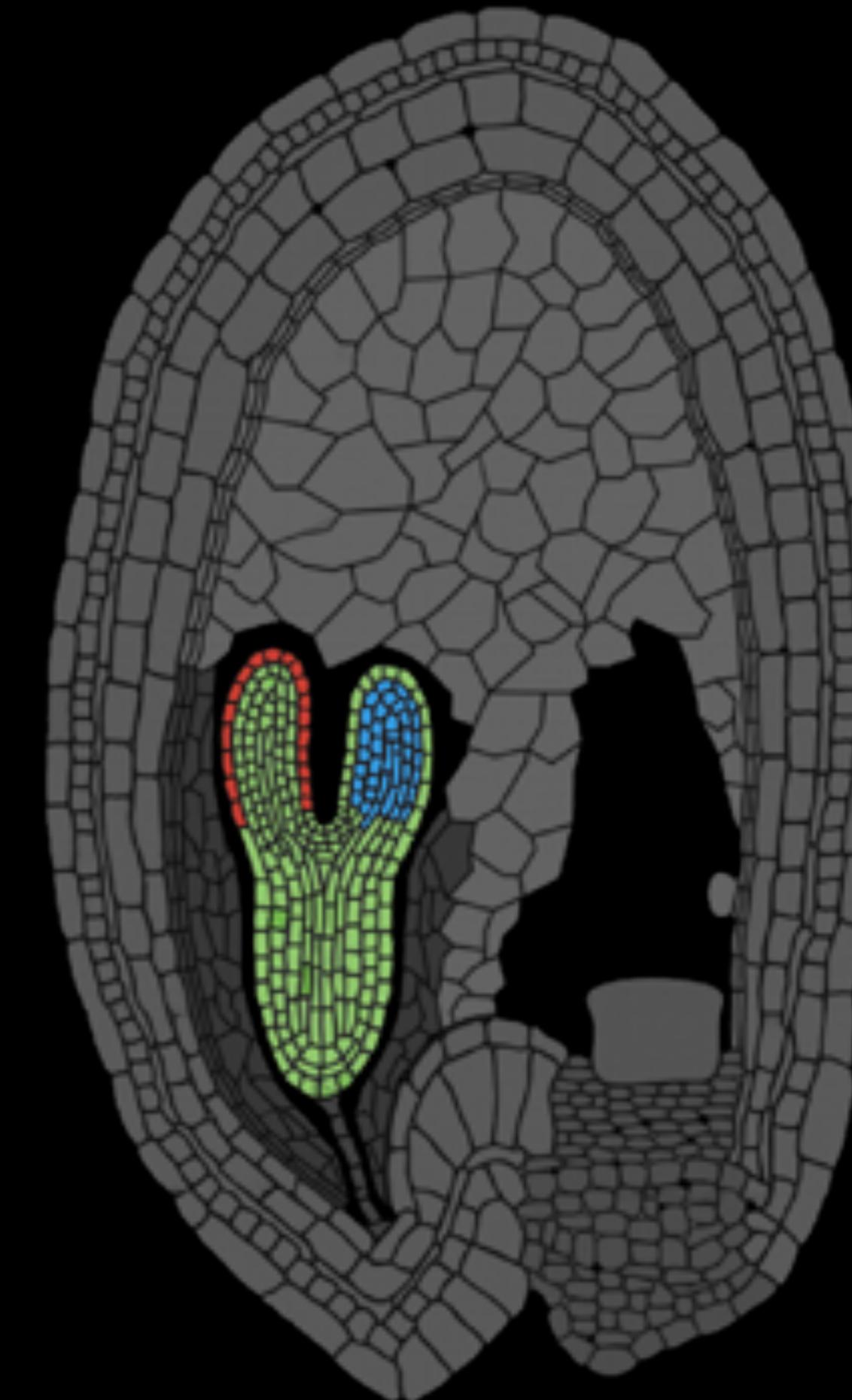


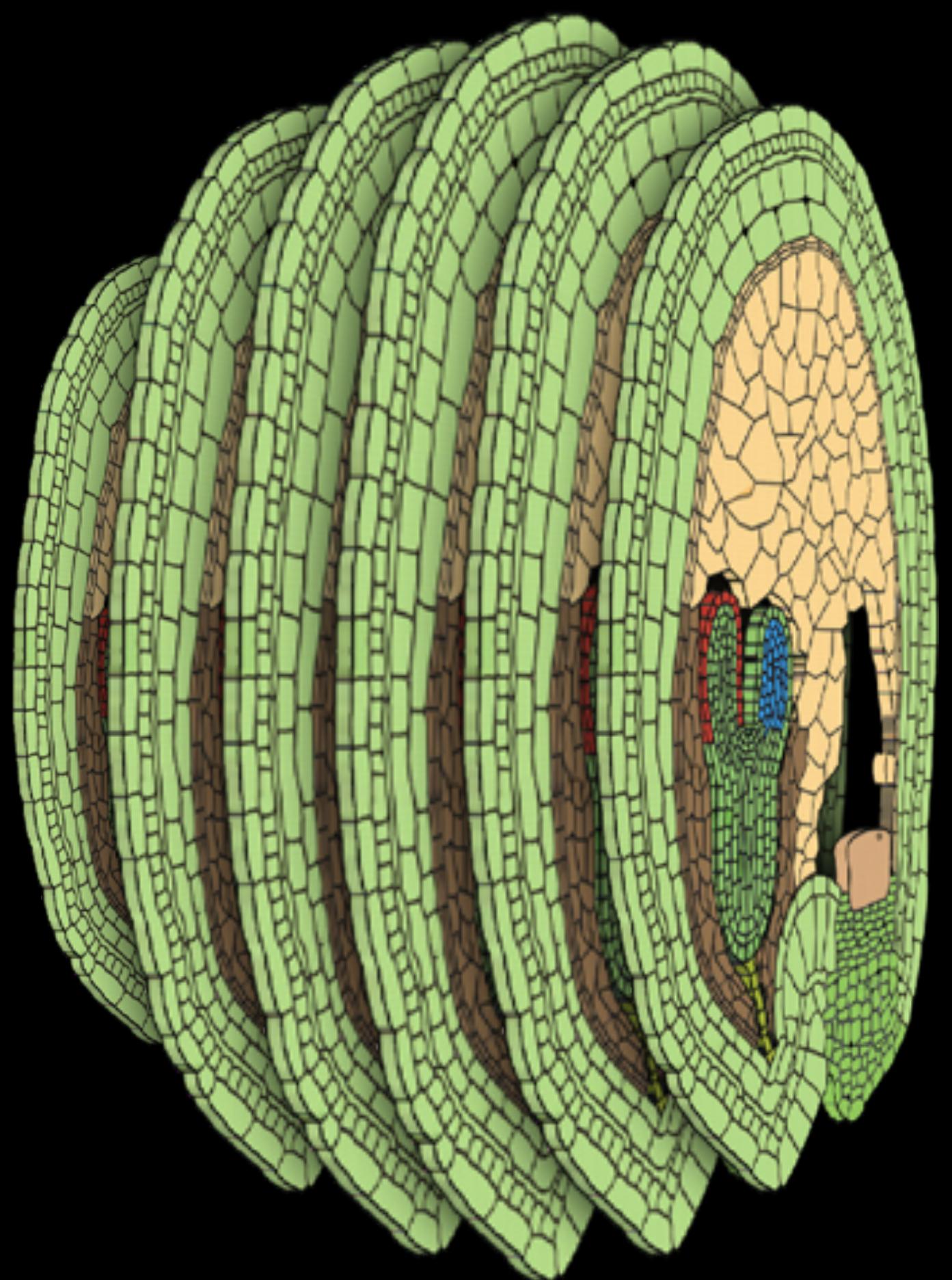
0.25mm

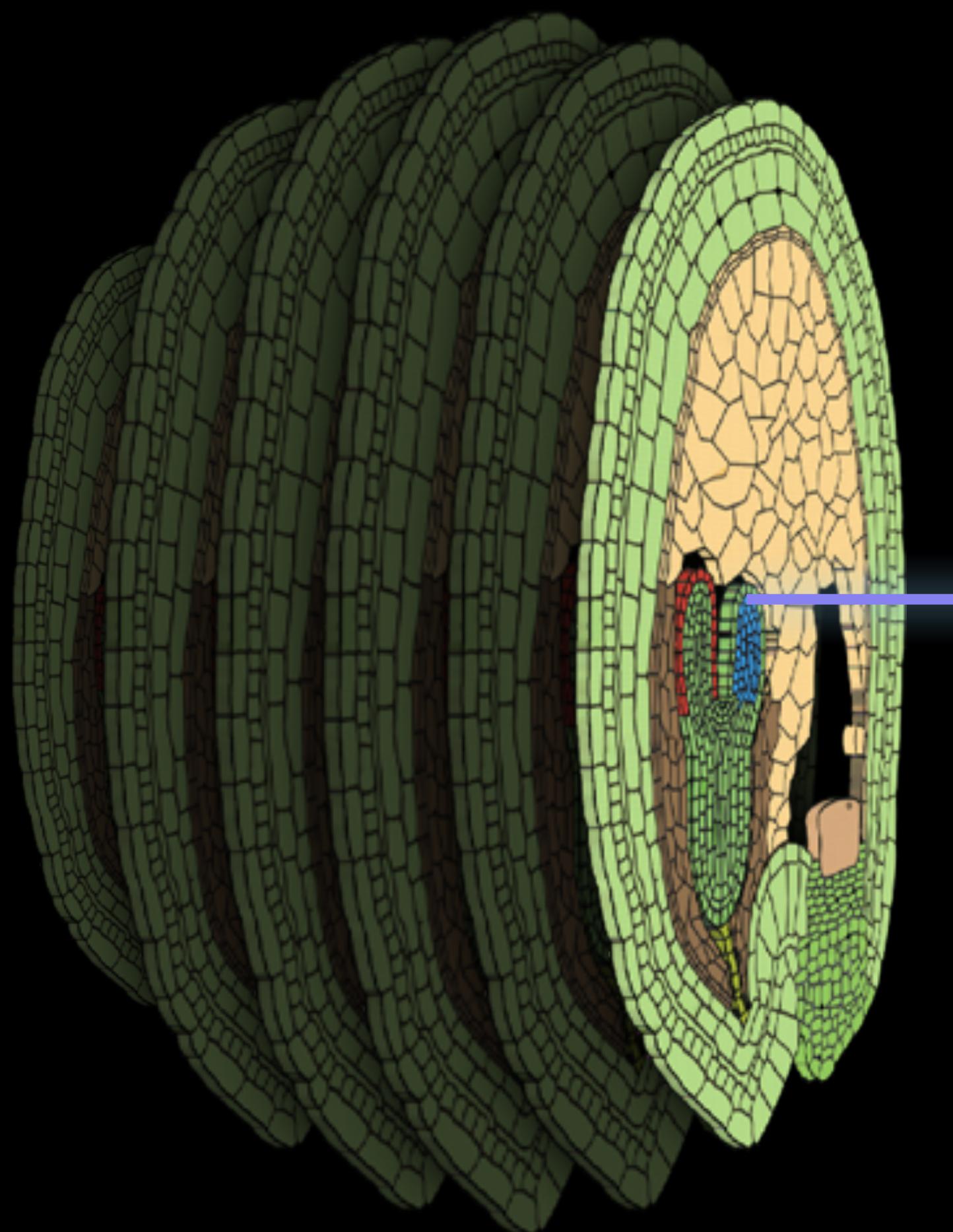
**Epidermis**



**Mesophyll**

**0.25mm**







**RNA (ng)**

**5**

**0.10**

**0.75**

**0.05**

**0.25**

**0.01**

---

**Library  
size ( $10^6$ )**

53

38

25

32

33

30

**Genes  
detected**

17,400

19,300

19,500

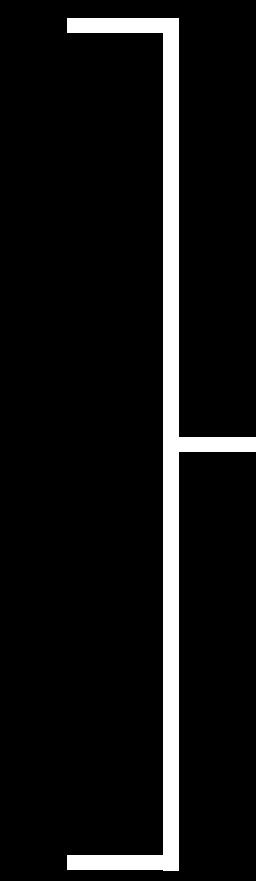
18,500

16,800

15,900

*"As low as 0.05ng of total RNA can be used ... to perform  
a comprehensive analysis of the expressed genome ...  
without compromising the number of genes detected."*

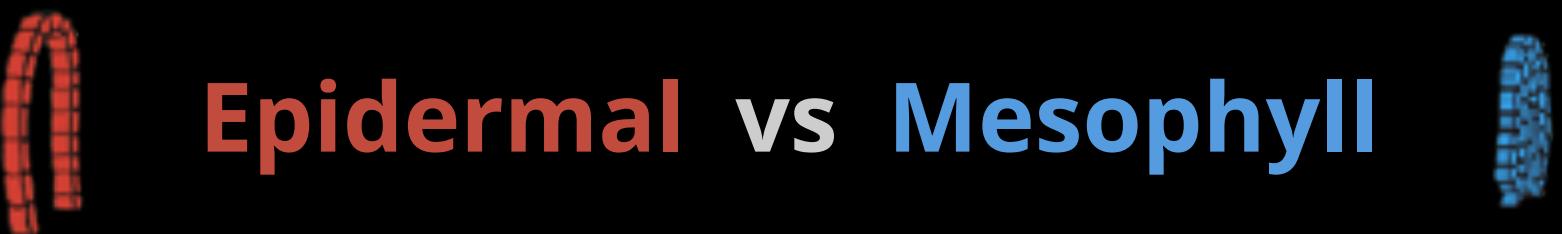
**0.10ng = 25x**



**4 biological replicates each  
10 min collection time each**

**0.10ng = 25x**





## Epidermal vs Mesophyll

CPM	No Significant Difference	Up	Down	Total DEGs
0.3	17817	201	111	312
1.0	16943	144	94	238

*“Of our genes, 571 had a log ratio > 1”*

Gene	Published logFC	My logFC	My adj. p-value
<b>MYB56</b>	>10	8.76	<b>0.019</b>
<b>NAC2</b>	9.70	8.87	<b>0.029</b>
<b>WOX8</b>	9.09	7.43	<b>0.126</b>
<b>AT4g31680</b>	6.50	6.60	<b>0.195</b>
<b>ULT2</b>	5.23	6.97	<b>0.111</b>
<b>bHLH</b>	5.08	5.95	<b>0.315</b>
<b>MYB118</b>	5.03	4.61	<b>0.304</b>