

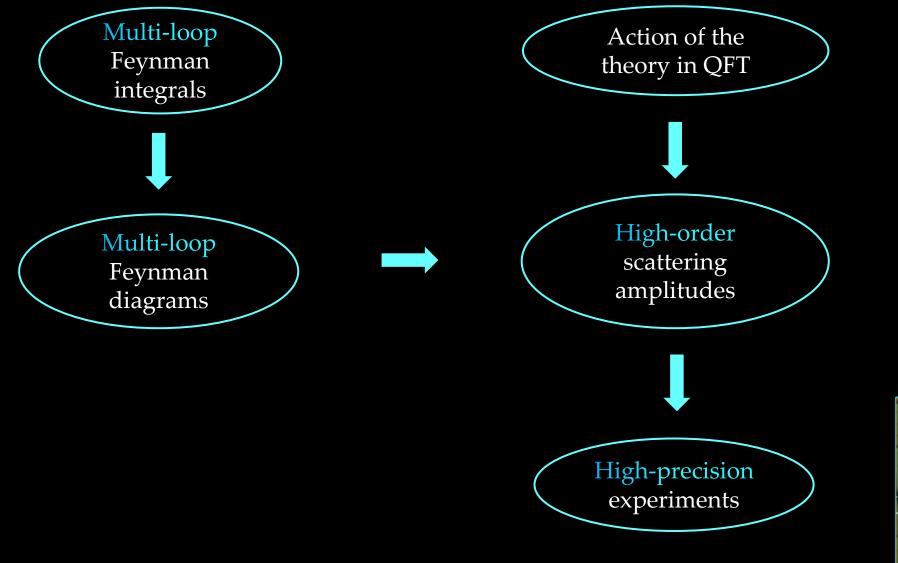
## **Recent developments of NeatIBP**

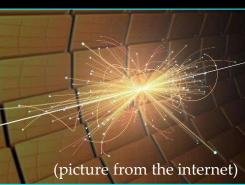
A package generating small sized integration-by-parts relations for Feynman integral reduction

#### Zihao Wu

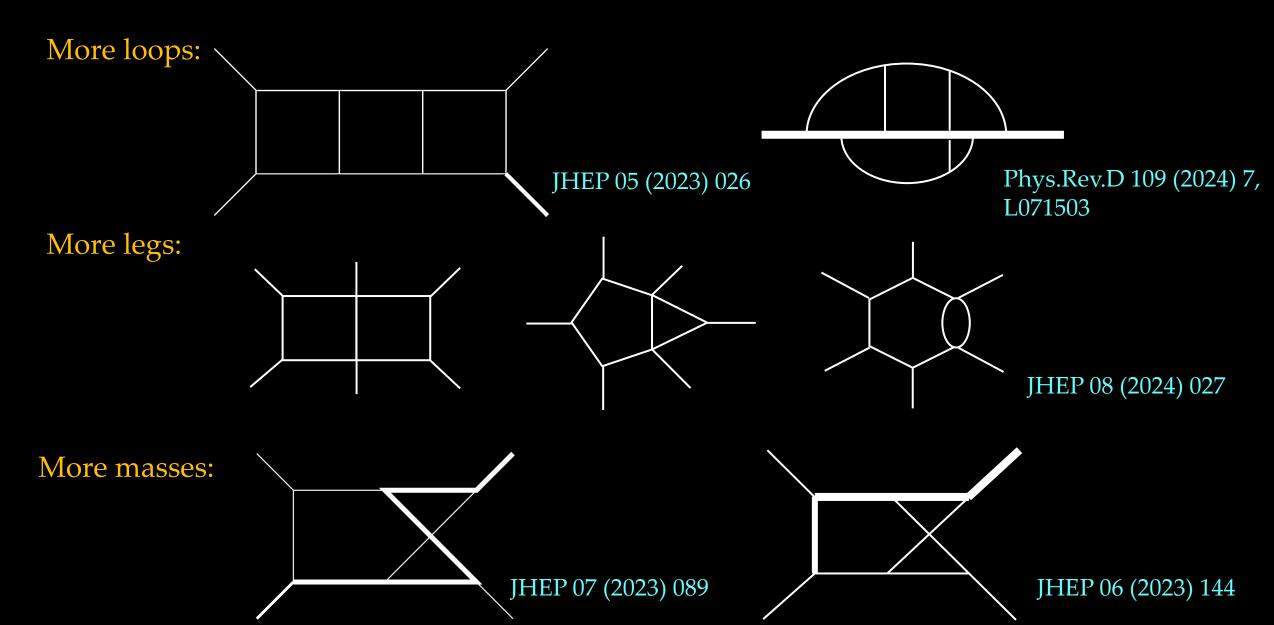
Based on: Rourou Ma, Johann Usovitsch, ZW, Yingxuan Xu, Yang Zhang, 24xx.xxxx, related previous work: ZW, Janko Boehm, Rourou Ma, Hefeng Xu, Yang Zhang, Comput.Phys.Commun. 295 (2024) 108999 and important discussions with: David Kosower, Fabian Lange, Roman Lee, Yan-Qing Ma, Simone Zoia.

### Multi-loop Feynman integrals in particle physics

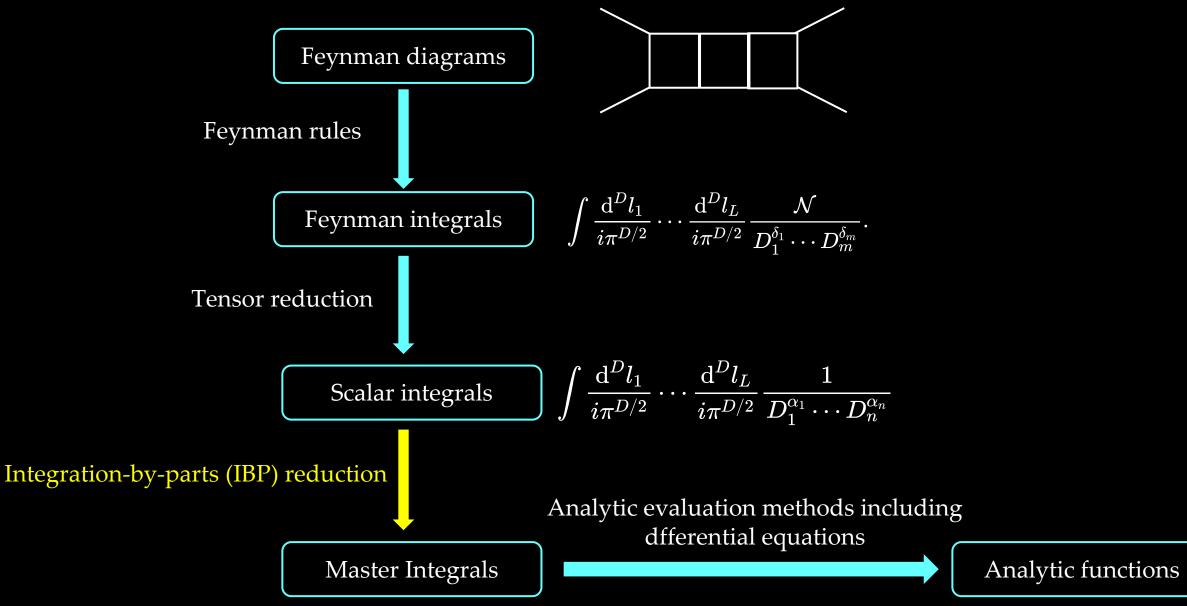




# Multi-loop Feynman integrals, frontiers



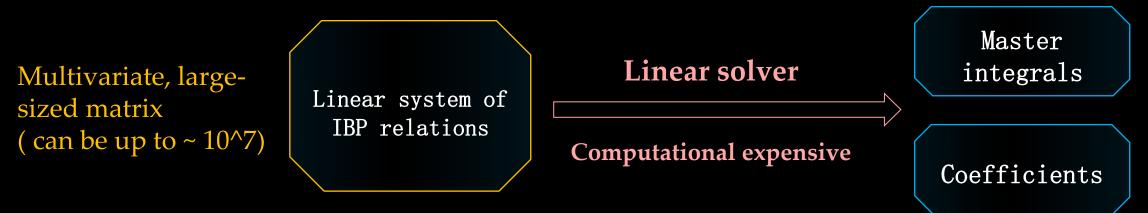
### Feynman integral reduction



A linear independent basis of the scalar integrals

### Computational bottle necks in integration-by-parts (IBP) reductions

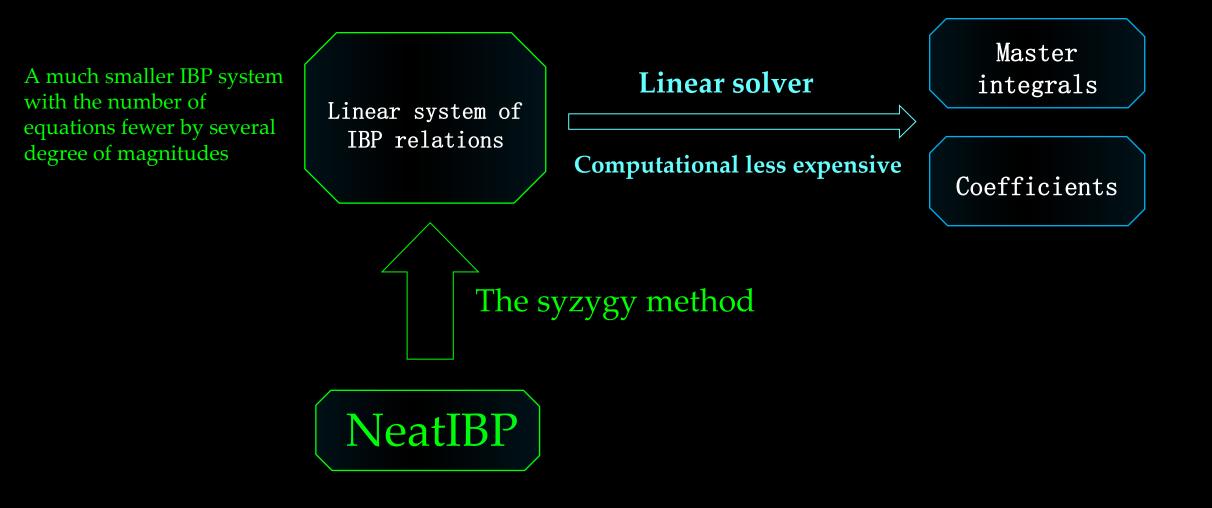
A traditional IBP reductio method such as Laporta's algorithm Int. J. Mod. Phys. A, 2000, 15: 5087-5159. Computational bottle necks in frontier problems



Costing days/weeks on clusters for a cutting-edge problem

Costing hundreds of GB / several TB RAM or more

### Generating small-size IBP system using NeatIBP



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Progress of NeatIBP from its v1.0.2.4 to the latest version, v1.0.5.1, at 6<sup>th</sup> Sep. 2024.

1. The spanning cuts method

2. The Kira+NeatIBP interface

3. Simplification of syzygy vectors using maximal cut

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## IBP relations in with multiple propagators

 Target integrals

  $G[1, 1, 1, \dots, 1, -1, -4]$ 
 $G[1, 1, 1, \dots, 1, -5, 0]$ 

$$\begin{array}{c} \text{IBP-Related integrals} \\ G[1,1,0,\cdots,1,-1,-2] & G[0,1,1,\cdots,2,0,-1] \\ G[2,1,0,\cdots,1,-1,-1] & G[0,1,1,\cdots,0,0,-2] \\ G[1,1,0,\cdots,1,-3,-1] & G[1,2,0,\cdots,1,-3,0] \\ \dots \end{array}$$

 $egin{array}{c} Master integrals \ G[1,1,1,\cdots,1,-1,0] \ G[1,1,1,\cdots,1,0,0] \ \ldots \end{array}$ 

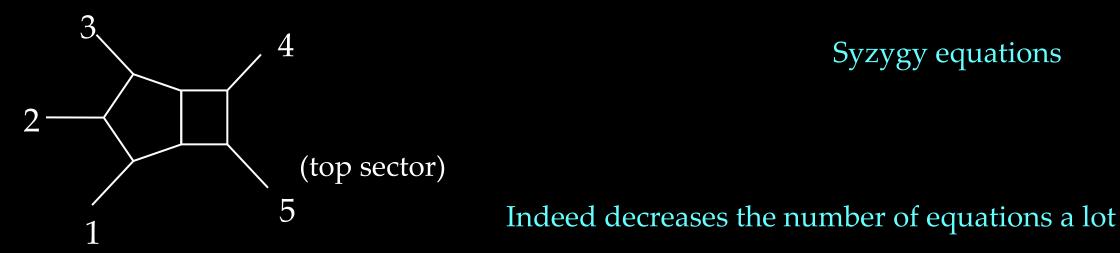
Contains **redundant integrals** with denominator indices lifted

$$0 = \int \frac{\mathrm{d}^{D} l_{1}}{i\pi^{D/2}} \cdots \frac{\mathrm{d}^{D} l_{L}}{i\pi^{D/2}} \frac{\partial}{\partial l_{k}^{\mu}} \frac{v^{\mu}}{D_{1}^{\alpha_{1}} \cdots D_{n}^{\alpha_{n}}} = \int \frac{\mathrm{d}^{D} l_{1}}{i\pi^{D/2}} \cdots \frac{\mathrm{d}^{D} l_{L}}{i\pi^{D/2}} \frac{\frac{\partial v^{\mu}}{\partial l_{k}^{\mu}} - v^{\mu} \sum_{i=1}^{n} \frac{\partial D_{i}}{\partial l_{k}^{\mu}} \frac{\alpha_{k}}{D_{i}}}{D_{1}^{\alpha_{1}} \cdots D_{n}^{\alpha_{n}}}$$
Introducing multiple propagators

### IBP relations from syzygy method

$$0 = \int \frac{\mathrm{d}^{D} l_{1}}{i\pi^{D/2}} \cdots \frac{\mathrm{d}^{D} l_{L}}{i\pi^{D/2}} \frac{\partial}{\partial l_{k}^{\mu}} \frac{v^{\mu}}{D_{1}^{\alpha_{1}} \cdots D_{n}^{\alpha_{n}}} = \int \frac{\mathrm{d}^{D} l_{1}}{i\pi^{D/2}} \cdots \frac{\mathrm{d}^{D} l_{L}}{i\pi^{D/2}} \frac{\frac{\partial v^{\mu}}{\partial l_{k}^{\mu}} - v^{\mu} \sum_{i=1}^{n} \frac{\partial D_{i}}{\partial l_{k}^{\mu}} \frac{\alpha_{k}}{D_{i}}}{D_{1}^{\alpha_{1}} \cdots D_{n}^{\alpha_{n}}}$$
Introducing multiple propagators

Janusz Gluza, Krzysztof Kajda, David A. Kosower: Phys.Rev.D 83 (2011) 045012 What if we find a good combination of *v* such that the additional *Di* cancels?



### NeatIBP: syzygy method in Baikov representation

Feynman integrals in momentum space:

$$I[lpha_1, \cdots, lpha_n] = \int rac{\mathrm{d}^D l_1}{i\pi^{D/2}} \cdots rac{\mathrm{d}^D l_L}{i\pi^{D/2}} rac{1}{D_1^{lpha_1} \cdots D_n^{lpha_n}}$$
 $\int \int \mathrm{Variable\ transformation}$ 

Baikov representation: integrates directly over propagators *zi* 

$$I[lpha_1,\cdots,lpha_n]=C\int\mathrm{d} z_1\cdots\mathrm{d} z_nP(z)^lpharac{1}{z_1^{lpha_1}\cdots z_n^{lpha_n}}$$

## IBP relations in Baikov representation

#### Polynomial equations in Baikov IBP relations

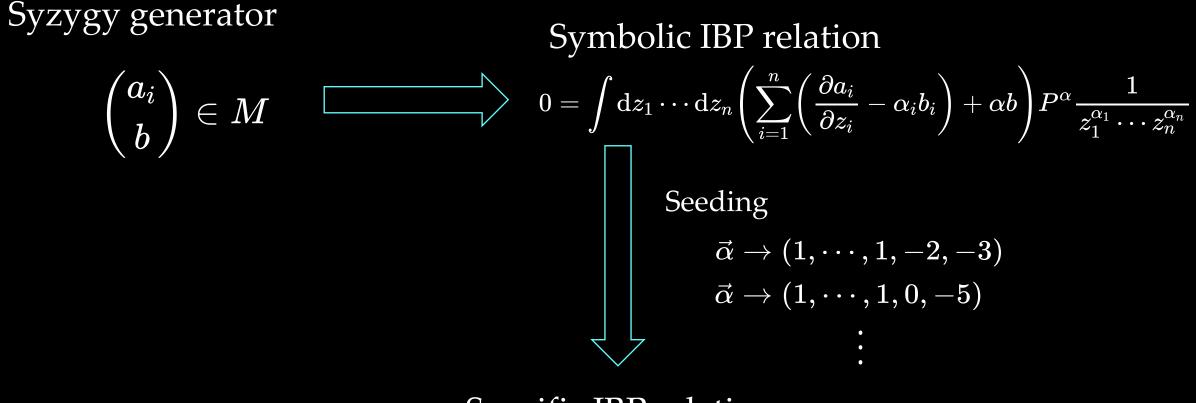
$$\left(\sum_{i=1}^n a_i(z) rac{\partial P}{\partial z_i}
ight) + b(z)P = 0$$

$$a_i(z)=b_i(z)z_i \ \ ext{for} \ i\in \{j|lpha_j>0\}$$



### Generators of the solution module

$$egin{pmatrix} a_i \ b \end{pmatrix} \in M \ = < f_1, f_2, \cdots >$$



Specific IBP relation

(# generators) × (# seeds) = (# IBP relations)

**IBP** relation selection

An enough IBP system  $0 = \sum_{j} c_{ij} I_j$ Column reduction (numeric + finite field)

Linearly independent system

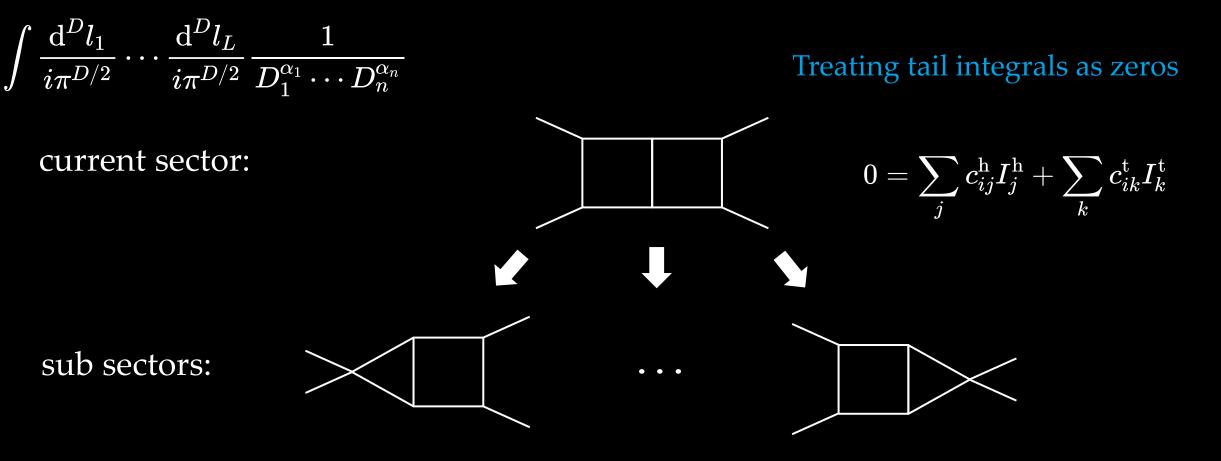
$$0=\sum_j { ilde c}_{ij} I_j$$

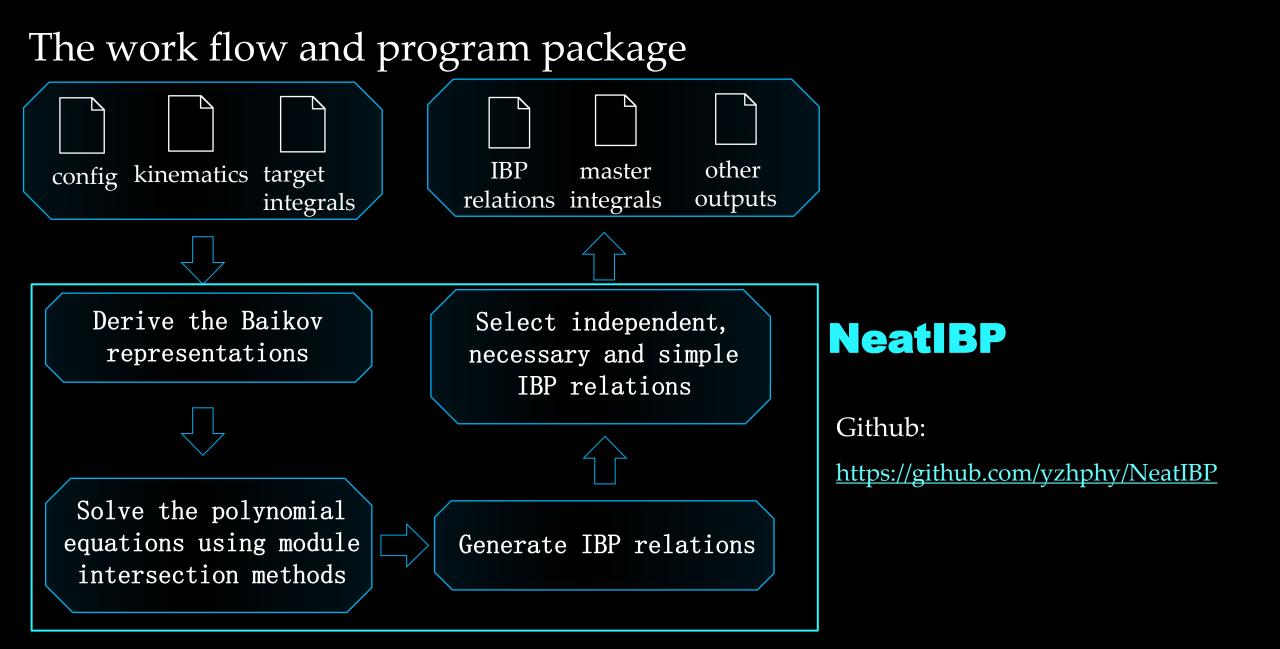
Row reduction (numeric + finite field)  $R_{ik} = L_{ij} \tilde{c}_{jk}$ 

Remove the unneeded relations for reducing the targets

Small-size IBP system minimally needed

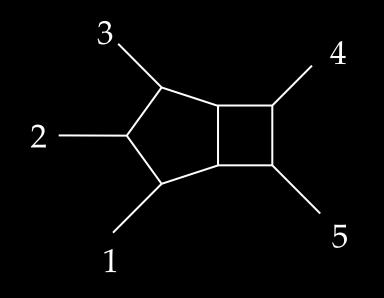
## Tail mask strategy





#### **Performance of NeatIBP**

2L5P Example



Target integrals with high-degree numerators Quantity: 2483

# of IBP (FIRE6): **11207942** 

Max numerator degree: 5 Max denominator power: 1 # of MI: 61 # of IBP: **14120** 

Time used: 27m at



CPU threads: 20 RAM: 128GB

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Generalized unitarity cuts in Baikov representation

$$I_{lpha_1,\cdots,lpha_n}|_{\mathcal{C}- ext{cut}} \propto \oint_0 \prod_{i\in\mathcal{C}} \mathrm{d} z_i \int \prod_{i
otin\mathcal{C}} \mathrm{d} z_i P^lpha rac{1}{z_1^{lpha_1}\cdots z_n^{lpha_n}}$$

Cuts change the integrals but preserve IBP relations

$$\sum_i c_i I_i = 0 \quad \longrightarrow \quad \sum_i c_i (I_i|_{\mathcal{C}- ext{cut}}) = 0$$

Cuts increases the number of *zero sectors* in a family

$$\mathcal{C} 
ot \subseteq S \Rightarrow I|_{\mathcal{C}- ext{cut}} = 0, orall I \in ext{Sector } S$$

## Spanning cuts method

A spanning cuts  $\{C_i\}$  is defined so that for all nonzero sector S in a family,  $\exists C_i$ , such that  $C_i \subseteq S$   $\{1,3\} - \operatorname{cut} \{2,4\} - \operatorname{cut}$ 

Step 1: Run NeatIBP on each cut

Current version of NeatIBP avoids cutting multiple propagators. Thus, for integrals such that  $lpha_i=1, i\in \mathcal{C}$ ,  $\mathcal{C} ext{-cut}$  means

 $P o P|_{z_i o 0, i \in \mathcal{C}}$ 

Step 2: Reduce the IBP system on each cut

Step 3: Merge the reduced IBP system of all cuts

#### Benefit:

1.Σ difficulty of evaluation on
spanning cuts < evaluation on the</li>
family as a whole
2. Parallelizable

### Merging the spanning cut systems

$$\begin{split} I|_{\{1,3\}\text{-cut}} &= \sum_{i} c_{\{1,3\},i}^{1234} I_{i}^{1234}|_{\{1,3\}\text{-cut}} + \sum_{i} c_{\{1,3\},i}^{13} I_{i}^{13}|_{\{1,3\}\text{-cut}} \\ I|_{\{2,4\}\text{-cut}} &= \sum_{i} c_{\{2,4\},i}^{1234} I_{i}^{1234}|_{\{2,4\}\text{-cut}} + \sum_{i} c_{\{2,4\},i}^{24} I_{i}^{24}|_{\{2,4\}\text{-cut}} \\ & \swarrow \\ & \swarrow \\ & \text{merge} \end{split}$$

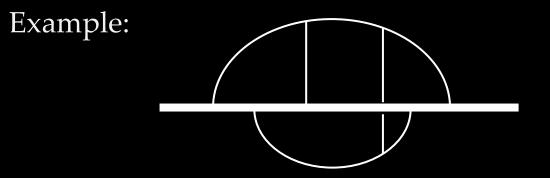
$$I = \sum_i c_i^{1234} I_i^{1234} + \sum_i c_i^{13} I_i^{13} + \sum_i c_i^{24} I_i^{24} ~~ (c_i^{13})^{13}$$

Currently, NeatIBP uses a "direct" merging strategy:

$$\left\{ \begin{array}{l} c_i^{13} = c_{\{1,3\},i}^{13} \\ c_i^{24} = c_{\{2,4\},i}^{24} \\ c_i^{1234} = c_{\{1,3\},i}^{1234} = c_{\{2,4\},i}^{1234} \end{array} \right.$$

NeatIBP checks consistency conditions before merging the spanning cut systems. This step cannot be omitted because the condition is **NOT** always guaranteed according to our observation.

### Automated evaluation on spanning cuts in NeatIBP



Phys.Rev.D 109 (2024) 7, L071503

Plainly running NeatIBP:

No, I cannot finish this computation. I stuck at syzygy vectors evaluation for days....  $\mathfrak{S}$ 

Running NeatIBP with spanning cuts CutIndices = "spanning cuts";

*Yes, after splitting the problem into 29 pieces of different cuts, I finished the IBP relation generation on spanning cuts in 10+ hours.* <sup>(C)</sup>

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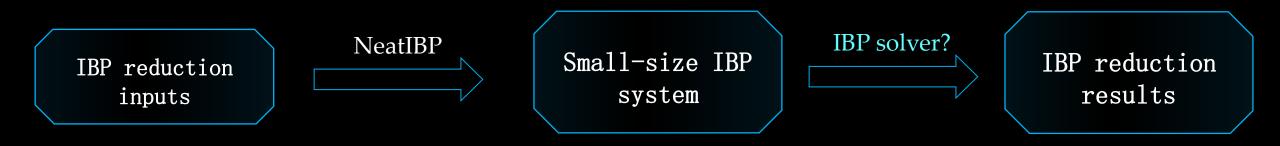
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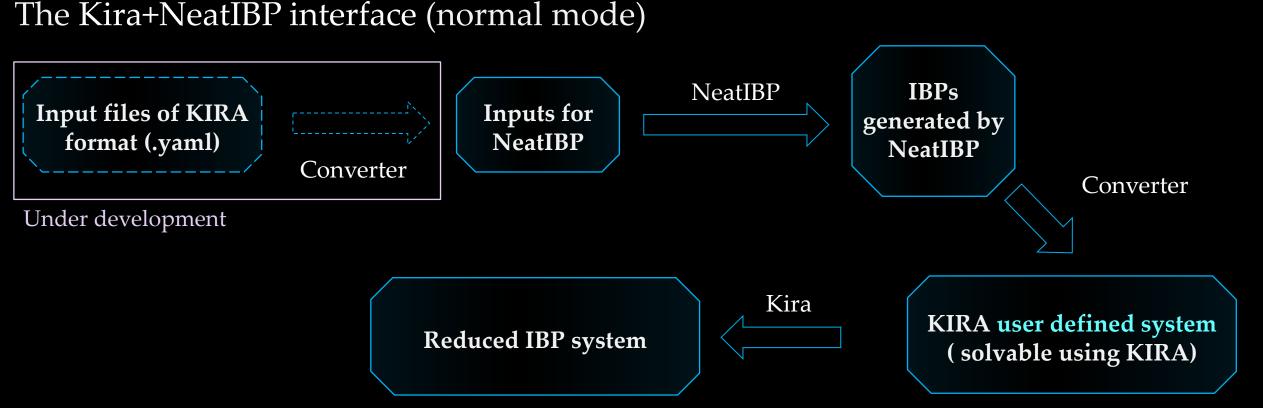
#### Currently, NeatIBP itself dose not perform IBP reduction



KIRA Comput.Phys.Commun. 230 (2018) 99-112, Comput.Phys.Commun. 266 (2021) 108024, etc. as a popular integral reduction software, is a very nice IBP solver for NeatIBP output.

Kira allows user to feed in linear systems of IBP relations (called user defined system) and then reduce them .

The Kira+NeatIBP interface is included in the latest version of NeatIBP (v1.0.5.1, at 6<sup>th</sup> Sep. 2024)

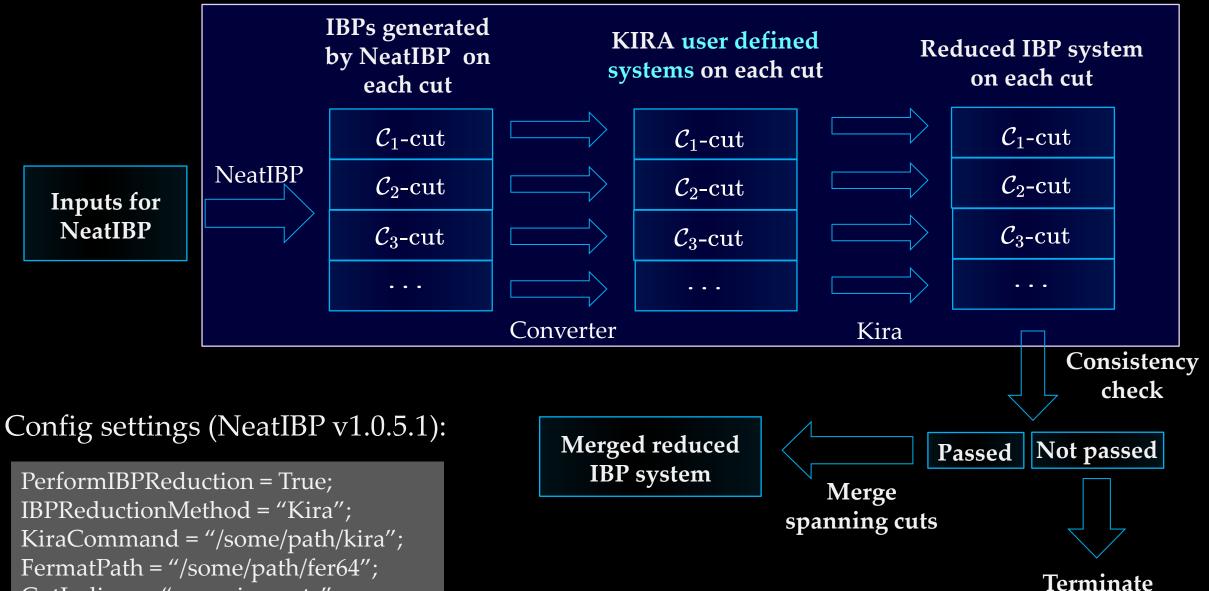


In NeatIBP (since v1.0.5.0), to use Kira to reduce the IBP generated by NeatIBP, add the following commands into NeatIBP "config.txt"

PerformIBPReduction=True; IBPReductionMethod="Kira"; KiraCommand="/some/path/kira" FermatPath="/some/path/fer64"

#### The Kira+NeatIBP interface (spanning cuts mode)

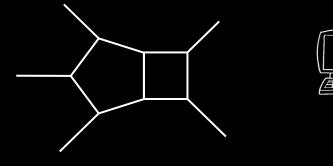
Parallelized by default

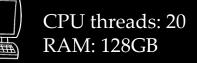


CutIndices = "spanning cuts";

A baby example of Kira+NeatIBP performance (with spanning cuts)

Two-loop five-point, with numerator degree 3





Running NeatIBP + Kira in spanning cuts mode:

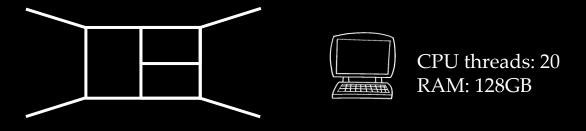
- 1. NeatIBP: Generates equations on 10 spanning cuts, using 5m
- 2. Number of equations: vary from 130 ~ 624, on different cuts
- 3. Kira running time: 10m (including converting, checking consistency, reducing and merging, in parallel mode)

Running NeatIBP + Kira in normal mode:

- 1. NeatIBP: Generates 1358 equations spanning cuts, using 16m
- 2. Kira reduction time used: 27m

Another example of Kira+NeatIBP performance (with spanning cuts)

Three-loop four-point, with numerator degree 5



Running NeatIBP + Kira in spanning cuts mode:

- 1. NeatIBP: Generates equations on 22 spanning cuts, using 6h21m
- 2. Number of equations:  $2k \sim 10k+$  or so, at most 35k, on different cuts
- 3. Kira running time: 7h40m (including converting, checking consistency, reducing and merging, in parallel mode)

Running NeatIBP + Kira in normal mode:

1. NeatIBP: Generates 114k equations spanning cuts, using 18h44m

2. Kira reduction: Testing in progress. Timing unknown yet. My 128GB RAM limit on the current machine is exceeded.

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Generator vector of the solution module of the syzygy equations

Symbolic IBP relation

$$egin{aligned} egin{aligned} egin{aligned} a_i \ b \end{pmatrix} \in M \end{aligned} egin{aligned} & 0 = \int \mathrm{d} z_1 \cdots \mathrm{d} z_n igg( \sum_{i=1}^n igg( rac{\partial a_i}{\partial z_i} - lpha_i b_i igg) + lpha b igg) P^lpha rac{1}{z_1^{lpha_1} \cdots z_n^{lpha_n}} & & \ & \ & & \ & & \ & & \ & & \ & & \ & & \ & & \ & & \ & \ & \ & \ & & \ &$$

Specific IBP relation

(# generators)  $\times$  (# seeds) = (# IBP relations)

Generator vector of the solution module of the syzygy equations

Formal IBP relation

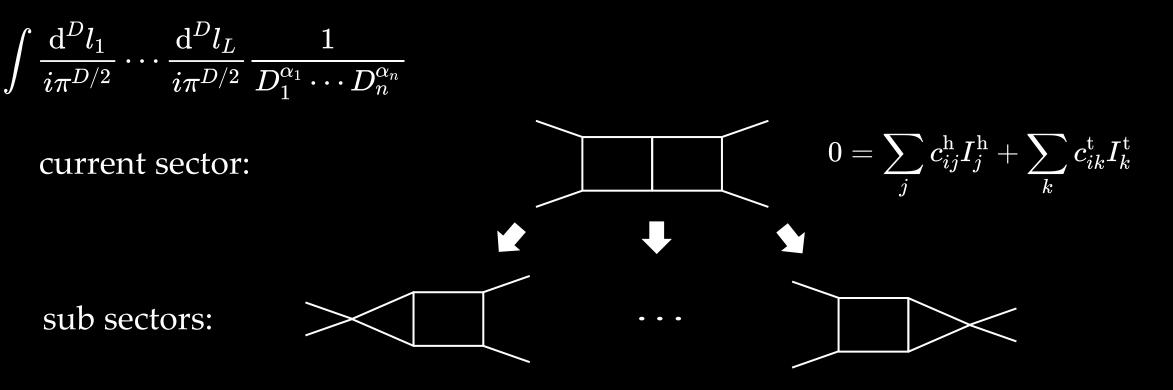
$$egin{aligned} egin{aligned} egin{aligned} a_i \ b \end{pmatrix} \in M \end{aligned} egin{aligned} 0 &= \int \mathrm{d} z_1 \cdots \mathrm{d} z_n igg( \sum_{i=1}^n igg( rac{\partial a_i}{\partial z_i} - lpha_i b_i igg) + lpha b igg) P^lpha rac{1}{z_1^{lpha_1} \cdots z_n^{lpha_n}} \ &igg[ Seeding & ec{lpha} o (1, \cdots, 1, -2, -3) & ec{lpha} o (1, \cdots, 1, 0, -5) & ec{lpha} o (1, \cdots, 1, 0, -5) & ec{ec{lpha}} \end{array} \end{aligned}$$

Specific IBP relation

(# generators) × (# seeds) = (# IBP relations)

~ hundreds large binomial numbers Seeding cost  $\uparrow$ 

## Tail mask strategy



Treating tail integrals as zeros:

- 1. IBP relations containing only sub sector integrals will be discarded
- 2. This is equivalent to maximal cut (mc):

$$ext{ for } I \in ext{ sector } S\,,\,\, \left. I 
ight|_{ ext{mc}} := I|_{S ext{-cut}}$$

## Maximal cut

Consider IBP relations with "no-multiple-propagator" integrals in a sector *S* such that  $\begin{cases} \alpha_i = 1, & i \in S, \\ \alpha_i < 0, & i \notin S, \end{cases}$ 

Define maximal cut on a syzygy vector

$$f:=egin{pmatrix} a_i\ b\end{pmatrix}\in M \qquad \qquad fert_{\mathrm{mc}}:=fert_{z_i o 0,i\in S}$$

From the IBP formula

$$0=\int \mathrm{d} z_1\cdots \mathrm{d} z_n igg(\sum_{i=1}^n igg(rac{\partial a_i}{\partial z_i}-lpha_ib_iigg)+lpha bigg) P^lpha rac{1}{z_1^{lpha_1}\cdots z_n^{lpha_n}}$$

We have

 $\|f\|_{
m mc}=0 \quad \Leftrightarrow \quad$  Corresponding IBP relation contains only integrals in sub sectors

### Simplification of syzygy generators via maximal cut

Delete generators that dose not change the "maximal cut" module  $M|_{mc}$  defined as

$$M|_{\mathrm{mc}}:=\langle f_1|_{\mathrm{mc}},f_2|_{\mathrm{mc}},\cdots
angle \hspace{1.5cm} ext{for}\hspace{1.5cm}M=\langle f_1,f_2,\cdots
angle$$

Step 1: Using "maximal cut" Groebner basis

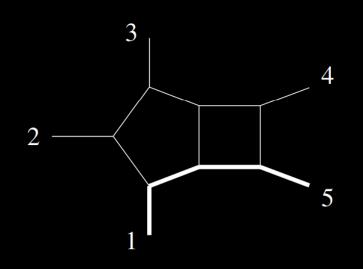
$$\left| G 
ight|_{
m mc} := {
m GB}(M|_{
m mc}) = \langle g_1, g_2, \cdots 
angle \qquad ext{ lifting } g_i = \sum_j c_{ij} f_i |_{
m mc}$$

Delete  $f_j$  in generators of M if  $c_{ij} = 0, \forall i$ Because this means  $f_j|_{
m mc}$  does not contribute to  $M|_{
m mc}$ 

Step 2: Scanning

After step 1, sort the remaining generators from complex to simple  $M' = \langle f'_1, \dots, f'_n \rangle$ Scanning *i* from 1 to *n*, if deleting  $f'_i$  does not change the GB of maximal cut module, then delete it. Example

Top sector of a pentagon-box diagram



Before simplification: (# generators) = 374

After simplification: (# generators) = 14

~ 25 times shortened

### More general cases

$$0 = \int \mathrm{d} z_1 \cdots \mathrm{d} z_n igg( \sum_{i=1}^n igg( rac{\partial a_i}{\partial z_i} - lpha_i b_i igg) + lpha b igg) P^lpha rac{1}{z_1^{lpha_1} \cdots z_n^{lpha_n}}$$

Consider IBP relations with generic integrals in a sector *S* such that  $\begin{cases} lpha_i > 0, & i \in S \\ lpha_i \leq 0, & i \notin S \end{cases}$ 

modification

$$f:=egin{pmatrix} a_i\ b\end{pmatrix}\in M \quad \longrightarrow \quad f:=egin{pmatrix} ilde{a}_i\ b\end{pmatrix} ext{ where } egin{pmatrix} ilde{a}_i:=egin{pmatrix}rac{a_i}{z_i}, & i\in S\ a_i, & i
otin S \end{pmatrix}$$

## Summary

The integration-by-parts (IBP) reduction is one of the bottle-neck steps in the evaluation of Feynman integrals.

**NeatIBP** is a automated program generating small-size IBP system. It helps to reduce the computation cost of IBP reduction.

We have introduced some new features of NeatIBP till its latest version. They include:

- 1. The spanning cuts method : Splitting the hard problem into simpler pieces.
- 2. The Kira interface: Providing automated IBP solver.
- 3. Simplification of syzygy generators using maximal cut method: Making the seeding process computational cheaper.