

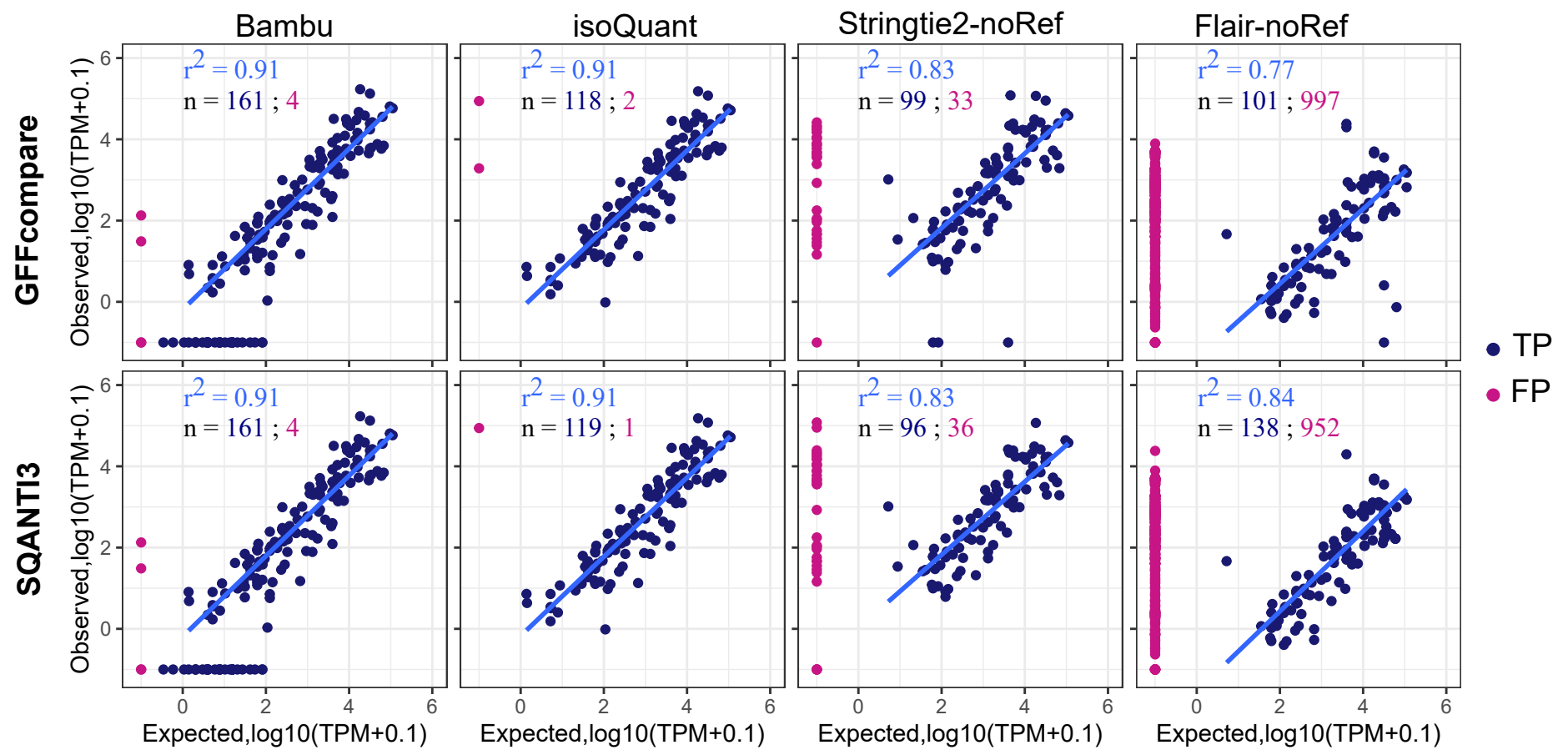
Methodology

Assembly  
Original .fastq reads

Alignment  
Minimap2

Quantification  
Salmon

LSK114



RNA004

