

Validate Power: d3.1

December 27, 2021

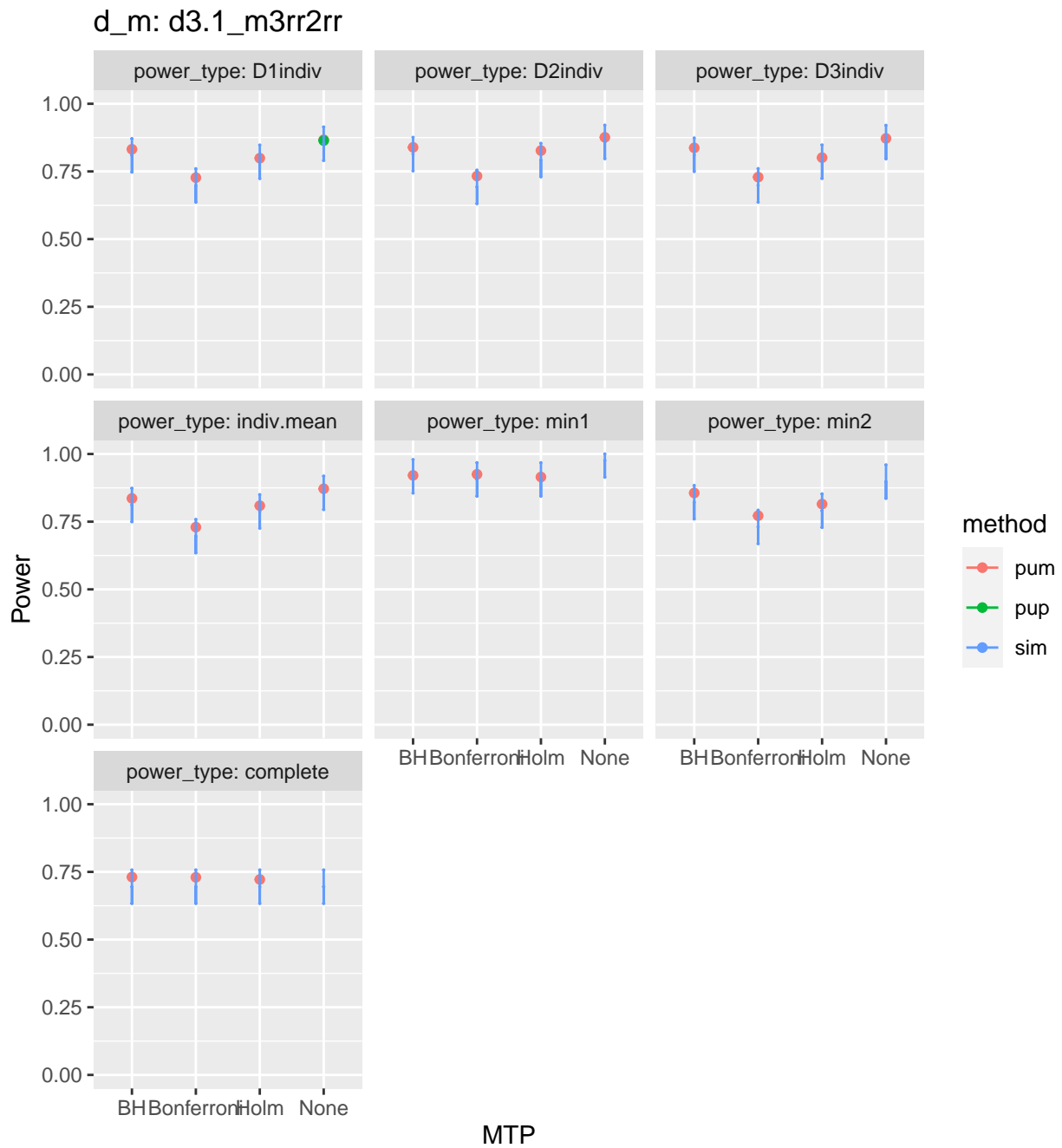
Design: Blocked RCT, with 3 levels, and randomization done at level 1 (individual level).

Models: random treatment effects.

- $M = 3$
- $J = 30$
- $K = 15$
- $\bar{n} = 100$ (unless otherwise noted)
- rho: $\rho = 0.5$
- MDES = 0.125, 0.125, 0.125
- R2: $R_1^2 = 0.1, 0.1, 0.1$
- ICC: $ICC_2 = 0.2, 0.2, 0.2, ICC_3 = 0.2, 0.2, 0.2$
- Omega: $\omega_2 = 0.1, 0.1, 0.1, \omega_3 = 0.1, 0.1, 0.1$

Power Validation

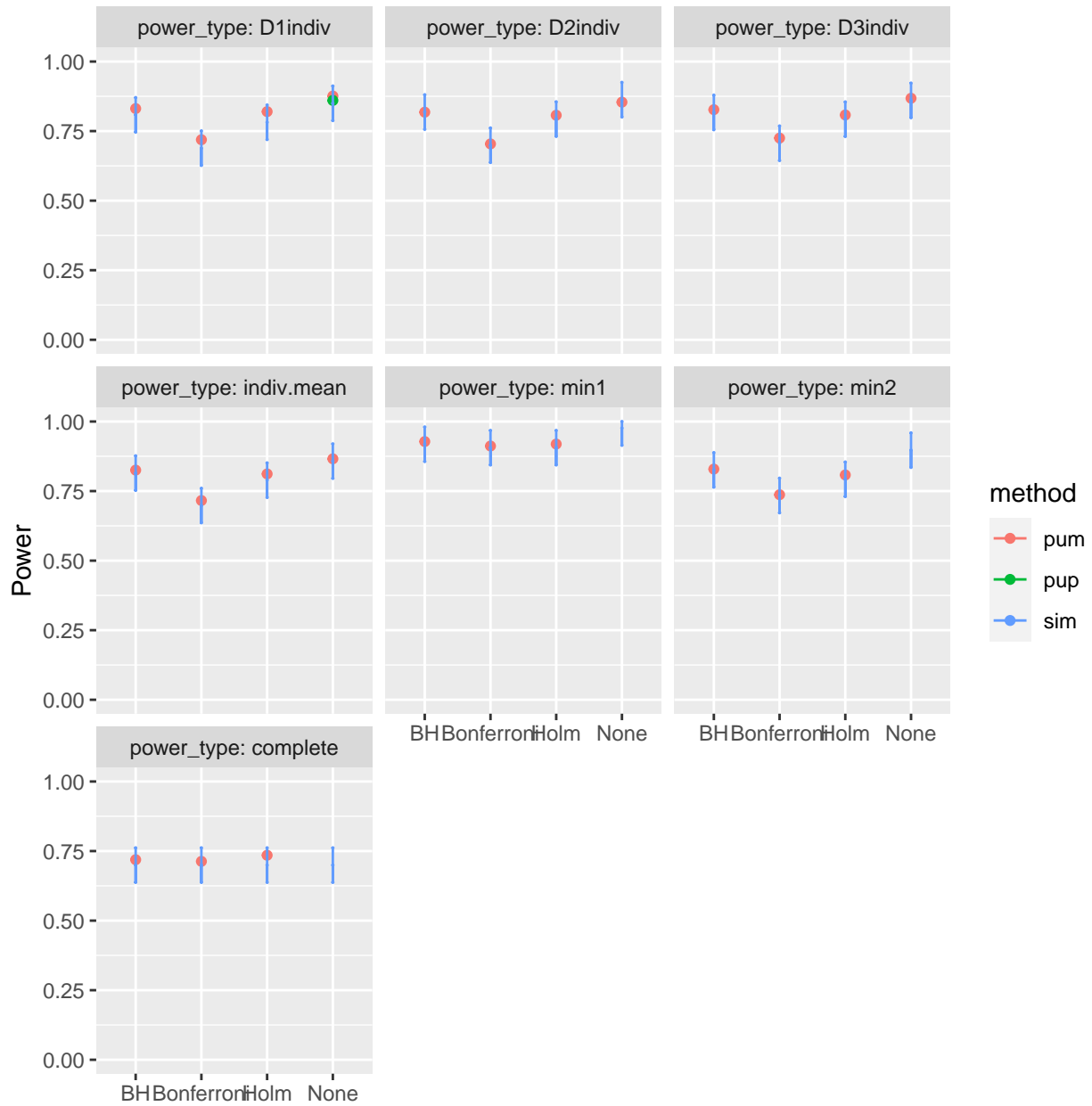
Base case



Varying school size

$\bar{n} = 75$

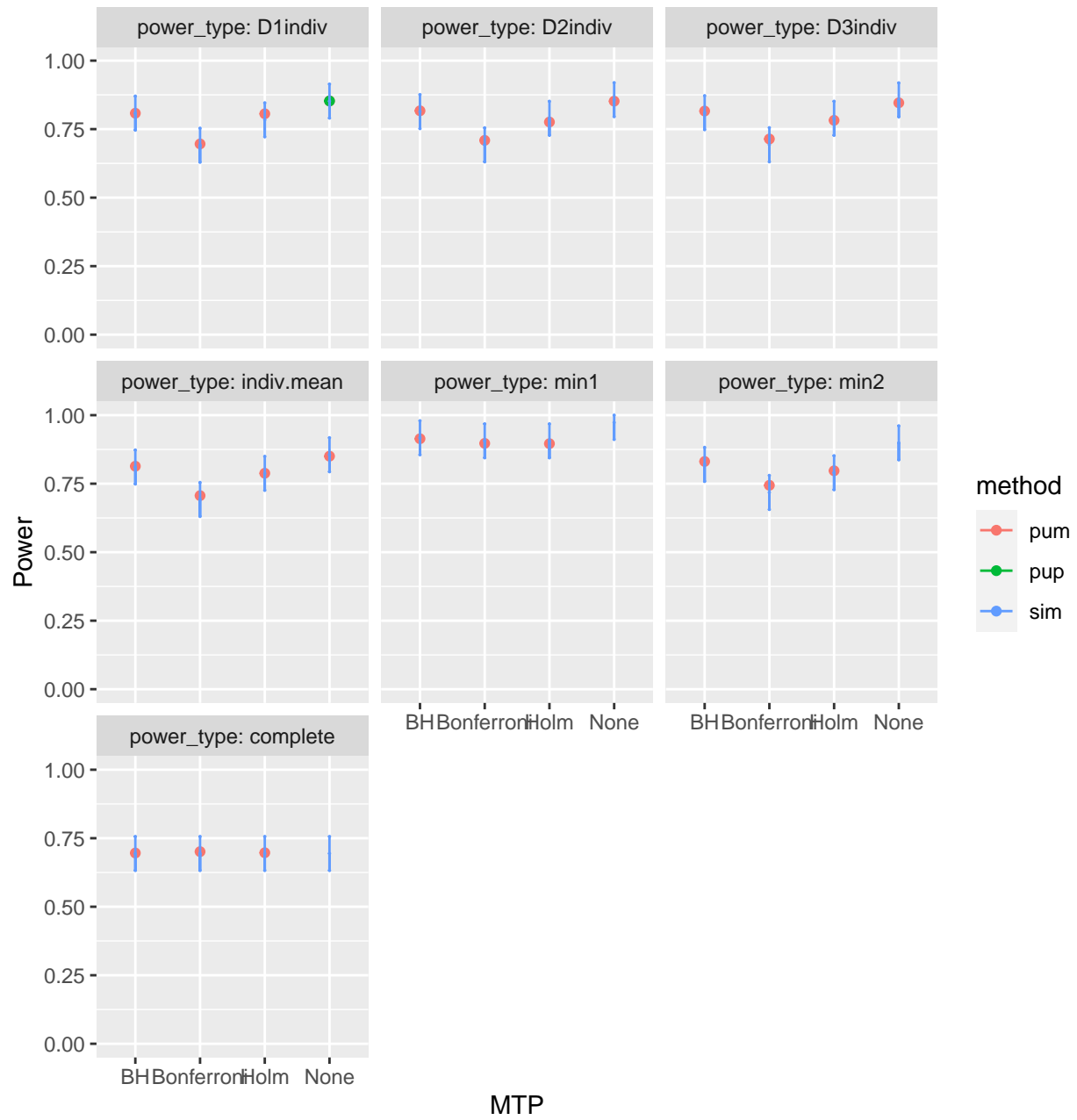
d_m: d3.1_m3rr2rr



MTP

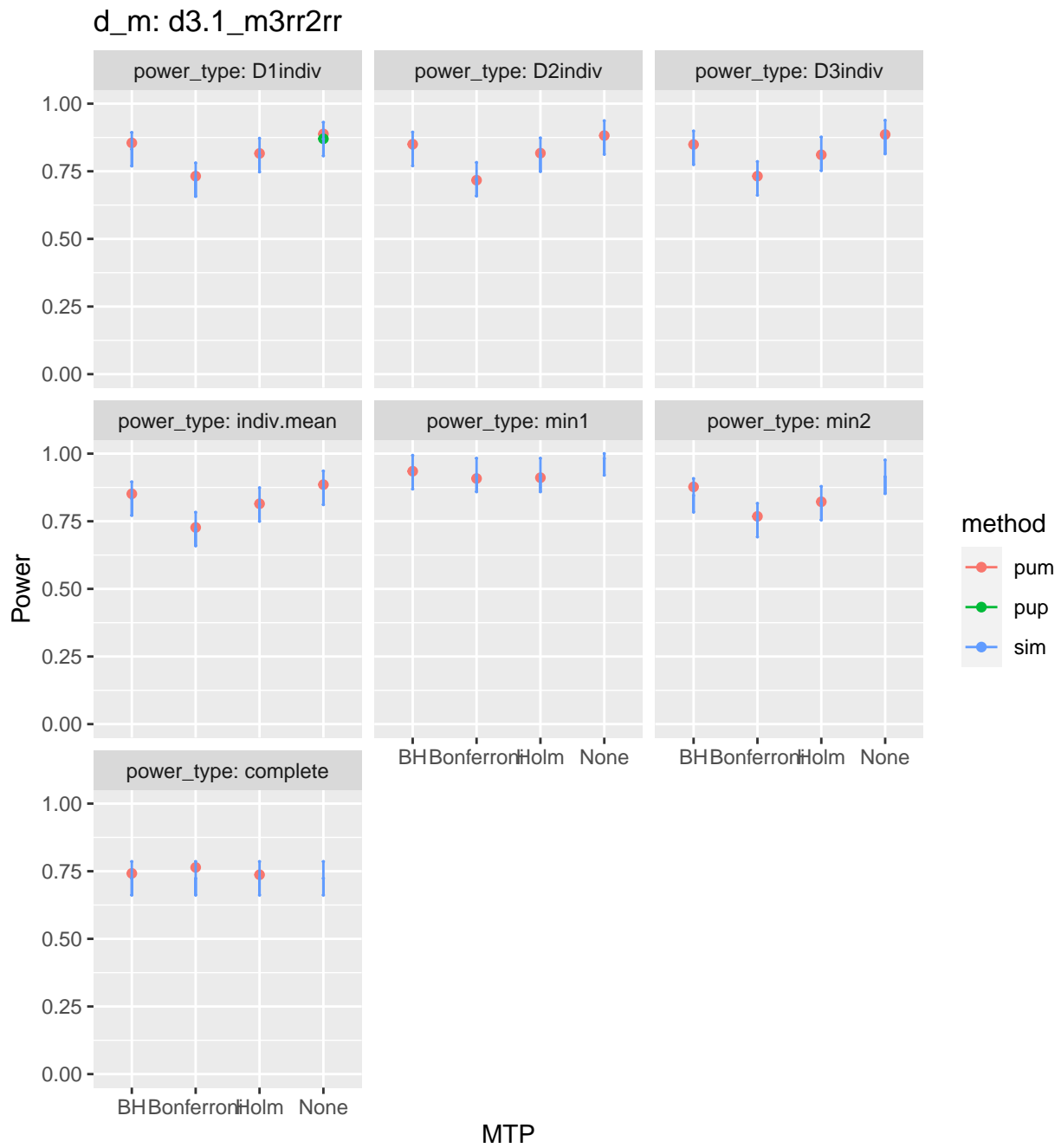
$\bar{n} = 50$

d_m: d3.1_m3rr2rr



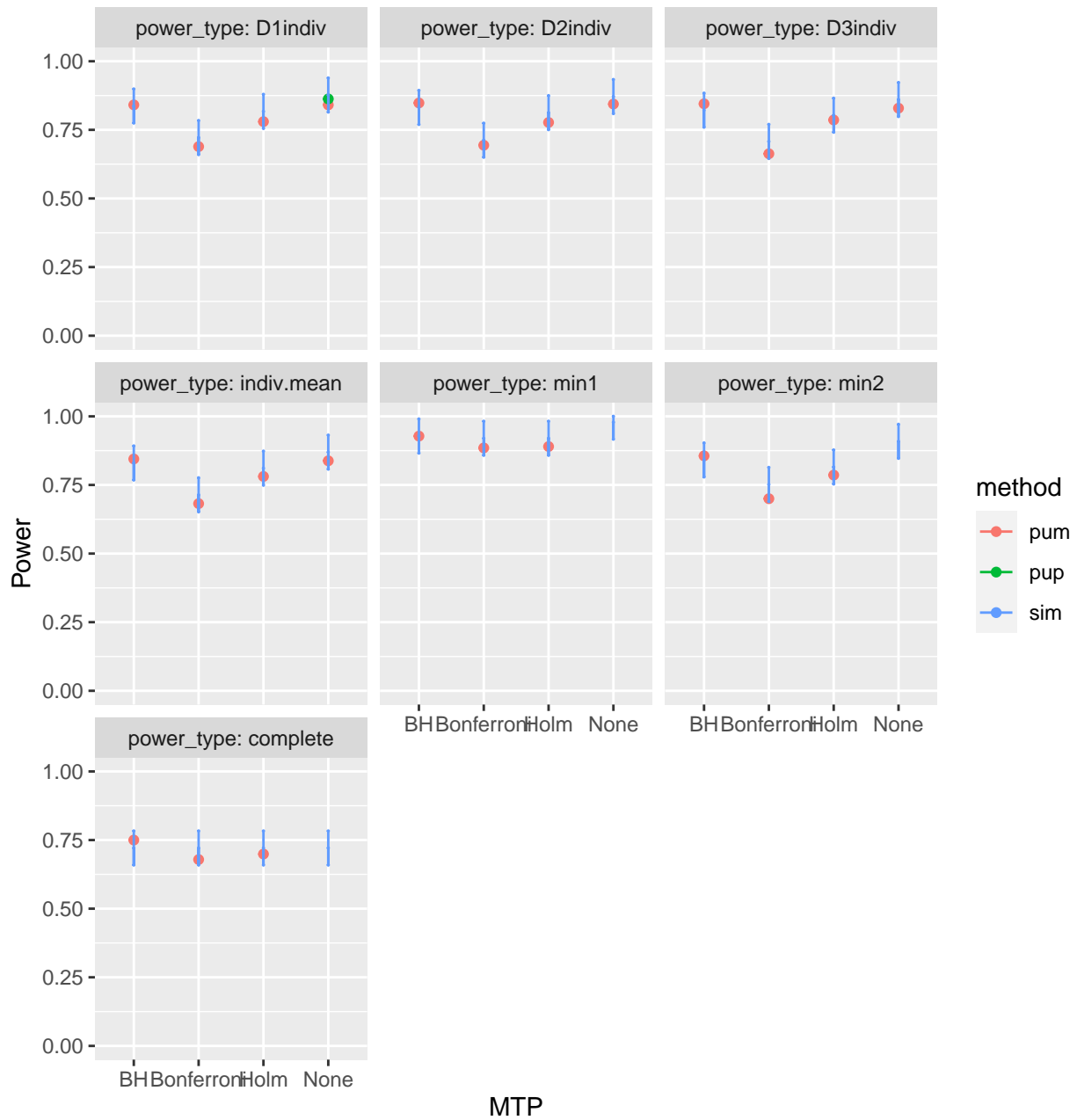
Varying R2

$R_1^2 = 0.6, 0.6, 0.6$



$$R_1^2 = 0, 0, 0$$

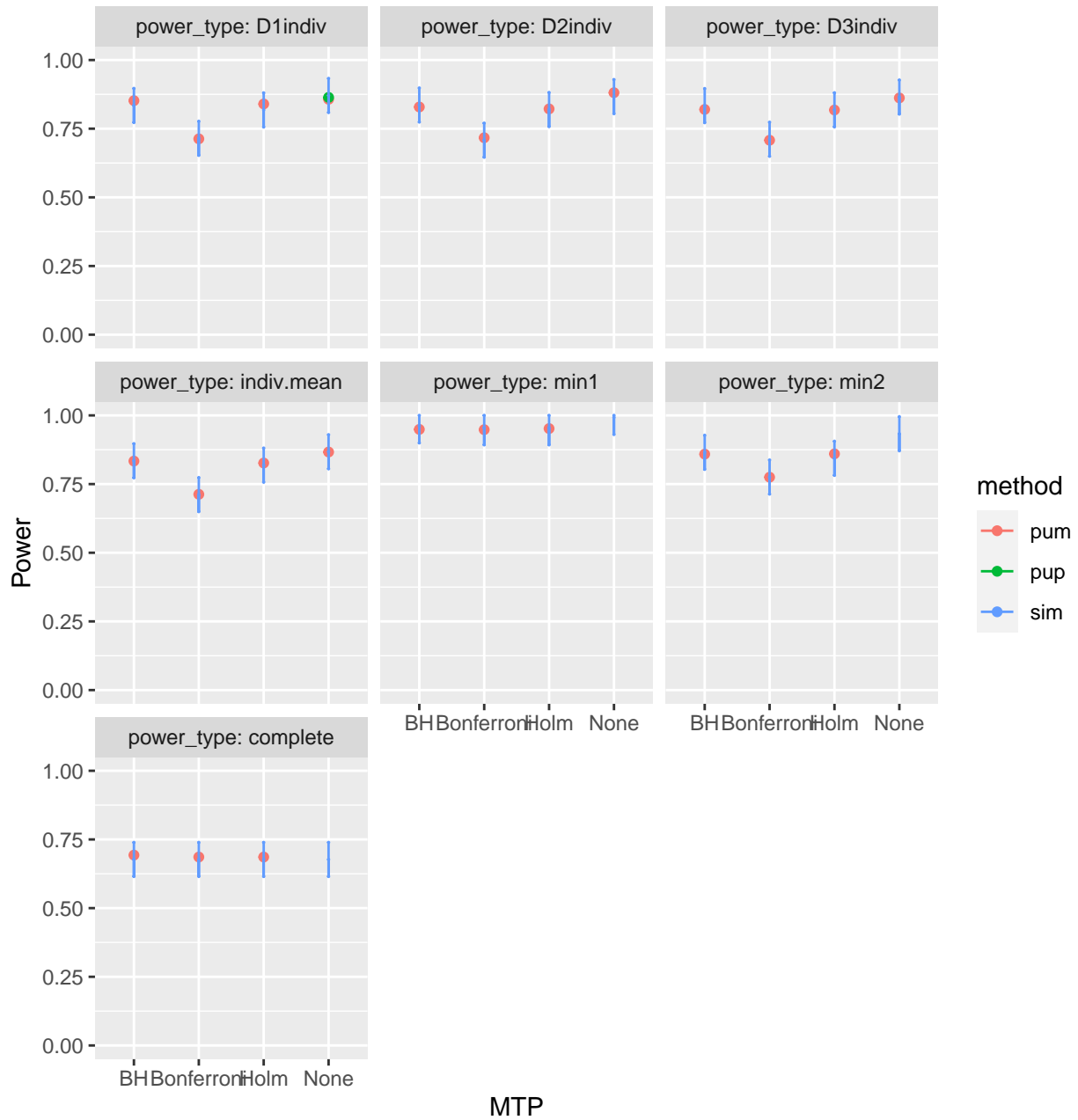
d_m: d3.1_m3rr2rr



Varying rho

$\rho = 0.2$

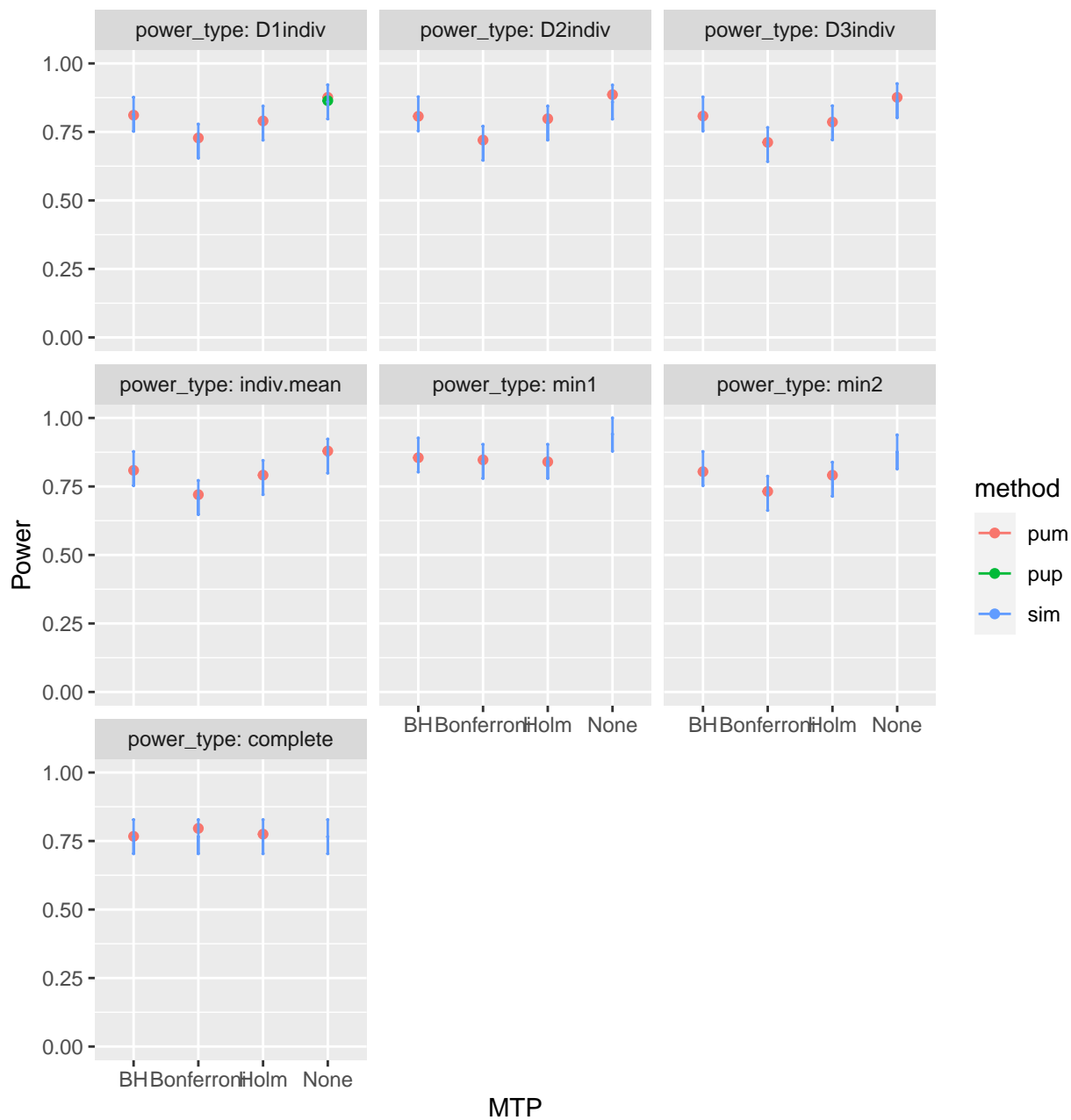
d_m: d3.1_m3rr2rr



MTP

$\rho = 0.8$

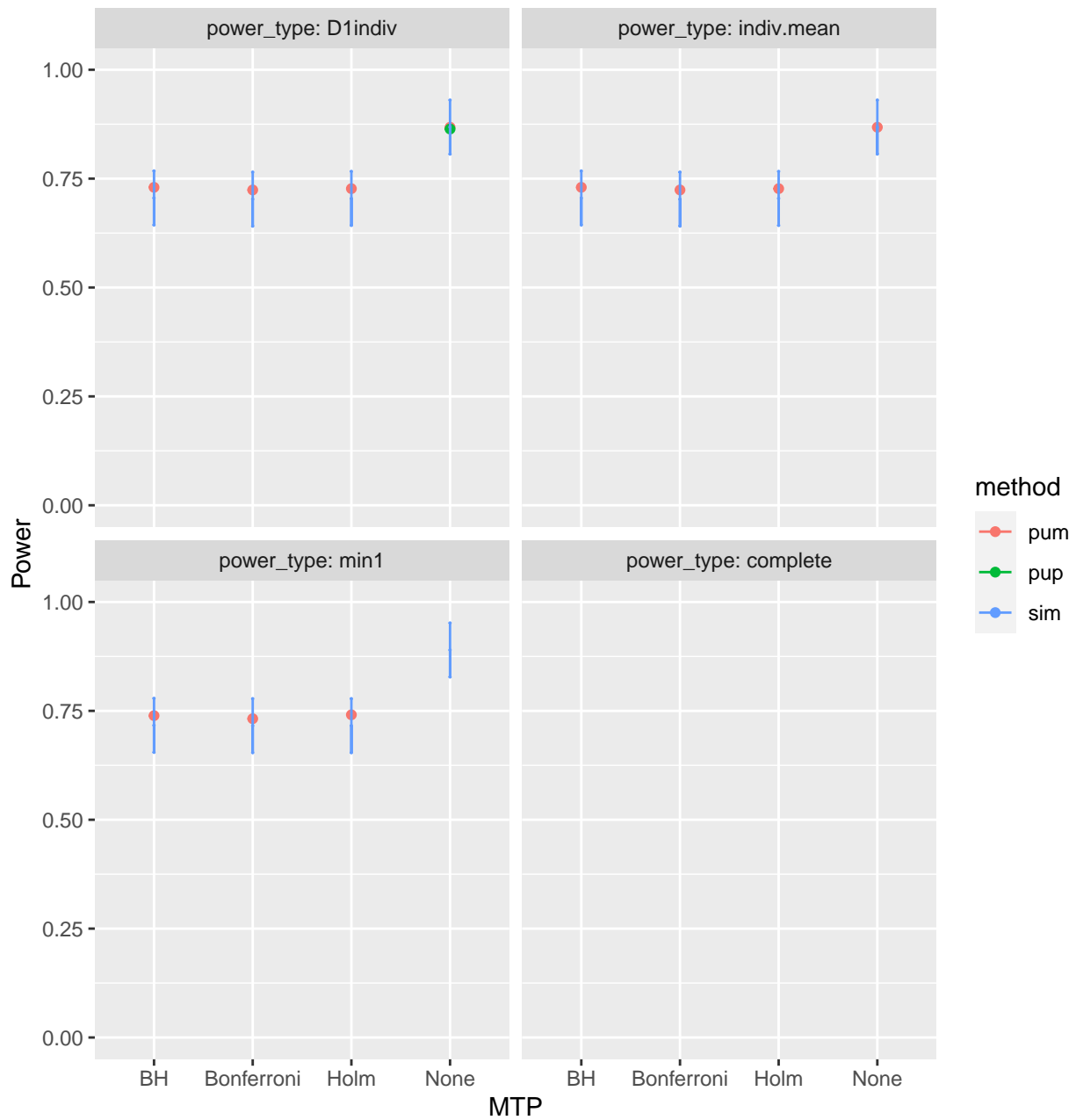
d_m: d3.1_m3rr2rr



Varying true positives

MDES = 0.125, 0, 0

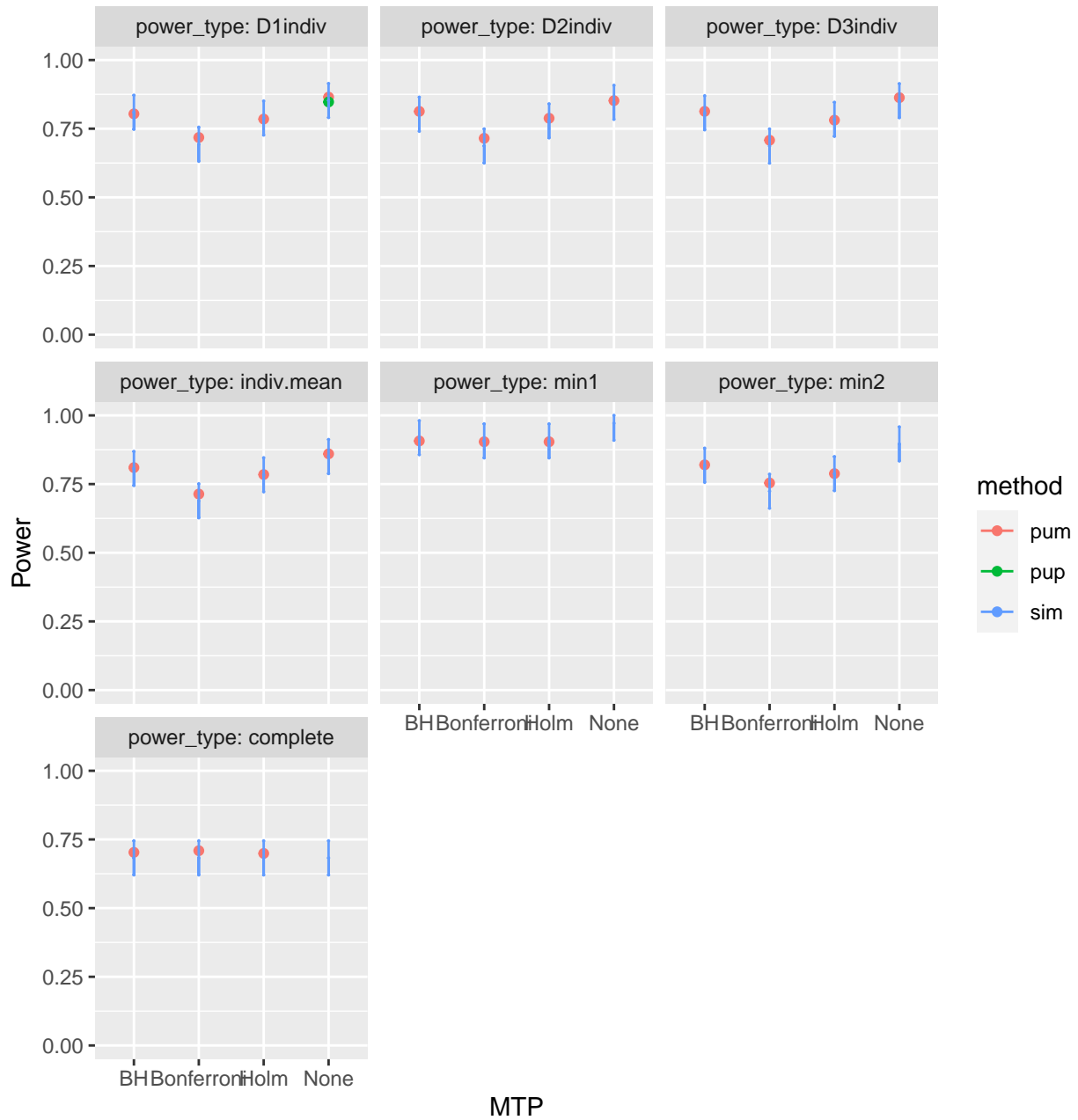
d_m: d3.1_m3rr2rr



Varying ICC

$ICC_2 = 0.7, 0.7, 0.7$

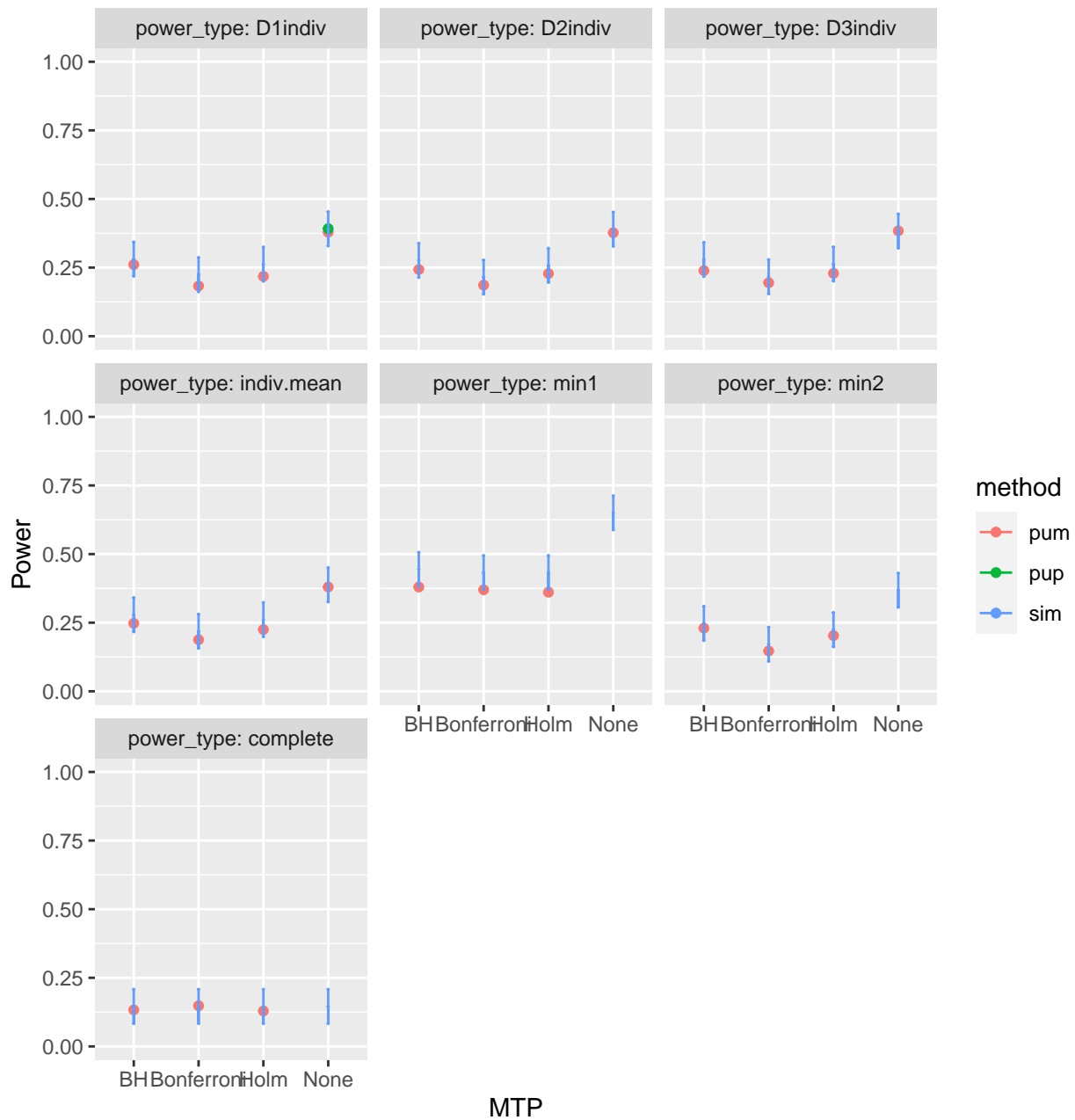
d_m: d3.1_m3rr2rr



MTP

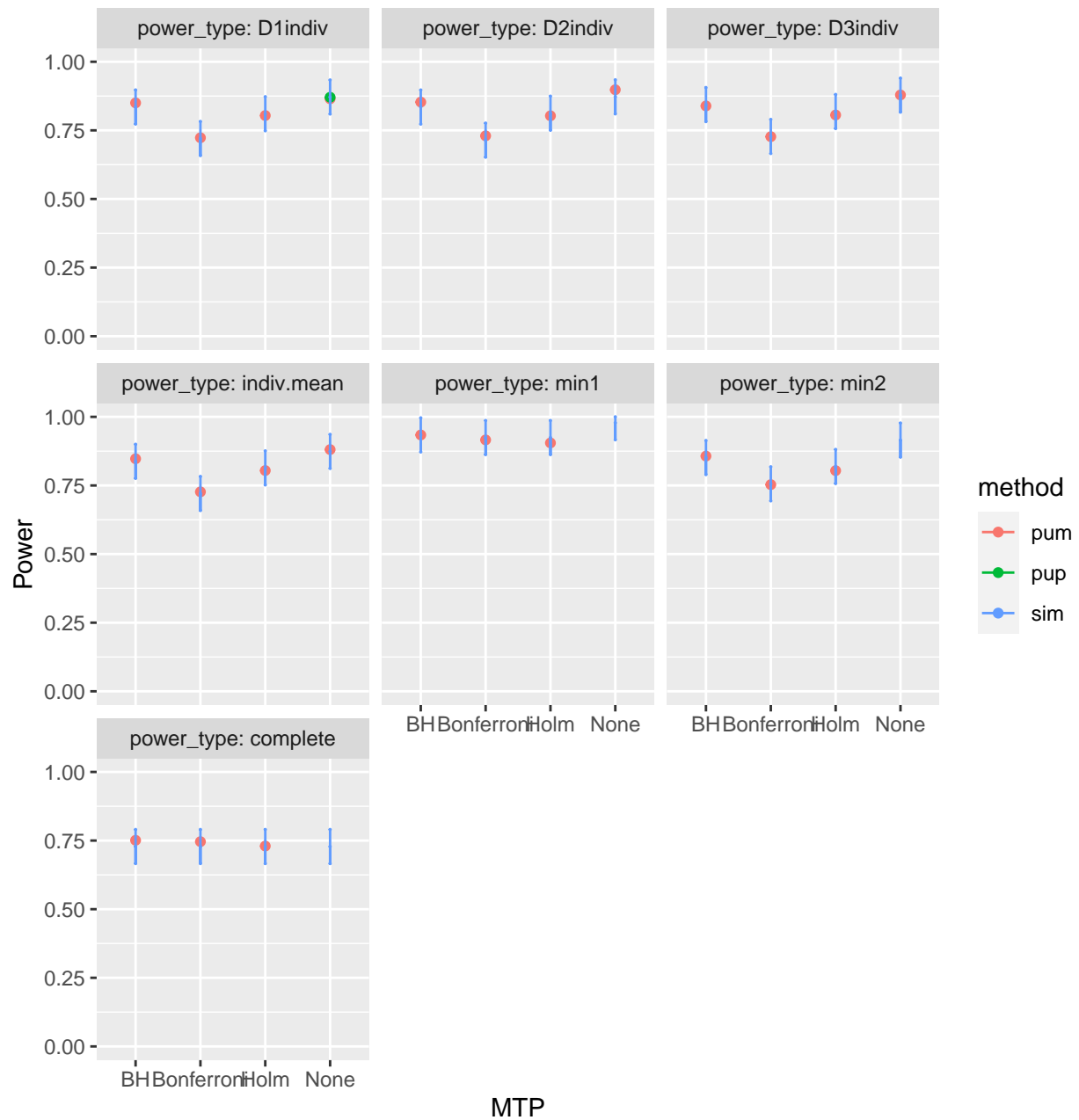
$ICC_3 = 0.7, 0.7, 0.7$

d_m: d3.1_m3rr2rr



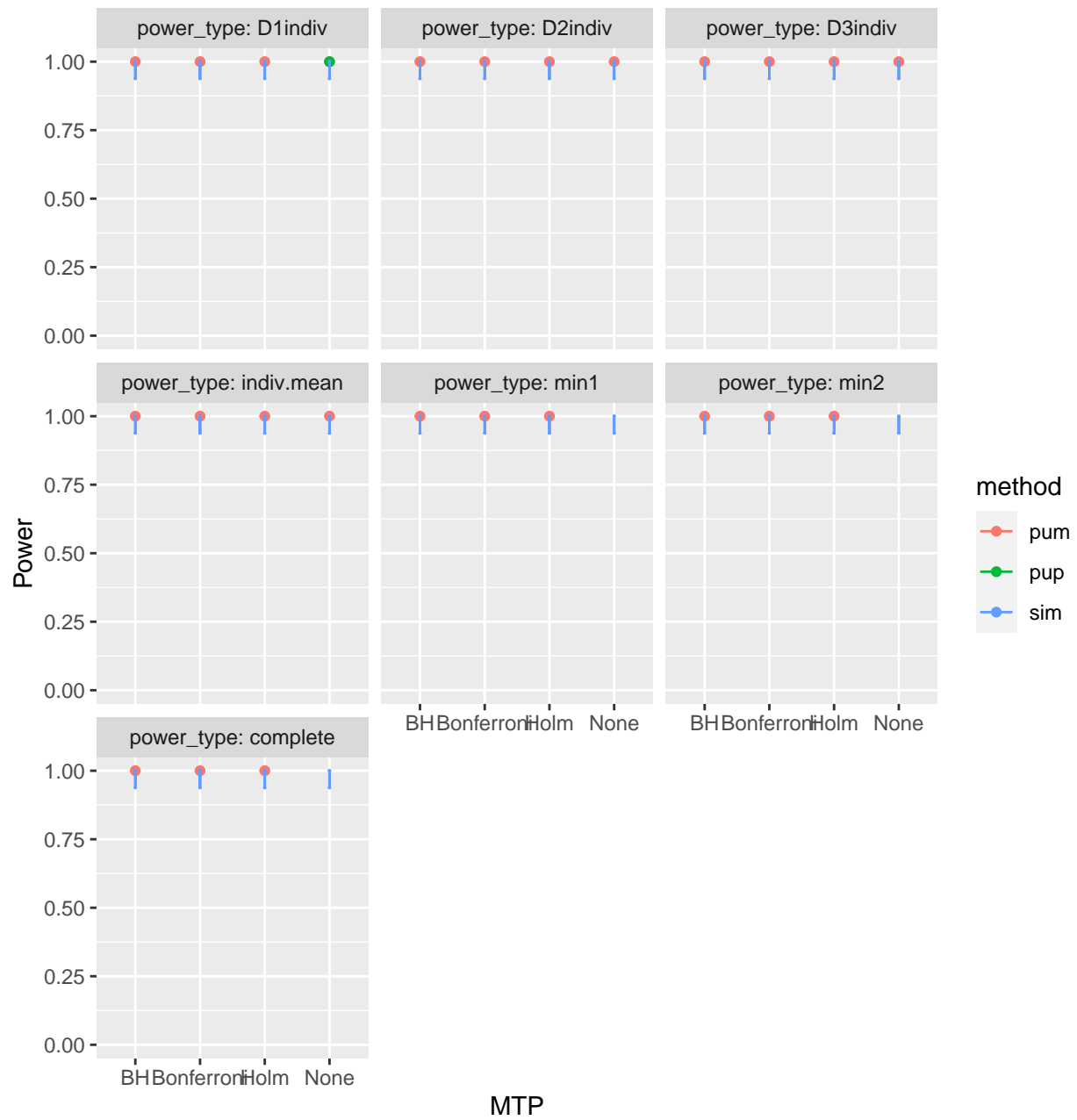
ICC₂ = 0, 0, 0

d_m: d3.1_m3rr2rr



$ICC_2 = 0.2, 0.2, 0.2$

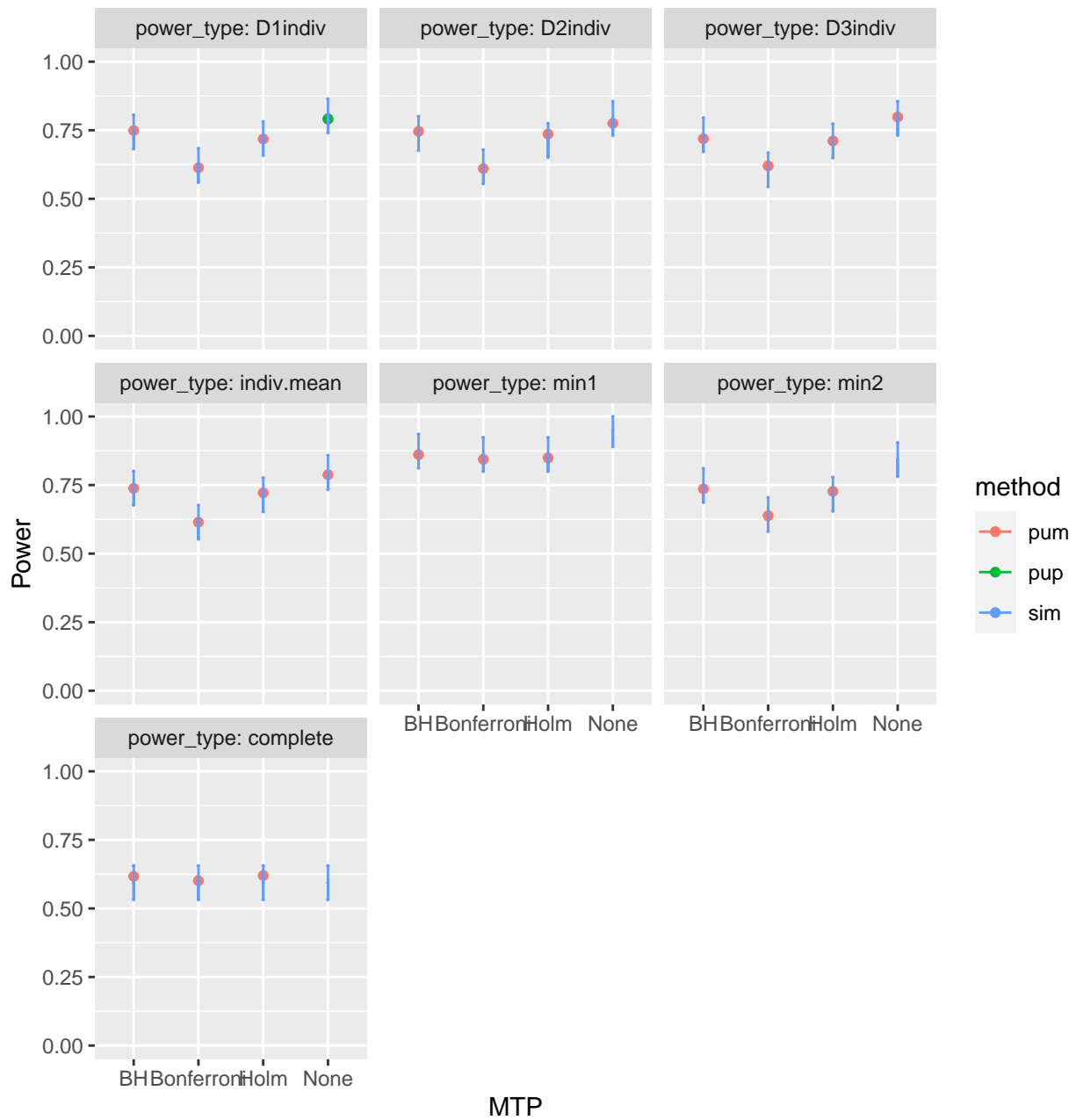
d_m: d3.1_m3rr2rr



Varying Omega

$\omega_2 = 0.8, 0.8, 0.8, \omega_3 = 0.1, 0.1, 0.1$

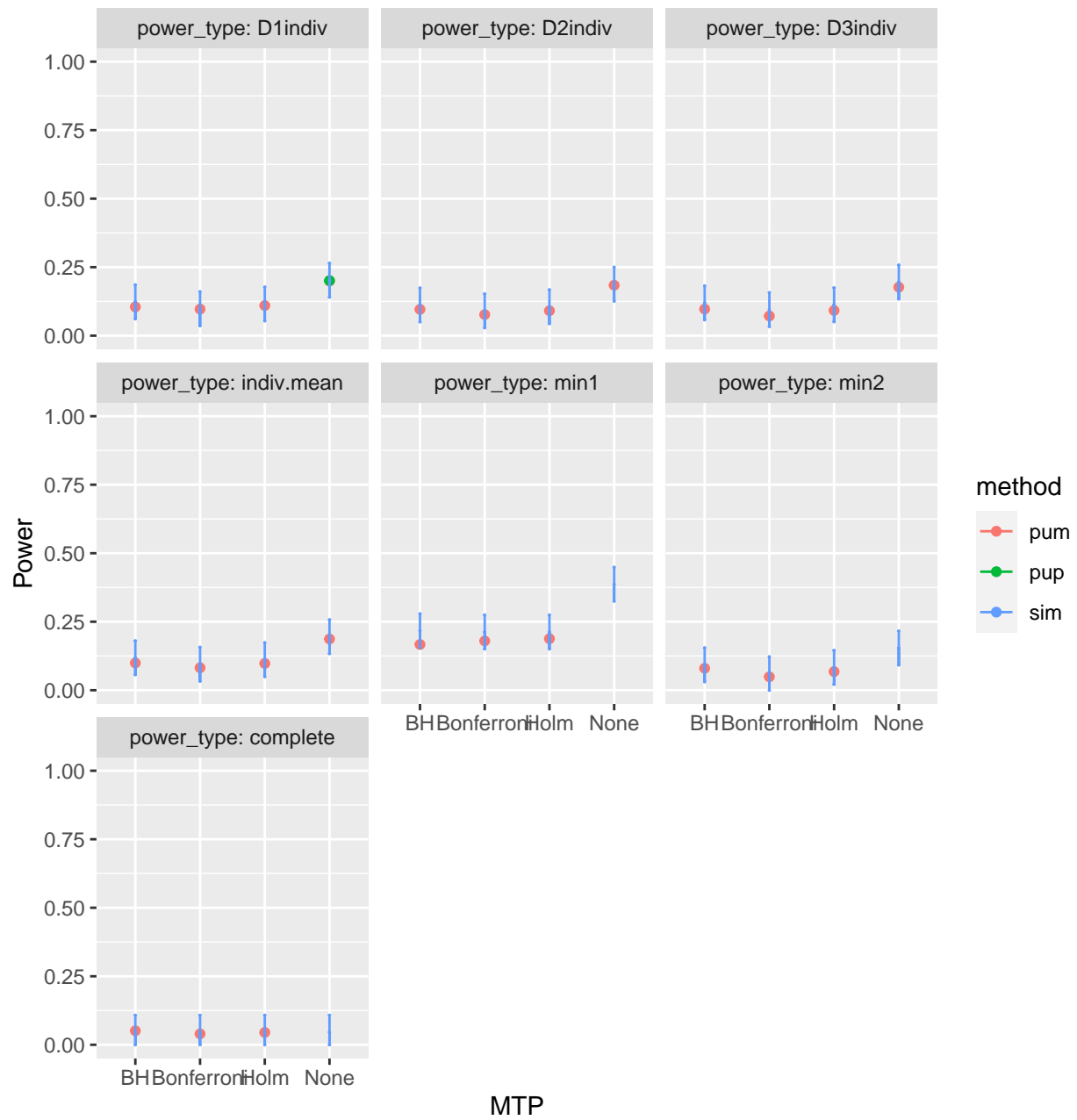
d_m: d3.1_m3rr2rr



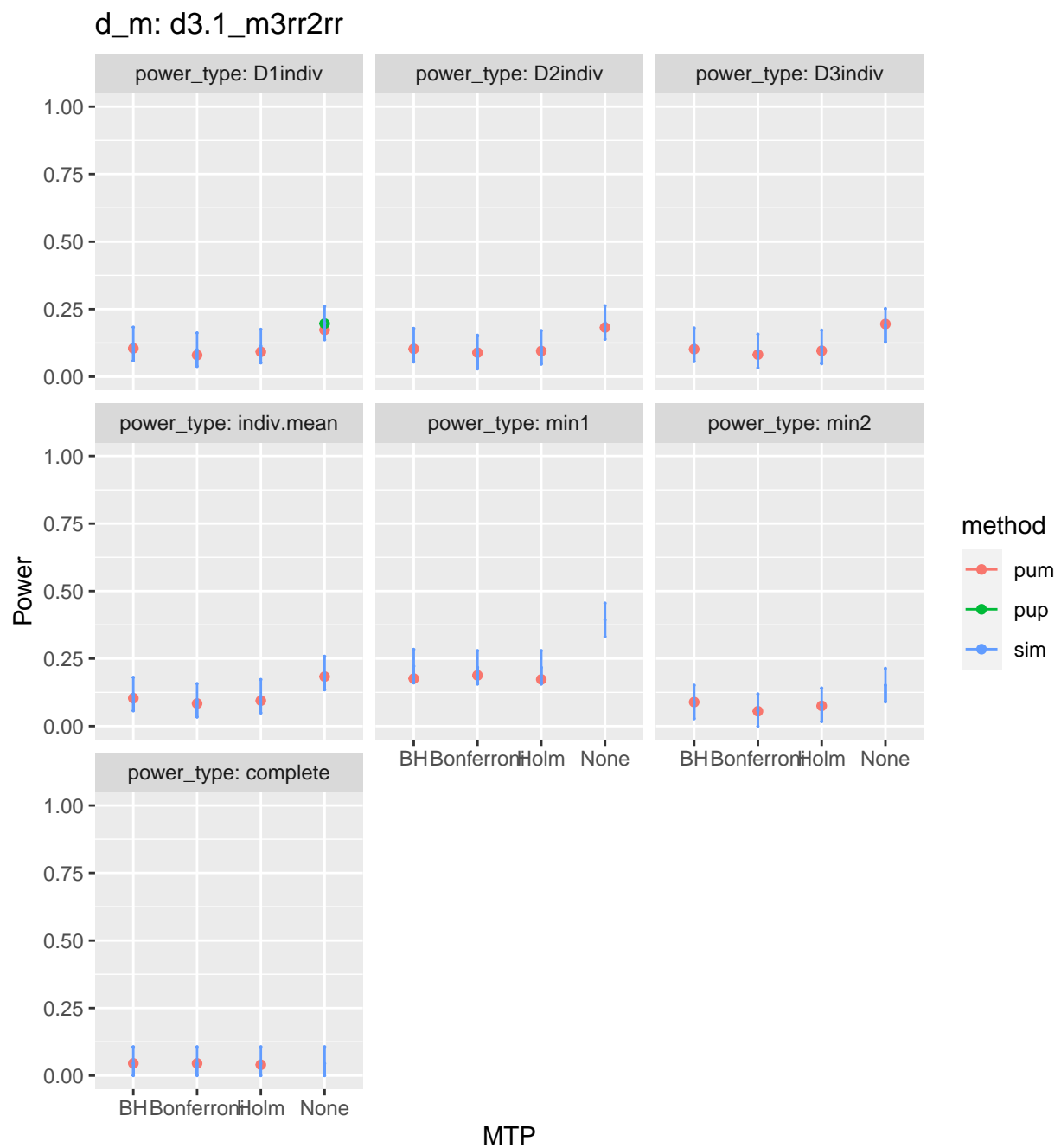
MTP

$\omega_2 = 0.1, 0.1, 0.1, \omega_3 = 0.8, 0.8, 0.8$

d_m: d3.1_m3rr2rr

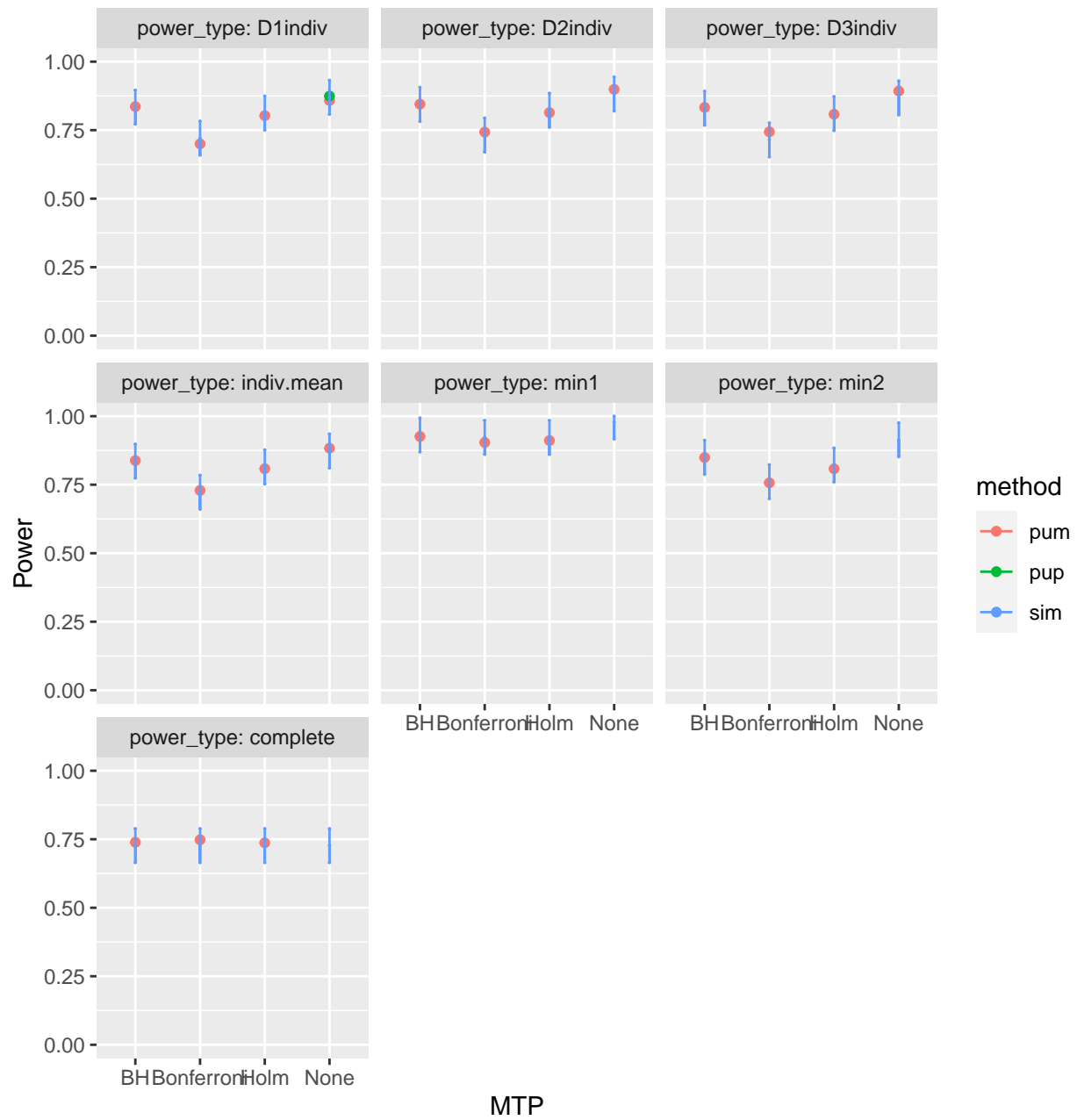


$\omega_2 = 0.8, 0.8, 0.8, \omega_3 = 0.8, 0.8, 0.8$



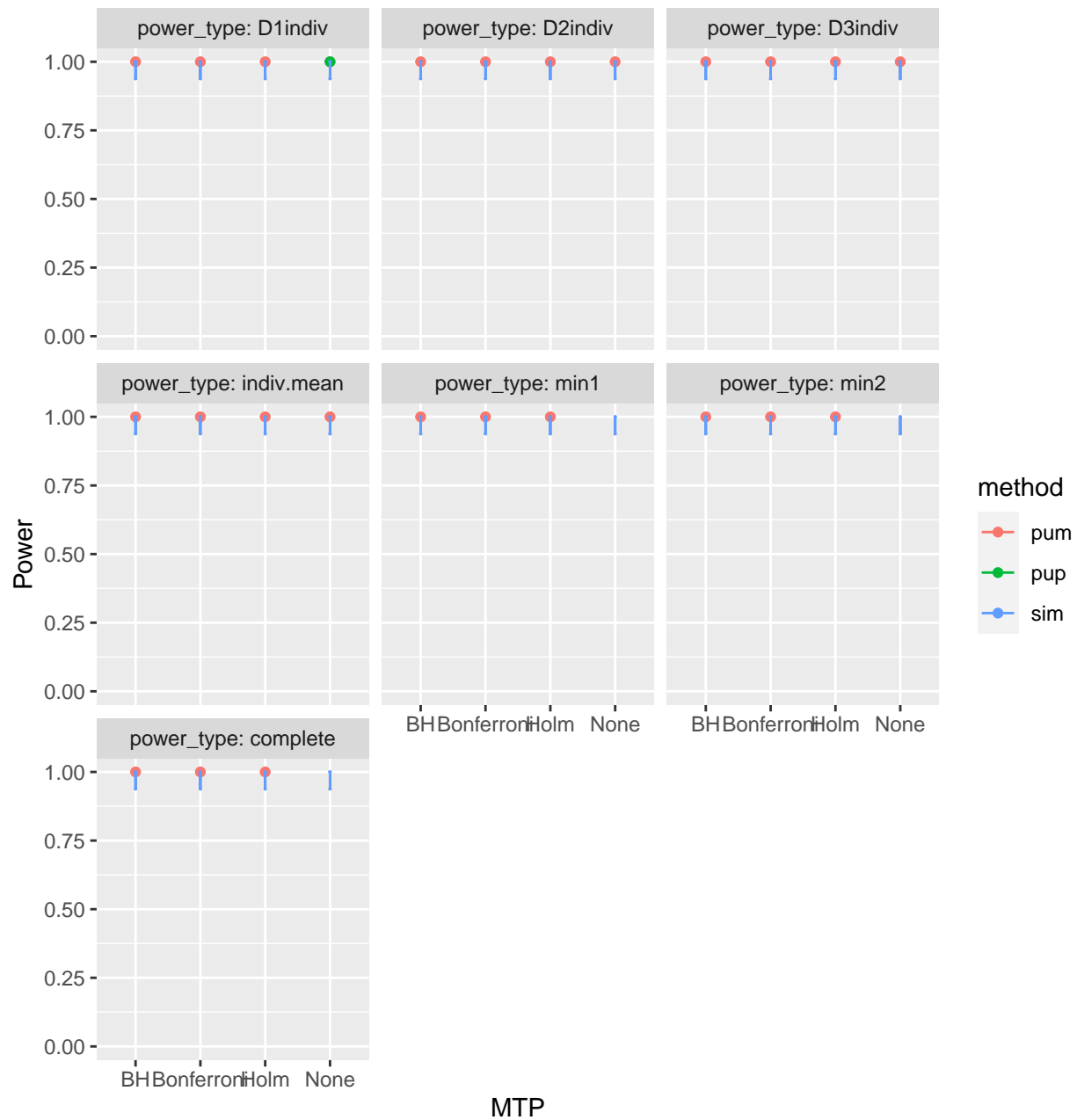
$\omega_2 = 0, 0, 0, \omega_3 = 0.1, 0.1, 0.1$

d_m: d3.1_m3rr2rr



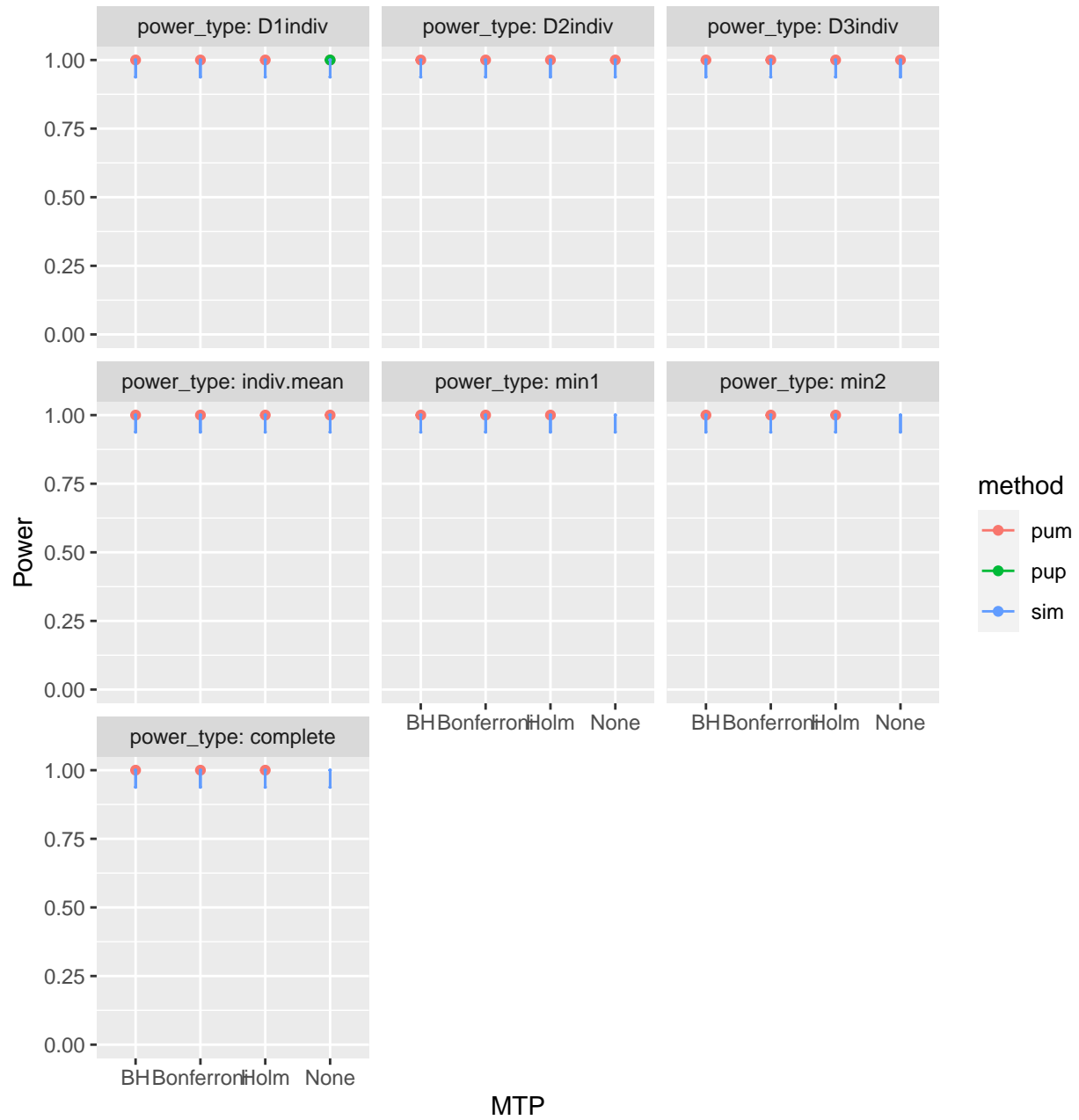
$\omega_2 = 0.1, 0.1, 0.1, \omega_3 = 0, 0, 0$

d_m: d3.1_m3rr2rr



$\omega_2 = 0, 0, 0, \omega_3 = 0, 0, 0$

d_m: d3.1_m3rr2rr



MDES validation

Target value: 0.125

```
##
##
## +-----+-----+-----+-----+
## |      MTP      | Adjusted MDES | D1indiv Power | Target MDES |
## +=====+=====+=====+=====+
## | Bonferroni |      0.126    |      0.727    |      0.125    |
## +-----+-----+-----+-----+
## |      BH      |      0.124    |      0.826    |      0.125    |
## +-----+-----+-----+-----+
## |      Holm     |      0.123    |      0.796    |      0.125    |
## +-----+-----+-----+-----+
##
## Table: d3.1_m3rr2rr
```

Sample size validation

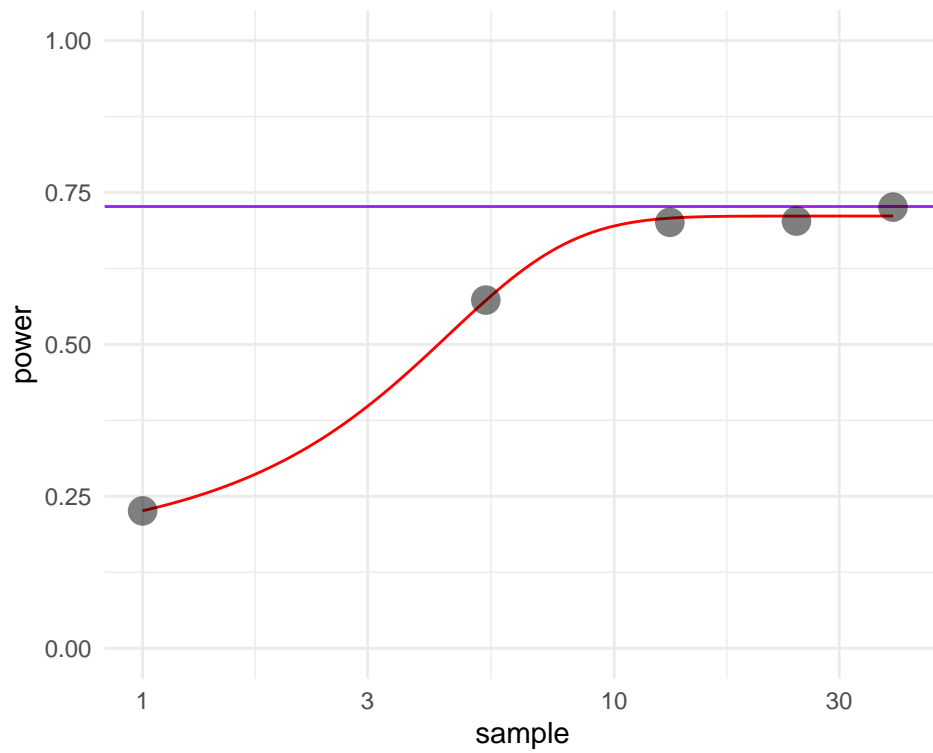
Target value: 15

```
##
##
## +-----+-----+-----+-----+
## |      MTP      | Sample.type | Sample.size | D1indiv.power |
## +=====+=====+=====+=====+
## | Bonferroni |      K      |      15     |      0.727     |
## +-----+-----+-----+-----+
## |      BH      |      K      |      15     |      0.829     |
## +-----+-----+-----+-----+
## |      Holm     |      K      |      15     |      0.795     |
## +-----+-----+-----+-----+
##
## Table: d3.1_m3rr2rr
```

Target value: 30

```
##
##
## +-----+-----+-----+-----+
## |      MTP      | Sample.type | Sample.size | D1indiv.power |
## +=====+=====+=====+=====+
## | Bonferroni |      J      |      39     |      0.727     |
## +-----+-----+-----+-----+
## |      BH      |      J      |      27     |      0.826     |
## +-----+-----+-----+-----+
## |      Holm     |      J      |      NA     |      NA        |
## +-----+-----+-----+-----+
##
## Table: d3.1_m3rr2rr
```

Note: particularly flat power curves results in discrepancy for J.



Target value: 100

```
##
##
## +-----+-----+-----+-----+
## |      MTP      | Sample.type | Sample.size | D1indiv.power |
## +=====+=====+=====+=====+
## | Bonferroni |    nbar    |    186.2    |    0.727    |
## +-----+-----+-----+-----+
## |      BH      |    nbar    |    1120     |    0.842     |
## +-----+-----+-----+-----+
## |      Holm     |    nbar    |     58      |    0.797     |
## +-----+-----+-----+-----+
##
## Table: d3.1_m3rr2rr
```

Note: particularly flat power curves results in discrepancy for `nbar`.

