Dichotomous

· TIPdates.

· functions.

→ MCMC
- .Cleveland_Plot.BMDdichotomous_MA. → ① Show only - Posterior Probability > 0.05 ... Pending.
  → ② If model yield NA then remove. - Done.
- . Plot.density.BMDdichotomous_MA. → Change Color of model average Part.

→ Laplace.
- · Cleveland_Plot.BMDdichotomous_MA. → (Same as MCMC Case.)
  x We don't need to create Plot density Case here.

→ Base Plot
- . Plot.BMDdichotomous_MA → ① # of Variable should be adjusted dynamically based on valid fitting models.
  ② Change base Plot's object as ggplot2 instead of Plotly one.] Done.
  ② MA_Maximized Case should be Considered. Currently defined function seems wrong.
  ~ Combine it as . Plot.BMDdichotomous_MA
- . Plot.BMDdich_fit_maximize
- . Plot.BMDdich_fit_MCMC

→ Single model fit