

Nextflow: difference between the original syntax and DSL2

```
input_ch = Channel.fromPath("*.txt")
```

- All "from" and "into" are removed
- Channel - process relationships are now specified through workflow
- The output channel is referred to as p.out

```
process p1 {  
    input:  
    file(x) from input_ch  
  
    output:  
    file("head.txt") into output_ch1  
    file("tail.txt") into output_ch2  
  
    """  
    head $x > head.txt  
    tail $x > tail.txt  
    """  
}
```

```
process p2 {  
    input:  
    file(y) from output_ch2  
  
    output:  
    val(true) into is_finished  
  
    """  
    gzip $y  
    """  
}
```

```
nextflow.enable.dsl=2
```

```
input_ch = Channel.fromPath("*.txt")
```

```
workflow {  
    input_ch | p1  
    p1.out.output_ch2 | p2  
    p2.out | p3  
}
```

- When there are multiple output channels, and need to specify which one, use "emit" to give the channel a name, and in workflow refer to it as .out.name

```
process p1 {  
    input:  
    file(x)  
  
    output:  
    file("head.txt")  
    file("tail.txt"), emit: output_ch2  
  
    """  
    head $x > head.txt  
    tail $x > tail.txt  
    """  
}
```

```
process p2 {  
    input:  
    file(y)  
  
    output:  
    val(true)  
  
    """  
    gzip $y  
    """  
}
```

Practical steps:

```
input_ch = Channel.fromPath("*.txt")
```

Step 1:

- Write the skeleton of workflow based on channel and process relationships.
- When process has single output channel, can pipe directly into the next process: "input_ch | p1 | p2 | p3"

```
process p1 {
```

```
  input:
  ... file(x) from input_ch
  output:
  ... file("head.txt") into output_ch1
  ... file("tail.txt") into output_ch2
  ...
  head $x > head.txt
  tail $x > tail.txt
  ...
}
```

```
process p2 {
```

```
  input:
  ... file(y) from output_ch2
  output:
  ... val(true) into is_finished
  ...
  gzip $y
  ...
}
```

Step 2:

- Make sure each process only takes 1 input channel.
- Remove all "from" and "into".
- When there are multiple output channels, name them as needed with "emit".

DSL2

```
nextflow.enable.dsl=2
```

```
input_ch = Channel.fromPath("*.txt")
```

```
workflow {
  ... input_ch | p1
  ... p1.out.output_ch2 | p2
  ... p2.out | p3
}
```

Step 3:

- In workflow, combine input channels as needed since each process takes only 1 input channel (see cheatsheet). May need to go back to process definition to make sure the channel inner structures agree.
- Fill in the output names as used in "emit".

```
process p1 {
```

```
  input:
  ... file(x)
  output:
  ... file("head.txt")
  ... file("tail.txt"), emit: output_ch2
  ...
  head $x > head.txt
  tail $x > tail.txt
  ...
}
```

```
process p2 {
```

```
  input:
  ... file(y)
  output:
  ... val(true)
  ...
  gzip $y
  ...
}
```