

# **Evaluating Single-Level and Hierarchical Maximum Likelihood Estimation in Shifted-Wald Models**



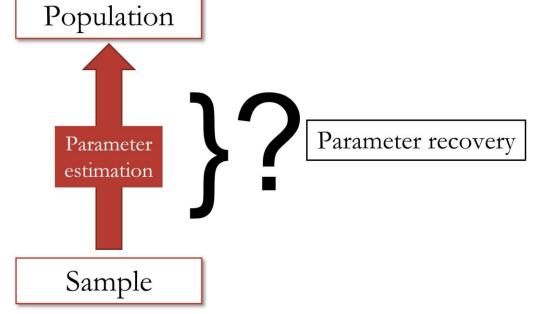
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### **Parameter Recovery**

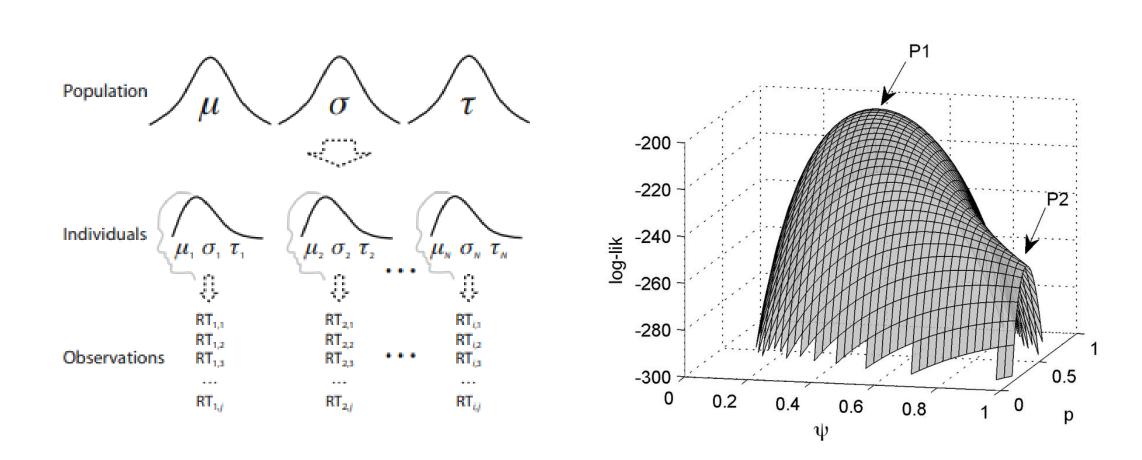
Response times are a crucial component of understanding and measuring cognitive processes. By fitting a model to a distribution of observed response times, we can estimate population parameters that could have potentially generated the observed sample. But, how do we know that the estimated parameters are representative of the true population parameters? To assess the validity of our estimation techniques, we must conduct a parameter recovery study.



## **Estimation Methods**

This study focused on two methods for estimating parameters:

- Classical (single-level) maximum likelihood estimation (CMLE)
- Hierarchical maximum likelihood estimation (HMLE)

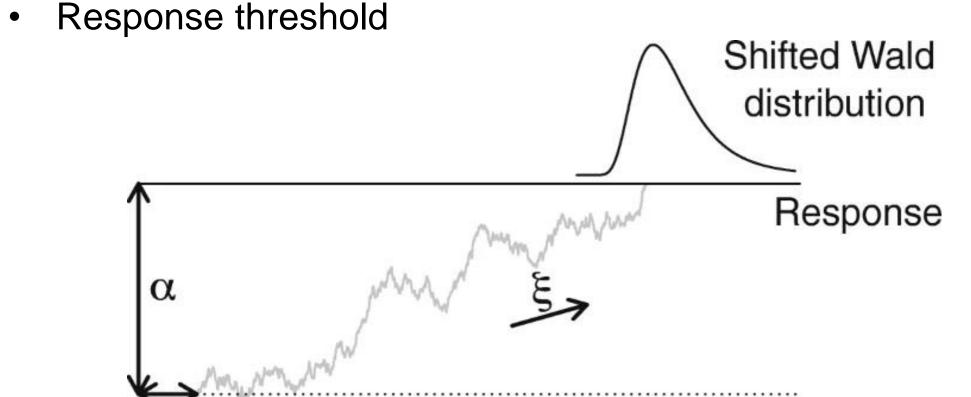


#### **Shifted-Wald Models**

For this study we used a model of response times known as the shifted-Wald model, which is composed of three parameters:

Time →

- Shift
- Drift rate



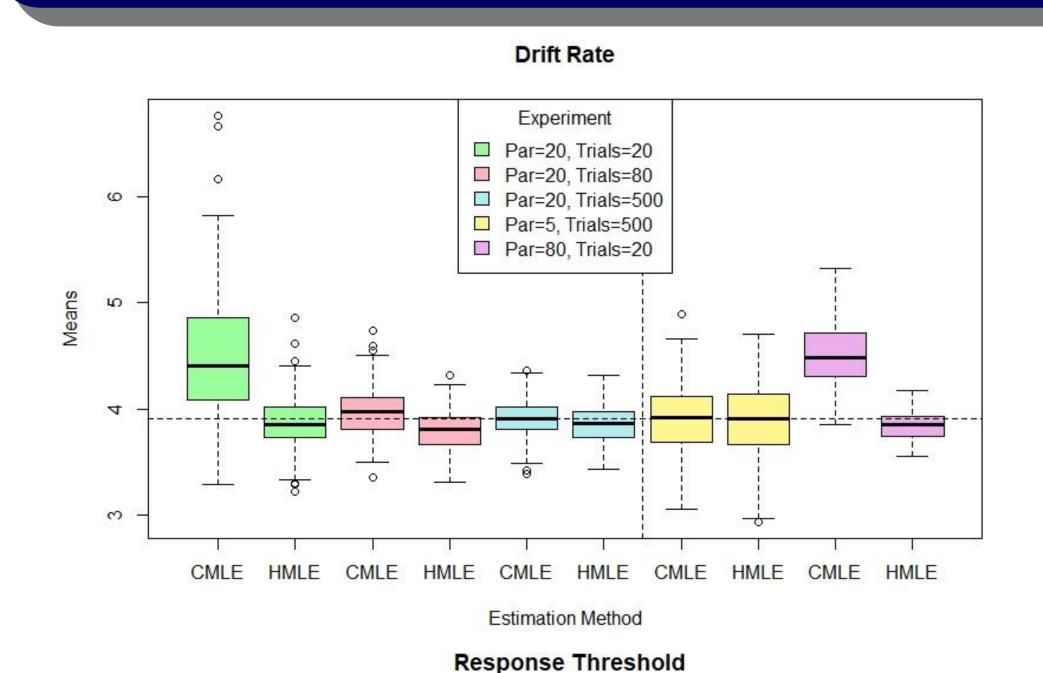
#### Methods

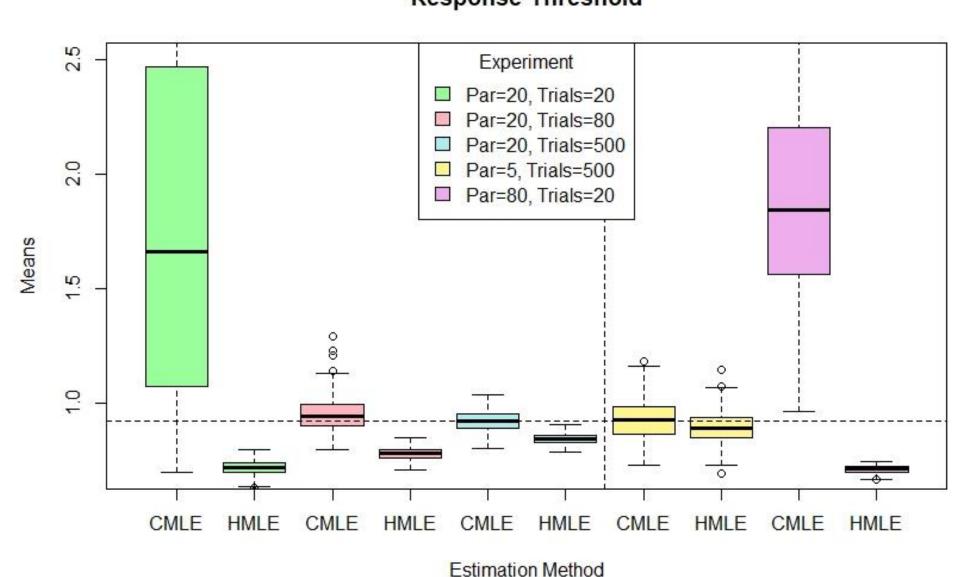
For this study, we simulated data using shifted-Wald parameter targets reported by Faulkenberry et al. (2018). This was composed of four main steps:

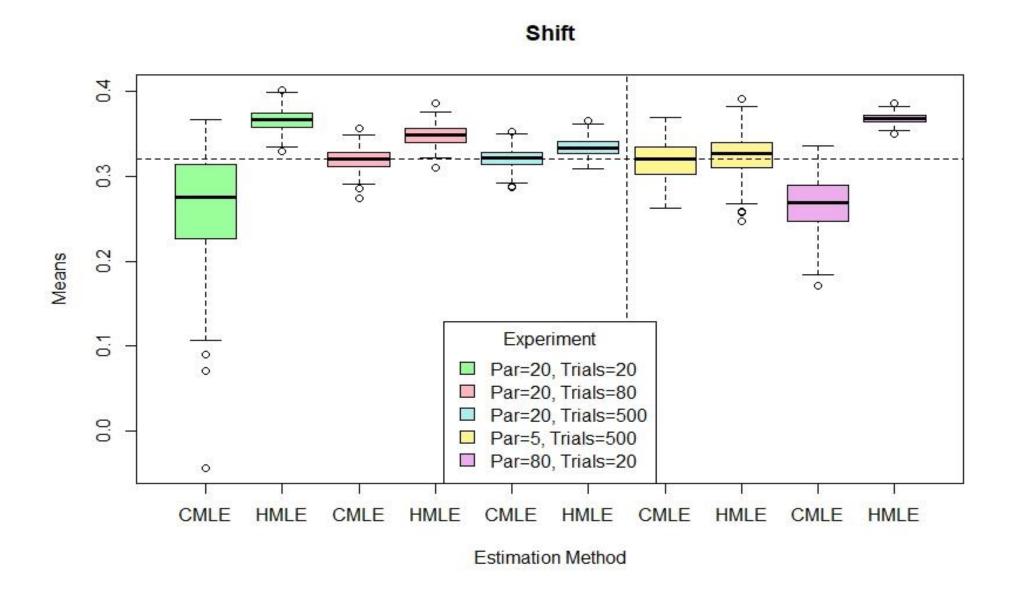
- Generate 'artificial' people from parent population
- Generate response times for artificial people
- Fit shifted-Wald model to RT distribution with CMLE and HMLE
- Compare estimates to original target parameters

This process was applied to 5 sub-experiments in a design used by Farrell and Ludwig (2008). These sub-experiments had 5, 20, or 80 participants, with 20, 80, or 500 trials per participant.

## **Results: Parameters**







#### **Results: Error and Bias**

Root Mean Square Error Root Mean Square Error in Seconds						
	γ	α	θ			
Par = 20, Trials = 20						
HMLE Group Level	0.198	0.202	0.046			
HMLE Individual Level	0.673	0.226	0.046			
CMLE Individual Level	2.419	3.474	0.236			
Par = 20, Trials = 80						
HMLE Group Level	0.175	0.139	0.028			
HMLE Individual Level	0.401	0.157	0.029			
CMLE Individual Level	0.657	0.296	0.036			
Par = 20, Trials = 500						
HMLE Group Level	0.144	0.074	0.014			
HMLE Individual Level	0.187	0.101	0.017			
CMLE Individual Level	0.236	0.095	0.012			
Par = 5, Trials = 500						
HMLE Group Level	0.275	0.060	0.019			
HMLE Individual Level	0.364	0.115	0.016			
CMLE Individual Level	0.231	0.092	0.012			
Par = 80, Trials = 20						
HMLE Group Level	0.107	0.209	0.048			

CMLE Individual Level 2.647 4.171

Note. CMLE = classical maximum likelihood estimation;

HMLE Individual Level

HMLE = hierarichal maximum likelihood estimation

#### Mean Bias n Bias (B) and Standard Doviation(SD) in Seconds

0.232

0.048

0.268

Mean Bias (B) and Standard Deviation(SD) in Seconds								
γ		α		θ				
В	SD	В	SD	В	SD			
-0.045	0.251	-0.202	0.032	0.046	0.013			
-0.050	0.203	-0.200	0.042	0.046	0.007			
0.578	0.581	0.961	0.975	-0.053	0.062			
-0.111	0.188	-0.139	0.026	0.028	0.012			
-0.100	0.114	-0.137	0.032	0.028	0.005			
0.064	0.148	0.034	0.073	-0.001	0.008			
-0.061	0.171	-0.074	0.026	0.013	0.010			
-0.062	0.053	-0.078	0.020	0.014	0.004			
0.008	0.052	0.003	0.021	0.000	0.003			
-0.014	0.342	-0.023	0.070	0.005	0.023			
-0.008	0.149	-0.033	0.054	0.004	0.009			
0.017	0.102	0.006	0.042	0.000	0.006			
-0.066	0.117	-0.209	0.017	0.048	0.006			
-0.075	0.100	-0.211	0.021	0.048	0.003			
0.615	0.308	0.980	0.476	-0.051	0.036			
	B -0.045 -0.050 0.578 -0.111 -0.100 0.064 -0.061 -0.062 0.008 -0.014 -0.008 0.017 -0.066 -0.075 0.615	B       SD         -0.045       0.251         -0.050       0.203         0.578       0.581         -0.111       0.188         -0.100       0.114         0.064       0.148         -0.062       0.053         0.008       0.052         -0.014       0.342         -0.008       0.149         0.017       0.102         -0.066       0.117         -0.075       0.100         0.615       0.308	B         SD         B           -0.045         0.251         -0.202           -0.050         0.203         -0.200           0.578         0.581         0.961           -0.111         0.188         -0.139           -0.100         0.114         -0.137           0.064         0.148         0.034           -0.061         0.171         -0.074           -0.062         0.053         -0.078           0.008         0.052         0.003           -0.014         0.342         -0.023           -0.008         0.149         -0.033           0.017         0.102         0.006           -0.066         0.117         -0.209           -0.075         0.100         -0.211           0.615         0.308         0.980	B         SD         B         SD           -0.045         0.251         -0.202         0.032           -0.050         0.203         -0.200         0.042           0.578         0.581         0.961         0.975           -0.111         0.188         -0.139         0.026           -0.100         0.114         -0.137         0.032           0.064         0.148         0.034         0.073           -0.061         0.171         -0.074         0.026           -0.062         0.053         -0.078         0.020           0.008         0.052         0.003         0.021           -0.014         0.342         -0.023         0.070           -0.008         0.149         -0.033         0.054           0.017         0.102         0.006         0.042           -0.066         0.117         -0.209         0.017           -0.075         0.100         -0.211         0.021           0.615         0.308         0.980         0.476	B         SD         B         SD         B           -0.045         0.251         -0.202         0.032         0.046           -0.050         0.203         -0.200         0.042         0.046           0.578         0.581         0.961         0.975         -0.053           -0.111         0.188         -0.139         0.026         0.028           -0.100         0.114         -0.137         0.032         0.028           0.064         0.148         0.034         0.073         -0.001           -0.061         0.171         -0.074         0.026         0.013           -0.062         0.053         -0.078         0.020         0.014           0.008         0.052         0.003         0.021         0.000           -0.014         0.342         -0.023         0.070         0.005           -0.008         0.149         -0.033         0.054         0.004           0.017         0.102         0.006         0.042         0.000           -0.066         0.117         -0.209         0.017         0.048           -0.075         0.100         -0.211         0.021         0.048			

likelihood estimation

likelihood estimation

- There is a trade-off in estimation parameters
- If a study is constrained to a smaller trial size, utilizing HMLE could ensure more accurate estimation

## Discussion

There are still many questions left to be answered:

- How would an alternative approach, (i.e., Bayesian) change our results?
- Is there an optimum number of participants or trials?
- How can we utilize these results to inform participant/ trial size decisions?



For references, R script, and more, please see QR code.