	1	2	3	4	5	6	7	8
Cohort		1.8E−18	9.0E−05		3.9E−21	2.3E−03	3.0E−17	7.7E−13
Cell/Tissue	2.2E−27	5.7E−12	2.4E−06	3.8E−09	1.1E−22	1.7E−06	1.5E−22	6.6E−12
Age	1.0E−04	1.0E−04	1.0E−04	1.0E−04	1.0E−04			
Gender			3.2E−03		2.3E−33		5.4E−11	
Sentrix ID	8.6E−06	2.2E−35	3.8E−75	6.9E−36	6.7E−27	2.2E−75	8.7E−59	8.8E−81
Sentrix Position		1.8E−10		2.1E−07	5.6E−06	9.9E−05		
Cell Type	1.7E−05	4.4E−03	4.6E−04	4.7E−05	9.9E−06	3.0E−04	4.2E−05	9.8E−04
Age Group	5.8E−16	1.4E−34	3.0E−31	1.5E−27	2.9E−07			
Predicted Male Probability			5.4E−03		1.0E−04		1.0E−04	
Predicted Gender			3.5E−03		2.9E−33		1.1E−10	
Genome−wide methylation	1.0E−04	1.0E−04	1.0E−04	1.0E−04	1.0E−04	2.5E−03	1.6E−03	
predicted_ages	1.0E−04	1.0E−04	1.0E−04	1.0E−04				
age_increase			1.0E−04			1.0E−04	3.0E−04	1.0E−04
Immune Cell Content (LUMP)		1.5E−03	1.0E−04	1.0E−04		2.0E−04		

Principal component