



UNIVERSITY OF
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Probing Inflation with Precision Bispectra

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St. John's

This dissertation is submitted for the degree of Doctor of Philosophy

Declaration

This dissertation is the result of my own work and includes nothing which is the outcome of work done in collaboration except as declared in the Preface and specified in the text. It is not substantially the same as any that I have submitted, or am concurrently submitting, for a degree or diploma or other qualification at the University of Cambridge or any other University or similar institution except as declared in the Preface and specified in the text. I further state that no substantial part of my dissertation has already been submitted, or is being concurrently submitted, for any such degree, diploma or other qualification at the University of Cambridge or any other University or similar institution except as declared in the Preface and specified in the text. This dissertation does not exceed the prescribed limit of 60 000 words.

Philip Clarke
July, 2021

Abstract

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My abstract ...

Acknowledgements

My acknowledgements ...

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Chapter 1

Introduction to cosmology/inflation

Calculating the primordial bispectrum predicted by a model of inflation and comparing it to what we see in the sky is very computationally intensive, necessitating layers of approximations and limiting the models which can be constrained. Exploiting the inherent separability of the tree level in-in formalism using expansions in separable basis functions provides a means by which to obviate some of these difficulties. Here, we develop this approach further into a practical and efficient numerical methodology which can be applied to a much wider and more complicated range of bispectrum phenomenology, making an important step forward towards observational pipelines which can directly confront specific models of inflation. We describe a simple augmented Legendre polynomial basis and its advantages, then test the method on single-field inflation models with non-trivial phenomenology, showing that our calculation of these coefficients is fast and accurate to high orders.

1.1 General intro

The primordial bispectrum is one of the main characteristics used to distinguish between models of inflation. While it is well known that the physics of inflation must have been extremely close to linear, and the initial seeds of structure it laid down very close to Gaussian, there is expected to have been some level of coupling between the Fourier modes of the perturbations. In the simplest example of an inflation model this is expected to be unobservable [1], but the possibility remains that inflation was driven by more complex physics that may have left an observable imprint on our universe today. Some models of inflation have interactions that predict non-Gaussian correlations at observable levels. Ways this can happen include self-interactions [2, 3], interactions between multiple fields [4], sharp features [5] and periodic features [6]. However, constraining such imprints is extremely difficult observationally. Even once the data has been obtained, using existing methods it is extremely computationally intensive to translate this into constraints on

specific inflation scenarios. Much progress has been made by course-graining the model space into a small number of approximate templates, and leveraging the simplifying characteristic of separability with respect to the three parameters of the bispectrum [7, 8].

The primordial bispectrum is the Fourier equivalent of the three-point correlator of the primordial curvature perturbation. If this field is Gaussian, the bispectrum vanishes, so it is a valuable measure of the interactions in play during inflation. If some inflation model predicts a bispectrum that is sufficiently well approximated by the standard separable templates, the constraints on those standard templates can be translated into constraints on the parameters of the model. The fact that all primordial templates estimated thus far from the CMB are consistent with zero has already provided such constraints in certain scenarios [9, 10]. With this high-precision *Planck* data, and data from forthcoming experiments such as the Simons Observatory (SO) [11] and CMB-S4 [12], robust pipelines must be developed to circumvent the computational difficulties and extract the maximum amount of information possible. Due to the nature of bispectrum estimation in the CMB and LSS [13, 14, 15, 16] constraining an arbitrary template is difficult. Our aim in this work is to develop the inflationary part of a pipeline to allow to efficiently test a much broader range of models. In this work, we explore shapes arising from tree-level effects in single field models. We do this numerically, allowing quantitative results for a broad range of models, and avoiding extra approximations. Our general aim is to apply the modal philosophy of [17, 18, 19] to calculating primordial bispectra. This modal philosophy is a flexible method that has broadened the range of constrained bispectrum templates, by expanding them in a carefully chosen basis. The Modal estimator is thus capable of constraining non-separable templates, while the KSW estimator cannot. In this work we exploit the intrinsic separability of the tree-level in-in formalism to apply these methods at the level of inflation. Expressing the primordial bispectrum in a separable basis expansion leads to vast increases in efficiency both at the primordial and late-universe parts of the calculation. The main advantage is that expressing the primordial shape function in this way reduces the process of bispectrum estimation in the CMB to a cost which is large, but need only be paid once per basis, not per scenario. A proof of concept of this approach at the primordial level was presented in [20], and the details of the bispectrum estimation part will be detailed in [21]. We go beyond the work of [20] both in developing the choice of basis (the feasibility of the method depending vitally on the chosen basis achieving sufficiently fast convergence in a broad range of interesting models) and in the methods we use to allow us to go to much higher order in our modal expansion, allowing us to apply the method to feature bispectra for the first time.

The paper is organised as follows. In section ?? we present brief reviews of the various parts of the pipeline that connects inflation scenarios to observations through the bispectrum. We review the usual paradigm of bispectrum estimation in the CMB, and

the motivation for separable bispectra. We review the in-in formalism, for calculating the tree level bispectrum for a given model of inflation. We review $P(X, \phi)$ models of inflation as an example, and some of the usual approximate bispectrum templates that we aim to bypass. We will draw our validation scenarios from these models. We discuss previous numerical codes for calculating the primordial bispectrum k -configuration by k -configuration, which contrasts our separable basis expansion. We review the previous work in achieving separability through modal expansions in [20], and we discuss methods of testing numerical bispectrum results, defining our relative difference measurement. In section ?? we present our methods. Since the paradigm we aim to present is only viable if we can find a basis that can efficiently represent a wide variety of bispectra, we begin with this vital discussion. We discuss the effects of the non-physical k -configurations on the convergence of our expansion on the tetrapyd, and present an efficient basis. Then, we recast the usual in-in calculation into an explicitly separable form, in terms of an expansion in an arbitrary basis, and detail our methods for carefully calculating the coefficients to high order. In section ?? we validate our methods and implementation on inflation scenarios with varied features from the literature, and we finish with a discussion of future work in section ??.

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Chapter 2

Decomposing primordial shapes

2.1 Pulling out the k -dependence

2.1.1 Set up formalism.

Given its separable form, the tree-level in-in formalism is amenable to more efficient calculation using separable modes, as first mentioned in [20]. That work extended the separable methodology previously implemented for the CMB bispectrum [19]. Our goal in this work is the efficient calculation of more general bispectra which may have significant (possibly oscillatory) features, requiring searches across free parameter dependencies. To achieve this, we represent the shape function (??) using a set of basis functions as

$$S(k_1, k_2, k_3) = \sum_n \alpha_n Q_n(k_1, k_2, k_3), \quad (2.1)$$

where the basis functions $Q_n(k_1, k_2, k_3)$ are explicitly separable functions of their arguments. Translating this result into a constraint from the CMB will require a large once-off computational cost, paid once per set of basis functions Q_n , not per scenario (encoded in α_n). The details of this once-per-basis calculation will be presented in [21]. As such, while the general computational steps we describe will be independent of the basis, it is vital we explore possible sets of basis functions $Q_n(k_1, k_2, k_3)$ and their effects on convergence; we do this in section ?? . In section ?? we set the notation we will use to recast the standard numerical in-in calculation into a calculation of α_n , and sketch the steps involved. In section ?? we outline the details of the interaction Hamiltonian, including accounting for the spatial derivatives in our final result. In section ?? we make precise the numerical considerations of the calculation, especially our methods of dealing with the high-frequency oscillations at early times.

2.1.2 Forced to decompose on the cube.

2.1.3 Defining the notation for the two main time integrals.

In this section we set up the notation, and sketch the steps required to calculate the coefficients α_n in (2.1). The values of these coefficients will depend on the choice of basis, but the description of the methods below will remain mostly basis agnostic. Our aim will be to separate out the dependence on k and τ_s , without losing information, except in the sense that is controlled by p_{\max} . We will set up an efficient numerical implementation of the calculation, a necessary consideration to allow this method to be useful in exploring parameter spaces in primordial phenomenology. Throughout we will see that we are able to preserve the separability of the dependence on k_1 , k_2 and k_3 .

The tree-level in-in formalism for the bispectrum (??) is inherently separable given the form of the cubic interaction Hamiltonian H_{int} . Consider indexing with $i = 1, 2, 3 \dots$ the interaction vertices in H_{int} , so then the bispectrum (??) can be expressed as a sum over separable contributions of the form:

$$\begin{aligned}
 S(k_1, k_2, k_3) &= \sum_i I^{(i)}(k_1, k_2, k_3) \\
 &= \sum_i \left[v^{(i)}(k_1, k_2, k_3) \int_{-\infty(1-i\varepsilon)}^0 d\tau w^{(i)}(\tau) F^{(i)}(\tau, k_1) G^{(i)}(\tau, k_2) J^{(i)}(\tau, k_3) \right. \\
 &\quad \left. + \text{cyclic perms} \right] \tag{2.2}
 \end{aligned}$$

where $w^{(i)}(\tau)$ is a function of the scale factor and the other background parameters (??) for the i -th interaction vertex, while the terms $F^{(i)}, G^{(i)}, J^{(i)}$ are given by the Fourier mode functions $k^2 \zeta_{\mathbf{k}}(0) \zeta_{\mathbf{k}}^*(\tau)$ or their time derivatives $k^2 \zeta_{\mathbf{k}}(0) \dot{\zeta}_{\mathbf{k}}^{*'}(\tau)$. Spatial derivative terms, such as $\partial_i \zeta \partial_i \zeta \rightarrow (\mathbf{k}_2 \cdot \mathbf{k}_3) \zeta_{\mathbf{k}_2}^*(\tau) \zeta_{\mathbf{k}_3}^*(\tau)$ also separate because of the triangle condition (??) giving $\mathbf{k}_2 \cdot \mathbf{k}_3 = (k_1^2 - k_2^2 - k_3^2)/2$, yielding a sum of separable terms. These time-independent contributions are contained in $v^{(i)}(k_1, k_2, k_3)$, as they do not force us to compute extra time integrals. Note that $v^{(i)}(k_1, k_2, k_3)$ need not be symmetric in its arguments.

The terms contained in $v^{(i)}(k_1, k_2, k_3)$ depend on the structure of the spatial derivatives in the interaction Hamiltonian, but not the specific scenario. These terms are separable; for details, see section ??. We include their contribution to the final result after the time integrals have been computed. The factors which depend only on time, $w_i(\tau)$, depend on the scenario but do not need to be decomposed in k . The remaining factors have both k and time dependence; they must be decomposed in k at every timestep. These terms look like $F^{(i)}(k, \tau) = k^2 \zeta_{\mathbf{k}}(0) \zeta_{\mathbf{k}}^*(\tau)$ (or $k^2 \zeta_{\mathbf{k}}(0) \dot{\zeta}_{\mathbf{k}}^{*'}(\tau)$), where k^2 comes from using the weighting

of the scale-invariant shape function (??). This could be absorbed into $v^{(i)}(k_1, k_2, k_3)$, but we have the freedom to keep it here to aid convergence.

If the expressions being expanded have some known pathology in their k -dependence, we can then see two ways of dealing with this. The basis can be augmented to efficiently capture the relevant behaviour (see section ??) or the behaviour can be absorbed into the analytic prefactor, $v^{(i)}(k_1, k_2, k_3)$.¹ We use the former, as the numerics of the latter are less transparent and less physically motivated.

The internal basis used for the decomposition at each timestep need not match that which is used for the final result, and indeed in dealing with the spatial derivatives in section ?? we will find it useful to change to a different basis than the one used to perform the time integrals of the decompositions.

Using the approximate mode functions (??), an explicit example for the first interaction term in (3.4), i.e. $H_{\text{int}}^{(1)} = \zeta'^2 \zeta$, takes the form

$$F^{(1)}(\tau, k) = G^{(1)}(\tau, k) = c_s k^2 \tau \frac{H^2}{4\epsilon c_s} e^{ic_s k \tau}, \quad J^{(1)}(\tau, k) = (1 - ik c_s \tau) \frac{H^2}{4\epsilon c_s} e^{ic_s k \tau}. \quad (2.3)$$

In the simple mode approximation (??), such terms in (2.2) are straightforward to integrate analytically (using the $i\epsilon$ prescription), provided the time-dependence of the slow-roll parameters and the sound speed is neglected [1]. However, for high precision bispectrum predictions we must incorporate the full time-dependence, while solving (??) to find accurate mode functions $\zeta_{\mathbf{k}}(\tau)$ numerically. Obtaining the full 3D bispectrum directly is computationally demanding at high resolution because it requires repetitive integration of (2.2) at each specific point for the wavenumbers (k_1, k_2, k_3) , a problem which is drastically compounded by bispectrum parameter searches e.g. for oscillatory models.

Consider representing the primordial shape function $S(k_1, k_2, k_3)$ in (2.2) as a mode expansion for each interaction term $I^{(i)}(k_1, k_2, k_3)$ as

$$S(k_1, k_2, k_3) = \sum_i I^{(i)}(k_1, k_2, k_3) = \sum_i \sum_n \alpha_n^{(i)} Q_n(k_1, k_2, k_3), \quad (2.4)$$

where $Q_n(k_1, k_2, k_3)$ is separable, built out of some orthonormal set $q_p(k)$ as in (2.20). Armed with this set of modes, we can expand any of the interaction terms $F^{(i)}(\tau, k)$,

¹At early times the modes are highly oscillatory in both k and τ_s , which certainly requires special attention. We discuss this in section ??.

$G^{(i)}(\tau, k)$, $J^{(i)}(\tau, k)$ in (2.2) as:

$$F^{(i)}(\tau, k) = \sum_p f_p^{(i)}(\tau) q_p(k), \quad (2.5)$$

$$\text{where } f_p^{(i)}(\tau) = \int_{k_{\min}}^{k_{\max}} dk F^{(i)}(\tau, k) q_p(k). \quad (2.6)$$

Note that in the simple mode approximation, as in (2.3), we must expand $e^{ic_s k \tau}$ in the terms of the $q_p(k)$. At early times τ is large, so in k this is highly oscillatory. This creates two problems. Firstly, this seems to require many samples in k to accurately calculate each $f_p^{(i)}(\tau)$, adding more modes that must be evolved in time. To bypass this, we extract the oscillatory part at early times, reducing the number of needed k -samples; see section (??) for details. Secondly, it forces us to calculate $f_p^{(i)}(\tau)$ up to very high p if we want to accurately converge to $F^{(i)}(\tau, k)$, for sets of basis functions such as the Legendre polynomials. In fact, obtaining a convergent final bispectrum result does not require calculating the full convergent sum for $F^{(i)}(\tau, k)$ in (2.5), as the highly oscillatory parts will cancel in the time integrals for any sufficiently smooth $S(k_1, k_2, k_3)$.

Substituting these expansions into (2.2), we obtain the following decomposition for the i -th vertex contribution,

$$\begin{aligned} I^{(i)}(k_1, k_2, k_3) &= v^{(i)}(k_1, k_2, k_3) \int d\tau w^{(i)}(\tau) \sum_p f_p^{(i)}(\tau) q_p(k_1) \sum_r g_r^{(i)}(\tau) q_r(k_2) \sum_s h_s^{(i)}(\tau) q_s(k_3) + \text{cyclic perms} \\ &= v^{(i)}(k_1, k_2, k_3) \sum_{prs} \left(\int d\tau w^{(i)}(\tau) f_p^{(i)}(\tau) g_r^{(i)}(\tau) h_s^{(i)}(\tau) \right) q_p(k_1) q_r(k_2) q_s(k_3) + \text{cyclic perms}. \end{aligned}$$

For the sake of compactness we use P to stand for the triplet p, r, s and \tilde{P} to stand for the triplet $\tilde{p}, \tilde{r}, \tilde{s}$. Writing $q_P(k_1, k_2, k_3) = q_p(k_1) q_r(k_2) q_s(k_3)$, we continue,

$$\begin{aligned} I^{(i)}(k_1, k_2, k_3) &= v^{(i)}(k_1, k_2, k_3) \sum_P \tilde{\alpha}_P^{(i)} q_P(k_1, k_2, k_3) + \text{cyclic perms} \\ &= \sum_P \tilde{\alpha}_P^{(i)} \sum_{\tilde{P}} V_{P\tilde{P}}^{(i)} q_{\tilde{P}}(k_1, k_2, k_3) + \text{cyclic perms} \\ &= \sum_{\tilde{P}} \alpha_{\tilde{P}}^{(i)} q_{\tilde{P}}(k_1, k_2, k_3) + \text{cyclic perms}, \end{aligned} \quad (2.7)$$

where we have written

$$\tilde{\alpha}_P^{(i)} = \tilde{\alpha}_{prs}^{(i)} = \int d\tau w^{(i)}(\tau) f_p^{(i)}(\tau) g_r^{(i)}(\tau) h_s^{(i)}(\tau), \quad (2.8)$$

and included the time-independent k -prefactors from the interaction Hamiltonian by

writing

$$v^{(i)}(k_1, k_2, k_3) q_P(k_1, k_2, k_3) = \sum_{\tilde{P}} V_{P\tilde{P}}^{(i)} q_{\tilde{P}}(k_1, k_2, k_3), \quad (2.9)$$

and

$$\alpha_P^{(i)} = \sum_{\tilde{P}} \tilde{\alpha}_{\tilde{P}}^{(i)} V_{\tilde{P}P}^{(i)}. \quad (2.10)$$

We connect to the coefficients of the ordered, symmetrised basis in (2.4) by taking the symmetry factor into account,

$$\alpha_n^{(i)} = \frac{3!}{\Xi_P} \alpha_P^{(i)}. \quad (2.11)$$

The numerical calculation of $V_{P\tilde{P}}^{(i)}$ (as defined by (2.9)) is highly efficient as $v^{(i)}(k_1, k_2, k_3)$ is a sum of separable terms. The details of these terms depend only on the spatial derivatives in the interaction Hamiltonian, not the scenario being considered, so the matrix can be precomputed and stored. Note that this is not the only way one can organise this calculation to explicitly preserve the separability. One could also include the contributions coming from the spatial derivatives first, decomposing (as in (2.5)) not only terms like $k^2 \zeta_{\mathbf{k}}(0) \zeta_{\mathbf{k}}^*(\tau)$, but also terms that include each power of k_1 , k_2 or k_3 that appears in $v(k_1, k_2, k_3)$. The index i in the sum in (2.4) would then run over not only each vertex in the interaction Hamiltonian, but also each separable term within those vertices. We do not choose this path as, for the sake of efficiency, we wish to minimise the number of time integrals of the form (2.8) we need to calculate.

Note the basis sets on the left and right hand side of (2.9) need not match. In fact, if those two basis sets do match, then generically information will be lost—for example, if the basis set on the left is \mathcal{P}_0 , then terms in $v^{(i)}(k_1, k_2, k_3)$ with positive powers will introduce higher order dependencies on k , and negative powers will introduce $1/k$ behaviour. In practice, to prevent this loss of information, we take the basis set on the right hand side of (2.9) to be an expanded version of that on the left. For example, if the left hand basis was \mathcal{P}_0 of size p_{\max} , the right hand basis would be \mathcal{P}_1 of size $p_{\max} + 3$.

We can see from (2.8) that the number of time integrals needed is controlled by $N_V \times p_{\max}^3$ ², where N_v is the number of interaction vertices and p_{\max} is the size of the final basis. Since the calculational cost of doing the internal decompositions depends only linearly on the size of internal basis, improvements there are dwarfed by improvements

²In fact the number is not quite p_{\max}^3 . Since we have extracted the spatial derivatives, the only remaining possible source of asymmetric k -dependence comes from ζ^3 , $\zeta^2 \zeta'$, $\zeta'^2 \zeta$ or ζ'^3 so the time integral in (2.8) will always be (at least) symmetric in p and r .

gained from reducing the number of terms needed in the final basis.

We will calculate the contribution of each H_{int} vertex separately, indexing the vertices as above by (i) , so the overall shape function (2.2), (2.4) is then simply

$$S(k_1, k_2, k_3) = \sum_n \left(\sum_i \alpha_n^{(i)} \right) Q_n(k_1, k_2, k_3) = \sum_n \alpha_n Q_n(k_1, k_2, k_3). \quad (2.12)$$

Depending on the scenario, some vertex contributions will converge faster than others or be completely negligible; for efficiency the maximum modal resolution defined by p_{\max} can be allowed to be different for each vertex.

The *raison d'être* for this approach is that all time integrals (2.8) are now independent of the k -configuration³. In a configuration-by-configuration method one improves the precision by decreasing the spacing which defines the density of the grid of points within the tetrapyd. Instead, in the modal approach, we increase precision by adding more modes to the shape function expansion (2.12) until the result converges at high precision. At first sight, this appears to increase the dimensionality of the calculation. Directly integrating the in-in formalism requires one time integration for each k -configuration, i.e. N_k^3 integrals, ignoring symmetry. The method detailed here requires decomposing the modes, then a time integral for every coefficient, i.e. p_{\max}^3 integrals (again ignoring symmetry) plus the decomposition. However for every model we have explored from the literature, our expansion in p_{\max} converges far faster than in the number of k -modes that would be required to have confidence in a sampled bispectrum. This is clear in smooth bispectra such as (??) and (??), but is also true of bispectra with complicated features, as seen in section ??.

To be efficiently connected to a late-time observable a sampled bispectrum would have to be fit by a smooth template, a complication that is automatically taken care of in this formalism. We note that while the primordial basis is chosen for computational speed and convenience, it can be independent of the final bispectrum basis employed for observational tests; a change of basis $Q_n \rightarrow \tilde{Q}_n$ can be achieved through a linear transformation Γ with the new expansion coefficients given by $\tilde{\alpha}_m = \Gamma_{mn} \alpha_n$.

Discussion of convergence in this work is considered only at the primordial level, with no concept of the signal to noise of an actual experiment. There could be a basis that converges faster in some observationally weighted sense, efficiently describing the primordial modes which will matter most at late times. We leave discussion of this point to a later work, as converting between the two, after the in-in computation is completed, is trivial.

Having now set our notation and outlined the calculation, in the following sections we

³They are not independent of k_{\min} and k_{\max} which define the domain of interest, which is analogous to the coefficients of a Taylor expansion depending on its expansion point.

discuss the actual numerical implementation of these methods.

2.2 Inner product choice

The inner product of two bispectrum shape functions is given by

$$S_1 \cdot S_2 = \langle S_1, S_2 \rangle = \int_{T_k} d^3k S_1(k_1, k_2, k_3) S_2(k_1, k_2, k_3), \quad (2.13)$$

where T_k refers to the tetrapyd, the region of the cube $[k_{\min}, k_{\max}]^3$ that obeys the triangle inequality. Following [22] we define the two correlators:

$$\mathcal{S}(S_1, S_2) = \frac{S_1 \cdot S_2}{\sqrt{(S_1 \cdot S_1)(S_2 \cdot S_2)}}, \quad \mathcal{A}(S_1, S_2) = \sqrt{\frac{S_1 \cdot S_1}{S_2 \cdot S_2}}. \quad (2.14)$$

Here, we refer to $\mathcal{S}(S_1, S_2)$ as the shape correlator between the two bispectra; $\mathcal{A}(S_1, S_2)$ is the amplitude correlator. In principle, we could add some observationally motivated weighting to the above measure, as considered in [17, 18, 19], but in this work we restrict ourselves to accurately calculating the full primordial bispectra, weighting each configuration equally.

Writing $|S|^2 = S \cdot S$, we can then re-express a measure of the relative error between one bispectrum template and another:

$$\begin{aligned} \mathcal{E}(S_1, S_2) &= \sqrt{\frac{|S_1 - S_2|^2}{|S_2|^2}} = \sqrt{\frac{|S_1|^2 - 2S_1 \cdot S_2 + |S_2|^2}{|S_2|^2}} \\ &= \sqrt{\mathcal{A}(S_1, S_2)^2 - 2\mathcal{A}(S_1, S_2)\mathcal{S}(S_1, S_2) + 1}. \end{aligned} \quad (2.15)$$

This error measure takes into account differences in overall magnitude as well as shape. If we are only interested in comparing the differences coming from the shape, we can scale the bispectra so that $\mathcal{A}(S_1, S_2) = 1$ and so

$$\mathcal{E}(S_1, S_2) = \sqrt{2(1 - \mathcal{S}(S_1, S_2))}. \quad (2.16)$$

With this measure of relative difference, a shape correlation of 0.9 corresponds to an error of 45%, a shape correlation of 0.99 corresponds to an relative difference of 14%, a shape correlation of 0.999 corresponds to an relative difference of 4%. Thus this more exacting measure \mathcal{E} from [22] is a far better representation of actual convergence between two shape functions than the correlation used in [20]. We will use this measure to test the accuracy and efficiency of our basis expansion in reconstructing the standard templates, and later to quantify the convergence of our validation examples in section ???. In that section we

also plot residuals on slices through the tetrapyd, relative to the representative value

$$\sqrt{\frac{S \cdot S}{\int_{T_k} d^3 k}}. \quad (2.17)$$

The squeezed limit of canonical single-field bispectra will not cause observable deviations from a Gaussian universe, due to a cancellation when switching to physical coordinates [23]. Here, we will only consider primordial phenomenology in comoving coordinates, so despite this cancellation, the squeezed limit is still a useful validation test of our results, using the standard single-field squeezed limit consistency condition [24, 25]. With $\mathbf{k}_S \equiv (\mathbf{k}_2 - \mathbf{k}_3)/2$:

$$S(k_1, k_2, k_3) = - \left[(n_s - 1)|_{k_S} + \mathcal{O}\left(\frac{k_1^2}{k_S^2}\right) \right] P_\zeta(k_1) P_\zeta(k_S), \quad k_1 \ll k_S \quad (2.18)$$

where $S(k_1, k_2, k_3)$ is again our dimensionless shape function. That the error in the consistency relation decreases at least quadratically in the long mode was shown in [25].

2.2.1 Easier to interpret, more stringent.

2.2.2 Notes on observable convergence being the deciding factor.

2.3 Template testing

2.3.1 Need to understand expected convergence before validation on numerical results.

2.4 Decomposing shapes on the cube vs tetrapyd

We begin our methods discussion by exploring possible sets of separable basis functions $Q_n(k_1, k_2, k_3)$ for use in the expansion (2.1). Whether the goal is to explore primordial phenomenology or for direct comparison with observations, the convergence of our basis set will determine the efficiency and practicality of our methods. We shall consider constructing the separable basis functions $Q_n(k_1, k_2, k_3)$ out of symmetrised triplet products of normalized one-dimensional modes $q_p(k)$ as

$$Q_n(k_1, k_2, k_3) \equiv \Xi_{prs} q_p(k_1) q_r(k_2) q_s(k_3). \quad (2.19)$$

Here, n labels the ordered integer triplet $n \leftrightarrow \{prs\}$ in an appropriate manner (see some ordering alternatives in [19]), while the symmetrised average of all $\{prs\}$ permutations is

$$q_{(p}q_rq_s) \equiv (1/3!) \sum_{\text{perms}} q_p q_r q_s \quad \text{and} \quad \Xi_{prs} = \begin{cases} 1, & p = r = s \text{ all equal,} \\ \sqrt{3} & \{prs\} \text{ any two equal,} \\ \sqrt{6} & \{prs\} \text{ all different.} \end{cases} \quad (2.20)$$

Unless stated otherwise, the $\{prs\}$ triples for each permutation set n in (2.20) are represented by the coefficient with $0 \leq p \leq r \leq s$, that is, $\alpha_n = \alpha_{prs} \equiv \alpha_{(prs)}$. This modal expansion is terminated at some p_{\max} for which $\max(p, r, s) < p_{\max}$. Given the basis-agnostic methods we shall outline in the following sections ??, ?? and ??, we are free to choose our set of basis functions to optimise for efficient convergence, ensuring our results are useful for comparison with observations. There are a wide variety of options available, such as polynomial bases or Fourier series, that can be chosen for the $q_p(k)$. While not strictly necessary for the method, it is more convenient if the resulting 3D basis functions $Q_n(k_1, k_2, k_3)$ are orthogonal on the cubic region of selected wavenumbers, making it much more straightforward to obtain controlled convergence. Overall, then, rapid convergence is the key criterion in choosing the basis functions $q_p(k)$ in (2.20), thus determining the nature of the numerical errors in the calculated bispectrum. However, since we are going beyond the featureless examples of [20] this matter deserves considerable care and close attention. Ideally we would have a three-dimensional basis that can efficiently capture a wide variety of shapes on the tetrapyd, with relatively few modes. In this work we aim for basis functions that work well in a wide variety of scenarios, so we endeavour to use as little specific information as possible (e.g. guessing the frequency of bispectrum oscillations from the power spectrum of a given scenario), though we will allow ourselves to use a representative value of the scalar spectral index, n_s^* . It is worth emphasising that a major advantage of the flexibility of the basis in the methods detailed in the following sections is the ease with which the basis can be modified to yield drastic increases in the rate of convergence at the primordial level, for the purposes of exploring primordial phenomenology.

In this section we will use some standard templates to investigate different possible sets of basis functions. An important issue is that when leveraging the separability of the in-in formalism, we are essentially forced to expand the shape function on the entire cube $[k_{\min}, k_{\max}]^3$. This is because the only decomposition we actually perform is a one-dimensional integral over $[k_{\min}, k_{\max}]$ (as we will see in (2.6)). With a uniform weighting, this integral does not know anything about the distinction between the tetrapyd and the cube. This is important as it means the non-physical configurations outside the tetrapyd will affect the convergence of our result on the tetrapyd, the region where we require

efficient convergence. To mimic this in testing our sets of basis functions, each shape will be decomposed on the entire cube, but the quoted measures of convergence will be between the shape and its reconstruction on the tetrapyd only (unless stated otherwise).

For a shape like (??) the non-physical off-tetrapyd configurations will not have a large effect, as the bispectrum on the faces of the cube is comparable to the bispectrum in the squeezed limit of the tetrapyd. On the other hand, for a shape of the equilateral type such as (??), this effect can be disastrous if not handled properly. This can be easily seen from (??), in the limit of small k_3 . The triangle condition in that limit enforces $(k_2 - k_1)^2 \leq k_3^2$. This implies that $0 \leq k_3^2 - (k_2 - k_1)^2 \leq k_3^2$, forcing the shape to go to zero in that limit despite the k_3 in the denominator. On the non-physical part of the face, $k_2 - k_1$ is not small, and so the shape is boosted by $1/k_3$ relative to the equilateral configurations. These regions then dominate any attempted basis expansion. To overcome this problem, as we shall discuss, we will extend our basis to explicitly include this $1/k$ behaviour⁴.

To date the most useful starting choice for modal bispectrum expansions has been shifted Legendre polynomials $P_r(x)$:

$$q_r(k) = \left(\frac{2r+1}{k_{\max} - k_{\min}} \right)^{1/2} P_r(\bar{k}), \quad (2.21)$$

with a rescaling of the argument \bar{k} to ensure the wavenumber k falls within the chosen (observable) domain $k_{\min} < k < k_{\max}$, that is,

$$\bar{k} = \frac{2k - k_{\max} - k_{\min}}{k_{\max} - k_{\min}}. \quad (2.22)$$

We shall label as \mathcal{P}_0 the basis function set of pure Legendre polynomials in (2.21), with $r = 0, 1, \dots, p_{\max}-1$. These were considered also in [20], however, while they prove to be particularly functional building blocks for other modal applications, in the context of the in-in formalism they converge so slowly even for simple shapes as to be inadequate when taken on their own. This poor rate of convergence with \mathcal{P}_0 for two local- and equilateral-type shapes is shown in figure 2.1. It is due to the $1/k$ behaviour inherent in these shapes, which is compounded in the equilateral models by pathologies exterior to the tetrapyd, as we have discussed. We can mitigate against this by including a basis function

⁴There are results in the literature that describe generic $K = k_1 + k_2 + k_3$ poles in correlators—see for example [26]. A simple example can be understood by recalling that in standard calculations using the in-in formalism, the $i\varepsilon$ prescription is used to damp out contributions in the infinite past. This does not work for $K = 0$. While the resulting divergence (in K) is clearly outside the physical region of the tetrapyd, we will see its effects in the physical configurations. Given that this three-dimensional behaviour is generic, one might worry that we should take more care in building it into our one-dimensional basis. However, the excellent convergence in section ?? shows that \mathcal{P}_1 and $\mathcal{P}_{01}^{n_s}$ can capture this behaviour well, and that this worry is unwarranted. In fact, since this behaviour comes from the oscillations at early times, observing this behaviour is a useful check on our results.

to capture this $1/k$ behaviour,

$$q_{p_{\max}}(k) = \text{Orth}[1/k], \quad \text{with} \quad \mathcal{P}_1 = \{q_r(k) \mid r = 0, 1, 2, \dots, p_{\max}\}, \quad (2.23)$$

where Orth represents the projection orthogonal to the original Legendre polynomial basis \mathcal{P}_0 . As we see in figure 2.1, the convergence properties for the augmented basis \mathcal{P}_1 are dramatically improved.

The two basis function sets actually used in [20] to calculate primordial bispectra were as follows. The first was the Legendre polynomials taken with a log-mapping between k and the polynomial argument as

$$q_r(k) = \left(\frac{2r+1}{\ln k_{\max} - \ln k_{\min}} \right)^{1/2} P_r(\overline{\ln k}) \quad \text{with} \quad \overline{\ln k} = \frac{2 \ln k - \ln k_{\max} - \ln k_{\min}}{\ln k_{\max} - \ln k_{\min}}. \quad (2.24)$$

The second basis was implicitly mentioned in a reference to the possibility of multiplying the functions to be decomposed by k , and dividing that factor out when evaluating the result. In our language, this is equivalent to working with an unnormalised basis set of the Legendre polynomials divided by k :

$$q_r(k) = \left(\frac{2r+1}{k_{\max} - k_{\min}} \right)^{1/2} \frac{P_r(\bar{k})}{k}, \quad (2.25)$$

where the rescaled \bar{k} is defined in (2.22). This can also be thought of as expanding the bispectrum $k_1 k_2 k_3 S(k_1, k_2, k_3)$ in \mathcal{P}_0 , instead of the shape function $S(k_1, k_2, k_3)$ itself. The consequence is that neither (2.24) nor (2.25) are orthogonal with respect to the flat weighting of the inner product (2.13). However, as shown in [20], these two basis sets (2.24) and (2.25) are able to approximate the three canonical bispectrum shapes. Nevertheless, our aim is to go beyond the featureless examples investigated in [20], so we require a basis that can capture many different forms of bispectrum features. To this end, we prefer not to weight the large or small wavelengths in our fit, as is done in (2.24) and (2.25). The deciding factor for which weighting is optimal to include in the primordial inner product is information about which configurations are most important for observables, that is, the expected signal-to-noise. We will not discuss this matter in detail here, except to note that motivated by the form of (??), we will take as our aim the accurate calculation of the primordial shape function with a flat weighting. Based on this motivation, we will not pursue (2.24) and (2.25) any further.

One could certainly also consider sets of basis functions more tailored to a particular example, or indeed even use power spectrum information to, on the fly, generate a basis tailored to a rough form of the expected bispectrum features. We save this possibility for

future work. In the following we will perform a more general exploration of orthogonal sets of basis functions that can efficiently describe the necessary $1/k$ behaviour. In addition to using the Legendre polynomials as building blocks, we will also consider a Fourier basis for the purposes of comparison.

Our general strategy will be to augment these basic building blocks with a small number of extra basis elements, while retaining orthogonality, using the standard modified Gram-Schmidt process. If we want to use some function f to augment a given set of orthogonal functions q_r , with $r = 0, \dots, p_{\max}-1$, then we define

$$\tilde{f}(k) = \text{Orth}[f(k)] \equiv f(k) - \sum_{r=0}^{p_{\max}-1} \frac{\langle f, q_r \rangle}{\langle q_r, q_r \rangle} q_r(k) \quad (2.26)$$

and add \tilde{f} to our basis set, now of size $p_{\max} + 1$. We note that the inner product here $\langle f, g \rangle$ is the 1D integral of the product $f(k)g(k)$ from k_{\min} to k_{\max} . The resulting basis is orthogonal, provided sufficient care is taken to avoid numerical errors.

In addition to our Legendre basis functions, pure \mathcal{P}_0 and augmented \mathcal{P}_1 , we will also introduce a Fourier series basis denoted by \mathcal{F}_0 and defined by

$$q_0(k) = 1, \quad q_{2r-1}(k) = \sin(\pi r \bar{k}), \quad q_{2r}(k) = \cos(\pi r \bar{k}), \quad 1 \leq r \leq (p_{\max}-3)/2 \quad (2.27)$$

$$q_{p_{\max}-2}(k) = \bar{k}, \quad q_{p_{\max}-1}(k) = \bar{k}^2. \quad (2.28)$$

Here, even the basic Fourier series have to be augmented by the linear k and quadratic k^2 terms (for a total size of p_{\max}), in order to satisfactorily approximate equilateral shapes (reflecting in part the preference for periodic functions). As with \mathcal{P}_1 defined in (2.23), we will similarly create an augmented Fourier basis \mathcal{F}_1 by adding the inverse $1/k$ term to the \mathcal{F}_0 basis, i.e. using $q_{p_{\max}}(k) = \text{Orth}[1/k]$ with (2.26). When we refer to convergence, we mean in increasing number of Legendre polynomials (or sines and cosines) within the initial set. The total size of the set will always be referred to as p_{\max} .

Notation	Building Blocks	Augmented by	Definition
\mathcal{P}_0	Legendre polynomials		(2.21)
\mathcal{F}_0	Fourier Series	k, k^2	(2.27)
\mathcal{P}_1	Legendre polynomials	k^{-1}	
\mathcal{F}_1	Fourier Series	k, k^2, k^{-1}	
$\mathcal{P}_1^{n_s}$	Legendre polynomials	$k^{-1+(n_s^*-1)}$	(2.29)
$\mathcal{P}_{01}^{n_s}$	Legendre polynomials	$k^{n_s^*-1}, k^{-1+(n_s^*-1)}$	(2.30)

Table 2.1: Basis summary—the augmentation of the basis is done using (2.26). The size of each basis is referred to as p_{\max} .

In order to compare the efficacy of these four different basis function sets ($\mathcal{P}_0, \mathcal{F}_0, \mathcal{P}_1$

and \mathcal{F}_1), we have investigated their convergence on Maldacena’s shape (??) and the DBI shape (??). To mimic the in-in calculation, we expand the shape on the cube, but test the result on the tetrapyd using (2.15). The results are shown in figure 2.1 where we find that the Legendre polynomials basis set \mathcal{P}_0 converges so slowly as to be unusable (with the Fourier modes \mathcal{F}_0 worse and not plotted). However, the augmented Legendre basis \mathcal{P}_1 (including $1/k$) leads to rapid convergence with an improvement of four orders of magnitude at $p_{\max} = 15$. The augmented Fourier basis \mathcal{F}_1 also converges quickly relative to \mathcal{P}_0 , but is outdone by \mathcal{P}_1 . Though we do not show the convergence on the cube, we find that for Maldacena’s template this is of the same order of magnitude as the error on the tetrapyd. For the DBI shape, however, the fit on the tetrapyd lags significantly behind, due to the effect of the large non-physical configurations. This explains the order of magnitude difference between the convergence at each p_{\max} for the two shapes in figure 2.1.

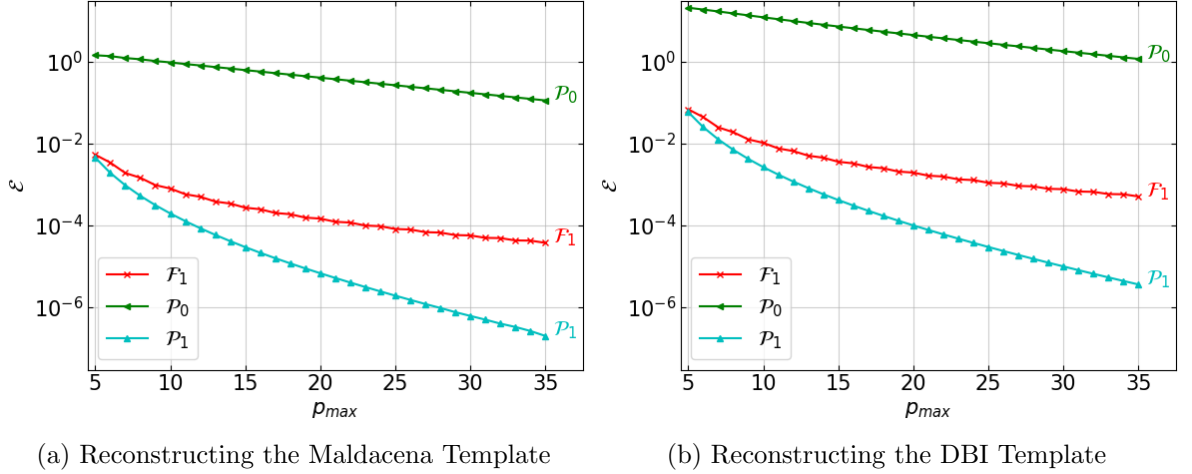


Figure 2.1: Convergence comparisons for the Legendre and Fourier basis functions for (a) the Maldacena template (??) and (b) the DBI template (??). The pure Legendre \mathcal{P}_0 basis requires many terms to fit the $1/k$ behaviour in both Maldacena’s template (??) and the DBI template (??). In contrast, the \mathcal{P}_1 basis (with an orthogonalised $1/k$ term) mitigates this dramatically, with the error already reduced by a factor of 100 at $p_{\max} = 5$. The Fourier \mathcal{F}_1 basis performs well, but converges more slowly than the \mathcal{P}_1 basis. Note that the convergence errors for (??) are larger than (??) because of the larger contributions outside the tetrapyd dominating the fit.

Next, we investigate oscillatory model templates. The simple feature model (??) and the resonance model (??) have scale dependence, but no shape dependence (in that they only depend on the perimeter of the triangle, $K = k_1 + k_2 + k_3$). We test our sets of basis functions on these two shapes, and also when they are multiplied by (??) to obtain a feature template with both shape and scale dependence. As shown in figure 2.2, \mathcal{F}_0 naturally outperforms \mathcal{P}_0 for a pure oscillation, but when the equilateral-type DBI template (??) is superimposed, even the augmented Fourier modes \mathcal{F}_1 converge poorly

and the Legendre modes \mathcal{P}_1 clearly offer a better more robust option. In figure 2.3 we see that for a logarithmic oscillation, \mathcal{P}_1 always converges in the fewest modes.

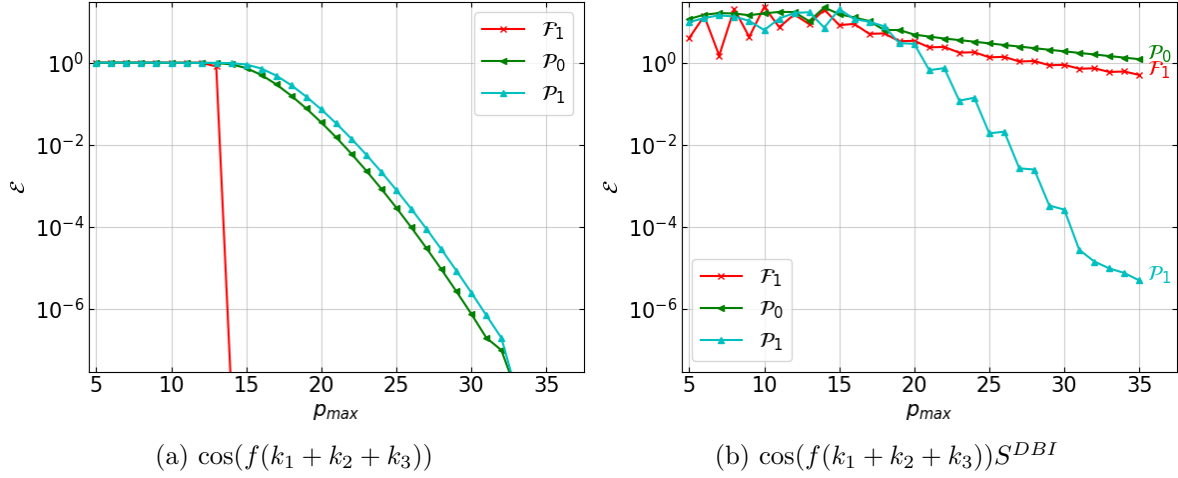


Figure 2.2: Convergence comparison for oscillatory models. (a) As expected, the \mathcal{F}_1 basis fits an oscillation with no shape dependence (??) (that is periodic in the k -range) perfectly. For this special case, the \mathcal{P}_0 and \mathcal{P}_1 sets of basis functions require more modes to accurately describe the shape. (b) However, moving to the more complex and realistic case of a feature with scale and shape dependence (in this case the product of (??) and (??)), we see that again \mathcal{P}_1 converges with the fewest modes. Note that before the expansion has fully converged, the fit on the tetrapyd can actually degrade slightly when the basis set is extended. This is an artifact of fitting on the cube and restricting (2.15) to the physical configurations on the tetrapyd; when considered over the entire cube the fit improves monotonically.

Finally, we consider convergence in the light of the more subtle scale-dependence due to the spectral index n_s of the power spectrum. The simple canonical examples in figure 2.1 had shape dependence and no scale dependence, but this would only be expected of scenarios unrealistically deep in the slow-roll limit. When we include this scale dependence, using (??) with n_s , it proves very useful to include these deviations from integer power laws in the basis functions. We consider two cases, first augmenting \mathcal{P}_0 by a scale-dependent $1/k$ term using the orthogonalisation procedure (2.26),

$$q_{p_{\max}}(k) = \text{Orth} [k^{-1+(n_s^*-1)}], \quad (2.29)$$

which we refer to as $\mathcal{P}_1^{n_s}$. Secondly, we also augment \mathcal{P}_0 with an additional scale-dependent constant term as

$$q_{p_{\max}}(k) = \text{Orth} [k^{(n_s^*-1)}], \quad q_{p_{\max}+1}(k) = \text{Orth} [k^{-1+(n_s^*-1)}], \quad (2.30)$$

which we refer to as $\mathcal{P}_{01}^{n_s}$. As we see in figure 2.4, for equilateral type shapes even a small overall scale dependence causes significant degradation in the convergence of the original

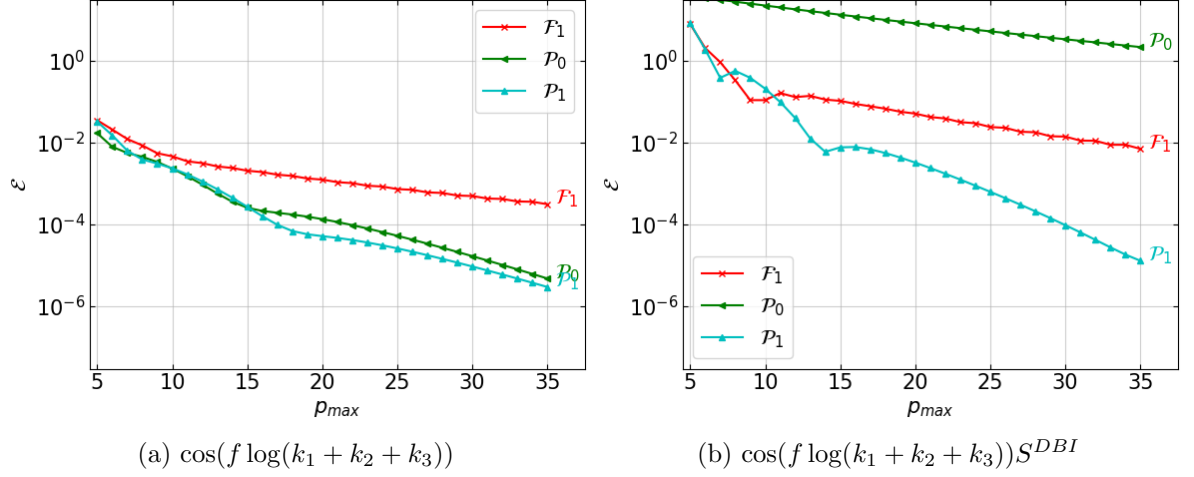


Figure 2.3: (a) The convergence for a log oscillation model (??) with no shape dependence. For this type of feature, the \mathcal{P}_0 and \mathcal{P}_1 sets of basis functions require fewer modes to accurately describe the shape than the \mathcal{F}_1 basis. (b) For the more complex and realistic case of a feature with scale and shape dependence (in this case the product of (??) and (??)), we see that again \mathcal{P}_1 converges with the fewest modes, though in this case \mathcal{F}_1 outperforms \mathcal{P}_0 .

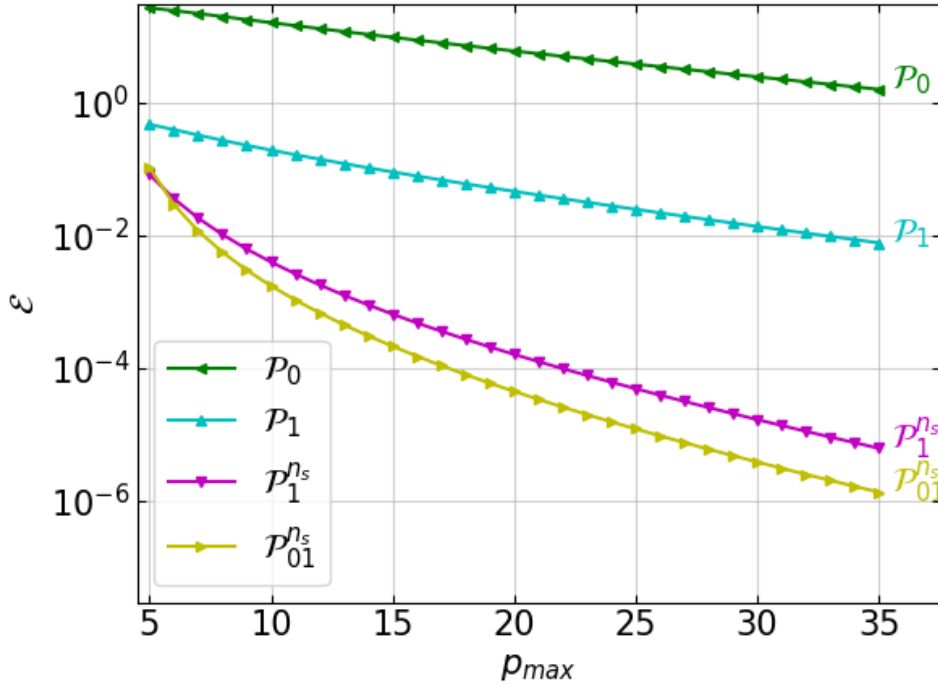


Figure 2.4: For the scale-dependent DBI template (??), by including a minimal amount of power spectrum information using (2.29) and (2.30) (here with $n_s^* = 0.97$), we can push the errors to less than 0.1% at $p_{\max} = 15$, allowing us to work with a basis that can efficiently capture the expected scale dependence of the shape function.

augmented Legendre basis \mathcal{P}_1 . However, incorporating the spectral index n_s into the basis functions $\mathcal{P}_1^{n_s}$ and $\mathcal{P}_{01}^{n_s}$ results again in rapid convergence to the scale-dependent DBI template, which can be accurately approximated with a limited number of modes. We conclude that augmenting the basis functions with terms incorporating the expected dependence on the spectral index enables the efficient approximation of high precision primordial bispectra.

2.4.1 Large non-physical contributions.

Words

2.4.2 FIGURE: DBI on cube vs tetra.

Words

2.5 Setting up a basis: augmentation

2.5.1 Using modified GS.

Words

2.5.2 Legendre and Fourier as building blocks

Words

2.6 Basis choice matters!

2.6.1 Why the basic basis expansion is so bad.

Words

2.6.2 My new basis sets, and their dramatic improvement.

2.6.3 Nice table with descriptions and some single-number comparisons Wordson on examples.

Words

2.6.4 FIGURES: Recon for Malda, DBI, scale-inv, with P0, P1, F0, F1 vs P_{max} .

Words

2.6.5 FIGURES: Recon for cos, cosDBI, with P0, P1, F0, F1 vs P_{max} .

Words

2.6.6 FIGURES: Recon for cos-log, cos-logDBI, with P0, P1, F0, F1 vs P_{max} .

Words

2.6.7 FIGURES: Recon for scaled-DBI, with P0, P1, P1ns, P01ns, P_{log} vs P_{max} .

Words

2.7 Not obvious, but k_{ratio} matters!

Words

2.8 The P_{max} - k_{ratio} tradeoff (and DBI vs Equil).

Words

2.9 Log basis for basis without PS info.

Words

2.10 Loginv basis for resonant shapes.

Words

2.11 Factor basis?

Words

2.12 Conclusion

2.12.1 Basic Legendres/Fourier are not sufficient.

Words

2.12.2 Need to pay attention to the cube.

Words

2.12.3 Including $1/k$ terms helps massively.

Words

2.12.4 The log basis performs excellently, without being tied to a specific n_s .

Words

Chapter 3

Methods and Validation

3.1 Numerics of mode evolution

3.1.1 τ_{u_s}

Words

3.1.2 When to set IC's for each mode.

Words

3.1.3 Swapping variables.

Words

3.1.4 Freeze-out.

Words

3.2 Integration choices

3.2.1 The $i\varepsilon$ prescription

Words

3.2.2 Expand on what F&RP wrote.

Words

3.2.3 Suppression from plane wave expansion (intuition).

Words

3.3 Starting the integration with a pinch

The previous two sections detailed the methods required to calculate the coefficients α_n in (2.1), obtaining an explicitly separable expression for the shape function. In this set-up, there are two kinds of integrals we must compute: integrals over time of the form (2.8), and integrals over k of the form (2.6). The first is done once per coefficient for each vertex, the second is a decomposition done once for ζ_k and ζ'_k each, every timestep. In this section we detail how to numerically evaluate these integrals accurately and efficiently.

Since calculating each point in the time integrand requires a decomposition (2.6), which is highly oscillatory at early times, it is worthwhile to consider how to perform the time integral efficiently. From the form of the Bunch-Davies mode functions, we expect the dominant frequency (in τ_s) to be $3k_{\max}$. Assuming we are earlier than any features that might change this, we can use this knowledge to sample the integrand at a far lower rate, building the oscillation into our quadrature weights. A second important consideration comes from how early we sample the integrand. We can of course only obtain a point in the time integrand after our mode functions have burned in from their set initial conditions to their true attractor trajectory. This means that sampling earlier in the time integrand requires us to set the initial conditions for the mode functions deeper in the horizon, a regime in which they are expensive to evolve.

The integrals of the form (2.8) that we must calculate have $\tau = -\infty$ as their lower limit. The highly oscillatory nature of the mode functions in these early times ($|k\tau_s| \gg 1$) suppresses the coefficients of our basis expansion by a factor of $1/\tau_s$. As noted in [20], this means that we do not need to explicitly use the $i\varepsilon$ prescription to force the integrals to converge. In the case of using the Legendre polynomials as our basis, we can see this more precisely by considering the plane wave expansion (e.g. [27]):

$$e^{-i\bar{k}(k_{\max}-k_{\min})\tau/2} = \sum_{n=0}^{\infty} (2n+1)i^n P_n(\bar{k}) j_n(-(k_{\max}-k_{\min})\tau/2) \quad (3.1)$$

for \bar{k} in $[-1, 1]$. When $(k_{\max}-k_{\min})\tau/2$ is large, the spherical Bessel functions oscillate with an amplitude $\propto \frac{1}{\tau}$. Thus, the initial conditions (??) expanded in Legendre polynomials (and similar) give us suppression of $1/\tau^3$ in (2.8).

While our method has extra suppression compared to configuration-by-configuration methods (and thus does not need the $i\varepsilon$ prescription to converge) it still converges rather slowly, as we push the lower limit to earlier times. This expensive sampling can be wasteful

of resources, especially in a feature scenario where we know this region will not contribute to the final result. Care is required however, as starting the integration in the wrong way can easily lead to errors which can completely swamp the result, since higher order modes are more sensitive to early times. The authors of [28] used an artificial damping term to smoothly “turn on” their integrand. The point at which this is done can then be pushed earlier to check for convergence. However they found that the details of the damping needed to be carefully set to avoid underestimating the result. In [29] they replaced this method by a “boundary regulator”; they split the integral into early and late parts and used integration by parts to efficiently evaluate the early time contribution. As our integrand already has extra suppression compared to the configuration-by-configuration integrands considered in [28, 29], we can safely use the simpler first method.

We understand this situation by taking advantage of asymptotic behaviour of highly oscillatory integrals (for a review see [30]). Since the leading order term depends on the value of the non-oscillatory part only at the endpoints, and the next-to-leading order correction depends on the derivative only at the endpoints, we can approximate the integral $\int_{-\infty}^T f(\tau_s) e^{iw\tau_s} d\tau_s$ by replacing the non-oscillatory part $f(\tau_s)$ with a function with matching value and derivative at $\tau_s = T$, but which converges far faster. We use

$$f(\tau_s) e^{-\beta^2(\tau_s - T)^2}, \quad (3.2)$$

for $\tau_s < T$. In this way, for sufficiently large T , we obtain the accuracy of the first two terms of the asymptotic expansion ($O(\beta^2/w^2)$, $w = 3k_{\max}$) without needing to explicitly calculate the derivative at T , or needing any phase information (as one would need to accurately impose a sharp cut on the integrand).

We use a damping of the form $e^{-\beta^2(\tau_s - T)^2}$ for $\tau_s < T$ to smoothly set the integrand to zero before a certain initial time, T . As long as T is sufficiently early and β is not too large their precise values have no significant effect on the final result. For definiteness, we take $\beta/w = 1 \times 10^{-4}$, small enough that the integrand has many oscillations while it is “turning on”, so matches the contribution of an infinite limit to high accuracy. We demonstrate this in figure 3.1 for a toy $H_{int} = (-1/\tau)\dot{\zeta}^3$, as in (??).

To obtain a k -sample we must evolve a Fourier mode from Bunch-Davies initial conditions deep in the horizon until it becomes constant after horizon crossing. We denote by N_k the number of Fourier modes we evolve. Different choices of distributing the k -samples are possible; for example, one could distribute them with an even spacing, log-spacing or cluster them more densely near k_{\min} and k_{\max} . The k -integrals themselves can be computed quite efficiently since at every timestep the integral is over the same sample points. One can therefore calculate and store the values of the basis functions at each of these points, along with the integration weights which will depend on the distribution of k -samples. The actual integration at each timestep, the calculation of the coefficients

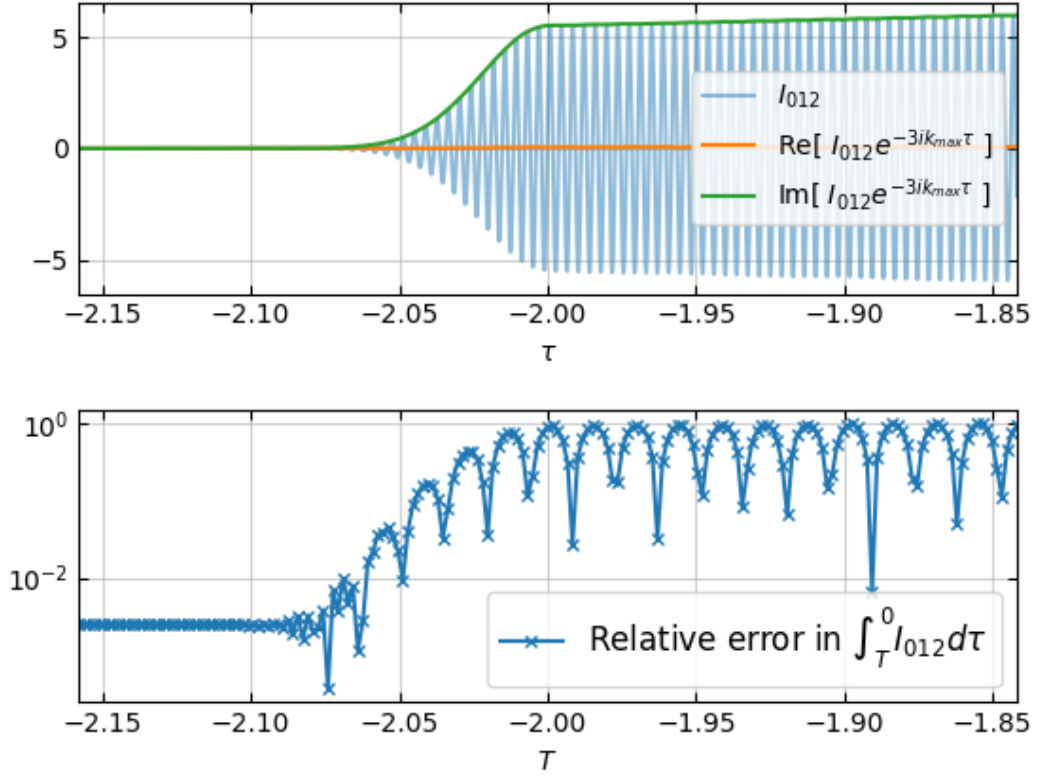


Figure 3.1: A toy example demonstrating the considerations involved in performing the time integrals (2.8). By carefully starting the time integrations, using the form (3.2), we can avoid errors that would otherwise swamp our result. The coefficient being calculated is the α_{012} coefficient of the \mathcal{P}_0 expansion of (??).

in (2.6), then becomes nothing more than a dot product of a time-independent array with the numerically evolved mode functions, for each order up to p_{\max} . We have found the best convergence results from distributing the k -samples according to the prescription of Gauss-Legendre quadrature.

To calculate the basis expansion of the bispectrum using the in-in formalism we must first calculate the basis expansion of the mode functions at each timestep (2.6). At early times the mode functions are highly oscillatory, taking the form $z_k e^{-ik\tau_s}$ for some much smoother z_k . Directly decomposing this would require evolving more ζ_k samples than is practical. We want an expansion of the form

$$z_k e^{-ik\tau_s} = \sum_{n=0}^{\infty} \alpha_n q_n(k). \quad (3.3)$$

We can obtain this by using standard oscillatory quadrature, if the τ_s dependence of the weights does not add too much overhead. We can also use an expansion of $e^{-ik\tau_s}$ with a known explicit time dependence, for example the expansion (3.1).

To use this second method, the first (smooth) factor z_k can be expanded in whatever basis we are working in, $q_n(k)$, and the second factor (highly oscillatory in k) is expanded in some convenient basis $\tilde{q}_n(k)$ (e.g. \mathcal{F}_0 , or \mathcal{P}_0 using the analytic form (3.1)). Then by precomputing $q_a(k)\tilde{q}_b(k)$ as a linear combination of the set of basis functions $q_c(k)$ all we need calculate at each timestep is the coefficients of the smoother $z_k(\tau_s)$, which we then convert to the coefficients of $F^{(i)}(\tau, k)$. In this way we can retain flexibility in our bispectrum basis, as well as efficiency and precision in the calculation. In the case of using \mathcal{P}_0 for the $\tilde{q}_n(k)$, assuming the expansion in (2.12) converges, we need only compute the expansion for $e^{-ik\tau_s}$ to enough terms that the first p_{\max} of the coefficients in the expansion (2.5) of the $F^{(i)}(\tau, k)$ converge, not until the actual sum (2.5) converges, since for high enough orders the integrals in (2.8) will integrate to zero.

Clearly, once τ_s becomes small enough these considerations will no longer be necessary and we can simply decompose the mode function directly. We do this around the horizon crossing of the geometric mean of k_{\min} and k_{\max} . If there is an extreme feature which causes a large deviation from the usual slow-roll form this switch will need to be made sooner. Also, this method would need to be adapted for non-Bunch-Davies initial conditions. Since anything related to the basis but independent of the scenario can be precomputed, certain parts of this calculation do not hurt the efficiency of this method in the context of, for example, a parameter scan. Using the methods outlined above, (2.8) and (2.6) can be computed precisely and efficiently in a mostly basis-agnostic context allowing us to (i) preserve the intrinsic separability of the tree-level in-in formalism and (ii) do so in a way that allows easy exploration of possible sets of basis functions, to find a set that converges quickly enough to be useful in comparison with observation.

3.3.1 When to start.

Words

3.3.2 Show how easy it is to swamp the result with a sharp start.

Words

3.3.3 **FIGURE:** Relative error for simple π^3 and for Reso, in (β, T) .

Words

3.4 Oscillatory weights

3.4.1 When to swap.

Words

3.4.2 Validation.

Words

3.5 Stopping the integration

3.5.1 A note on boundary terms and difficult (time-dep) cancellations.

Words

3.6 The interaction Hamiltonian

The methods detailed in the previous section depend on the separability of the third-order interaction Hamiltonian, H_{int} , and the possibility of including the spatial derivatives in a numerically accurate and efficient way. To make precise how our methods take into account the details of H_{int} , we will take $P(X, \phi)$ inflation as an example. The full cubic

interaction Hamiltonian, not neglecting boundary terms, can be calculated as [2, 31, 32]

$$\begin{aligned}
H_{int}(t) = \int d^3x \Big\{ & -\frac{a^3\varepsilon}{Hc_s^4} \left(1 - c_s^2 - 2c_s^2 \frac{\lambda}{\Sigma}\right) \dot{\zeta}^3 + \frac{a^3\varepsilon}{c_s^4} (3 - 3c_s^2 - \varepsilon + \eta) \zeta \dot{\zeta}^2 \\
& - \frac{a\varepsilon}{c_s^2} (1 - c_s^2 + \varepsilon + \eta - 2\varepsilon_s) \zeta (\partial\zeta)^2 \\
& - \frac{a^3\varepsilon^2}{2c_s^4} (\varepsilon - 4) \dot{\zeta} \partial\zeta \partial(\partial^{-2}\dot{\zeta}) - \frac{a^3\varepsilon^3}{4c_s^4} \partial^2\zeta (\partial(\partial^{-2}\dot{\zeta}))^2 \Big\} \quad (3.4)
\end{aligned}$$

with $\Sigma = \frac{H^2\varepsilon}{c_s^2}$ and $\lambda = X^2 P_{,XX} + \frac{2}{3} X^3 P_{,XXX}$. See [2] for further details.

This is commonly quoted with a term proportional to the equation of motion, but this will never contribute [2, 33, 34, 35]. We do not need to make a slow-roll approximation (the quantities defined in (??) are not required to be small, except in that we wish to have a successful inflation scenario), nor do we need to neglect any terms in the interaction Hamiltonian. We do no field redefinition, so do not need to add a correction to the final bispectrum. Following the calculation of [2] (see also [33, 34, 35]) we do not work with any boundary terms. Numerically this is preferable to forms with boundary terms, whether they come from undoing a field redefinition or from integration by parts. Since the boundary term contribution will depend on the choice of when to end the integration, its time dependence must cancel with a late-time time-dependent contribution of some vertex, requiring us to track the necessary quantities much longer than otherwise needed to obtain the desired precision.

Schematically, the correction from a field redefinition would look like

$$\langle \zeta_{\mathbf{k}_1} \zeta_{\mathbf{k}_2} \zeta_{\mathbf{k}_3} \rangle = \langle \tilde{\zeta}_{\mathbf{k}_1} \tilde{\zeta}_{\mathbf{k}_2} \tilde{\zeta}_{\mathbf{k}_3} \rangle + \lambda \langle \tilde{\zeta}_{\mathbf{k}_1} \tilde{\zeta}_{\mathbf{k}_2} \rangle \langle \tilde{\zeta}_{\mathbf{k}_1} \tilde{\zeta}_{\mathbf{k}_3} \rangle + \text{cyclic} \quad (3.5)$$

where λ is some function of the slow-roll parameters. The correction terms will have a time dependence from λ , so the $\langle \tilde{\zeta}_{\mathbf{k}_1} \tilde{\zeta}_{\mathbf{k}_2} \tilde{\zeta}_{\mathbf{k}_3} \rangle$ term must have some late time contribution to cancel it. To obtain an accurate result, care would need to be taken with this cancellation, an unnecessary complication.

By integrating by parts and using the equation of motion, the interaction Hamiltonian can be rewritten without picking up boundary terms [36]. Using (3.7) from [36], with $f = -\varepsilon/(c_s^2 H)$, we obtain the following form:

$$\begin{aligned}
H_{int}(t) = \int d^3x \Big\{ & -\frac{a^3\varepsilon}{Hc_s^4} \left(-c_s^2 - 2c_s^2 \frac{\lambda}{\Sigma}\right) \dot{\zeta}^3 + \frac{a^3\varepsilon}{c_s^4} (-3c_s^2) \zeta \dot{\zeta}^2 \\
& - \frac{a\varepsilon}{c_s^2} (-c_s^2) \zeta (\partial\zeta)^2 - \frac{a\varepsilon}{Hc_s^2} \dot{\zeta} (\partial\zeta)^2 \\
& - \frac{a^3\varepsilon^2}{2c_s^4} (\varepsilon - 4) \dot{\zeta} \partial\zeta \partial(\partial^{-2}\dot{\zeta}) - \frac{a^3\varepsilon^3}{4c_s^4} \partial^2\zeta (\partial(\partial^{-2}\dot{\zeta}))^2 \Big\}. \quad (3.6)
\end{aligned}$$

To leading order, this formulation is made up of terms that give equilateral shapes when

the slow-roll parameters are roughly constant. It was pointed out in [20] that using (3.4) in a scenario that results in an equilateral shape would require sensitive cancellations in the squeezed limit. Likewise, using (3.6) for a local scenario would require sensitive cancellations in the equilateral limit.

As mentioned in [20], the spatial derivatives can be manipulated into simple prefactors of k_i using the triangle condition ($\mathbf{k}_1 + \mathbf{k}_2 + \mathbf{k}_3 = 0$), and so preserve the separability of the result. To absorb these prefactors in our calculation, we precompute $k^p q_a(k)$ as a linear combination of the $q_a(k)$ for the relevant values of p , from which $V_{p\bar{p}}^{(i)}$ defined in (2.9) is built. For certain sets of basis functions this matrix can be calculated analytically, but it is simpler and more robust to numerically calculate the relevant integral directly. The processing cost this incurs is small, and must only be paid once per basis. We note especially that this means the matrix can be stored and efficiently used in many scenarios. To summarise, we calculate the bispectrum contribution from each vertex in H_{int} separately: we assemble the integrands, integrate them with respect to time, include the prefactors coming from the spatial derivatives, then sum the resulting sets of basis coefficients. Of course, these methods are not restricted to this example of H_{int} .

3.6.1 Using integration by parts from RP paper.

3.7 Set-up of DBI example scan

3.7.1 Matching A_s , n_s , r , N_* , using β_{IR} etc.

3.8 Going to high P_{max}

3.8.1 How high can I go, in terms of Planck w ?

3.8.2 How high can I go, in terms parameters that can't be compared to Planck?

3.9 Validation (on numerical results)

3.9.1 Validation methods

In this section we validate our implementation of our methods on different types of non-Gaussianity, sourced in different ways. While our actual results take the form of a set of mode expansion coefficients α_n , to make contact with previous results in the literature all of our validation tests take place on the tetrapyd, the set of physical bispectrum configurations.

We test that our results have converged using (2.15), between $p_{\max} = 45$ and $p_{\max} = 15$ for the featureless cases, and between $p_{\max} = 65$ and $p_{\max} = 35$ for the cases with features. We will refer to this as our convergence test. To verify that our results have converged to the correct shape, we perform full tetrapyd checks against known analytic results (where those are available, and in their regimes of validity) using (2.15), and point tests against the PyTransport code for the scenarios with canonical kinetic terms. Since all our scenarios are single-field, the most general test we have is the single-field consistency relation, which states that for small k_L/k_S , the shape function $S(k_S, k_S, k_L)$ must obey (2.18). The consistency condition should hold most precisely at the configurations with smallest k_L/k_S , the most squeezed being the three corners, $(k_{\max}, k_{\max}, k_{\min})$ and permutations. We want our test to be on an extended region of the tetrapyd however, so we choose the line

$$\frac{k_L}{k_S} = \frac{2k_{\min}}{k_{\max}}, \quad (3.7)$$

which connects $(k_{\max}, k_{\max}, 2k_{\min})$ to $(k_{\max}/2, k_{\max}/2, k_{\min})$. We will take $\frac{k_{\min}}{k_{\max}} = \frac{1}{550}$, so this is still sufficiently squeezed to be a stringent test.

First, we investigate convergence on simple featureless models, both local-type (??) and equilateral-type (??). We find that in our chosen basis $\mathcal{P}_{01}^{n_s}$ our results converge quickly and robustly as we increase the number of modes, where we quantify the convergence using (2.15). We compare the converged results against analytic templates (??) and (??), using the full shape information (2.15), finding them to match to high accuracy. Secondly we validate our methods on an example of non-Gaussianity from a feature: linear oscillations from a sharp step in the potential (??). The result converges robustly across the parameter range we explore. Throughout that range, we test the converged result using the squeezed limit consistency condition, and perform point tests against PyTransport, finding excellent agreement. For small step size we can further validate against the analytic template of [5], using the full shape information, finding agreement to the expected level given the finite width of the step. The final type of non-Gaussianity we use for validation on is the resonance type, logarithmic oscillations generated deep in the horizon (??). We test the converged result against the PyTransport code, by performing point tests on a slice. We also present a resonant DBI scenario, with out-of-phase oscillations in the flattened limit, as pointed out in [37], resulting from non-Bunch-Davies behaviour of the mode functions. We also test both resonant scenarios using the squeezed limit consistency condition.

We display the phenomenology of our various validation examples by plotting slices through the tetrapyd, as detailed in figure 3.2. Along with the phenomenology plots we plot the residual (with respect to the totally converged result) on the same slice, relative to the magnitude of the shape (2.17). We emphasise that while these plots display slices through the tetrapyd, our actual result describes the shape function on the entire three-dimensional

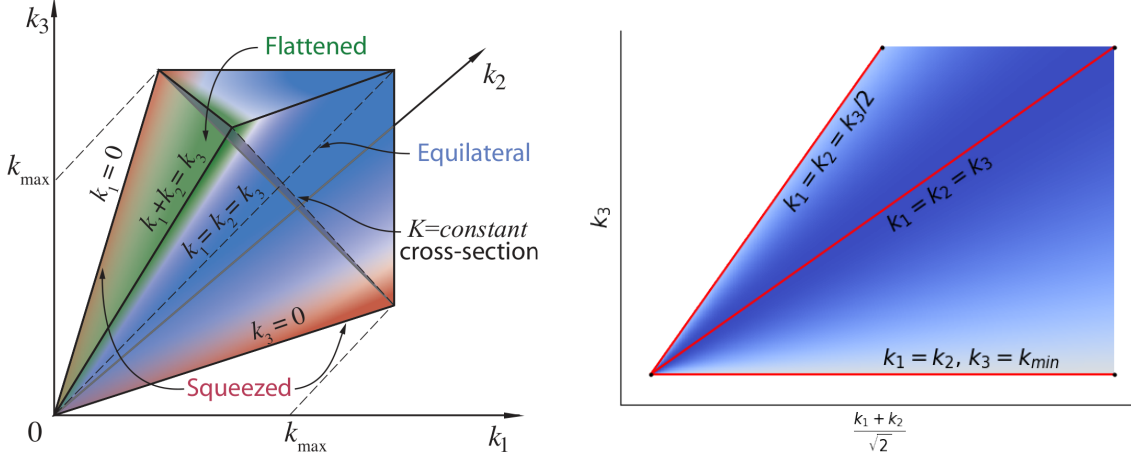


Figure 3.2: For ease of display, we will plot the two-dimensional $k_1 = k_2$ slice of the tetrapyd for each of our validation examples, as shown schematically here on the right. Horizontal lines on this plot have constant k_3 . The bottom edge is $k_3 = k_{\min}$, the top edge is $k_3 = k_{\max}$. The right edge is $k_1 = k_2 = k_{\max}$, the left edge is $k_1 = k_2 = k_3/2$, i.e. the limit imposed by the triangle condition. Plotted in red (in the right hand plot) from top-left to bottom-right, are the flattened, equilateral and squeezed limits. For comparison, half of the tetrapyd is shown in the three-dimensional plot on the left.

volume of the tetrapyd, and we measure our convergence over this whole space.

While one of the main advantages of this method is its direct link to the CMB, in this section we only concern ourselves with validating the code, not the observational viability of the scenarios considered. We focus on accurately and efficiently calculating the primordial tree-level comoving bispectrum, validating on models popular in the literature.

3.9.2 Quadratic slow-roll

The first model we will consider is slow-roll inflation on a quadratic potential (??). We consider two scenarios, both with $m = 6 \times 10^{-6}$. The first is deep in slow-roll, which we achieve by choosing $\phi_0 = 1000$; then, choosing ϕ'_0 according to the slow-roll approximation, we get $\frac{1}{2}\phi'^2 = \varepsilon \approx 0.2 \times 10^{-5}$. We can then choose the initial value for H to satisfy the Friedmann equation to sufficient precision. The second scenario is chosen to have a value for $n_s^* - 1$ consistent with the *Planck* result, by choosing $\phi_0 = 16.5$, so that $\varepsilon \approx 0.8 \times 10^{-2}$. The shapes are shown in figure 3.3.

We choose the first scenario to have such a small value of ε so that we can use Maldacena's shape (??) as a precision test. Indeed, we find that it has a scaled relative difference (2.16) of 2.7×10^{-5} with this shape, contrasting a scaled relative difference of 0.077 with the local template (??). This confirms that our methods and our implementation in code can accurately pick up this basic type of featureless non-Gaussianity.

For the second scenario, we cannot validate on Maldacena’s shape (??) or the local template (??), as for $\epsilon \approx 0.8 \times 10^{-2}$ we only expect these templates to match the true result to percent level accuracy. Indeed, we find that our result has a correlation of 0.998 with both (??) and (??), corresponding (in the sense of (2.16)) to a relative difference of 6%, as expected. Instead, we validate this model using the squeezed limit test described above, verifying our result to 0.05%.

This is a validation of the convergence of our basis, reaffirming the template decomposition results of figure 2.1 in the setting of the in-in formalism. It is also a stringent validation of our methods of including the higher-order coefficients, as insufficient care taken in the early-time sections of integrals (2.8), or in including the spatial derivatives from H_{int} , could have easily swamped the $p_{\max} = 45$ result.

3.9.3 DBI inflation

Next, we show results for a similar pair of scenarios for DBI inflation. We choose $V_0 = 5.2 \times 10^{-12}$ with $m = \sqrt{0.29V_0/3}$ in (??) and (??). We choose $\phi_0 = 0.41$, and then the starting condition for H according to the slow-roll approximation, allowing us to choose ϕ'_0 such that the Friedmann equation is satisfied to sufficient precision. The first scenario is deep in slow-roll, with $\lambda_{DBI} = 1.9 \times 10^{18}$, while the second scenario saturates the *Planck* limit on c_s , with $\lambda_{DBI} = 1.9 \times 10^{15}$. The resulting shapes are shown in figure 3.4.

The scenario deep in slow-roll has a error of 0.082% relative to the DBI shape (??), and 13% relative to the equilateral template (??). The second scenario has a relative error of 2.9% with the scale-invariant DBI shape, and 14% with the equilateral template. Including some scale dependence in the template, using (??), we get a relative error of 0.27%. On the line defined by (3.7), both scenarios have a sub-percent difference from the consistency condition, with respect to the equilateral configurations, which decreases when configurations with a larger k_S/k_L are considered.

Including the minimal information of an individual, approximately representative value of $n_s^* - 1$ in $\mathcal{P}_{01}^{n_s}$ allows us to converge to these smooth shapes quickly and robustly, overcoming the tetrapyd-vs-cube difficulties described in ??. Our accurate match to these shapes validates our implementation in code, and the ability of the method (and our basis in particular) to capture very different types of bispectrum shapes, local and equilateral.

3.9.4 Step features

Moving on from simple featureless bispectra, we present the results of our validation tests on non-Gaussianity coming from a sharp feature in the potential. We use the same parameters for the quadratic potential as in the second scenario in fig 3.3. In (??) we fix $d = 1 \times 10^{-2}$ and $\phi_f = 15.55$ (as with the second canonical quadratic example, $\phi_0 = 16.5$).

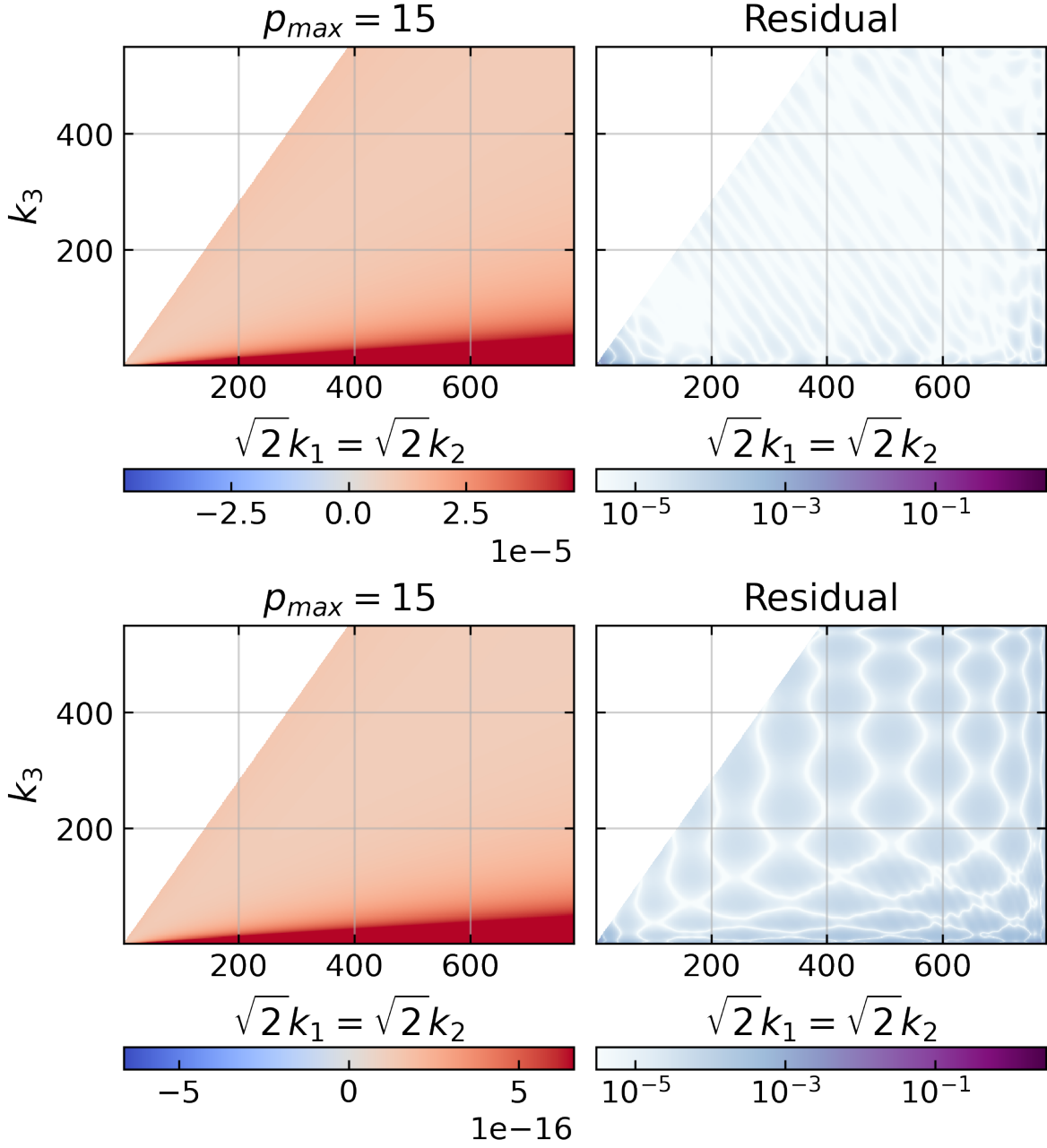


Figure 3.3: A canonical single-field model on a quadratic potential (??), slowly-rolling with $\varepsilon \approx 2 \times 10^{-6}$ in the top plot, and $\varepsilon \approx 0.8 \times 10^{-2}$ in the lower plot. This shape is dominated by its squeezed limit, and has a scale dependence determined by ε , very small in the top plot and “realistic” in the lower plot, relative to the *Planck* power spectrum. The first scenario converges well in the \mathcal{P}_1 basis, with a relative difference of 2.7×10^{-5} between $p_{\max} = 45$ and $p_{\max} = 15$. The second scenario converges well in the $\mathcal{P}_{01}^{n_s}$ basis (with $n_s^* - 1 = -0.0325$), with a relative difference of 7.9×10^{-5} between $p_{\max} = 45$ and $p_{\max} = 15$.

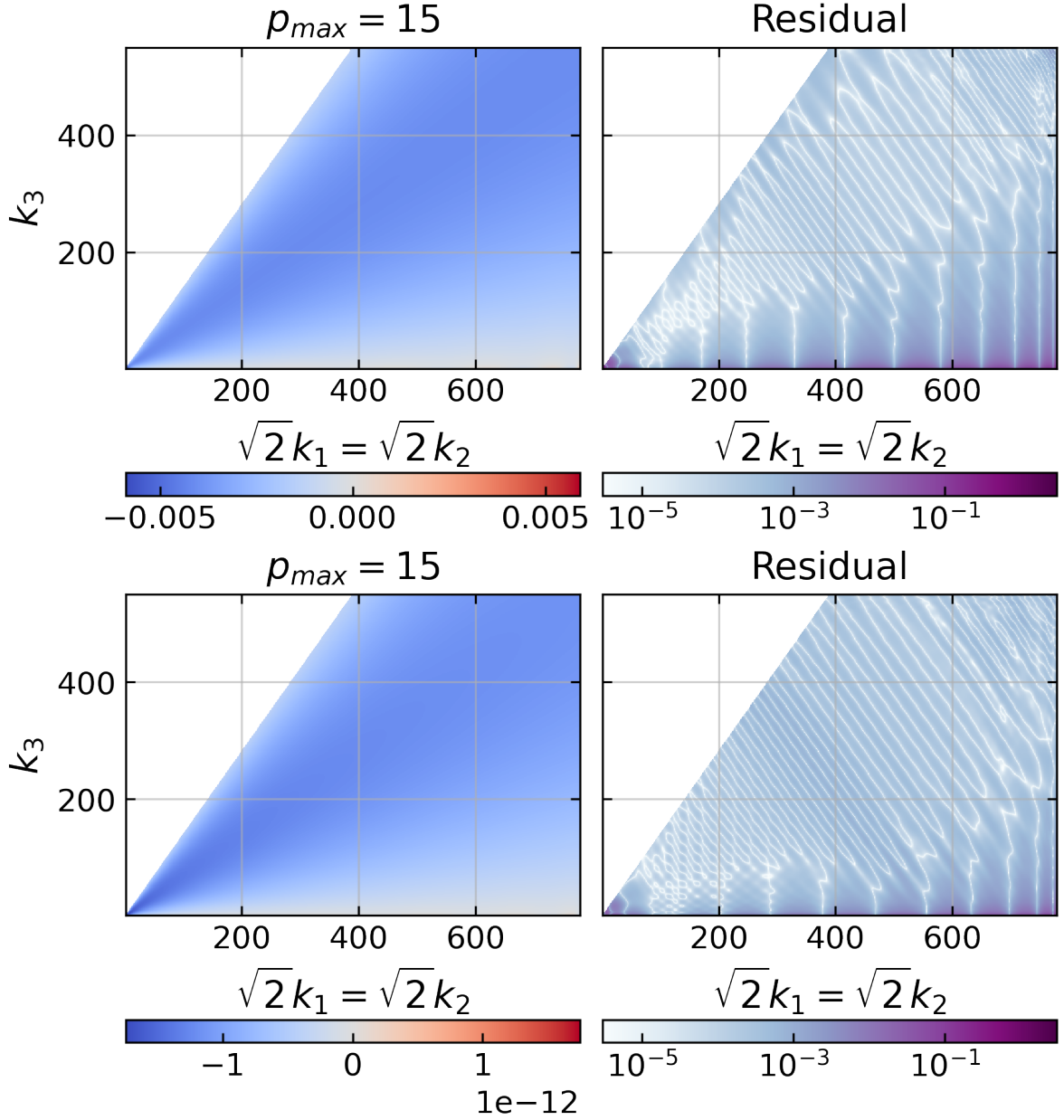


Figure 3.4: The upper plot shows the shape function for a DBI model deep in slow-roll. We set λ_{DBI} in (??) to 1.9×10^{18} , obtaining a scenario with $\varepsilon \approx 1.9 \times 10^{-6}$ and $c_s = 2.3 \times 10^{-3}$. This shape is dominated by its equilateral configurations, and has only a slight scale dependence. It converges well in the \mathcal{P}_1 basis, with a relative difference of 2.1×10^{-3} between $p_{\max} = 45$ and $p_{\max} = 15$. The lower plot shows a DBI model that saturates the *Planck* limit on c_s . We set λ_{DBI} in (??) to 1.9×10^{15} , obtaining a scenario with $\varepsilon \approx 8.0 \times 10^{-5}$ and $c_s = 8.0 \times 10^{-2}$. This shape is also dominated by its equilateral configurations, but has a scale dependence consistent with the measured power spectrum. It converges well in the $\mathcal{P}_{01}^{n_s}$ basis (with $n_s^* - 1 = -0.0325$), with a relative difference of 1.1×10^{-3} between $p_{\max} = 45$ and $p_{\max} = 15$.

Figure 3.5 shows results for the shape function for two step sizes, $c = 5 \times 10^{-5}$ and $c = 5 \times 10^{-3}$. The resulting shape for small step sizes contains simple oscillations, linear in $k_1 + k_2 + k_3$, whose phase is almost constant across the tetrapyd. When the step size is small, as expected, our result matches the analytic result of [5], presented there in equations (48), (54), (55). We plot a comparison of the result of [5] and our result in figure 3.7. For larger step size, we check the squeezed limit in figure 3.6, where we also show point tests against the PyTransport code. Across this range of step sizes, for the resulting shapes we obtain a full tetrapyd convergence test result (between $p_{\max} = 65$ and $p_{\max} = 35$) of between 0.17% and 0.15% and we verify the squeezed limit test to better than 0.5%.

These examples show the utility of our methods in calculating bispectra with non-trivial shape and scale dependence, going beyond the simple examples of [20]. They validate the calculation of the high order coefficients, and show that our code as implemented can handle sharp deviations from slow-roll, generating non-Gaussianity around horizon crossing.

3.9.5 Resonance features

Now we further validate our code against two resonance models. In contrast to the previous sharp kink, this feature is extended, requiring precision at earlier times. The first, shown in figure 3.8, is a model with a canonical kinetic term, on a quadratic potential with a superimposed oscillation (??). We take $bf = 10^{-7}$, and $f = 10^{-2}$. The resulting bispectrum has oscillations logarithmic in $k_1 + k_2 + k_3$. In figure 3.8 we see the excellent agreement between our result and the PyTransport result, once initial conditions in both codes are set early enough to achieve convergence. This validates the code on non-Gaussianity generated deeper in the horizon. Note the change of phase in the squeezed limit, though this is expected to be unobservable. We obtain a full tetrapyd convergence test result (between $p_{\max} = 65$ and $p_{\max} = 35$) of 0.93%, a squeezed limit test result of 1.1% (along the line defined by (3.7)), and a relative difference of 3.0% with respect to the PyTransport result, although this is only integrated over the two-dimensional slice presented in figure 3.8.

The time taken for the PyTransport code (per configuration) varies by a factor of around forty between the equilateral limit and the squeezed limit, as we show in figure 3.8. While the PyTransport code is extremely fast at calculating the shape function for a single k -configuration, to obtain this two-dimensional slice through the tetrapyd took around seven hours; to obtain the shape function on the full three-dimensional tetrapyd would take much longer. In contrast, our code took less than an hour on the same machine to calculate the full shape function, not limited to the shown slice. The overall speed increase is, therefore, a factor on the order of 10^2 to 10^3 for the full shape information, speaking only on the level of primordial phenomenology, in addition to the advantage that

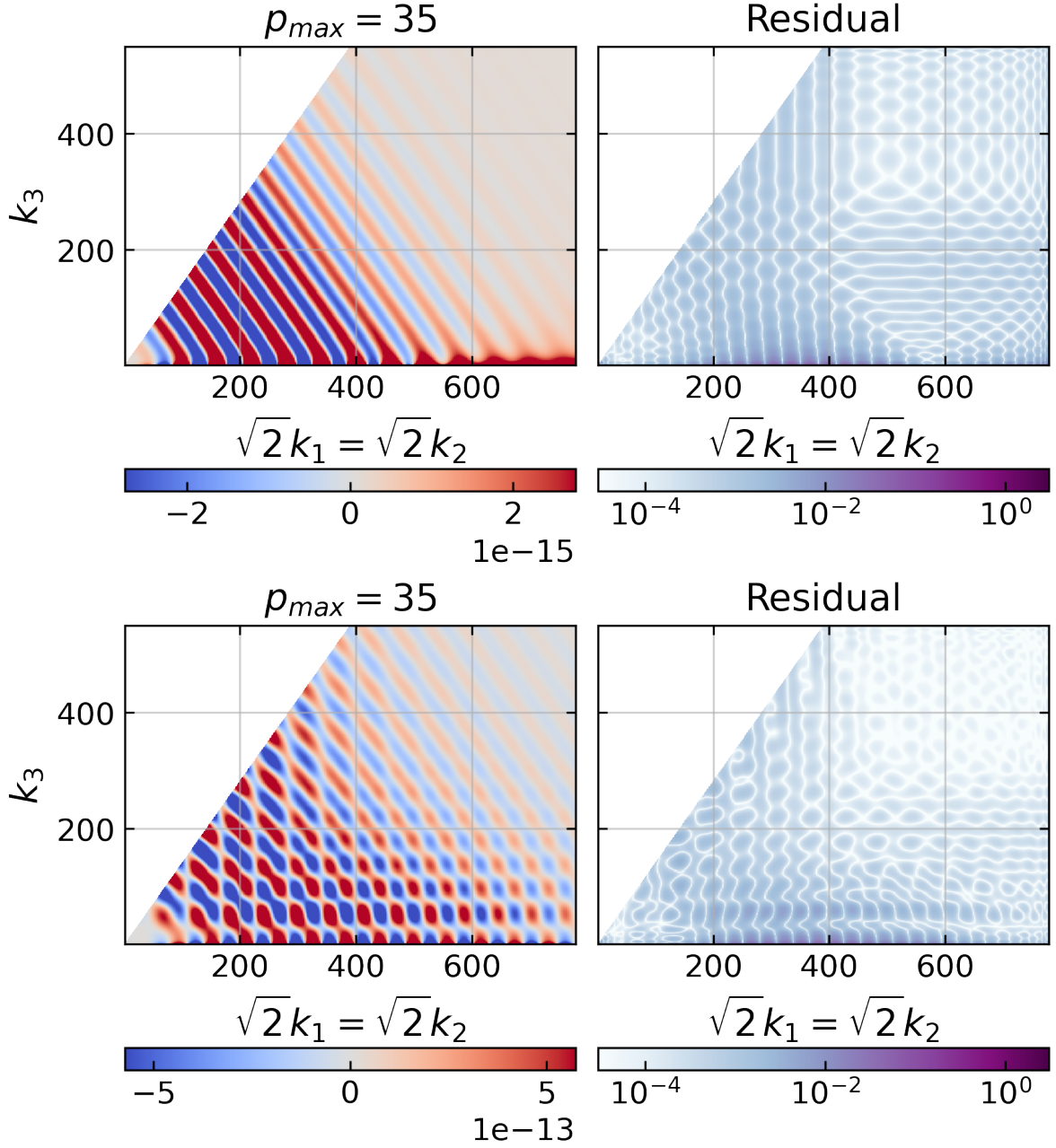


Figure 3.5: The tree-level shape function of a feature model (??), shown for step sizes of $c = 5 \times 10^{-5}$ (upper plot) and $c = 5 \times 10^{-3}$ (lower plot). The corresponding expansion parameter values of [5], $\mathcal{C} = 6c/(\varepsilon + 3c)$, are 0.035 and 1.3. For the smaller step size, the oscillations are almost entirely functions of $K = k_1 + k_2 + k_3$, except for a phase difference in the squeezed limit. The dependence is more complicated for $\mathcal{C} = 1.3$, however our result still converges well. In the $\mathcal{P}_{01}^{n_s}$ basis, with $n_s^* - 1 = -0.0325$, the results have a relative difference of 1.6×10^{-3} and 1.5×10^{-3} , respectively, between $p_{\max} = 65$ and $p_{\max} = 35$.

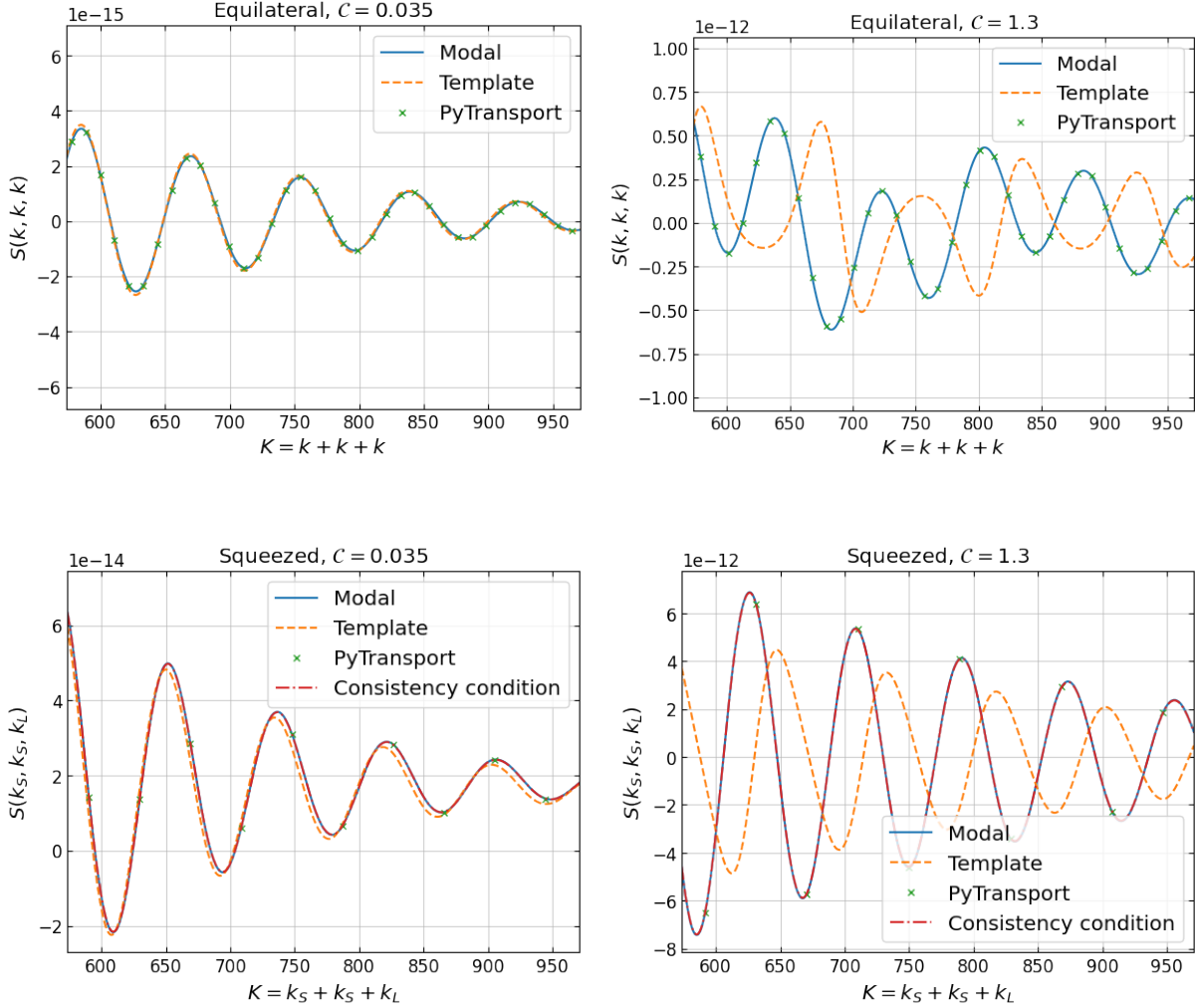


Figure 3.6: In the equilateral limit for the feature models (the top two figures) we validate our modal result against the PyTransport result. In the squeezed limit (the bottom two figures) we validate against PyTransport, and the consistency condition. In both limits, for both step sizes shown, we find excellent agreement. For the small step size (the two plots to the left), we additionally see a good match to the template of [5]. For the larger step size, the template amplitude is still accurate, but no longer captures the detailed shape information. This validates our code on non-Gaussianity generated by sharp features, and illustrates the general usefulness of our method. Our numerical results are accurate in a broader range than approximate templates, but are still smooth separable functions, unlike the results of previous numerical codes.

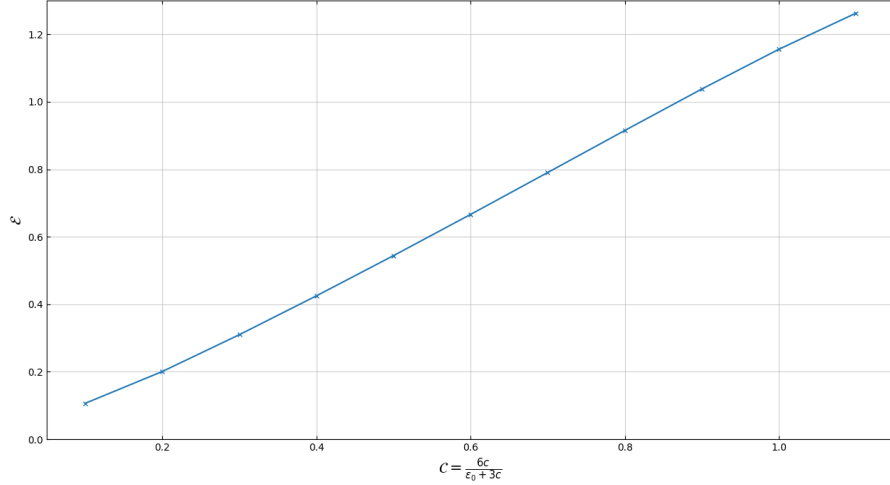


Figure 3.7: We sample more shapes with step sizes between the two feature models shown in figure 3.5. We plot the relative difference, integrated over the full tetrapyd in the sense of (2.15), between the modal result and the analytic template of [5], as a function of the template parameter $\mathcal{C} = \frac{6c}{\epsilon_0 + 3c}$ (where c is the step size and ϵ_0 is the value of the slow-roll parameter ϵ at ϕ_{step} when $c = 0$). We test our result by verifying the squeezed limit consistency condition to better than 1% throughout (not shown). The number of oscillations in the k -range is determined by the conformal time at which the kink in (??) occurs, which is kept constant across this scan. The width of the feature was also kept constant.

our result is in a form designed to be compared with observation. We expect that our implementation can be optimised beyond this.

The second scenario we consider here also has an oscillation superimposed on its potential, but this time is a non-canonical model, the DBI model. The resulting bispectrum is shown in figure 3.9. Note especially the out-of-phase oscillations in the flattened limit, which are potentially observable. For the purpose of displaying this phenomenology, we place a window on the oscillation in the potential, smoothing out the resulting oscillations in the shape at low $k_1 + k_2 + k_3$, to aid convergence. This validates our code on non-Gaussianity generated by deviations from Bunch-Davies behaviour [37, 38]. We obtain a convergence test result (between $p_{\max} = 65$ and $p_{\max} = 35$) of 0.15%, and a squeezed limit test result of 6.5%.

3.10 Speed comparison vs PyTransport. Something BINGO can't do?

Words

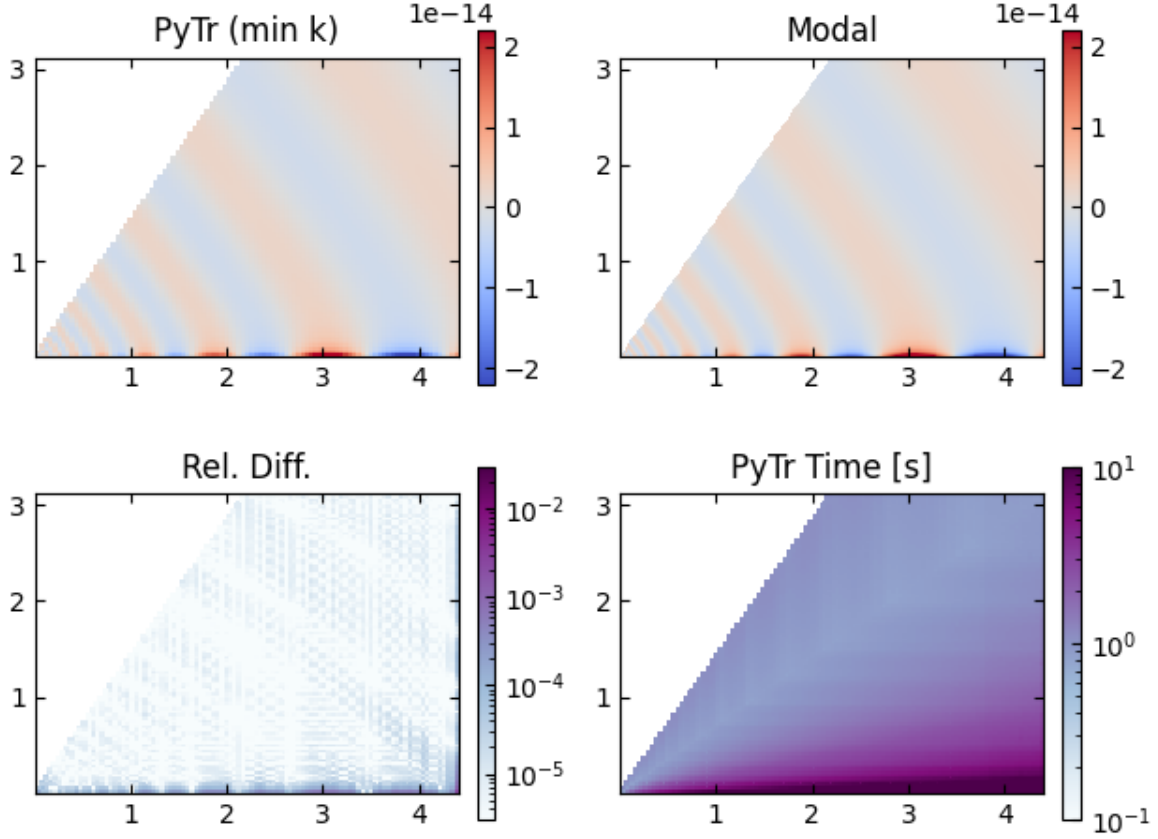


Figure 3.8: Resonance on a quadratic potential (??), testing our result using point tests against the PyTransport code. The logarithmic oscillations in the shape function are generated by periodic features deep in the horizon. The differences between our result and the PyTransport result are sufficiently small throughout that we can consider this a validation of our code on non-Gaussianity generated by periodic features deep in the horizon. In the $\mathcal{P}_{01}^{n_s}$ basis, with $n_s^* - 1 = -0.0325$, our result has a relative difference of 9.6×10^{-3} between $p_{\max} = 65$ and $p_{\max} = 35$.

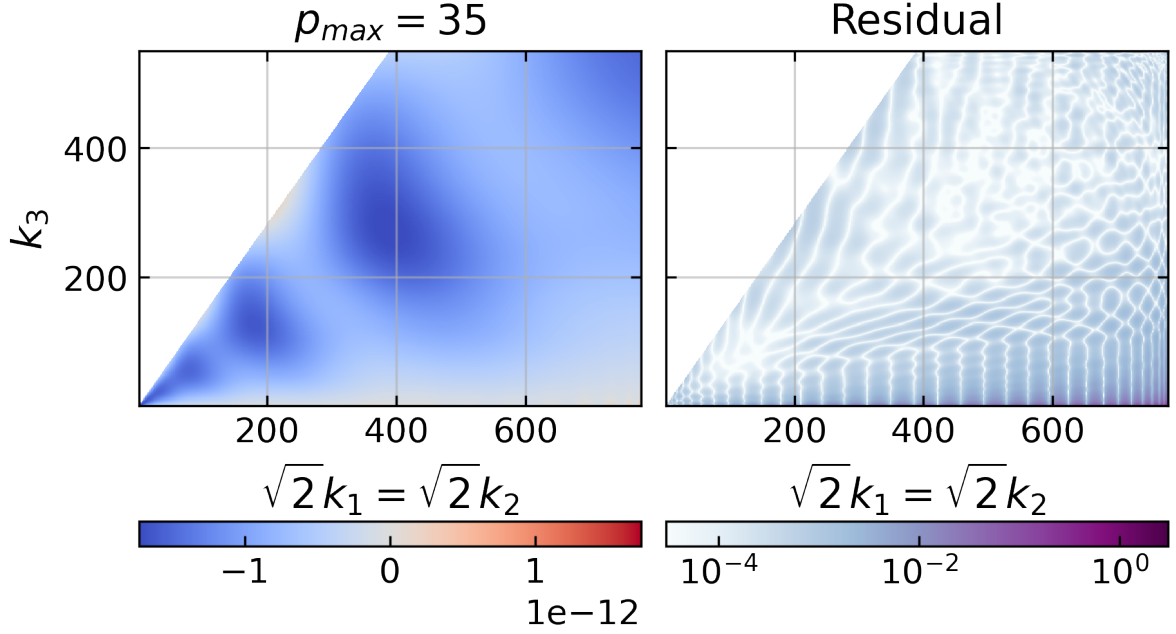


Figure 3.9: Non-Gaussianity generated by periodic features in a DBI model, including a phase difference in the flattened limit as described in [37]. For the purposes of demonstrating the phenomenology, we have placed an envelope on the oscillations in the potential to aid convergence. In the $\mathcal{P}_{01}^{n_s}$ basis, with $n_s^* - 1 = -0.0325$, the result has a relative difference of 1.9×10^{-3} between $p_{\max} = 65$ and $p_{\max} = 35$.

3.11 Map distinguishability of templates at primordial level

3.11.1 Determine where to usefully scan.

Words

3.12 Conclusion

3.12.1 My methods (and their implementation) have been validated on interesting examples.

Words

3.12.2 Have successfully developed separable methods for high orders and features for the first time.

Words

3.12.3 Can obtain full shape info far faster than previous numerical methods.

Words

Chapter 4

Validation

4.1 Abstract

4.2 High p_{\max}

4.2.1 How high can I push p_{\max} ?

Move section here.

Chapter 5

Constraints TO DO

5.1 With either CMB-BEst, or James' coeffs, or both.

Words

5.2 Compare convergence at primordial level to convergence at f_{NL} level

5.2.1 Talk about Wuhyun's work.

Words

5.3 Validate this convergence by reproducing Planck constraint using DBI template decomp.

Words

5.4 Then, check how the scaling from the real numerical result affects this.

Words

5.5 EFT stuff, build off Enrico's recent work.

Words

Chapter 6

Future Work/Conclusions

6.1 Further SF constraints

6.1.1 (Depending on what gets done.)

Words

6.2 Multi-field

6.2.1 Specific models, parameter goals.

Words

6.2.2 Notes on implementation.

Words

6.3 Factor basis to assess limits

6.3.1 (Unless this is already done)

Words

6.3.2 Map shape space?

Words

6.4 User guide for Primodal code, public release of Primodal

6.4.1 Not relevant to thesis, but could add some short notes on plans.

Words

6.5 Discussion

In this work we have extended the modal methods of [17, 18, 19] to recast the calculation of the tree-level primordial bispectrum (??) into a form that explicitly preserves its separability. We emphasise again that this work has two main advantages over previous numerical methods. The more immediate is that by calculating the primordial bispectrum in terms of an expansion in some basis, the full bispectrum can be obtained much more efficiently than through repetitive integration separately for each k -configuration. The second, more important, advantage is the link to observations. Unlike previous numerical and semi-analytic methods, once the shape function is expressed in some basis as in (??), the integral (??) and other computationally intensive steps involved in estimating a particular bispectrum in the CMB, can be precomputed. Since this large cost is only paid once per basis, once a basis which converges well for a broad range of models has been found, an extremely broad exploration of primordial bispectra becomes immediately feasible in the CMB. Making explicit the k -dependence in this way also opens the door to vast increases in efficiency in connecting to other observables, by precomputation using the basis set, then performing a (relatively) cheap scan over inflation parameters.

Our work here goes beyond that of [20] in that our careful methodology allows us to accurately and efficiently go to much higher orders, in particular our methods of starting the time integrals (2.8) and of including the spatial derivative terms in the calculation. This allowed us to present this method for feature bispectra for the first time, demonstrating the efficient exploration of much more general primordial bispectrum phenomenology. We have also identified and addressed the effects of the non-physical k -configurations on convergence within the three-dimensional tetrapyd. We explored, for the first time, possible basis set choices in the context of those effects. We showed rapid convergence on a broad range of scenarios, including cases with oscillatory features with non-trivial shape dependence, using our augmented Legendre polynomial basis, $\mathcal{P}_{01}^{n_s}$.

The immediate application of this work is the efficient exploration of bispectrum phenomenology, as our methods can much more quickly converge to the full shape information than previous numerical methods, which relied on calculating the shape function

point-by-point, for each k -configuration separately. We have implemented these methods for single field scenarios with a varying sound speed, scenarios which have a rich feature phenomenology. An important goal will be extending these methods to the case of multiple-field inflation.

The next immediate application will be to directly constrain parameters of inflationary scenarios through modal bispectrum results from the *Planck* satellite [10]. The details of the work required to directly connect our coefficients to the observed data, and the large but once-per-basis cost of this calculation, will be detailed in a forthcoming paper [21]. CMB and LSS data from forthcoming surveys will be able to use these separable primordial bispectra to even more precisely constrain the parameters of inflationary scenarios.

PhD project (1/2 page)

The standard cosmological history, the Λ CDM model, explains the structure we see in the distribution of matter (large-scale structure, LSS) and radiation (the cosmic microwave background, *CMB*) in our universe. However, this model still requires initial conditions. These initial conditions – adiabatic perturbations on a uniform background, close to Gaussian, with a power spectrum that is nearly (but not exactly) scale-invariant – are provided by an epoch of “inflation”, an epoch before the Big Bang of the Λ CDM model.

During inflation, the universe is supposed to have expanded rapidly, stretching quantum fluctuations from small scales to large. This seeded the formation of structure and encoded correlations that we would eventually measure today. Much information on fundamental physics has been gleaned from the two-point correlators of observations. For higher order correlators (which probe the non-linearities of interactions in the early universe) calculating the predictions of a model of inflation and comparing them to what we see in the sky is much more computationally intensive. Previous work, including that done by the Planck collaboration [10], has managed to constrain some models of inflation through their three-point correlations (or rather through their proxy, the bispectrum). The computational complexity was dealt with by making layers of approximations and the space of models which could be constrained was limited. One notable simplifying assumption is that of “separability”, which makes the process of *CMB* estimation much more tractable [7].

The formalism used to calculate inflationary non-Gaussianities within an inflation scenario is known as the in-in formalism [39]. In my PhD, I have exploited the inherent separability of the tree level in-in formalism, using expansions in separable basis functions. Following the modal philosophy of [19], and building on the basic idea of [20], this method obviates the requirement of considering an approximate separable template, which was a limitation of previous analyses. Instead, the goal is to work with the full numerically calculated tree-level inflationary bispectrum of the theory being considered—the output

is not a grid of points however, but a set of coefficients of an explicitly separable basis expansion. This preserves the separability and therefore allows us to skip the usual template-approximation step.

I developed this separable approach into a practical and efficient numerical methodology which can be applied to a wider and more complicated range of bispectrum phenomenology, making an important step forward towards observational pipelines which can directly confront specific models of inflation using the full bispectrum information in a template-free analysis.

The latter part of this bispectrum estimation pipeline (*CMB-BEst* [?]) was recently completed by my collaborators (Wuhyun Sohn, Dr James Fergusson). This code constrains the inflationary output of my “*Primodal*” code using *CMB* data.

In my recent paper (with E. P. S. Shellard) [40], which has been submitted to JCAP, we explored and described various basis choices and their advantages, and thoroughly validated the methods on single-field inflation models with non-trivial phenomenology, showing that our calculation of these coefficients is fast and accurate to high orders.

More recently I have reached the key exploitation phase of my research. My collaborators and I have tested the integration of the *Primodal* and *CMB-BEst* parts of the pipeline, and at present, my focus is on running *Primodal* on the Cambridge HPC facility, scanning models of inflation with the goal of obtaining some improved constraints on parameters of fundamental physics using our pipeline. Specifically, we are looking at models of DBI inflation [3] with sharp or oscillatory features, which are reviewed in, for example, [41].

Research proposal (1-2 pages)

General Aims

My research goals are to develop methods which will allow us to connect the fundamental physics of the early universe to observations. My most immediate goal is to connect single-field models of inflation with the full three-point correlations of the *CMB*. Beyond that, my goal is to generalise these methods to theories with multiple fields (which are well-motivated by “string swampland” considerations [42]), and later to connect the same theories to other observables coming from large-scale structure.

My PhD has focused on the first of these goals, calculating the bispectrum of the inflationary curvature perturbations, in a way tailored to be linked to observations. I have developed methods to do this in the case of single field inflation. A prime goal I would pursue in this RA position would be to extend these methods (and their implementation in code) to the case of multiple fields. This would open the path to placing constraints on this more general class of theories.

Readying *Primodal* for a public release is another immediate goal. This would benefit the community by providing it with a powerful tool that can capture the full shape information of a wide class of models far more efficiently than previous similar codes, as described in [40].

These goals support those laid out in the recent community white paper [43], by improving predictions of the phenomenology of non-linearities in the very early universe. The ability to efficiently compare the phenomenology of these scenarios to observation and thus constrain the space of theories is a very worthwhile goal, though the phenomenology is also of interest in its own right.

I will do this work in collaboration with Wuhyun Sohn, Dr James Fergusson and Prof. E.P.S.Shellard.

Specific Objectives

Non-linearities from inflaton self-interactions

If the fundamental physical model of inflation is of a single field, then the non-linearities of the self-interactions of that field may have produced observably large non-Gaussian signatures [44]. Detecting these signatures in the CMB bispectrum requires an understanding of their form [7], which motivates the study of these dynamics. Models of inflation that can produce observably large non-Gaussian signatures, and are therefore constrained by the current experimental bounds [10], include string theory-inspired models such as DBI inflation [45] and axion-monodromy [46, 47] (which gives large non-Gaussianity through the resonance mechanism). Also within reach with the current implementation of *Primodal* is the exploration of the effects of sudden features in the inflationary dynamics, and of more general time-dependence within the formalism of the effective field theory (EFT) of inflation.

Non-linearities from multi-field inflation

Another source of possibly observable non-Gaussian signatures can be found in inflation scenarios with more than one active field. In these models, which are motivated by string theory considerations [42], non-linearities of interactions between fields during inflation can generate observably large non-Gaussian signals. This is in part because they can evade the squeezed limit consistency condition [24]. The extra degrees of freedom in these scenarios can generate bispectrum shapes not usually seen in single-field models (for example, see recent work in [48]), making the exploration of the case of multiple fields a prime target.

Optimising the separable decomposition of primordial bispectra

The convergence of our separable expansion is a vital issue in determining the feasibility of our methods. This convergence depends centrally on the basis choice, as we explored in [40]. While in that work we described basis sets that could efficiently capture both the basic shapes and some more complicated oscillatory shapes, the question remains as to how to find the optimal separable description of a given bispectrum.

This is complementary to the philosophy of the main pipeline, which is to find a large general basis which covers a wide range of models. Instead, it may be possible to reduce the separable description of any given bispectrum to a sufficiently compact form that template-free estimation tailored to a scenario could be computationally tractable.

Additionally, the most relevant metric for convergence is the convergence of the observables, not the bispectrum at the end of inflation—exploring the relative efficiencies of different basis sets in this context could be very fruitful in expanding the range of models which could be constrained.

Public release of *Primodal*

The public release of *Primodal* is a clear and worthwhile deliverable of this project. While other publicly released software for the calculation of inflationary non-Gaussianities does exist, they are based on point-by-point methods, and are thus inefficient in their calculation and description of the smooth bispectrum. *Primodal*, which calculates the same quantity in an intrinsically smooth way, and whose result is tailored to comparison with observations, would thus fill an important gap.

Methodology

Non-linearities from inflaton self-interactions

Using the basis sets and the separable formulation of the in-in calculation developed in my PhD, We will reproduce and improve upon the established constraints on single-field inflation models. An initial target will be exploring how the scale-dependence of the numerically calculated DBI bispectrum affects the constraint on the sound speed, which was found in [10] using a scale-invariant approximate template. We will also explore oscillatory models, obtaining constraints using the full shape dependence, complementing the searches done in [10] which again used approximate templates (which exclude, for example, drifting frequencies).

Beyond this, we will expand the focus beyond DBI inflation, resonance features and sharp features and fully exploit the current implementation of my methods and code by comprehensively reviewing the single-field inflation models popular in the literature. We will explore the effects of including the full shape information in a template-free

analysis, improving bispectrum constraints on parameters of single-field inflation models. Categorising models into distinguishability classes would be a useful initial goal.

Non-linearities from multi-field inflation

While some aspects of multi-field scenarios have been previously explored numerically [?] using the transport method [49] we can go beyond these explorations in two ways. Firstly, the transport method is a point-by-point method of calculating the bispectrum, and is thus significantly less efficient than our modal expansion method, and less suited to parameter scans. Secondly, the resulting grid of points is not easily comparable to observations, which is the main motivator of the separability of our method.

I will extend the methods I have already developed to the case of multiple active fields during inflation, and implement them in the *Primodal* code. I will validate my established basis sets on standard templates for multi-field bispectra, and test their ability to capture the expected scale-dependence of the shapes, a property usually excluded in standard templates. Once these validations are passed, I will recast the multi-field in-in calculation into a separable form (as I have already done for the single-field case) and implement that calculation in code. Using that implementation, I will explore the phenomenology of multi-field inflation, understanding which theories and scenarios can be constrained through the bispectrum, and obtaining those constraints.

Optimising the separable decomposition of primordial bispectra

To pursue efficient separable decomposition, I intend to research techniques which have not previously been exploited in the setting of bispectrum calculation and estimation. One possible avenue is tensor rank decomposition techniques and other methods of tensor approximation, which are of use in, for example, signal processing.

By using these techniques to approximate the coefficient matrix (which is the output of *Primodal*) we can determine a linear combination of our basis functions which can more efficiently describe that particular bispectrum. I will apply these techniques to explore the general limits of separable approximations to primordial bispectra, with a focus on convergence at the level of observations. This will make clear the widest possible set of models that can be constrained through the bispectrum.

Public release of *Primodal*

My intention is to follow best practices when preparing *Primodal* for public release. I will document the code, prepare example test cases for validation by users on their local machines, and release the code within a container to package the code along with its dependencies, libraries, and system settings to ensure reproducibility.

Work plan (12 months)

- Oct-Dec
 - Obtain single-field constraints using the already-validated codes *Primodal* and *CMB-BEst*.
 - Exploit tensor rank decomposition for basis choice optimisation, and explore tools from other fields, e.g. signal processing.
 - Explore further applications to single-field models, for example time dependence within the EFT of inflation.
 - Prepare *Primodal* for public release in its single-field form.
- Jan-Mar
 - Test the basis sets that were used in the single-field case on multi-field templates. If needed, explore new basis sets to cover multi-field shapes.
 - Generalise the methods for separable in-in calculation to the multi-field case.
 - Prepare single-field constraint results for publication.
- Apr-Jun
 - Implement multi-field methods in code, extending *Primodal*.
 - Investigate multi-field phenomenology, with the goal of understanding which scenarios predict observable non-Gaussianity, and thus can be constrained.
 - Explore the application of these methods in other contexts, such as preparing non-Gaussian initial conditions for simulations of large scale structure evolution.
- Jul-Sep
 - Prepare the multi-field extension of *Primodal* for release.
 - Prepare multi-field results for publication.

Work plan (6 months)

If granted a six month award, I would adapt the work plan by prioritising the extension of our separable methods to multi-field dynamics, and delaying the public release of *Primodal*.

- Oct-Dec
 - Obtain single-field constraints using the already-validated codes *Primodal* and *CMB-BEst*.

- Test the basis sets that were used in the single-field case on multi-field templates. If needed, explore new basis sets to cover multi-field shapes.
 - Generalise the methods for separable in-in calculation to the multi-field case.
- Jan-Mar
 - Implement multi-field methods in code, extending *Primodal*.
 - Investigate multi-field phenomenology, with the goal of understanding which scenarios predict observable non-Gaussianity, and thus can be constrained.
 - Prepare single-field constraint results for publication.
 - Exploit tensor rank decomposition for basis choice optimisation, and explore tools from other fields, e.g. signal processing.

Career goals, benefits

Given the methods and expertise I have developed so far in my PhD, and the expertise in CMB bispectrum estimation in Cambridge, I am now ideally positioned to achieve my goal of obtaining new constraints, benefiting the department by producing leading results in an area with genuine discovery potential.

The award would benefit me by giving me the opportunity to pursue this project further into the exploitation phase of the single-field implementation, and the opportunity to expand the methods I have developed to the broader class of multi-field models of fundamental early-universe physics. Meanwhile, I would have the opportunity to further my computational and numerical skills, and expand my skills by learning new numerical techniques (such as tensor rank decomposition). Having the opportunity to pursue these goals and learn these skills will greatly benefit me in my preparation for further applications.

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Appendix A

Appendices

A.1 Efficient numerics

Efficient evaluation and numerical usage of the coefficients (vander, etc). Numeric choices that aren't specific to mode evolution, etc. Gauss-Legendre integration, fixed weights.

A.2 Tools used

Numpy, Scipy, Jupyter Notebooks, QUADPTS...?

A.3 Transport method for the Modal coeffs

Infinite hierarchy of coupled equations. Choose a basis to make it nice? Real vs Imag...? Sounds cool, not worth pursuing.

