

	cell_type	culture_cond	Size_Factor
cell_1			
cell_2			
cell_3			
cell_n			

colData(cds)

rowData(cds)

	biotype	gc_content
gene_1		
gene_2		
gene_3		
gene_n		

Console

Terminal x

Jobs x

~/Documents/agogate/monocle3/ ↗



> |

Console

Terminal ×

Jobs ×



~/Documents/agogate/monocle3/ 



> |

Console

Terminal ×

Jobs ×



~/Documents/agogate/monocle3/ ↗



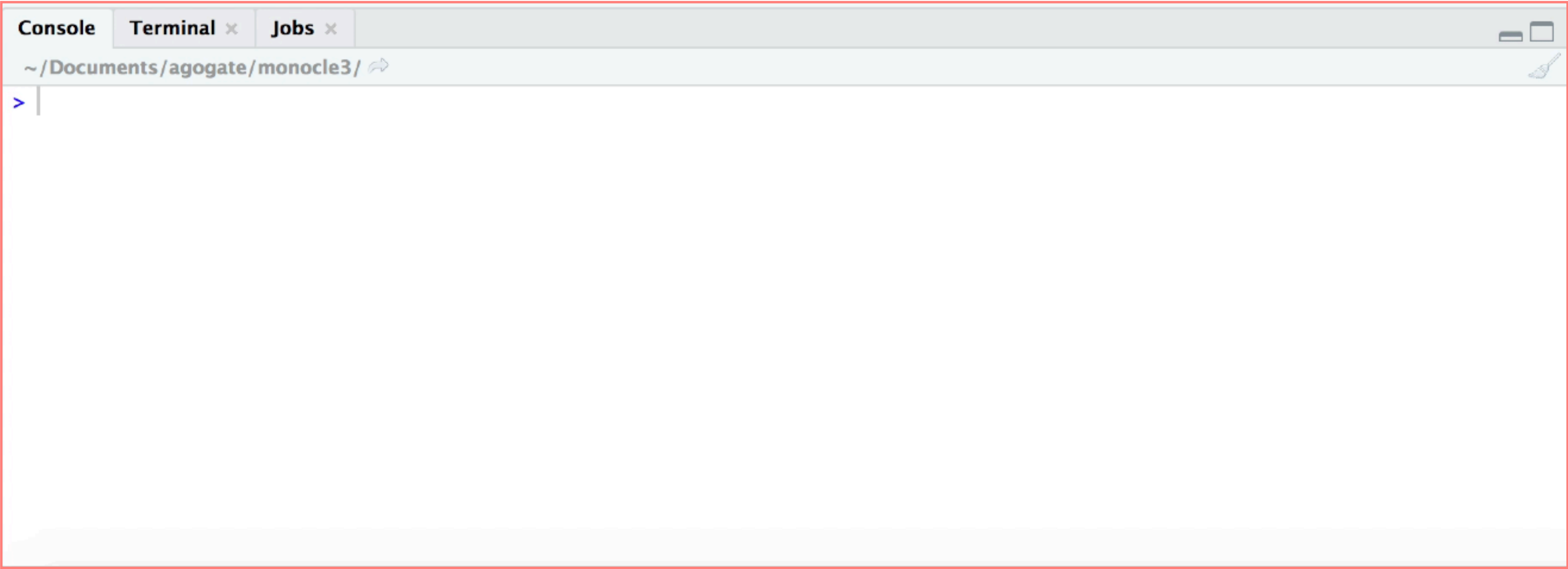
> |

colData(cds)

	cell_type	culture_cond	Size_Factor
cell_1			
cell_2			
cell_3			
cell_n			

rowData(cds)

	biotype	gc_content
gene_1		
gene_2		
gene_3		
gene_n		



Pre-process & batch

