Lineage Identifier Intersection

Comple ID	Principal Component		Lineage Alignment		
Sample ID	PC1	PC2	'A' Rate	'B' Rate	
CWMC_1.2	64.21193	-4.851574	0.64	0.552	
CWMC_1.4	63.69493	3.542804	0.652	0.561	Worms Identified as
CWMC_2.4	68.07215	-9.773377	0.585	0.509	lineage 'B',
CWMC_2.5	61.45939	11.193412	0.624	0.54	identified in the
CWMC_3.3	65.86026	-1.283664	0.643	0.553	
CWMC_3.6	64.74442	5.712063	0.641	0.553	same category in
CWMM_2.3	67.8455	-16.921207	0.644	0.557	both analyses
CWMM_3.9	61.909	9.868352	0.648	0.548	
CWMC_1.1	-19.29933	-30.1480973	0.878	0.46	
CWMC_1.3	-18.23981	-39.8997683	0.854	0.443	
CWMC_1.5	-17.71591	7.4527081	0.875	0.456	
CWMC_1.6	-15.10786	8.8624651	0.875	0.457	
CWMC_1.7	-14.14551	13.3799944	0.849	0.45	
CWMC_1.8	-18.07431	-9.4510231	0.887	0.468	
CWMC_2.1	-13.59617	-25.0566349	0.846	0.453	
CWMC_2.2	-16.52544	0.4291632	0.872	0.459	
CWMC_2.8	-16.10405	8.3065774	0.873	0.466	
CWMC_2.9	-15.01159	2.2175491	0.886	0.467	
CWMC_3.2	-14.30385	10.5853055	0.868	0.461	
CWMC_3.5	-17.19455	1.2274657	0.866	0.454	Worms Identified
CWMC_3.7	-18.34074	-3.4539132	0.878	0.457	as lineage 'A',
CWMC_3.8	-14.99761	6.068049	0.866	0.459	identified in the
CWMM_1.10	-16.52652	3.5011453	0.891	0.468	
CWMM_1.1	-18.11346	-14.3286941	0.894	0.469	same category in
CWMM_1.2	-15.34837	-36.6257094	0.87	0.473	both analyses
CWMM_1.3	-17.36745	8.9451205	0.876	0.459	
CWMM_1.4	-13.03161	12.9758288	0.876	0.465	
CWMM_1.5	-13.83473	18.1442847	0.867	0.463	
CWMM_1.7	-15.59906	13.236664	0.894	0.464	
CWMM_1.8	-17.29132	-17.6467317	0.883	0.463	
CWMM_1.9	-18.67533	0.3825125	0.879	0.457	
CWMM_3.1	-16.8167	12.3894567	0.884	0.461	
CWMM_3.2	-16.92392	9.6137812	0.884	0.466	
CWMM_3.3	-13.96111	8.1540588	0.862	0.453	
CWMM_3.4	-16.05824	9.0678384	0.857	0.449	
CWMM_3.7	-16.63925	2.2693346	0.863	0.458	