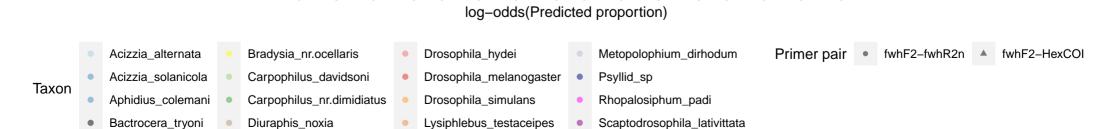
## Bias model fits complm lm metacal prop 0 DNA -4 -8 -12 log-odds(Observed proportion) DNEasy PCR -12 **-**0 QuickExtract -4 -8 -12 **-**-10 5 –10 5 –10 -5



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5

0

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