Regional_workflow

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Protocol 3A: Regional model workflow

In this notebook, we will create a diagram for the workflow that regional modellers should follow under FishMIP Protocol 3A.

Loading libraries

```
library(DiagrammeRsvg)
```

Designing workflow

```
grViz("digraph git_basics {
graph [
      margin = '0,0'
      newrank = true
      splines = false
      splines = compound
      nodesep = 0.3
      ranksep = 0.2
      overlap = true
      rankdir = TB]
node [
     shape = rectangle
      style = filled
      fillcolor = 'white'
      fontname = 'Helvetica, Arial, sans-serif'
      fontsize = 20
edge [
      fontname = 'Helvetica, Arial, sans-serif'
      fontsize = 30
      labelfontcolor = '#00000080'
      penwidth = 2
      shape = 'record']
```

```
Step_0[label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 0:</b> 
         Identify which climate variables<br/>
        to use and how these are <br/>
        implemented<br/>

     >]
Step 1 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 1:</b> 
         Provide shapefile of your model<br/>
        domain and fill out model<br/>
        template<br/>

     >]
Step_n [label = 'Is model spatial?' fillcolor = '#f6b979']
Step_2 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 2:</b> 
         Visualise and extract input<br/>
        variables to see if bias correction <br/>
        is needed<br/>

     >]
Step 3 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 3:</b> 
         Determine if further downscaling<br/>
         is needed <br/>>

     >]
Step_4 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 4:</b> 
         Match and extract fishing effort<br/>
        groupings to force your model<br/>

     >
     tooltip = 'Step 4:']
Step_5 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 5:</b> 
         Calibrate model with<br/>
        observational data for reference <br/>
        period<br/>

     >
     tooltip = 'Step 5:']
Step_6 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 6:</b><br/> 
         Set up model with forcings for<br/>
        each comparative experimental <br/>
        run<br/>

     >]
Step_7 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
          <b>Step 7:</b> 
         Output standard variables to<br/>
```

```
compare with observations and <br/>
           across models over time/space<br/>

       >]
Step_8 [label = <<table border='0' cellborder='1' cellspacing='0' cellpadding='28'>
             <b>Step 8:</b></br/> 
           Quality control checks and upload<br/>
          to FishMIP server<br/>

       >]
subgraph sub_1 {
 Step_0 -> Step_1
 Step_1 -> Step_n [style = dotted]
 Step_n -> Step_2 [label = 'No' fontcolor = '#329932' style = dotted color = '#329932'
                  fontname = 'bold Helvetica, Arial, sans-serif']
 Step_n -> Step_3 [label = 'Yes' fontcolor = '#d8ae2d' style = dotted color = '#d8ae2d'
                  fontname = 'bold Helvetica, Arial, sans-serif']
 {Step_2 Step_3} -> Step_4
 Step_4 -> Step_5 [weight = 10 splines = line];
 }
subgraph sub_2 {
 Step_5 -> Step_6 -> Step_7 -> Step_8
 }
subgraph sub_3 {
 graph [rank = same]
 edge [style = invis,
      weight = 1]
 Step_0 -> Step_5[weight = 1]
}
}",
width = "100%", height = "100%")
```

