# Title yannow

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### 1 Question 1

(A) a

```
print("hi")
## [1] "hi"
```

#### Cell state color key

#### early2cell 1. 2. mid2cell 3. late2cell 4. 4cell 5. 8cell 6. 16cell earlyblast 7. midblast 8. 9. lateblast

#### Cell state color key

1.	early2cell	
2.	mid2cell	
3.	late2cell	
4.	4cell	
5.	8cell	
6.	16cell	
7.	earlyblast	
8.	midblast	
9.	lateblast	

Supplemental Figure 1: I got twooooo figures.

## 2 Question 2

```
(A) print("bye")

## [1] "bye"
```

(B)

AML [1] GPLVM [2]

### References

- [1] M Ryan Corces, Jason D Buenrostro, Beijing Wu, Peyton G Greenside, Steven M Chan, Julie L Koenig, Michael P Snyder, Jonathan K Pritchard, Anshul Kundaje, William J Greenleaf, et al. Lineage-specific and single-cell chromatin accessibility charts human hematopoiesis and leukemia evolution. *Nature genetics*, 2016.
- [2] Neil D Lawrence. Gaussian process latent variable models for visualisation of high dimensional data. In *Advances in neural information processing systems*, pages 329–336, 2004.