Genotype	Dilution	Tech_rep	Count	True_count
WT	1000	00	37	37000
WT	1000	01	38	38000
WT	1000	02	43	43000
GFP	1000	00	33	33000
GFP	1000	01	40	40000
GFP	1000	02	46	46000

nanocount()
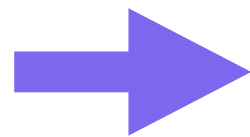
Genotype	Particle_N	Particle_count
WT	3	119000
GFP	3	118000

```
df %>%
    nanocount(Genotype,
             name = "Particle",
      param_var = True_count)
```