamental parameter estimates

The conversion of spectral line strengths to spectral type (Cannon & Pickering 1901)—and by extension line strengths to $T_{\rm eff}$ (Payne 1925)—has occupied a large chunk of stellar spectroscopy in the last century. The measurement of fundamental stellar properties remains one of the principal applications of stellar spectroscopy. Blasé as it currently stands does not output fundamental properties, and in fact it takes them for granted: you must specify e.g. the $T_{\rm eff}$, $\log g$, [Fe/H], and $[\alpha/{\rm Fe}]$ of your PHOENIX template as Step Zero before that template gets cloned and warped to match data. So obtaining a $T_{\rm eff}$ estimate from blasé is impossible, at least as disscussed so far.

The clearest way forward would be to calibrate the blasé-derived line properties. Simply put, we seek a set of functions, f, that relate the line properties back to fundamental properties:

$$f_j(a_j, \sigma_j, \gamma_j) \to (T_{\text{eff}}, \log g, [\text{Fe/H}])$$

There are $N_{\rm lines}$ such functions—one for each spectral line—because each line has its own temperature, gravity, and metallicity dependence. For visualization simplicity purposes, we have assumed the PHOENIX grid is merely 3-dimensional ($T_{\rm eff}$, $\log g$, [Fe/H]), but the same logic applies to 4-D and higher, and it works for sparse and irregularly sampled grids too.

Fascinatingly, there are two related ways forward towards this function. They both involve ensembles of spectra—such as a spectral atlas, spectral sequence, or library—and they both involve first obtaining the inverse function:

$$g_i \equiv f_i^{-1} \tag{13}$$

$$g_j \equiv f_j^{-1}$$
 (13)
$$g_j(T_{\text{eff}}, \log g, [\text{Fe/H}]) \rightarrow (a_j, \sigma_j, \gamma_j)$$
 (14)

A purely model-based approach would start by blasécloning every single PHOENIX spectrum in the grid dimensions-of-interest, over the wavelength range-ofinterest. The resulting product would be a line-by-line catalog of cloned spectral properties for each 3D coordinate on this grid. We could then assemble a heatmap of how each j^{th} line property changes across this heatmap, g_i . In this way, we are reverse-engineering the e.g. temperature dependence of each spectral line, as encoded by the PHOENIX atmosphere models. The final step could involve finding the neareast neighbor of each j^{th} observed spectral line to each j^{th} grid heatmap point; that is, finding a way to invert the function g_i to get f_i . The information-weighted mean, median, or mode of these nearest neighbors could then be reported as a revised "best-fit T_{eff} ", for instance.

A semi-empirical approach could improve on this purely model-based approach. We know that the PHOENIX line depths, widths, and shapes are imperfect, and so this $(T_{\text{eff}}, \log g, [Fe/H])$ heatmap will have large flaws in the line-by-line properties: the contours are systematically too bright or dark. There are many conceivable ways to quantify these heatmap flaws. Blasé provides an expedient route. One can pull the heatmap towards the locus of points established by running blasé on benchmark stars. The known $T_{\rm eff}$ of such systems would anchor the trend. We suspect the directionality of the purely model based heatmap must be correct, simply the slope, offset, and concavity of the trends may be wrong. With new information from each benchmark, the heatmap would get lifted like a central tentpole propping up an under-supported tent.

The construction and calibration of these ensembles of spectra represents a tremendous amount of work beyond the scope of this paper. But its creation could yield an extremely precise, fast, interpretable and reasonably accurate way to measure the properties of stars based solely on their high bandwidth échelle spectra. The overall accuracy hinges on the accuracy of the stellar benchmark labels. The mechanics of this approach could be adapted to fit within existing pipelines such as SAPP (Gent et al. 2022).

There exists a corrolary from the method described here to the approach of Czekala et al. (2015). There, a 3D heatmap was created for relating the eigenweights of a PCA basis back to the stellar fundamental properties, for spectrum emulation purposes. Here, the 3D heatmap is created on a line-by-line basis, and is therefore interpretable. The PCA eigenspectra were generally uninterpretable, at least not easily. So in the limiting case of obtaining a densely calibrated semi-empirical heatmap for all spectral lines across all grid dimensions, we will have achieved a spectral emulator that would obviate the need to clone spectra in the first place. We would simply start with this powerful line-by-line emulator as a forward model, and go directly to the extrinsic warping and data-model comparison.

8.3. Abundances

Stellar abundance work involves precisely measuring line strengths (often reported as EWs) of different chemical constituents evinced in a star or substar's photosphere. Relating those EWs back to physics can be done in a few ways. Most easily, trends and patterns in the EWs can be assembled, and metal-rich and metal-poor clusters can be identified. There exists a precision/accuracy tradeoff in the measurement process, usually stemming from the placement of the continuum, or assumptions about line blending. Blasé offers an immediate solution to these challenges, since it starts from our best-guess for how nearby lines may be shaping the continuum. Blasé appears to be a gateway to extremely fast and intervention-free industrial-scale abundances,

potentially useful for large surveys like APOGEE (Majewski et al. 2017), Gaia-ESO (Gilmore et al. 2012), RAVE (Steinmetz et al. 2006), and more.

8.4. Identifying missing lines

The automatic abundance determination spirit of blasé resembles ROBOSPECT (Waters & Hollek 2013), and other approaches aimed at automatically measuring equivalent widths for a large number of stars. One missing piece here is either to post-process the blaséoutput EW catalog with chemical labels (C, N, O, Si, Ba, Eu, etc.), or incorporate that information directly into the structure of blasé. The latter case is intriguing. For example, PHOENIX may not yet be aware of the existence of some obscure Lanthanum (La) line, possibly because the line was not yet understood and incorporated at the time of PHOENIX computation. Rather than wait for PHOENIX to receive the necessary update, blasé could instantiate one or more such lines in the locations guided by the atomic line list. This line could get initialized with some small amplitude, and would get added onto the underlying PHOENIX source as yet another freely tunable line. In this way, all the other nearby lines would be used for assigning continuum and blending, but this free-standing Lanthanum line would acquire a value properly taking into account the impact of its neighbors.

8.5. Doppler Imaging

The fixed $v \sin i$ approximation breaks down for stars with large-scale surface features. Doppler imaging attempts to reverse engineer the surface map from the extent to which observed line profiles depart from a pristine rotational broadening kernel. This reverse engineering step suffers from a vast number of geometrical degeneracies, but still provides useful constraints on stellar surfaces (Luger et al. 2021). We emphasize a distinction between A) longitudinally symmetric surface features, and B) longitudinally asymmetric surface features. Most radial velocity practitioners think about the latter, since longitudinally asymmetric surface features imbue changing-in-time skewness to the line profiles, causing radial velocity perturbations easily detectable in radial velocity time series. These confound exoplanet searches.

Longitudinally symmetric surface features—on the other hand—do not change as the star rotates on its axis. The existence of these features manifest as static-in-time *kurtosis* of the spectral line. For example, a hypothetical non-emitting (black) polar starspot exhibits a deficit in flux at the line core, resulting in less zero-velocity flux than its homogeneous counterpart. A dark

zonal band results in equal-sized bites out of the redand blue- sides of the line.

It is easier to reverse-engineer longitudinally asymmetric features than longitudinally symmetric ones, since we assume that we occasionally catch a glimpse of the spot-free limbs and their pristine line profiles. The latter requires exact knowledge of the underlying spectral template. Isolated, deep, well-calibrated spectral lines constitute the only practical scenario where exact knowledge can plausibly be claimed. Isolated spectral lines may be scarce or absent for M-dwarfs and brown dwarfs where lines blend ostensibly in an inseparable way, counfounding Doppler imaging.

blasé offers a new approach to Doppler imaging that may overcome these historical limitations by simultaneously fitting both the imperfections in the underlying spectrum and its line profile perturbations. This approach is analogous to the linearized model in Luger et al. (2021), but with the benefit of also handling nonlinear properties of the spectrum such as line widths, shapes, and locations, while also handling telluric contamination. Hypothetically the Luger et al. (2021) approach could be partially absorbed into blase, or vice versa, though such a merger may be complicated to implement.

8.6. Starspots and magnetic fields

One current assumption of starspot spectral decomposition is that the starspot spectrum itself resembles the stellar photosphere of a cooler star. This assumption appears adequate for detecting starspot spectra and measuring their physical properties (Gully-Santiago et al. 2017a). But to second order starspots should exhibit some spectral peculiarities that make them depart from a "normal" stellar photosphere. We may be probing deeper into the photosphere, and so the lines may experience higher pressure, with slightly different line widths. Or maybe the finite convective velocity shift can be directly seen as systematic shifts of the spectral lines (as stated in Section 8.1).

Rather than applying a mixture model of fixed PHOENIX templates, one could adopt a mixture model of pre-cloned blasé models. Then, the imperfections in the starspot spectrum can be learned alongside the filling factor of the starspots.

Often the most spotted stars are also rapidly rotating, and so the large $v\sin i$ blurs the starspot spectrum into the spot-free component of the photosphere. One experiment could measure the starspot spectral peculiarities more easily: obtainhigh resolution spectra of a pole-on (or nearly pole-on) young star where the lines will be

sharp and the two components of the spectra should be more easily discernable.

Relatedly, magnetic fields information could be incorporated into blasé as line-by-line Landé g-factors, to allow magnetic sensitive lines to exhibit either Zeeman broadening or Zeeman splitting. This labelling could be conducted in the same way as described in Section (?).

8.7. Circumstellar disk and accretion veiling

Circumstellar disk veiling suppresses the strengths of all spectral lines *en masse*, as the stellar photosphere gets outshined by a hot disk and/or envelope.

The bulk properties of that disk/envelope could be incorporated with a simple physical model: a black body of temperature $T_{\rm disk}$ and solid angle Ω_{disk} (Greene et al. 2018). Revealing the spectral shape of the veiling would require sufficiently high bandwidth spectra, such as Xshooter (Vernet et al. 2011) or possibly IGRINS. The extent of veiling would be fit alongside all the other stellar spectral lines. However, an MCMC approach may outperform blasé in accuracy, since the choice of picking an underlying spectral template is partially degenerate with the derived veiling.

Accretion veiling can be treated in an analogous way, but with some more complications. Emission lines can be easily incorporated as described in Section 2.5. These emission lines could be initialized with a line list of Hydrogen lines, forbidden lines, and other conspicuous features. The shapes of those lines can be affected by winds and other physical phenomena, yielding a variety of physically interpretable line shapes (Erkal et al. 2022). Blasé could incorporate those lineshapes as templates shared among several lines originating from similar physical environments, but scaled and shifted based on the details of each individual line's radiative transfer properties.

9. CONCLUSIONS

We have introduced an interpretable machine learning approach to forward modelling stellar, substellar, and telluric spectroscopic data. The line-by-line approach approach relies on a key enabling technology, automatic differentiation, that allows a nearly unlimited number of spectral lines to be forward modelled simultaneously. We initialize these lines to match precomputed synthetic stellar spectra, achieving excellent performance, and lending some confidence that the approach has a capacity to capture a tremendous amount of information at once.

We demo the framework on two sources: the K5 exoplanet host star WASP 69 using a precomputed PHOENIX model, and the T6 ultracool

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dwarf 2MASS J08173001-6155158 using a precomputed Sonora-Bobcat model. We discuss how blasé can be used to measure Equivalent Widths for thousands of lines automatically, understand line lists, measure rotation rates, generate surrogate models, and construct semiempirical models. This tool could readily have applications across stellar and substellar astronomy, including for PRV work, stellar compositions, Doppler imaging, and stellar activity.

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APPENDIX

A. LOG FLUX SCALING MODE

Here we illustrate how blasé gets altered when applying the logarithmic flux pre-processing step. First, we compute the natural log of the flux directly on the precomputed synthetic spectrum in its absolute flux scaling and native pixel sampling:

$$\ln \mathsf{S} = \ln \mathsf{S}_{\mathrm{abs}} - \ln \mathsf{B} - \mathsf{P} \tag{A1}$$

We simply "rebrand" P as residing in logarithmic flux units, and disregard it since it is largely a nuisance parameter anyways. We then treat the blasé clone model as a sum of opacities, retaining the Voigt profile:

$$\ln \mathsf{S}_{\text{clone}} = -\sum_{j=1}^{N_{\text{lines}}} a_j \mathsf{V}_j \tag{A2}$$

Here, the a_j 's have also been slightly rebranded from their meaning in Equation 2. We still want to enforce only absorption lines—and not spurious emission lines—so we use the sample trick of sampling the a_j 's in log and then exponentiating them to get guaranteed positive values. Note that Equations 2 and A2 carry modified meanings for the Voigt profile. Specifically, Equation 2 can be viewed as the Taylor Series expansion for A2 in the limit of small opacities:

$$e^{-a_j \mathsf{V}_j} \approx (1 - a_j \mathsf{V}_j) \tag{A3}$$

Both equations are approximate. A real stellar atmosphere's lineshape arises from a sum of disparate Voigt profiles weighted along a nonuniform column of gas, whereas here we have assumed the column of gas is approximated as a single uniform isothermal backlit layer. A sum of unlike-Voigt profiles is not exactly equal to any single Voigt profile. Theoreticians may resonate with this more "first principles" representation, while data practitioners may find Equation 2 more natural, so to some extent the choice is a matter of taste.

The sparse matrix gets rebranded as filled with opacity values, instead of log-fluxes, but operationally remains the same. All subsequent steps operate on the summed-and-exponentiated opacities, behaving identically to their linear counterparts. For example, we exponentiate before computing the residuals and data-model comparison, $R = e^{\ln S} - e^{\ln S_{\text{clone}}}$.

B. COMPARISON TO EXISTING SPECTROSCOPY FRAMEWORKS

Several astronomical spectral frameworks share similar aims as blasé. These existing frameworks will have enduring value for the wide range of problems in the field of stellar spectroscopy. Here we scrutinize the differences among some of these approaches to clarify how this work fits in.

The specmatch synthetic template matching tool produces noise-free nearest neighbor templates given an input spectrum (Petigura 2015). Several practical barriers limit the accuracy of using precomputed synthetic spectral models alone. First and foremost, real stars are usually more complicated than our simplified models of them. Real spectra often vary over more dimensions that our models do. Conspicuous examples of these hidden variables can be found in protostars: starspots, accretion veiling, dust extinction, and magnetic Zeeman splitting. Jointly modeling all of these phenomena alongside the intrinsic stellar photosphere is challenging.

The empirical version, specmatch-emp (Yee et al. 2017) matched spectra better than the synthetic templates, but is still too rigid for some applications and requires the assembly of hundreds of standardized high signal-to-noise-ratio templates, ideally with low intrinsic rotational broadening. Such a large number of high-quality templates with high resolving power and low $v \sin i$ has not yet been established in the near-infrared.

Table A1. Notation used in this paper

Symbol	Meaning
	Spectra
λ_S	Native wavelength coordinates of the precomputed stellar spectrum
$oldsymbol{\lambda}_T$	Native wavelength coordinates of the telluric spectrum
$oldsymbol{\lambda}_D$	Native wavelength coordinates of the data spectrum
S_{abs}	Flux values of the precomputed synthetic stellar spectral model λ_S
В	Blackbody of temperature $T_{\rm eff}$ to coarsely normalize $S_{\rm native}$
Р	Smooth polynomial to refine continuum-normalization
S	Continuum normalized augmentation of $S_{ m abs}$
T	Transmission values of the precomputed synthetic telluric model
D	The observed data spectrum flux values
ϵ	The estimated uncertainties in the data spectrum
$S_{ m clone}$	Evaluable and tunable cloned flux model of S
T_{clone}	Evaluable and tunable cloned transmission model of T
S_{ext}	An augmentation of S_{clone} with $v \sin i$ convolution and RV translation
M_{joint}	The joint stellar and telluric model: $S_{\mathrm{ext}} \odot T_{\mathrm{clone}}(\lambda_S)$
M	Joint model convolved with instrumental kernel and resampled to λ_D
R	The residual spectrum between a pair of inputs, e.g. $D - M$
$oldsymbol{v}$	The spectral coordinate axis λ expressed as a velocity difference
	Line properties
$\lambda_{c,j}$	Line center position of the j^{th} spectral line
a_j	Gaussian line profile amplitude of the j^{th} spectral line
σ_j	Gaussian line profile scale of the j^{th} spectral line
γ_j	Lorentzian line profile half width of the j^{th} spectral line
V_{j}	The Voigt profile of the j^{th} spectral line
$ec{ar{F}}$	The dense $(N_{\text{lines}} \times N_x)$ matrix of all line fluxes stacked vertically
$\hat{m{F}}$	The sparse $(N_{\text{lines}} \times N_{\text{sparse}})$ matrix of all line fluxes stacked vertically
ζ	The rotational broadening convolution kernel
g	The instrumental broadening convolution kernel, typically a Gaussian
	Scalars
$N_{ m lines}$	Number of spectral lines
N_x	Number of pixel coordinates in the precomputed spectrum λ_x
$N_{ m sparse}$	Number of non-zero pixels computed in the sparse implementation
$\pm \Delta \lambda_{ m buffer}$	Buffer exceeding the red and blue limits of the data spectrum
$P_{ m rom}$	The prominence threshold of spectral lines to include in cloning
$v \sin i$	Rotational broadening for stellar inclination i and equatorial velocity v
RV	Radial velocity of the star
R	Spectrograph resolving power $\lambda/\delta\lambda$
$\mathcal L$	The loss scalar, usually the sum of the squares of the residuals
	Operators
resample $[F(\lambda_x)]$	The resample operator, takes in a flux spectrum F evaluated at λ_x coordinates and returns the mean flux within the pixel boundaries of coordinate λ_z
*	The convolution operator
·	Hadamard product, an elementwise product of two same-length vectors

The wobble framework (Bedell et al. 2019) modernized the construction of high-SNR templates to account for temporally variable telluric lines. The tool requires dozens of high-SNR spectra acquired at a range of Barycentric Earth Radial Velocities (BERVs). The final telluric-free combined spectrum would still have to be compared to models for absolute calibration or can be used out-of-the-box for precision relative RVs. The wobble framework also pioneered the off-label application of automatic differentiation frameworks—in this case TensorFlow—towards their physically-motivated use in stellar spectra. blasé can be viewed as an evaluable and interpretable super-resolution version of wobble, that accepts more bias in the bias-variance tradeoff.

The starfish framework (Czekala et al. 2015) provides a robust likelihood function for data-model comparisons and retires many of the problems in this domain. starfish pioneered the use of whole-spectrum fitting with resilience to model imperfections by addressing the problem of what to do when the underlying atomic and molecular data was wrong or approximate or missing. It has been extended to inferring starspot physical properties (Gully-Santiago et al.

2017b), measuring veiling in Class 0 protostars (Greene et al. 2018), and quantifying imperfections in brown dwarf models (Zhang et al. 2021). The Spectral Inference Crank (sick, Casey 2016) shares similar aims as starfish, and provides additional useful grid search capabilities.

For very large bandwidths and very many spectral lines, the problem of identifying and cataloging line imperfections essentially becomes a book-keeping and continuum assignment problem. blasé and starfish provide different strategies for orchestrating the line-mismatch identification procedure, with each route having tradeoffs depending on the application.

C. CONCEIVABLE TECHNICAL IMPROVEMENTS

Blasé already performs very well under a wide range of cloning and transfer-learning tasks. However, some precision applications may demand even-more-strenuous performance than what the current implementation can accommodate. Here we describe some of these technical improvements, their design, and/or mention some science case they may unlock.

C.1. Exact instead of pseudo Voigt Profile

We currently employ the pseudo-Voigt profile for its low computational cost. We have a prototype exact-Voigt-Hjerting implementation following Kawahara et al. (2022). We coarsely estimate that moving to this exact-Voigt implementation could decrease some residual regions by $\sim 30\%$, while increasing the computational cost by more than $10\times$ over the existing pseudo-Voigt approximation. The exact-Voigt-Hjerting implementation still outbids the higher cost of a direct numerical convolution of a Gaussian and Lorentzian profile.

C.2. Addressing the pseudo-continuum with Gaussian Process regression

We currently assume the input spectra are adequately normalized to the continuum. We have a few options to relax this assumption. We could simply tune the P term that represents the wavelength-dependent pre-factor to Equation 2. Tuning P would correct for large-scale imperfections in the otherwise-fixed continuum flattening procedure. This change would be easy and effective, but has some challenge with model selection and flexibility: how to set the polynomial order to avoid over- and under-fitting. Gaussian Processes (GPs) offer many advantages for continuum fitting (Czekala et al. 2015). In short, a GP-likelihood relaxes the assumption that the continuum has been perfectly normalized, in favor of the more realistic statement "the continuum has been coarsely normalized, with some characteristic-but-as-yet-unknown correlation and scale length and amplitude of the imperfections". That statement translates to the following modification to Equation 3:

$$\mathcal{L} = \frac{1}{2} \mathsf{R}^{\intercal} \mathsf{C}^{-1} \mathsf{R} + \frac{1}{2} \ln \det \{ \mathsf{C} \}$$
 (C4)

where we introduce the covariance matrix C, with its associated kernel and collection of typically 2-3 parameters. We anticipate that this GP likelihood would have the greatest impact on stars with significant band-heads and line-blanketing: spectra with a so-called "pseudo-continuum". M-dwarfs and brown dwarfs fall into this challenging category.

The main demerit of moving to a GP-likelihood is computational cost. Fortunately, a few efficient autodiff-aware implementations of GPs exist. The celerite algorithm celerité (Foreman-Mackey et al. 2017) has an exact backpropation implementation (Foreman-Mackey 2018) that scales linearly with the number of data points. The celerité algorithm does not currently have a PyTorch implementation. The GPyTorch framework (Gardner et al. 2018) has a large category of approximate and exact GPs that could be straight-forwardly dropped into blasé. Even still these GPs could increase the computation cost by of order $10\times$.

C.3. Minibatches and Stochastic Gradient Descent

Currently, each training epoch sees the entire dataset, a setup dubbed *full-batch* gradient descent. An alternative scheme allows training with only a portion of the entire dataset at a time in *minibatches*. The massive data volumes in modern Neural Network applications cannot fit into the GPU memory, so minibatches are a necessity. Our meager 1 MB dataset can easily fit into the GPU memory, but our model can be large if we have a large number of pixels or lines or both. So while minibatches may not be required due to data size limitations they may be useful for particularly

large models. Minibatches also act as a form of regularization, the principal source of stochasticity in the Stochastic Gradient Descent algorithm, which tends to have better convergence than full-batch Gradient Descent (Ruder 2016).

We experimented with minibatches by assembling and evaluating only a portion of the dense \bar{F} matrix at a time, in minibatches. The choice to evaluate only a portion of lines at a time would mean the model is inaccurately evaluated at all wavelength pixels. Instead, we choose to evaluate all lines, but only on a random subset N_{batch} of the total pixels N_s , so that the model can eventually converge to exact at those points. All lines are allowed to update at each glimpse of a minibatch, but many lines with cores far from minibatch pixels will provide only weak information about how the loss scalar changes for their parameters.

Overall minibatches as implemented above performed worse than the sparse implementation, with both lower accuracy and slower computation time.

C.4. Broad lines and advanced lineshapes

Some lines—such as those arising from hydrogen, sodium, potassium, and others—have extremely broad line wings, approaching larger than the ~ 6000 pixels we allocate for the sparse implementation. These special lines should be handled separately from the weak lines, both from a computational performance perspective and an accuracy perspective.

Extremely broad lines will exhibit truncation effects if the sparse window is small compared to the line wing size. The truncation effects will look like tophat functions severing the asymptotic wings, imbuing artificial step function kinks in the emulated spectrum. We can afford to increase the sparse window on a few, say $N_{broad} \sim 20$ of the broadest spectral lines. We then construct and evaluate the entire dense matrix for those lines: $\sim 330~000 \times 20$. The number of FLOPS in each category scales as about 6 Million for the 20 broad and dense lines versus about 36 Million for the sea of about 7000 narrow and sparse lines, depending on the exact choices for wing cuts and the number of lines.

One could introduce advanced lineshapes for these ~ 20 broad lines, perturbing the Voigt line-wings with a smooth wavelength-dependent correction term G :

$$\tilde{\mathsf{V}}(\lambda) = \mathsf{V} \odot \mathsf{G} \tag{C5}$$

$$\mathsf{G} = 1 + (e^{\alpha_j} - 1) \cdot \mathcal{S}\left(\frac{|\boldsymbol{\lambda} - \lambda_{c,j}| - \lambda_{t,j}}{b_i}\right) \tag{C6}$$

where \odot is again the element-wise product (a.k.a Hadamard product), \mathcal{S} is the sigmoid function, and we have introduced three new tunable parameters for each of the j broad lines; λ_t is the truncation wavelength, b is a scale parameter for how slowly or how rapidly in wavelength-space the transition from non-Lorentzian proceeds, and α is a possibly negative stretch parameter that controls whether the line wing is sub- or super- Lorentzian.

This functional form has a few advantages. It is smooth. The smoothness of the transition is controlled by a tunable parameter, b. It can handle either sub- or super-Lorentzian shapes. In the limit $\lim_{a\to 0} \mathsf{G}$, the lineshape becomes exactly Lorentzian. The sigmoid is efficiently implemented in PyTorch. Finally, it enforces that the perturbation only produces absorption and not emission profiles.

C.5. Using native line lists rather than clones

For many practitioners, the choice to clone precomputed synthetic models in the first place may seem roundabout: "Why not just use the line lists?". Adopting the line lists would have many advantages: it would provide chemical and molecular provenance tags. Metadata associated with the quality of the atomic and molecular data could be used to assign physics-informed regularization. Many other benefits would effortlessly accrue from adopting the native line lists. The FAL project (Cargile et al. *in prep*) follows such a principled prescription.

As already emphasized, there exist at least a few demerits of adopting the line lists, and therefore supporting the blase strategy. First, these line lists need to undergo expensive multi-level radiative transfer calculations in order to obtain their amplitudes, so adopting the line lists would mean a laborious and computationally expensive pursuit simply to get close to what has already been computed. Second, as the effective temperature scales to ultracool dwarfs (Figure 1) the number of lines sky-rockets, tending towards the billions for T-dwarfs. The methane line list alone (Hargreaves et al. 2020) represents a prohibitive data volume. The ExoJAX and Radis (Pannier & Laux 2019; van den Bekerom & Pannier 2021) libraries offer a breakthrough solution to the voluminous line list problem. Even still, blase deals with the less pure but more practical "pseudo line" that gets closer to the astronomical observables anyways, and offers a middle ground between the extremes of interpretability and performance.

C.6. Wavelength dependent limb darkening

Currently the extrinsic model step possesses up to four parameters: the $v\sin i$ and RV, and 2 optional parameters for limb-darkening. These four parameters may adequately parameterize a star with a uniform stellar disk. Extremely high signal-to-noise-ratio spectra of rapidly rotating stars may require additional flexibility. The limb darkening is generally wavelength-dependent, and so a pan-chromatic spectrum may require a different limb darkening from the blue end to the red end. The limb darkening may instead depend on physical properties of the spectral line formation, such as physical depth of formation, and so the extent of limb-darkening may jump haphazardly from line-to-line, rather than as a predictably smooth function across wavelength. blasé could be built to handle such a seemingly pathological scenario by adding a vector of limb darkening parameters, one for each line. One would have to regularize the fits with some typical limb darkening and a heuristic penalty for departures from this mean.