

Power Validation

MDES validation

Sample size validation

Validate Power: blocked_c2_3fr

Code ▼

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Design: Blocked Cluster RCT, with 3 levels, and randomization done at level 2 (school level).

Models: random and fixed treatment effects.

Default parameters:

- $M = 3$
- $J = 20$
- $K = 10$
- $\rho = 0.5$
- ATE ES = 0.125, 0.125, 0.125
- R21, R22: $R_1^2 = 0.1, 0.1, 0.1, R_2^2 = 0.1, 0.1, 0.1$
- ICC2: $ICC_2 = 0.2, 0.2, 0.2$
- Omega: $\omega_2 = 0, \omega_3 = 0.1$

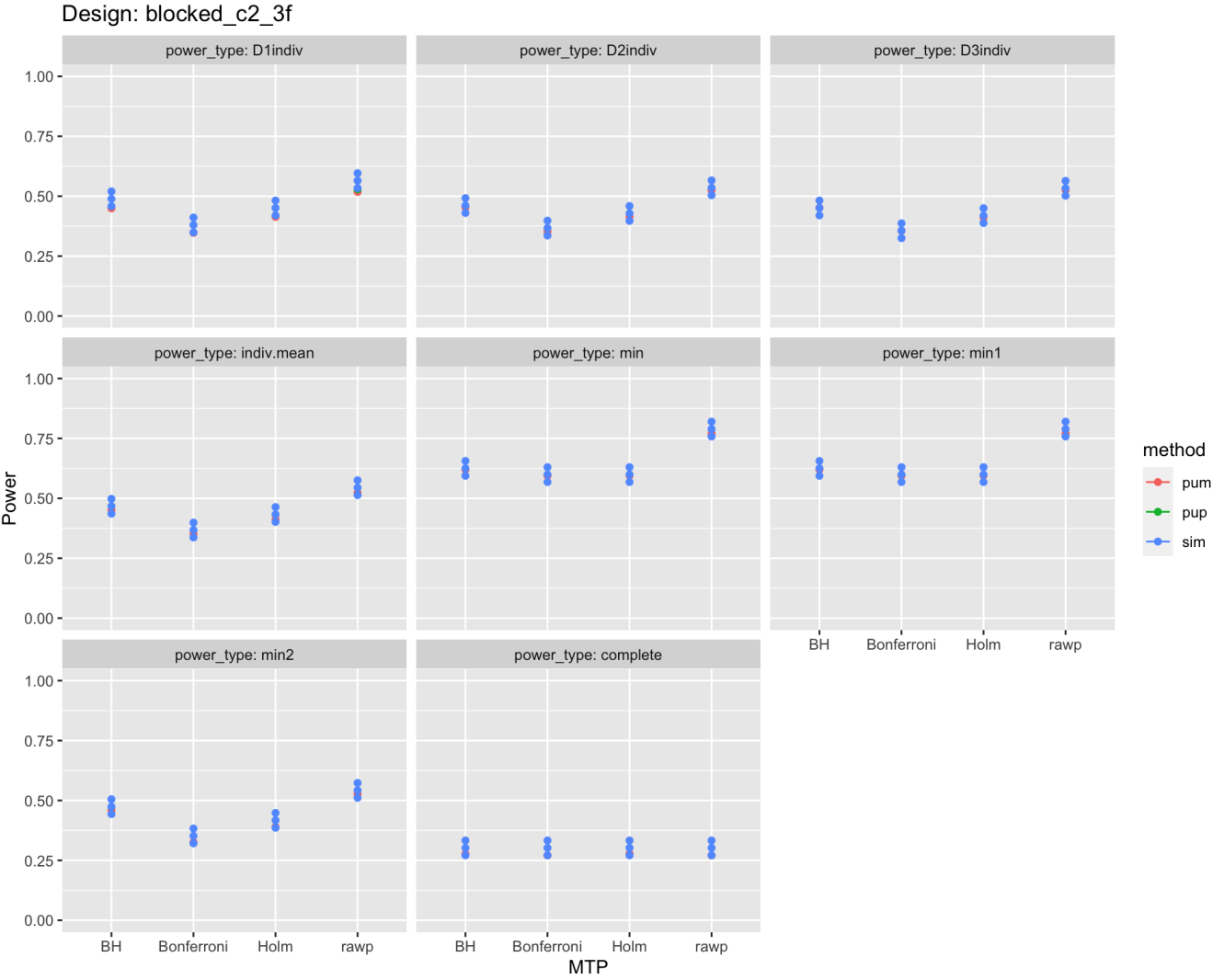
Parameters by model type:

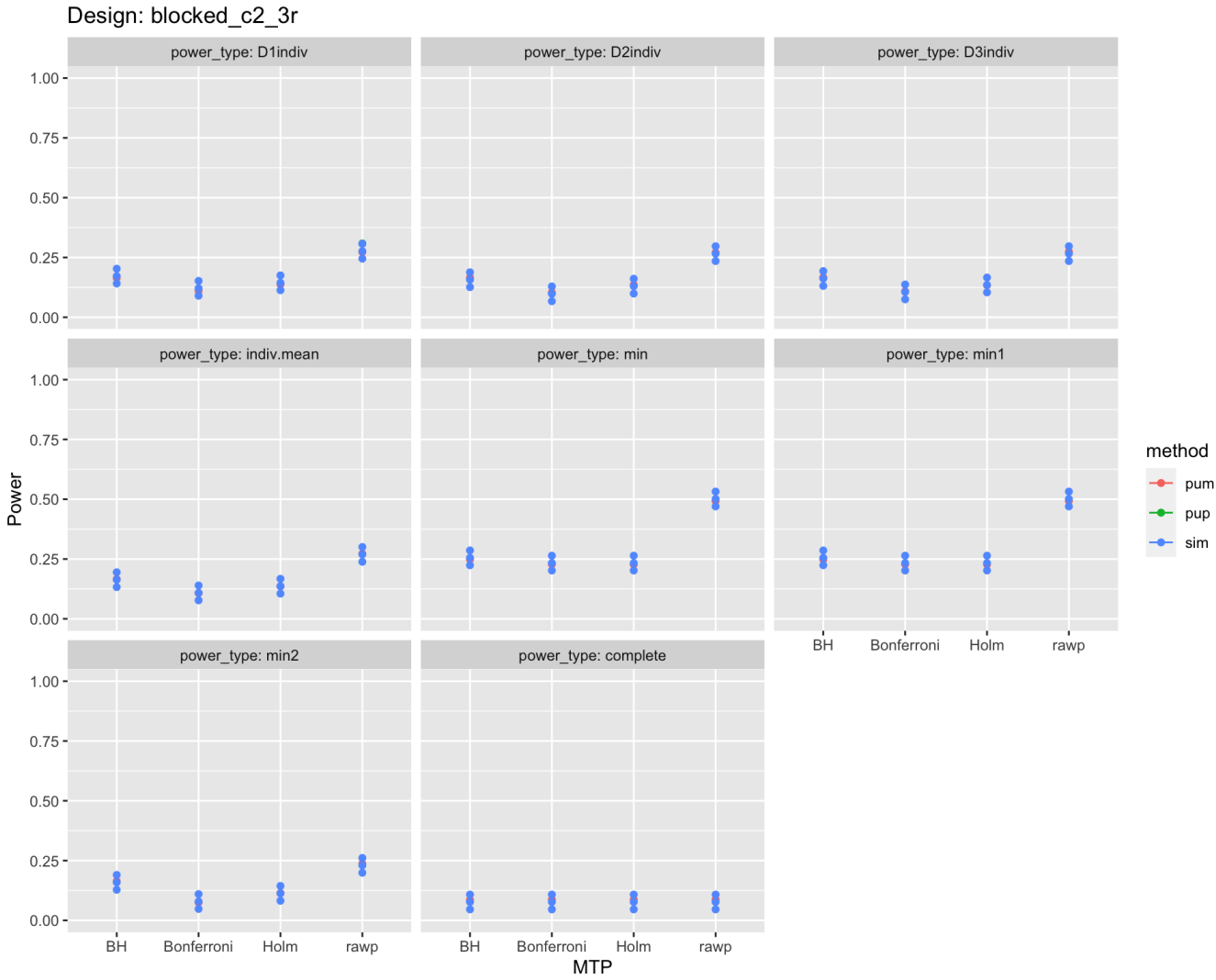
- R23: $R_3^2 = 0$ for fixed effects model, $R_3^2 = 0.1, 0.1, 0.1$ for random effects model
- ICC3: $ICC_3 = 0$ for fixed model, $ICC_3 = 0.2, 0.2, 0.2$ for random effects model

Power Validation

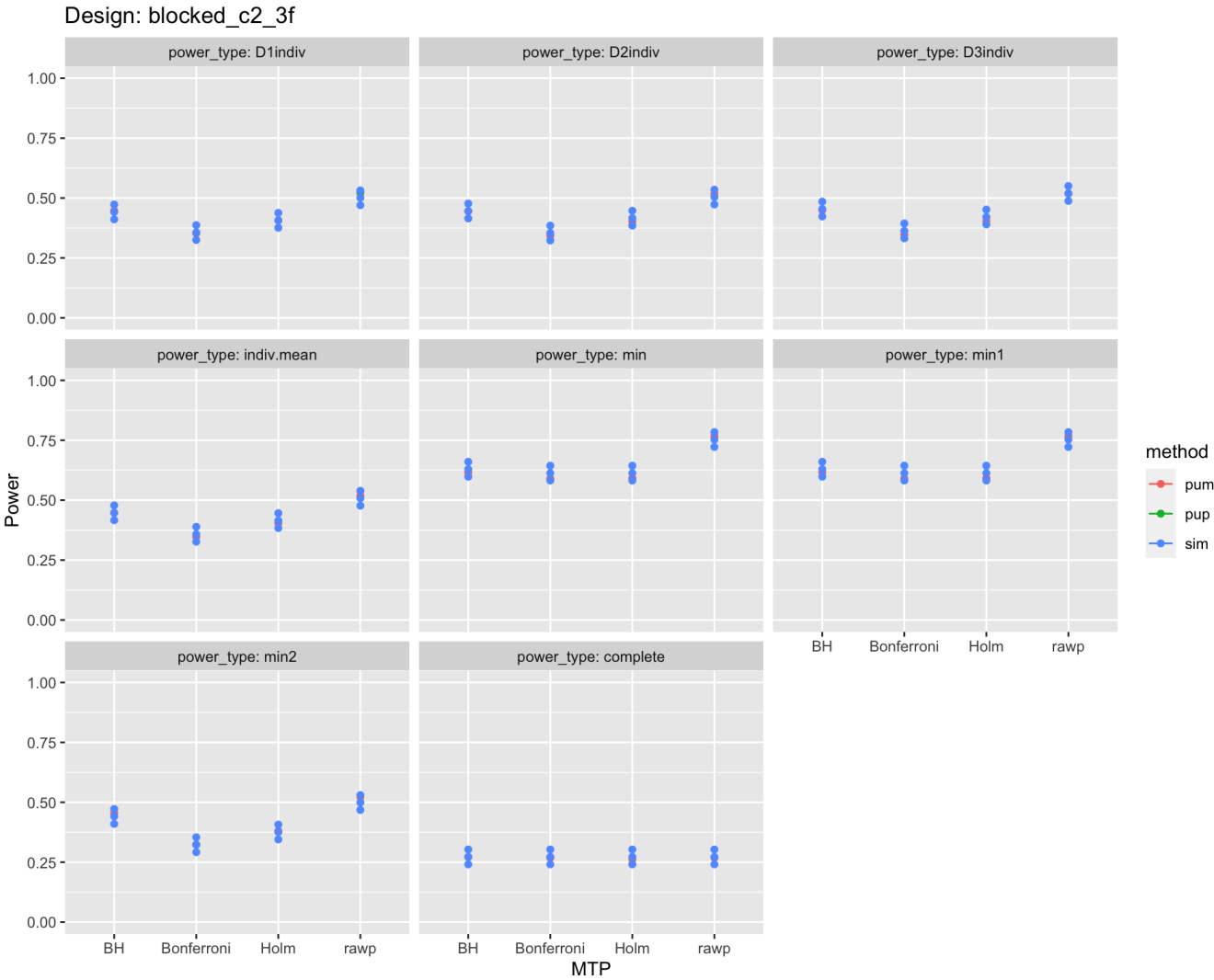
Varying school size

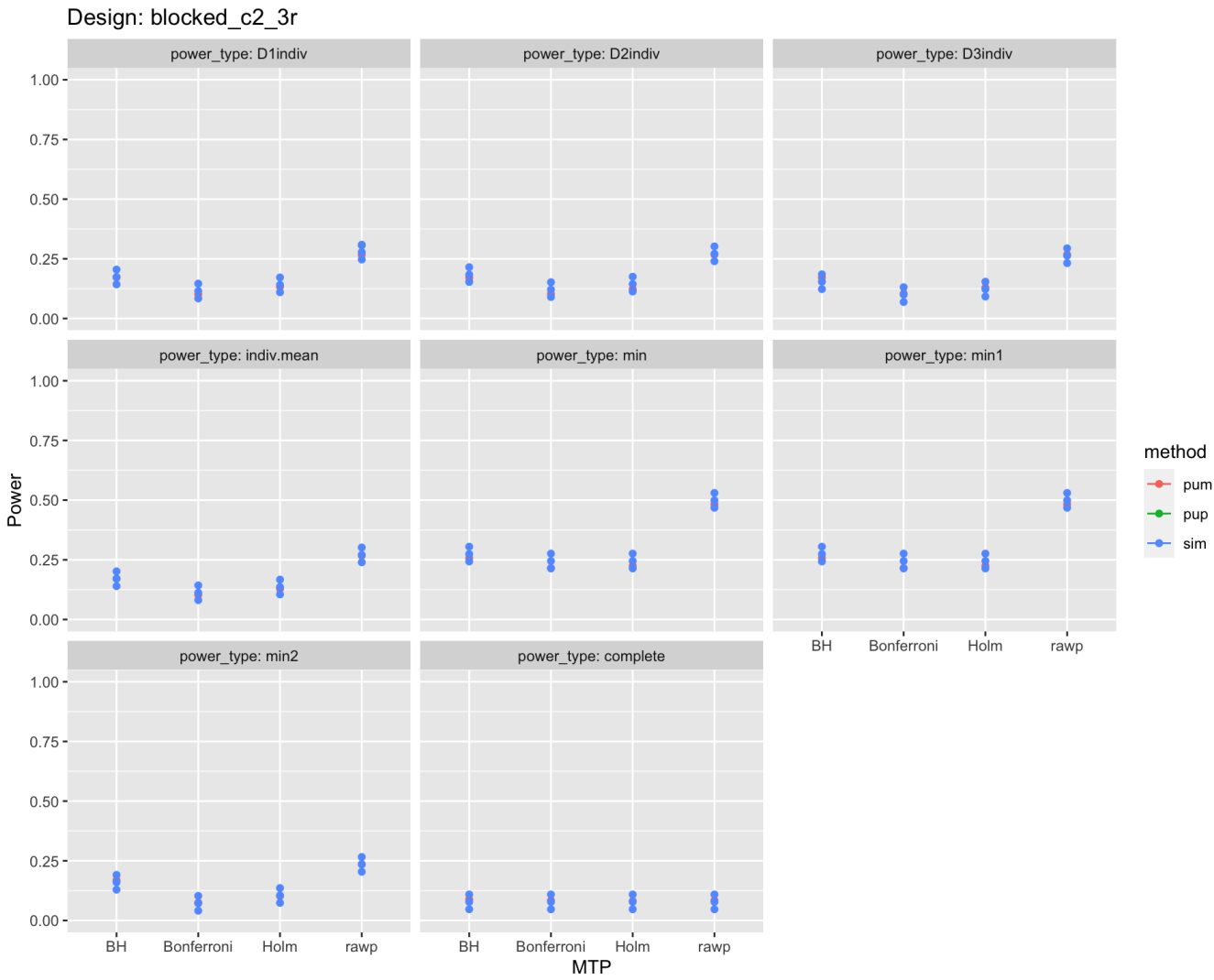
 $\bar{n} = 100$



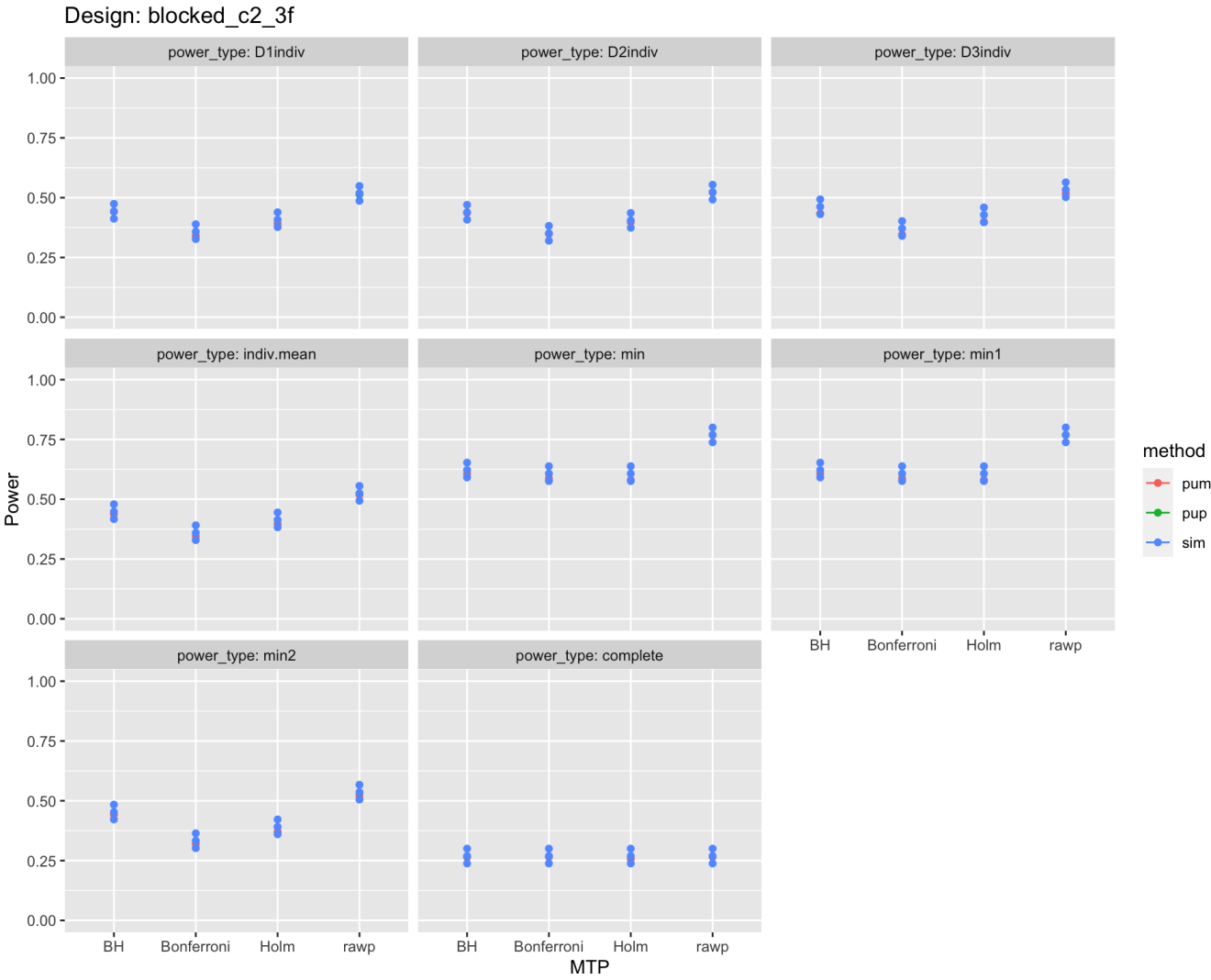


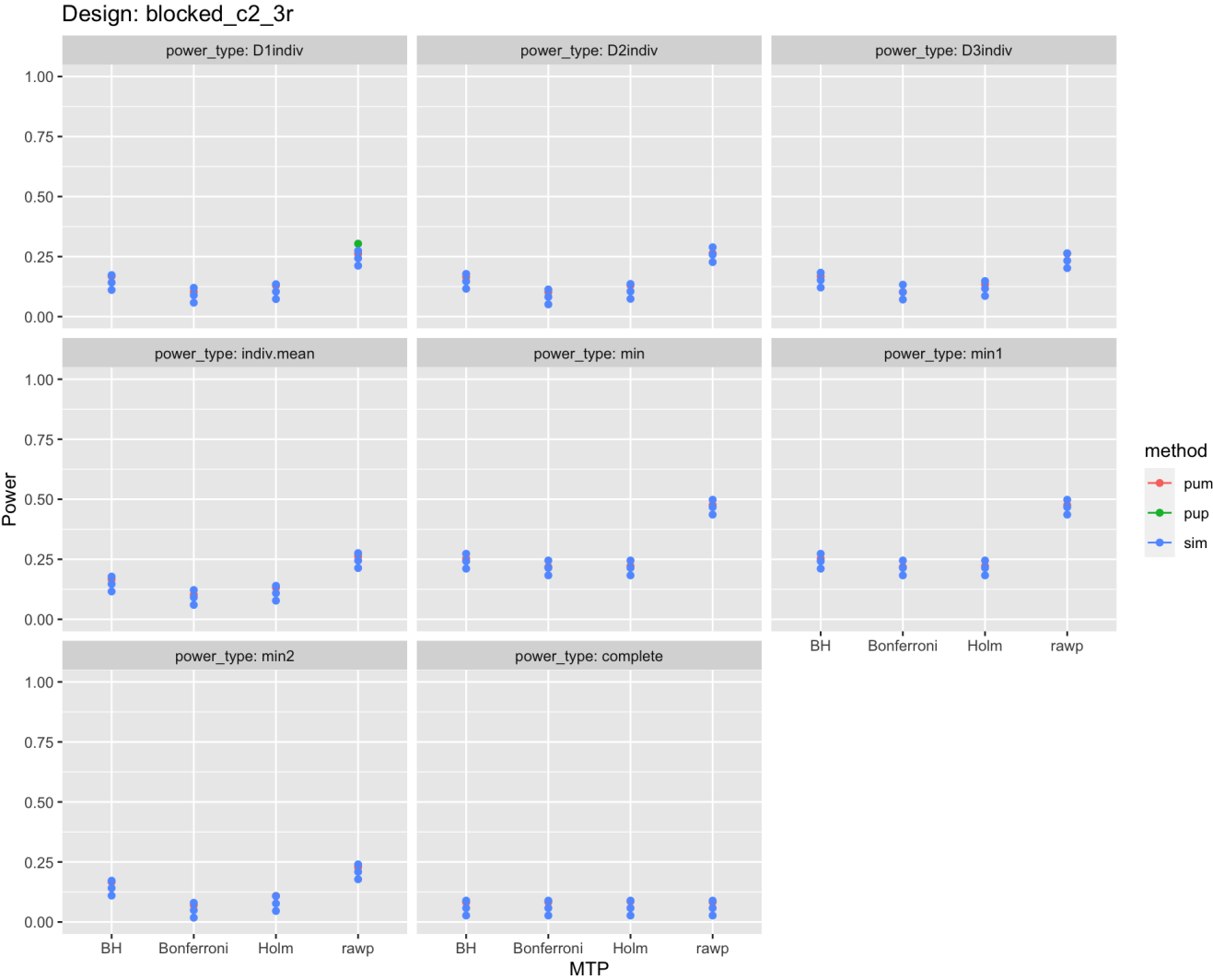
$\bar{n} = 75$





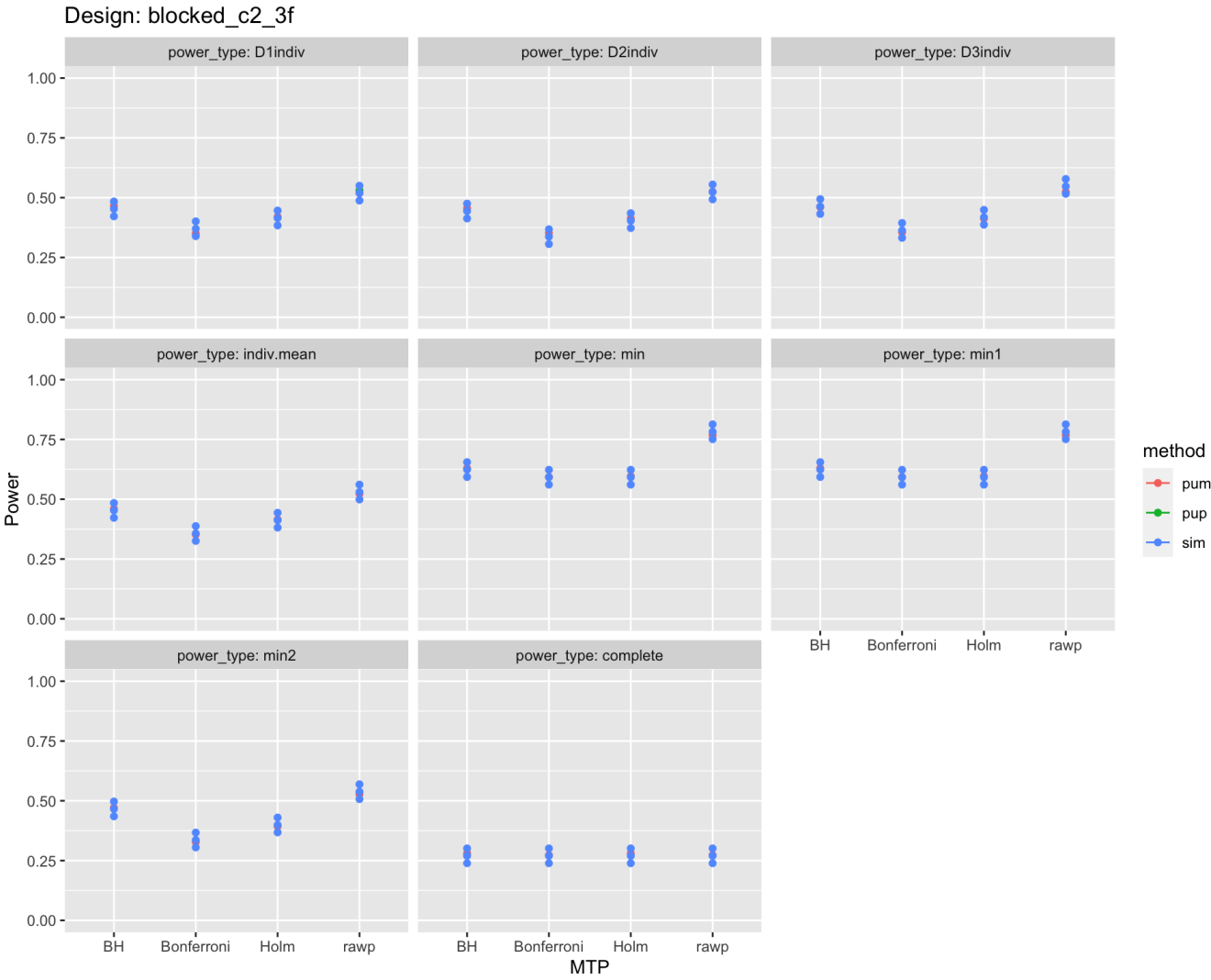
$\bar{n} = 50$

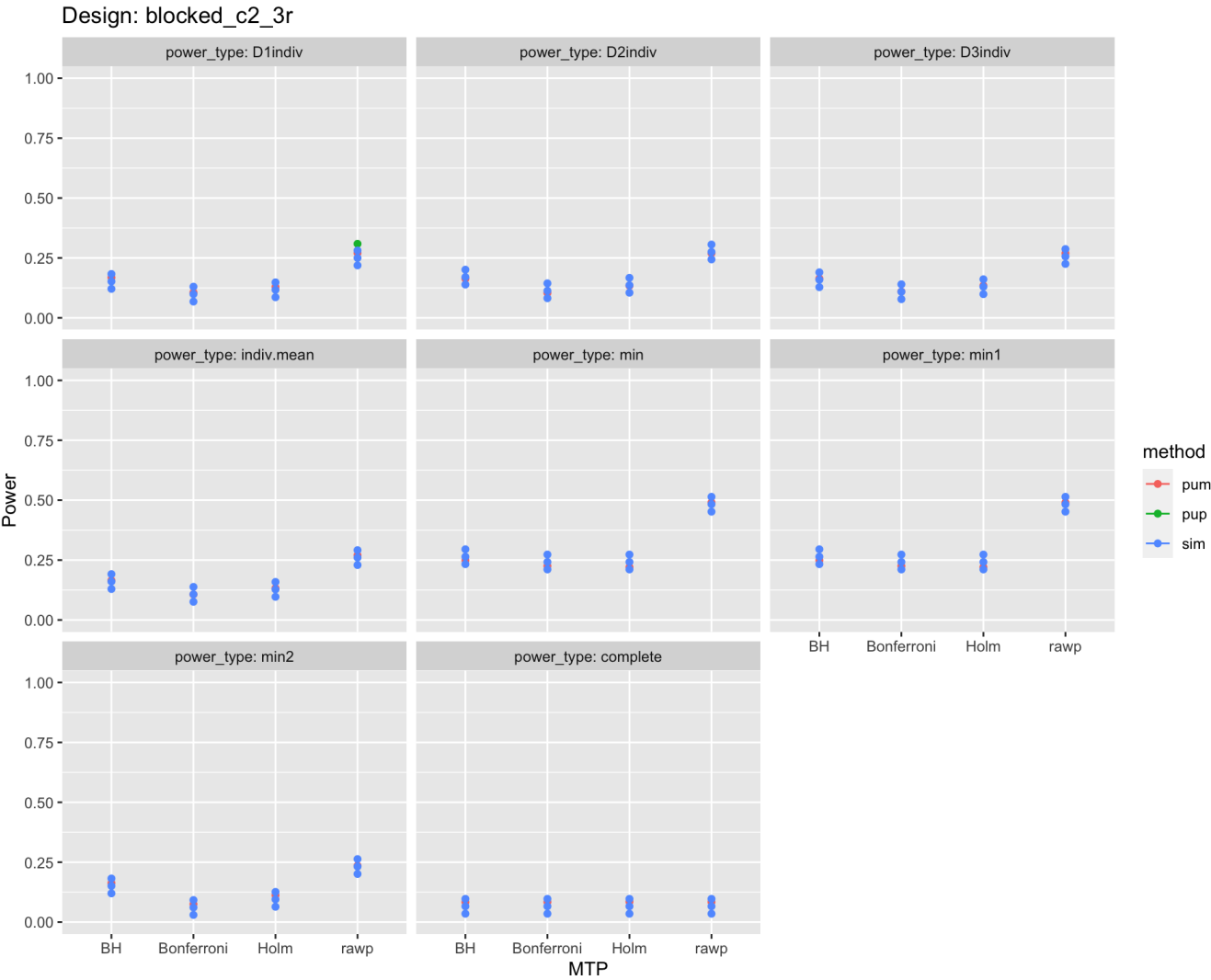




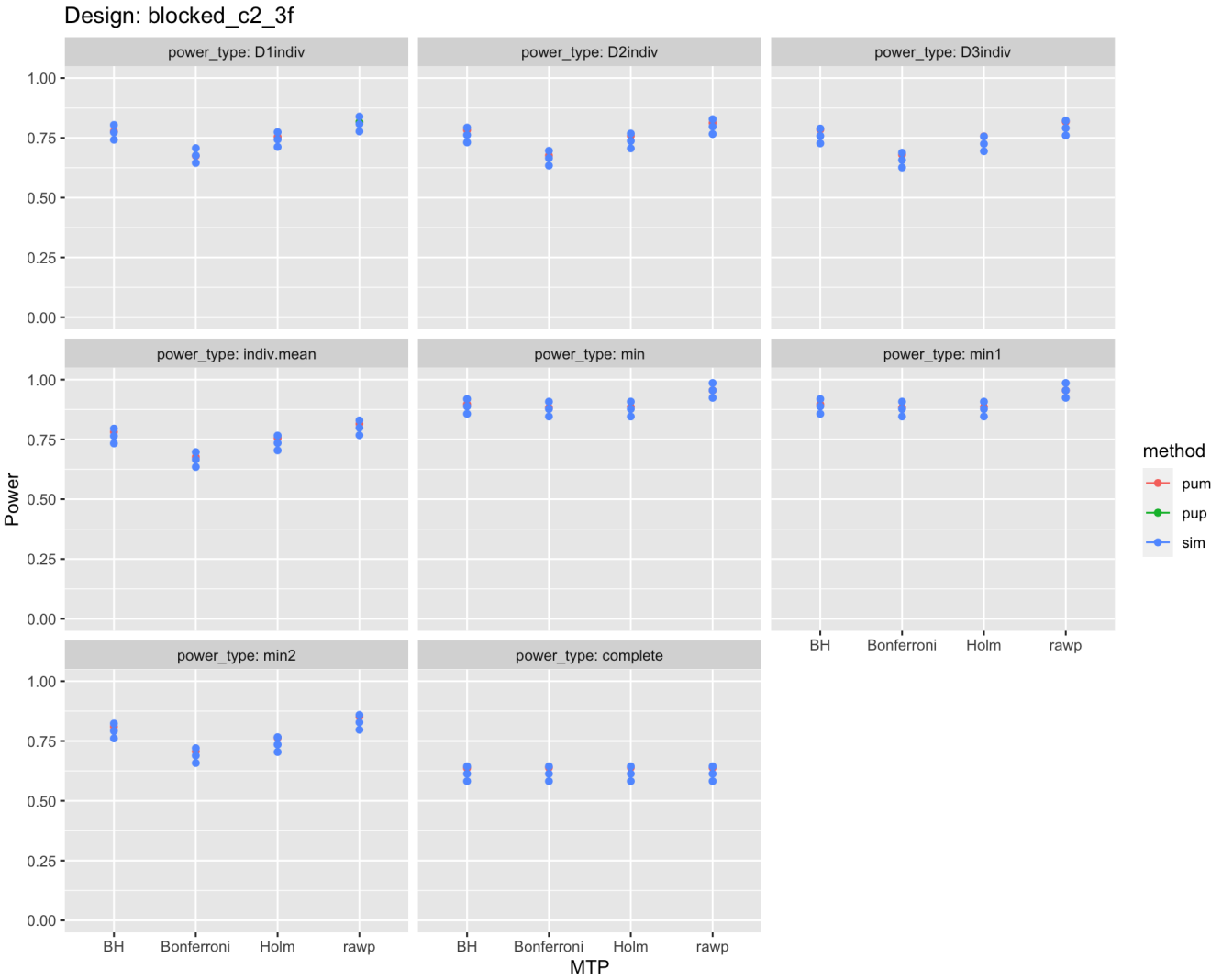
Varying R2

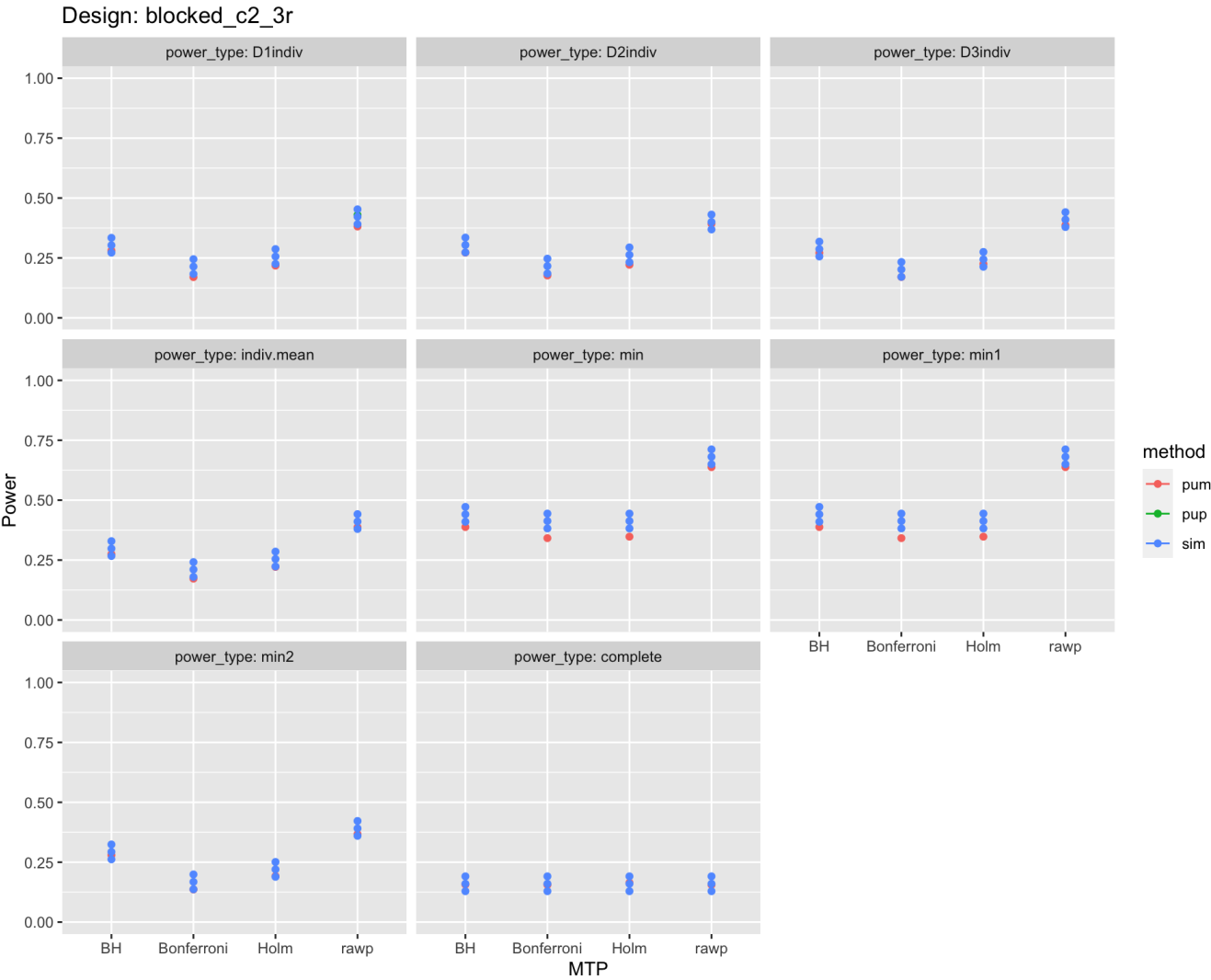
$R^2_1 = 0.6, 0.6, 0.6$



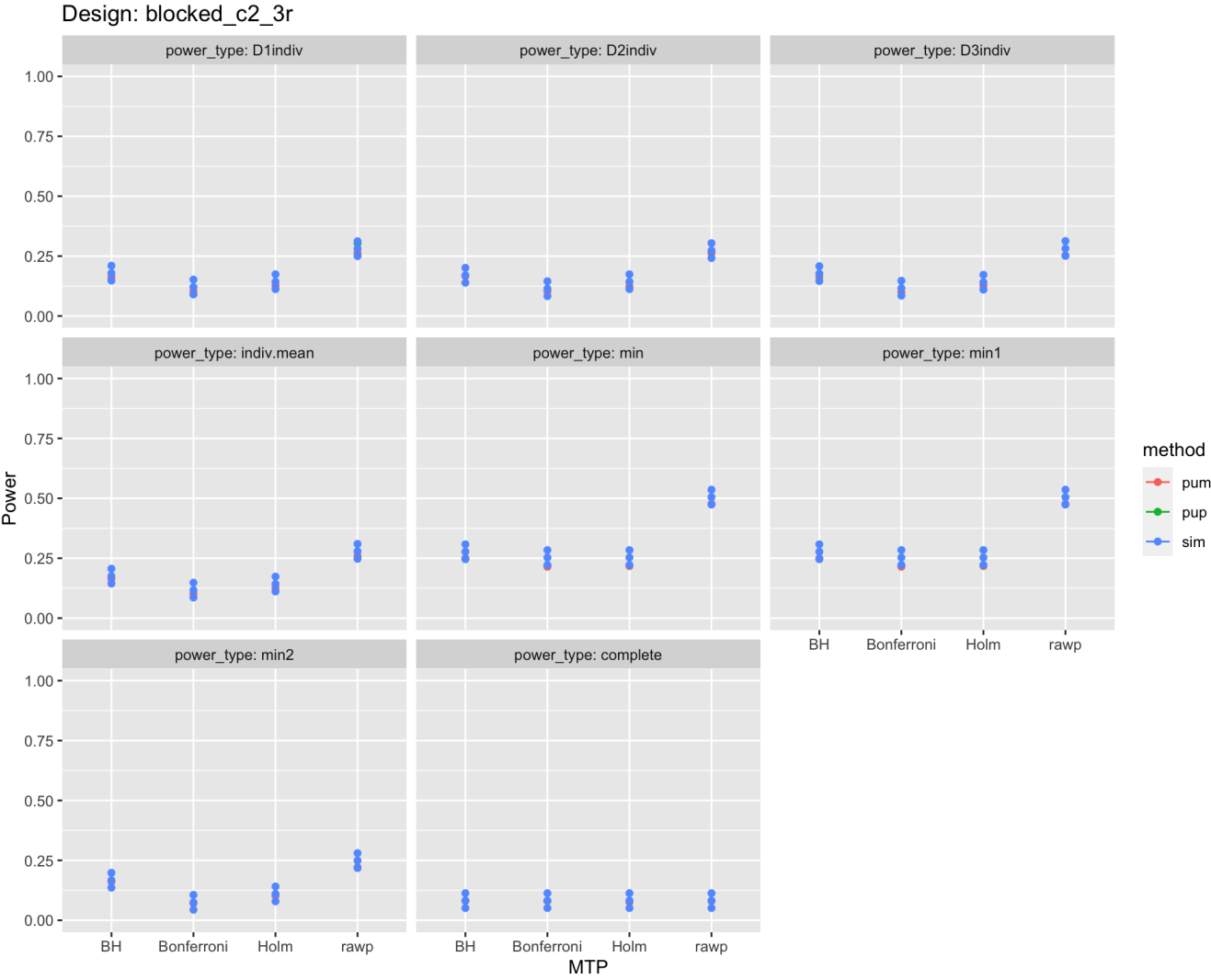


$R_2^2 = 0.6, 0.6, 0.6$

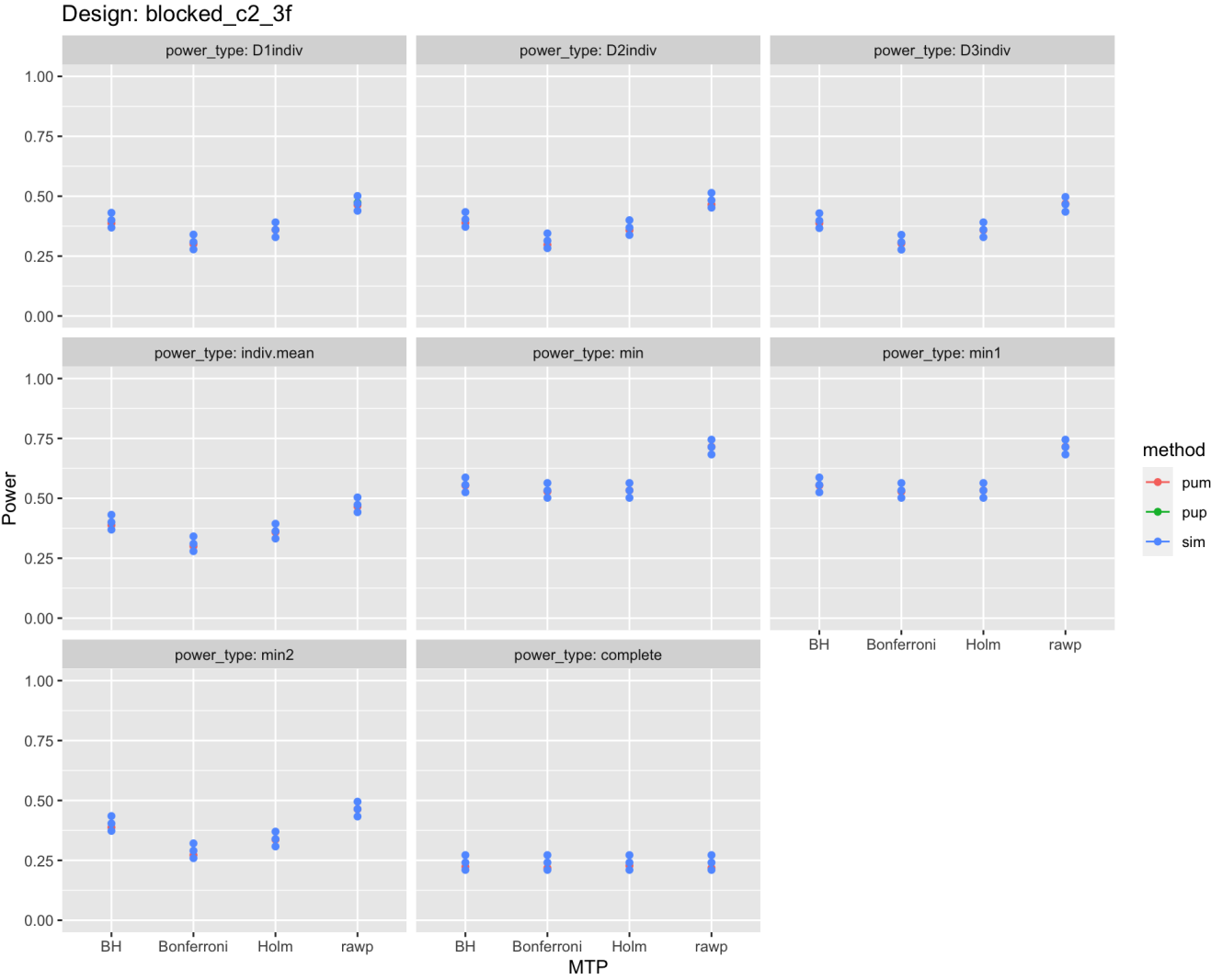


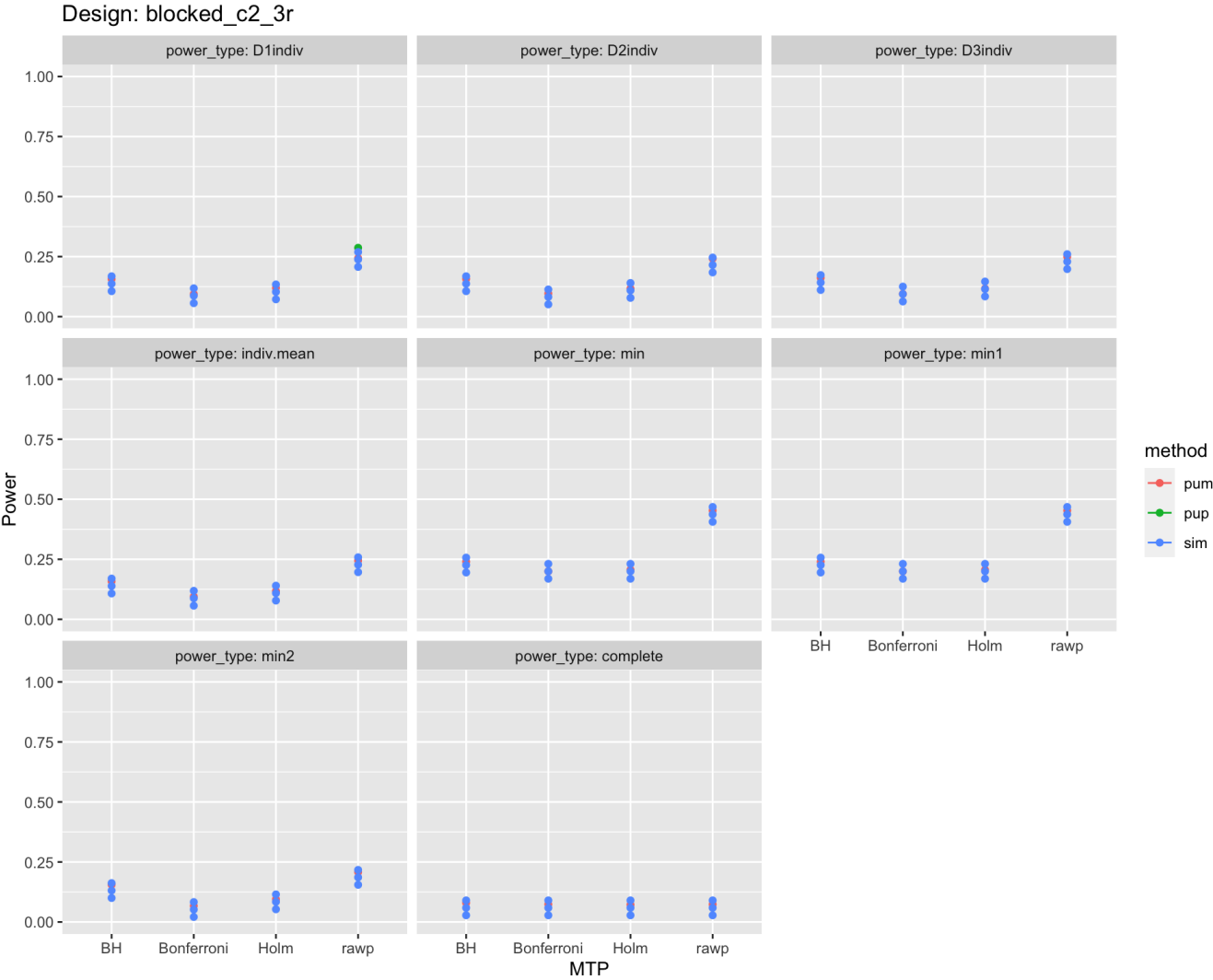


$R_3^2 = 0.6, 0.6, 0.6$



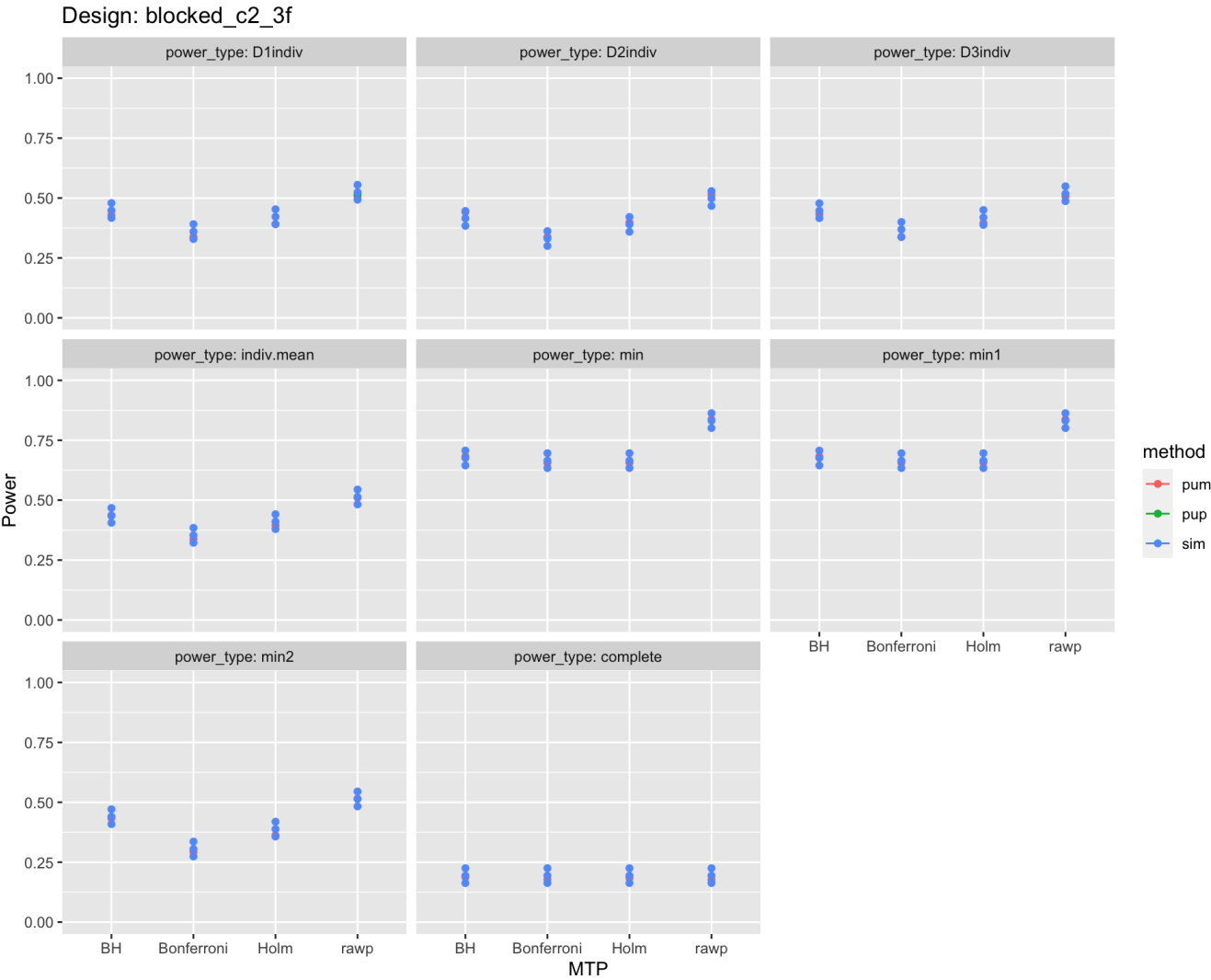
$R_1^2 = 0, 0, 0$ $R_2^2 = 0, 0, 0$ $R_3^2 = 0, 0, 0$



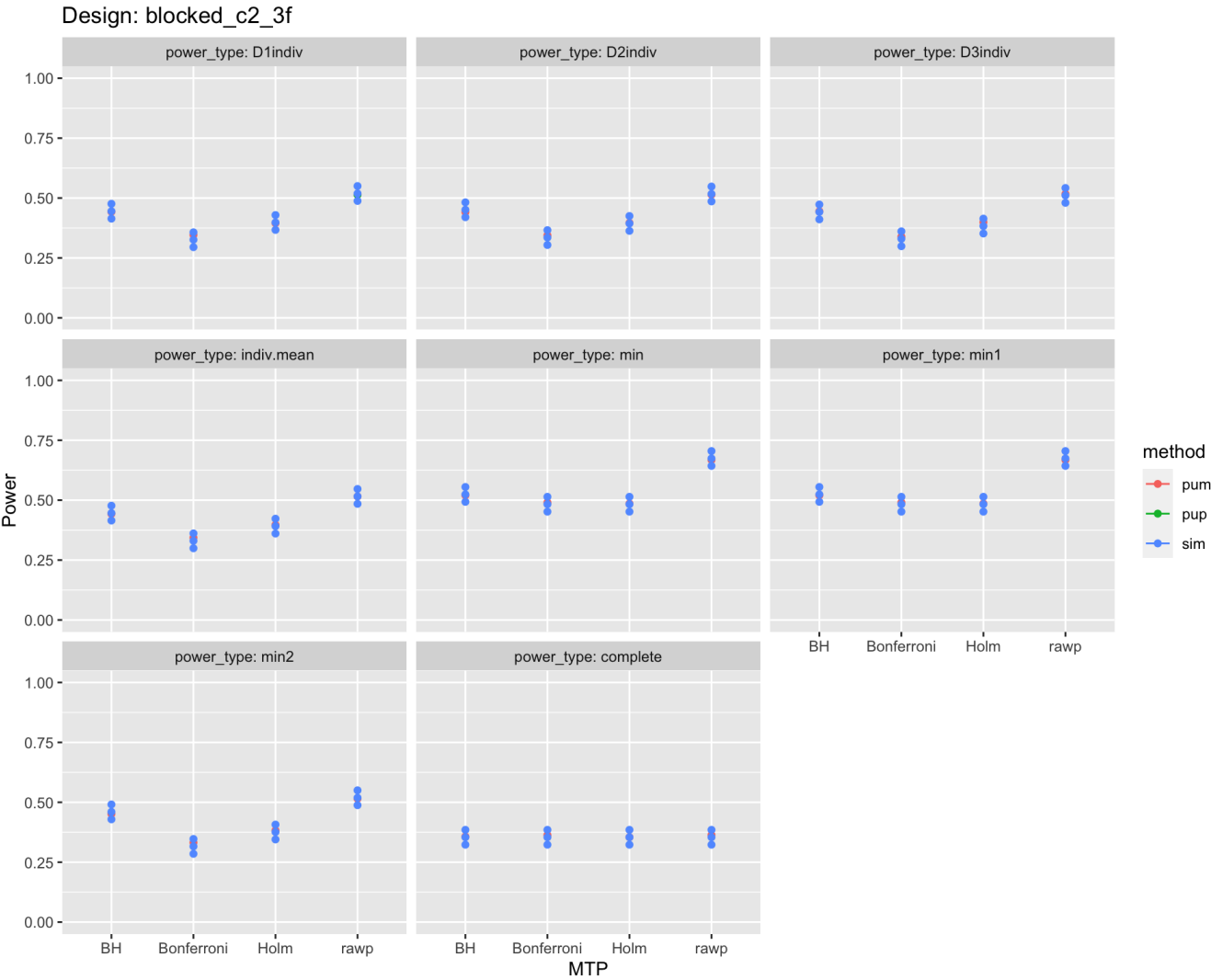


Varying rho

$\rho = 0.2$

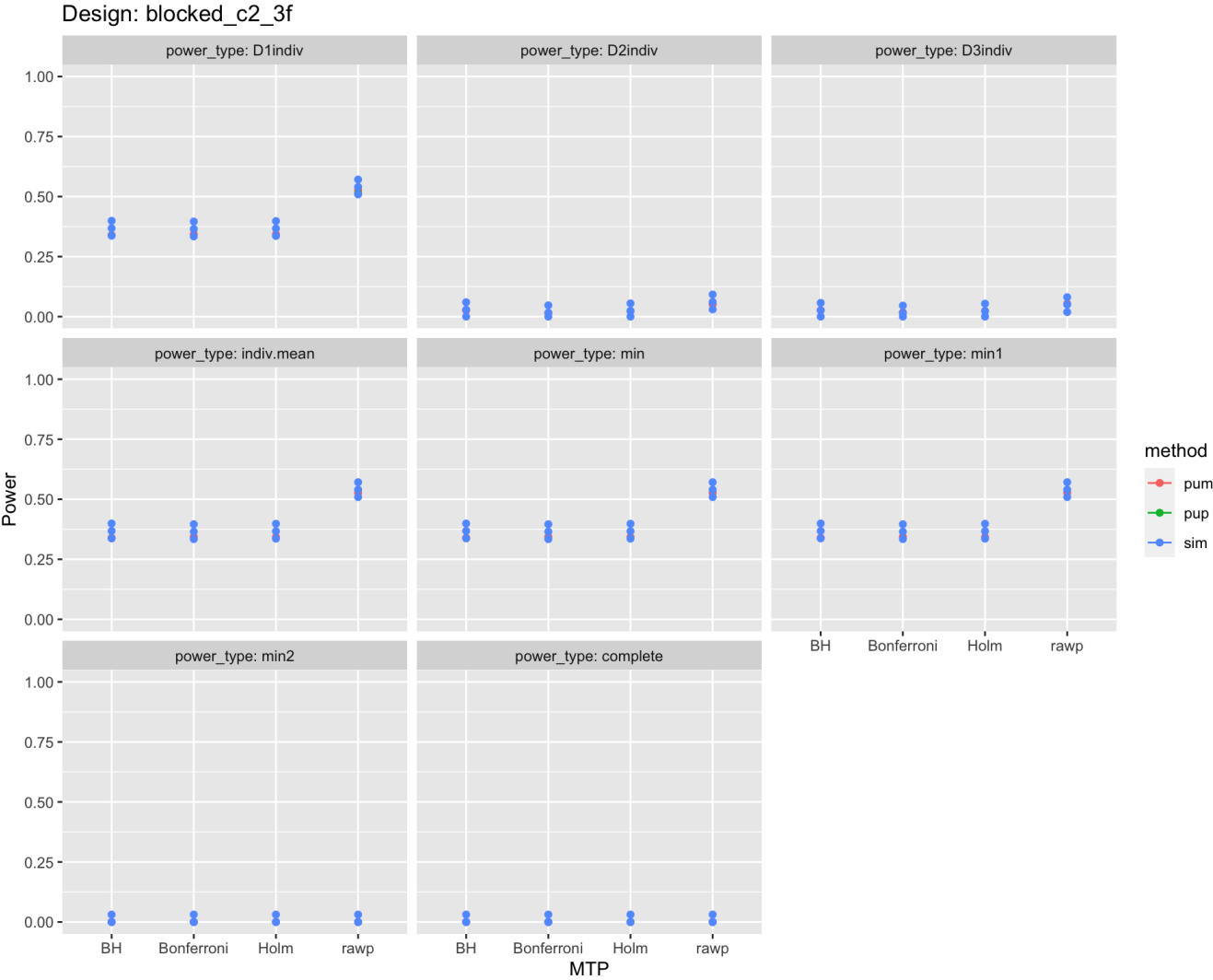


$\rho = 0.8$



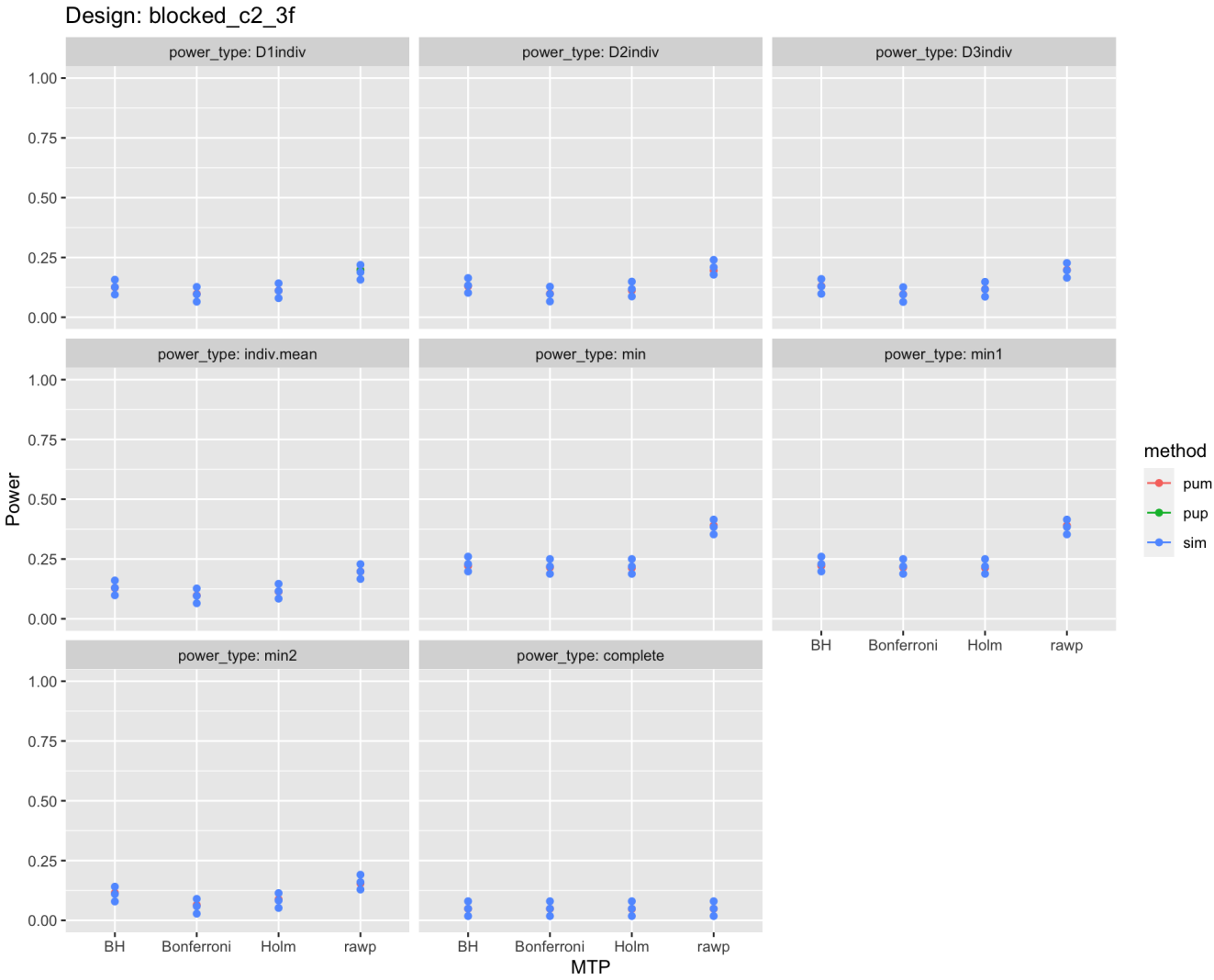
Varying true positives

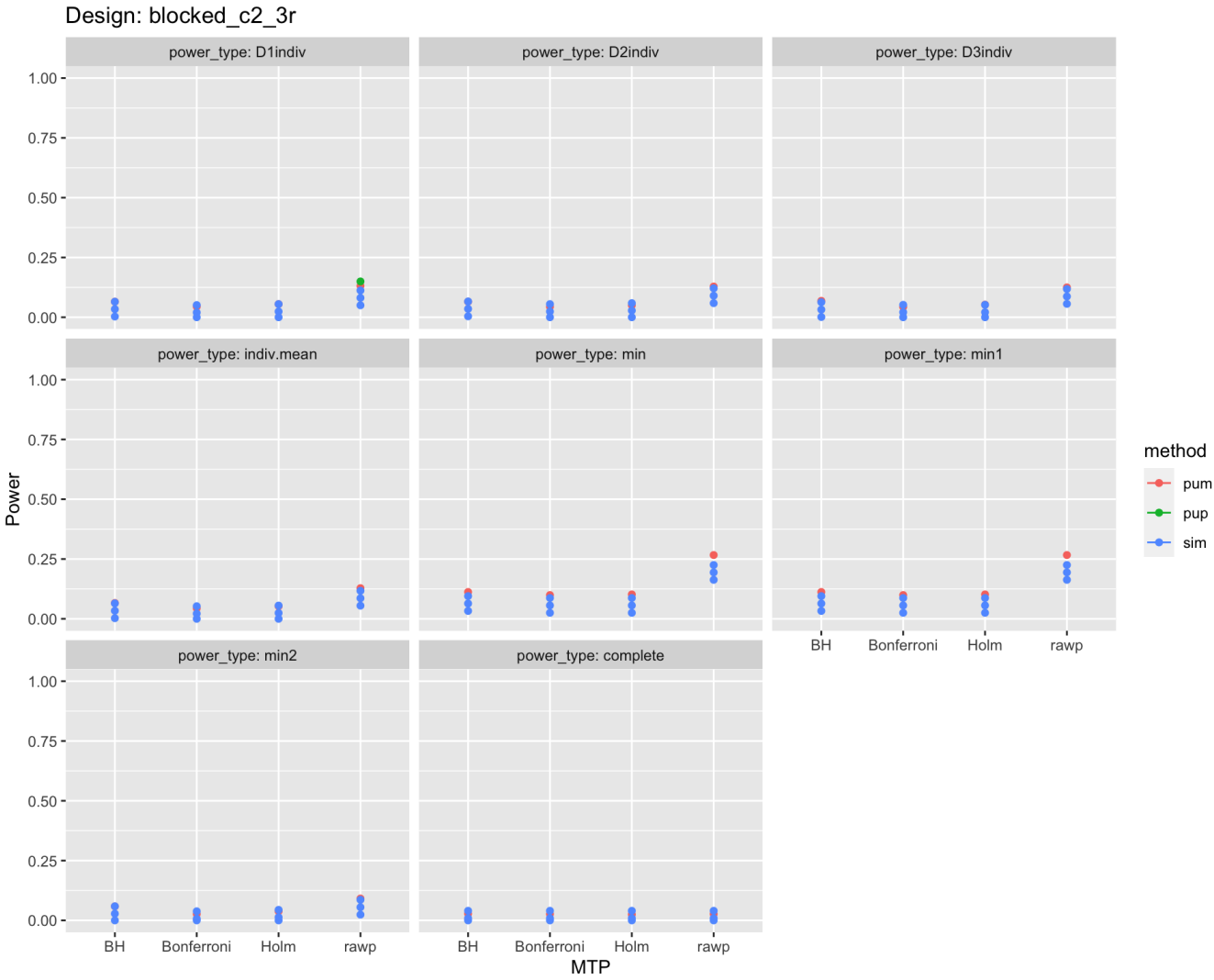
ATE (ES) = 0.125, 0, 0



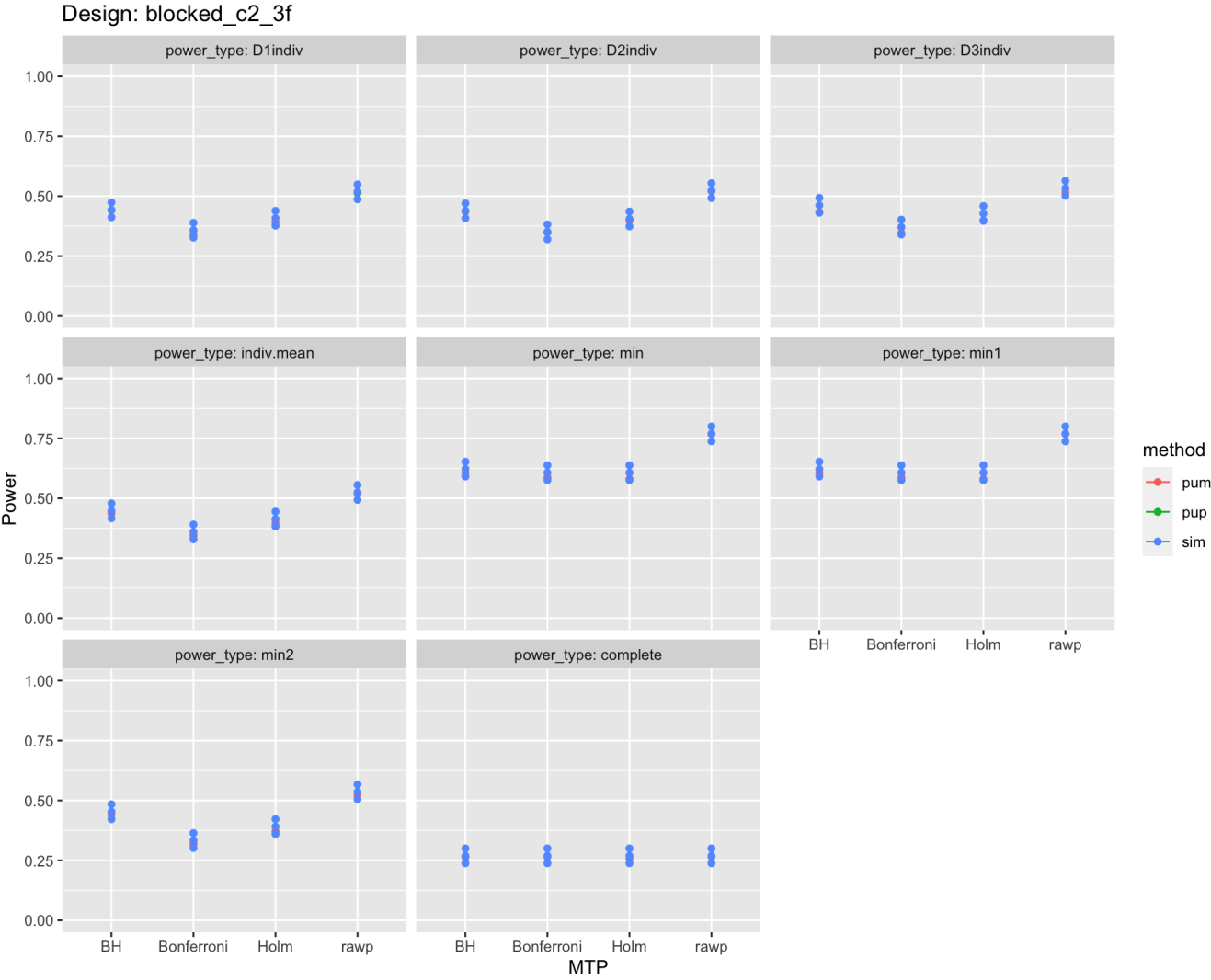
Varying ICC

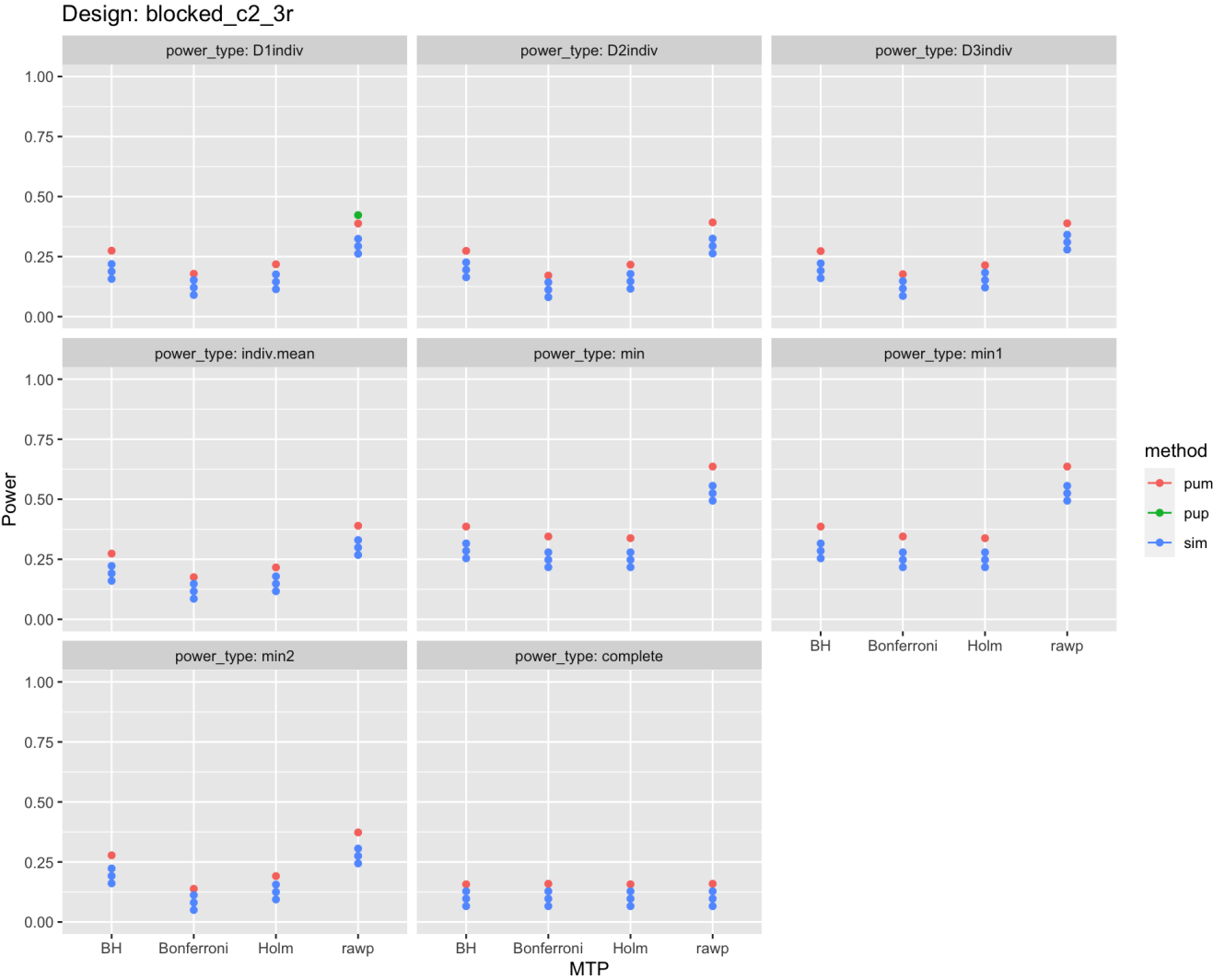
ICC₂ = 0.7, 0.7, 0.7





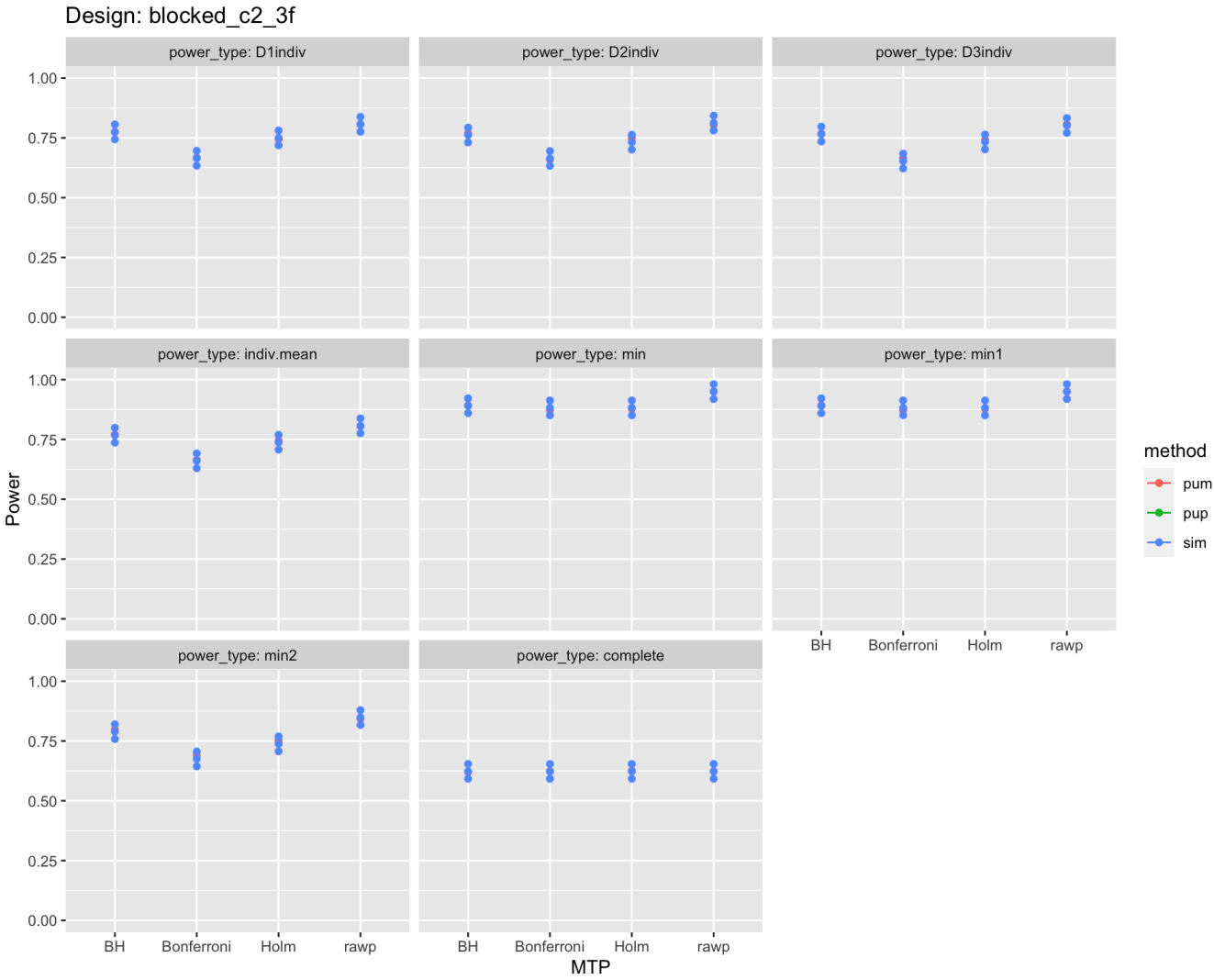
ICC₃ = 0.7, 0.7, 0.7

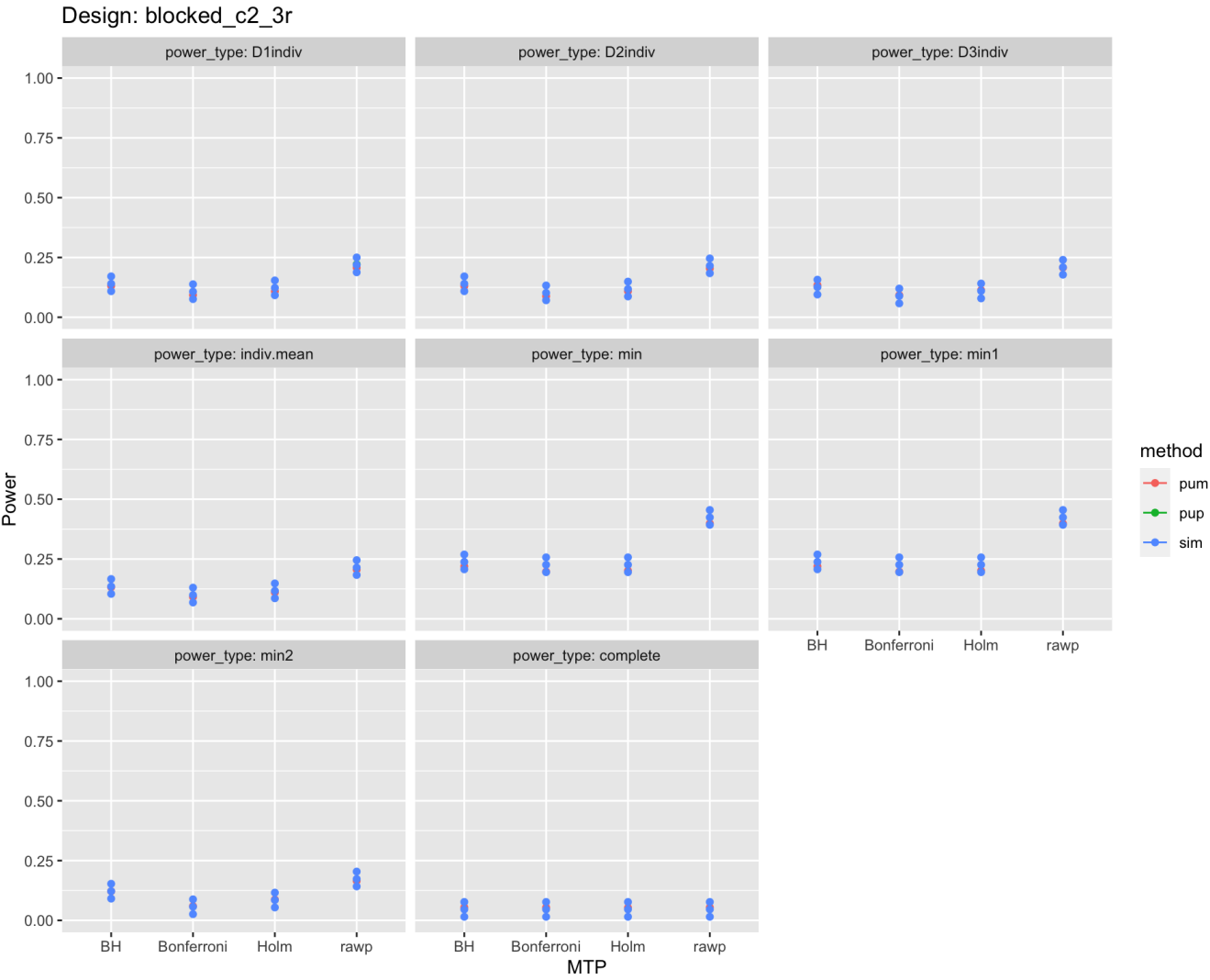




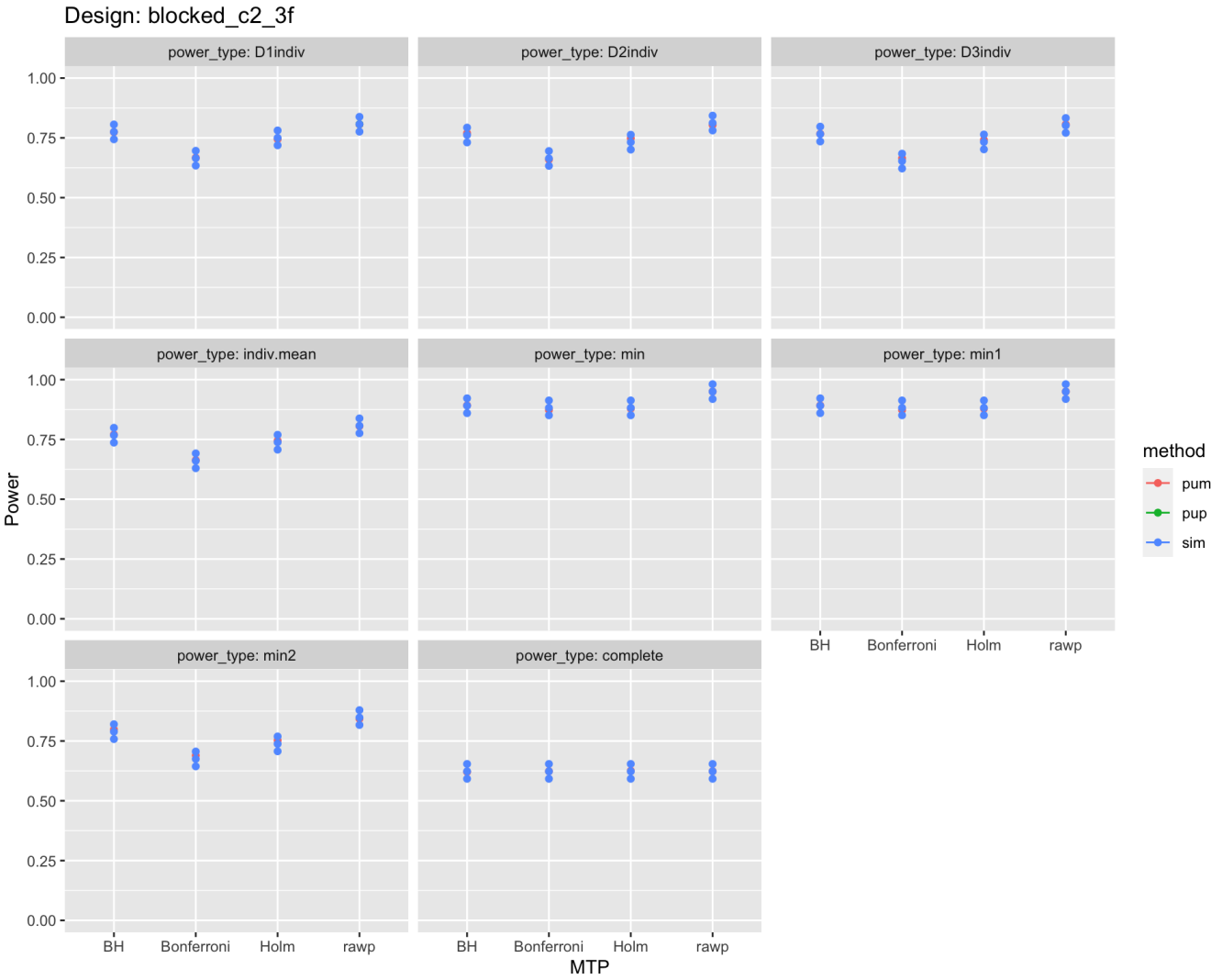
Varying Omega

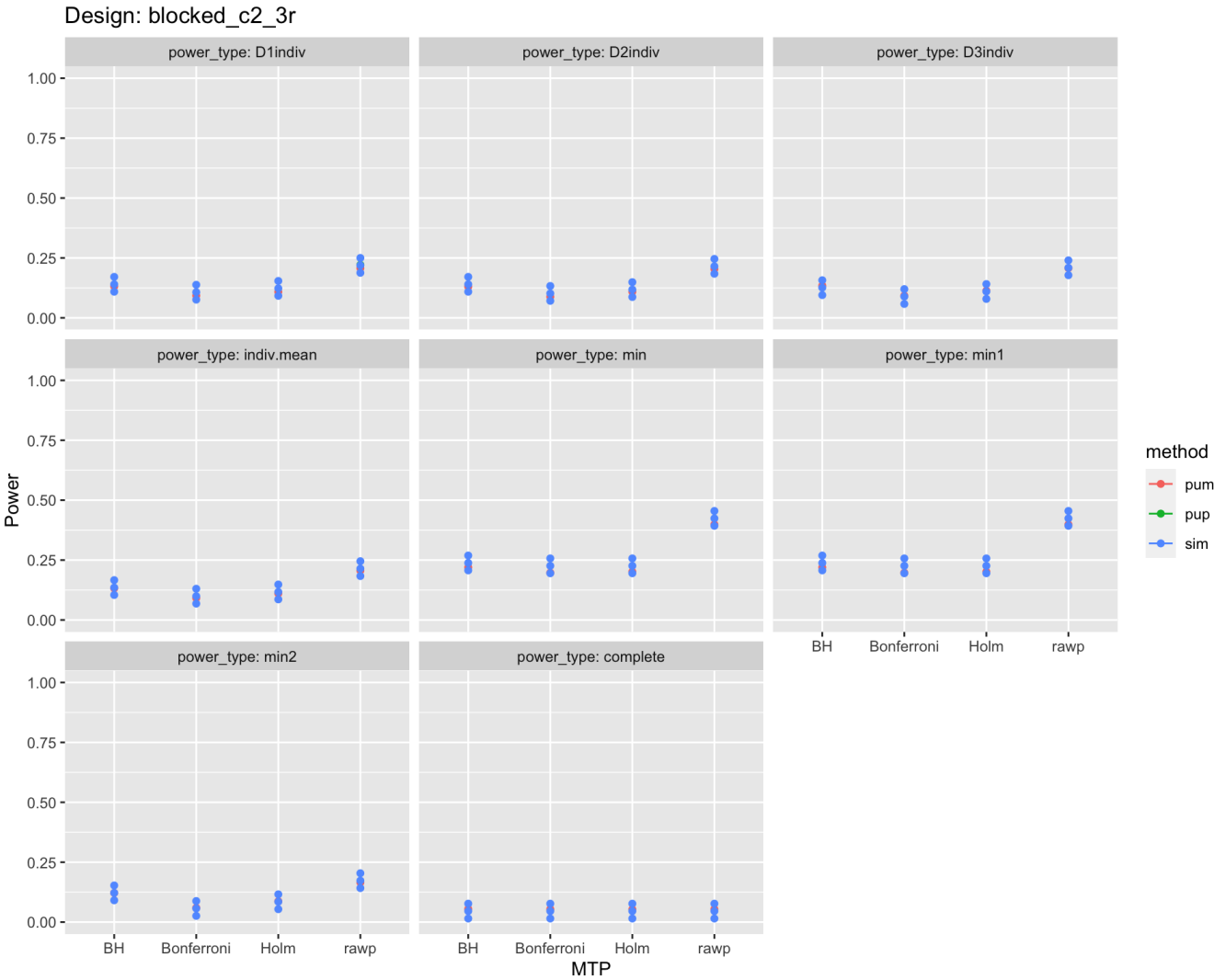
$\omega_3 = 0.8$





$\omega_3 = 0.8$ $ICC_3 = 0.7, 0.7, 0.7$





MDES validation

```
##
##
## +-----+-----+-----+-----+
## |      MTP      | Adjusted MDES | D1indiv Power | Target MDES |
## +=====+=====+=====+=====+
## |      rawp      |      0.126    |      0.517    |      0.125    |
## +-----+-----+-----+-----+
## | Bonferroni     |      0.125    |      0.34     |      0.125    |
## +-----+-----+-----+-----+
## |      BH        |      0.127    |      0.447    |      0.125    |
## +-----+-----+-----+-----+
## |      Holm      |      0.125    |      0.398    |      0.125    |
## +-----+-----+-----+-----+
##
##
## Table: blocked_c2_3f
```

```
##
##
## +-----+-----+-----+-----+
## |      MTP      | Adjusted MDES | Dlindiv Power | Target MDES |
## +=====+=====+=====+=====+
## |      rawp      |      0.125     |      0.261     |      0.125     |
## +-----+-----+-----+-----+
## | Bonferroni |      0.125     |      0.104     |      0.125     |
## +-----+-----+-----+-----+
## |      BH      |      0.124     |      0.161     |      0.125     |
## +-----+-----+-----+-----+
## |      Holm     |      0.129     |      0.139     |      0.125     |
## +-----+-----+-----+-----+
##
## Table: blocked_c2_3r
```

Sample size validation

```
##
##
## +-----+-----+-----+-----+-----+
## | MTP | Sample type | Sample size | Dlindiv power | Target sample size |
## +=====+=====+=====+=====+=====+
## | BH | K | 11 | 0.441 | 10 |
## +-----+-----+-----+-----+-----+
## | Holm | K | 10 | 0.386 | 10 |
## +-----+-----+-----+-----+-----+
##
## Table: blocked_c2_3f
```

```
##
##
## +-----+-----+-----+-----+-----+
## | MTP | Sample type | Sample size | Dlindiv power | Target sample size |
## +=====+=====+=====+=====+=====+
## | BH | K | 10 | 0.168 | 10 |
## +-----+-----+-----+-----+-----+
## | Holm | K | 10 | 0.125 | 10 |
## +-----+-----+-----+-----+-----+
##
## Table: blocked_c2_3r
```