

Cell/Tissue	2.0E-05												
Age													
Gender	2.3E-19	2.0E-05											
Senrix ID	2.0E-05	2.0E-05	8.3E-08	2.0E-05									
Senrix Position		8.2E-04											
Cell Type		2.0E-05			1.8E-04	2.0E-05							
Age Group			3.2E-62		4.0E-05								
Predicted Male Probability	3.8E-05			1.1E-130	4.5E-03								
Predicted Gender	2.3E-19	2.0E-05		5.9E-237	2.0E-05				5.3E-131				
Genome-wide methylation		4.0E-15			7.2E-37	5.6E-29	3.3E-05						
predicted_ages	6.3E-04		1.0E-04		6.6E-08			3.9E-62					
age_increase			1.0E-04		1.3E-19			4.9E-16				1.0E-04	
Immune Cell Content (LUMP)	9.1E-17	1.2E-09	2.0E-04		3.1E-97			1.6E-06				1.0E-04	1.0E-04
	Cohort	Cell/Tissue	Age	Gender	Senrix ID	Senrix Position	Cell Type	Age Group	Predicted Male Probability	Predicted Gender	Genome-wide methylation	predicted_ages	age_increase