# Using feynr

#### Peter Schmidt-Nielsen

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#### Installation

You can download feynr at http://github.com/petersn/feynr. All you need are the files feynr.sty and feynr\_render.py.

#### Best Practices Install

Mark feynr\_render.py as executable, and place it on your path. Add feynr.sty to your TrXpath, and rerun texhash. You are now set up.

#### Lazy (/Non-Root) Installl

Copy both feynr.sty and feynr\_render.py to the directory for your project. Instead of running feynr\_render.py source.tex to rebuild your diagrams you'll have to run python feynr\_render.py source.tex. C'est la vie.

### Overview

Include feynr with the usual \usepackage{feynr}. Place your diagram source code in a feynr environment to render a diagram. A minimalish example:

```
\documentclass{article}
\usepackage{feynr}
\begin{document}
\begin{feynr}
feynr diagram specification
\end{feynr}
\end{document}
```

Run feynr\_render.py on your LATEX file after each time you update a feynr diagram:

```
feynr_render.py source.tex
pdflatex source.tex
```

## Basic Usage

Anywhere in your preamble you may specify comma separated global options to feynr in the \feynroptions command. For example, \feynroptions{time-right} will override the default of time-up, and switch away from Feynman's original convention. A complete list of options can be found at the end of this document.

In feynr, particles are explicitly tracked. You can create an input particle to your diagram with the input command. For example, input a defines an input particle called a. Special flags can be passed to any feynr command with a pair of dashes. For example, to declare an input particle a that's an electron we may use: input a -- electron. More than one particle may be declared in a single command, and the flags passed apply to all particles. For example, to declare three positrons: input p1 p2 p3 -- positron. Particle types may be omitted if they can be unambiguously inferred from elsewhere in the diagram.

Particles can be made to interact with each other with the interact command.<sup>1</sup> For example, interact p1 p2 will cause the particles p1 and p2 to exchange a particle. Again, the interaction type may be unambiguous. For example, if p1 and p2 are photons, and p1 is already known to emit a positron (from another interact command), then the interaction is unambiguous, and an electron will be emitted from p1, and propagate to interact with p2. However, frequently you will have to specify what form the interaction takes. The flags electron, positron, and photon denote that the given particle propagated from the first argument to the second argument. For example, interact p1 p2 -- electron causes an electron to be emitted by p1 and absorbed by p2.<sup>2</sup> Note that as interact takes exactly two node arguments, the -- separating flags from node arguments is optional. You may write either interact a b -- flags or interact a b flags, these are equivalent. Just like input, there is a command output that can be used to disambiguate diagrams. For example, if, at the end of a sequence of interactions, it's ambiguous what type of particle a is, we can specify with, for example: output a -- photon. You can identify interactions in your diagram that couldn't have their type inferred – they will appear as red sine waves.

Finally, we need to give feynr some hints about the layout of our diagram. If you attempt to compile a diagram without any layout hints it will fail, as it can't figure out what angles the various lines should be drawn at. There are two default layout hints, Ospace and Otime. These hints indicate that the given interaction should be drawn perpendicular to the specified axis. The naming convention is that a Otime interaction has zero length in time, and a Ospace interaction has zero length in space. In other words, in a time-up Feynman diagram, Otime indicates that the interaction should be drawn horizontally. For example, we can write interact e1 e2 photon Otime to cause particles e1 and e2 to exchange a photon that is drawn perpendicular to the time axis.

The propagate command accumulates flags to be added on the next propagation

<sup>&</sup>lt;sup>1</sup>The command interact can also be abbreviated as a single dash.

<sup>&</sup>lt;sup>2</sup>Naturally, interact a b -- electron is equivalent to interact b a -- positron.

that a given particle makes. For example, if we would like the particle a to propagate straight along the time axis the next time it propagates, we can issue the command propagate a -- Ospace. Like all flags in feynr, they apply only to a single interaction/propagation of a particle, and are not permanent. Thus, order matters. In the sequence interact a b, propagate a -- flags, interact a c, the extra flags given in the propagate command affect how particle a gets from the interaction with b to the interaction with c.

Putting these all pieces together, let's look at some examples:

The first line defines an electron, and the second is a layout hint that it should propagate along the time axis.

Here two electrons are defined in the first line, and made to exchange a photon along the space axis in the second.

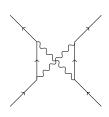
Not substantially more complicated than the previous example.

Here Delbrück scattering is concisely described in four lines. The third line (propagate p1 p2 -- Ospace) makes sure that the electrons propagate vertically in the diagram, forming a nice square box when used with the horizontal Otime interactions.

## Advanced Usage

Let's look how we might lay out a more complicated diagram.

```
input p1 p2 — electron skeleton p1 p2 photon 0time propagate p1 p2 — 0space skeleton p1 p2 photon 0time draw p1\sim2 p2\sim3 photon draw p2\sim2 p1\sim3 photon
```



Okay, there's a lot to break down here. We're using two new commands, skeleton and draw. The command skeleton is exactly equivalent to interact, only it automatically passes the flag hidden, a special flag that causes the interaction to not be rendered. That is to say, interact a b hidden is exactly equivalent to skeleton a b. Crucially, the interaction is still used for type-inference, and used for structural layout – thus skeleton. The purpose of the skeleton command is to allow you to quickly build the desired shape and types of your diagram without worrying too much about how you're going to fill in the interactions. Further, it allows for a tremendous amount of uniformity and control of the generated diagrams. You can produce a skeleton interaction, then later fill in multiple different kinds of interactions between the skeleton interacting particles without changing the layout of your diagram by a single pixel. What about draw? Just like skeleton, draw is a version of interact that passes an extra flag, in this case free. The free flag says that the given interaction is purely for display purposes, and does not affect type-inference or structural layout.

In the above diagram skeleton is necessary, and here's why: Unfortunately, we have a dependency loop: p1 is interacting with a future version of p2 and vice versa. There is nothing to be done about this, so we simply produce two skeleton interactions that form the same box as before with our Delbrück scattering. Finally, we use draw to add photons between p1 and p2, but we must specify the out-of-order interaction we want. Thus, we introduce nodes. Each particle produces a trail of nodes along its path. For example, p1 has four nodes: the four leftmost vertices in the above diagram. These vertices are automatically internally labeled  $p1\sim1$ ,  $p1\sim2$ ,  $p1\sim3$ , and  $p1\sim4$ . These correspond to p1's input node, two nodes connected to photons, and output node respectively. By issuing the command draw  $p1\sim2$   $p2\sim3$  photon we render a photon from the second node on p1's path to the third node on p2's path. This corresponds to the photon propagating up and to the right in the above diagram. In this way, every vertex in our diagram receives a node label, and we can add extra features to our diagram without too much effort.

When drawing Feynman diagrams we frequently want to add an emitted photon without affecting the layout of our diagram. Frequently we want this photon to be reabsorbed elsewhere in our diagram. Just like before where we used skeleton and draw to separate the layout and components of our diagram, here we can use draw to add additional details without distoring our diagram. Let's examine the following sort of diagram:

```
input e1 e2 — electron propagate e1 — no-arrow interact e1 e2 photon 0time propagate e1 — no-arrow draw e1\sim1:e1\sim2 e1\sim2:e1\sim3 photon arc
```



We've introduced two new flags, no-arrow and arc, and a new syntax which can be used in place of a node argument to draw. We've got almost the same diagram as in our very first example, except for the final draw command, which adds an arcing photon.

When specifying nodes in a draw command you may optionally use the syntax a:b to anchor one end of the drawn connection to half way between node a and b. For example, in this case the first end of our arcing photon is anchored half way between e1~1 and e1~2, namely the midpoint of the lower leftmost edge. The flag arc simply specifies that the drawn connection should be a segment of a circle rather than a straight connection – much nicer looking in this case. Compare to the unarced version to the right. Finally, we need to supress the arrows on the lines for e1, because they would overlap

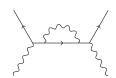


Figure 1: Ugly!

with the photon in an ugly way. We can do this by passing the no-arrow flag into propagate in the right two places, namely before and after the interaction.

Sometimes we may wish to render a diagram like the following, with a self-interaction on an edge. Here we're using the extended syntax proportion:a:b, which sets an

```
input p1 p2 — photon interact p1 p2 electron 0time long draw 0.8:p1\sim2:p2\sim2 0.2:p1\sim2:p2\sim2 photon arc
```



anchor point part way between two nodes. Here we draw a photon between two points 20% and 80% along the middle edge. If we had wanted the arc to be below the edge instead of above it we could add the flag flip. Additionally, I passed the argument long to make the middle edge 50% longer, giving some extra room for the photon semi-circle. However, if I were laying this diagram out as a fourth order upgrade to the second order version without the extra loop, I could choose to not pass long, and have a diagram that beautifully differs from the lower order version only in the extra looped photon.

We can also use this proportion feature to make room for arrows in the previous fourth order Coulomb repulsion diagram. It's up to you to decide which you think

```
input e1 e2 — electron
interact e1 e2 photon 0time
draw 0.25:e1\sim1:e1\sim2 0.75:e1\sim2:e1\sim3 photon arc
```



looks nicer – I really like putting the arrows back in. Using just these arc loops we can produce other useful features, such as circular electron propagation.

```
input e1 e2 — electron
skeleton e1 e2 photon 0time long
draw e1\sim2 0.3: e1\sim2: e2\sim2 photon
draw 0.3:e1\sim2:e2\sim2 0.7:e1\sim2:e2\sim2 electron arc
draw 0.7:e1\sim2:e2\sim2 0.3:e1\sim2:e2\sim2 electron arc
draw 0.7:e1\sim2:e2\sim2 e2\sim2 photon
```

There's a lot of stuff going on here, again using our powerful combination of skeleton and draw. It's a bit of a hack, but let's go through it. The skeleton command gives the type, length, and angle of the interaction, laying out the structure (same as always). The first and last draw commands draw little lengths of photons up along the first and last 30% of the middle edge. Finally, the middle two draw commands draw the two halfs of the circle. Note that the order of the arguments is swapped between them, which is why they aren't just drawn on top of each other. Of course, we could also have done it by using the positron and flip flags on one, and keeping the argument order the same. However, I think this looks cleaner. For convenience, the above four draw commands can be acheived on the edge between a and b with the single macro command photon-loop a b. For example:

```
input e1 e2 — electron
skeleton e1 e2 photon 0time long
propagate e1 e2 — 0space
skeleton e1 e2 photon 0time long
photon-loop e1\sim2 e2\sim2
photon-loop e1\sim3 e2\sim3
```

## Reference

#### Commands:

Command	Arguments
input	name1 name2 optionally: flag1 flag2
output	name1 name2 optionally: flag1 flag2
propagate	name1 name2 optionally: flag1 flag2
interact	node1 node2 optionally: flag1 flag2
-	node1 node2 optionally: flag1 flag2
skeleton	node1 node2 optionally: flag1 flag2
draw	location1 location2 optionally: flag1 flag2
photon-loop	node1 node2

In the above specifications, the field name can be any particle name, like xyz. The field node can be a particle name or particlar node of that particle, like xyz or xyz~2. Finally, the location field can any of the above plus the special syntaxes abc~1:xyz~2 or proportion:abc~1:xyz~2. Valid flags for nodes are:

electron, positron, photon, input, output

Valid flags for interactions are:

electron, positron, photon, hidden, free, Otime, Ospace, arc, flip, no-arrow, long

Valid options to pass to \feynroptions are:

Option	Description
time-up	Same as angle=0, flip=0
time-down	Same as angle=0, flip=1
time-left	Same as angle=270, flip=1
time-right	Same as angle=270, flip=0
flip=<0 or 1>	Sets whether the y-axis is mirrored.
angle= <degrees></degrees>	Rotate the diagram counter-clockwise.
photon-frequency= <num></num>	Change the frequency of photons.
photon-amplitude= <num></num>	Change the amplitude of photons.

The default is:

 $\verb| feynroptions{time-up, photon-frequency=1.0, photon-amplitude=1.0}|$