

## Measurement of pRSET-iGluSnFR DNA Concentration

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**Protocol:**

1. Cleaned NanoDrop spectrophotometer with DI water
2. Blanked NanoDrop with 2  $\mu\text{L}$  Elution Buffer
3. Cleaned NanoDrop before loading 2  $\mu\text{L}$  pRA
4. Repeated step 3 for all pRSET-iGluSnFR samples

**Results:**

Sample	DNA Concentration	Volume	Mass
pRA	148.6 ng/ $\mu\text{L}$	24 $\mu\text{L}$	3566.4 ng
pRB	139.8 ng/ $\mu\text{L}$	25 $\mu\text{L}$	3495 ng
pRC	158.7 ng/ $\mu\text{L}$	26.3 $\mu\text{L}$	4173.81 ng
pRD	164.4 ng/ $\mu\text{L}$	23.4 $\mu\text{L}$	3846.96 ng
pRE	173.1 ng/ $\mu\text{L}$	24.2 $\mu\text{L}$	4189.02 ng
pRF	113.2 ng/ $\mu\text{L}$	24.6 $\mu\text{L}$	2784.72 ng
pRG	139.5 ng/ $\mu\text{L}$	25.2 $\mu\text{L}$	3515.4 ng
pRH	140.6 ng/ $\mu\text{L}$	26 $\mu\text{L}$	3655.6 ng

**Conclusion:** We will select and digest one of these sample to generate template DNA for the insertion of the genes FadB (G4.1), FadA (G4.2), and yafH (G4.3).