

Models of disease transmission

- “Classical” assumption
- Simple dose-response model

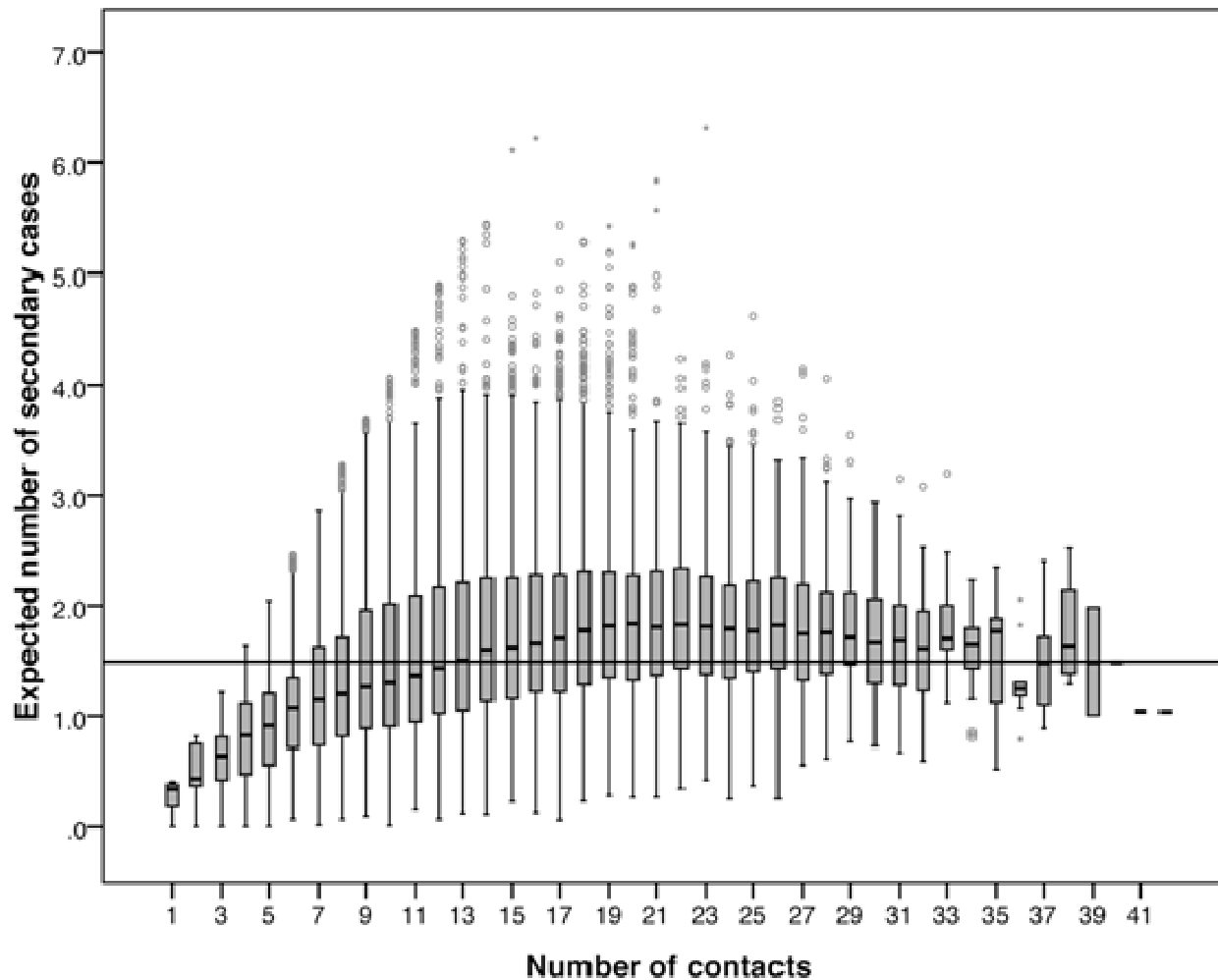
$$P_{nm} = \text{const.}$$

P_{nm} Infection probability of susceptible n
caused by infected m

$$P_n = 1 - \exp\left(-\sum_{m=1}^I q_m t_{nm}\right)$$

P_n Infection probability of susceptible n
 q_m Shedding rate of infector m
 t_{nm} Contact time of n and m

Secondary cases vs. number of contacts



Conclusions

- Focus not only on the number of contacts, but also on the quality of contacts (e.g. duration).
- Importance of highly connected individuals (“super-spreaders”) might be overestimated in classical models.
- Further research needed:
 - Is contact time a good proxy for exposure?
 - Which other factors are important to be measured for the quality of a contact?