

Ranges

	[,1]	[,2]	[,3]		[,19]	[,20]
[1,]	17	21	15	...	18	16
[2,]	5	4	1	...	4	5
[3,]	39	36	39	...	40	37
[4,]	39	39	34	...	32	35
[5,]	0	0	1	...	0	2
[6,]	21	17	27	...	19	22

[23053,]	10	3	6	...	8	5
[23054,]	31	37	49	...	48	46
[23055,]	23	15	17	...	11	10
[23056,]	32	26	38	...	38	37
	Samples					

Samples

Column (sample) Data			
DataFrame with 20 rows and 3 columns			
	Gender	Batch	Treatment
	<character>	<factor>	<factor>
[1]	Female	1	Control
[2]	Female	2	RNAi
[3]	Female	3	Control
...
[19]	Male	4	Control
[20]	Male	5	RNAi

Row (range) Data						
GRanges with 23056 ranges and 2 metadata columns:						
	seqnames	ranges	strand		SYMBOL	PValue
	<Rle>	<IRanges>	<Rle>		<character>	<numeric>
[1]	chr19	[58858172, 58874214]	-		A1BG	0.2784468
[2]	chr12	[9220304, 9268558]	-		A2M	0.6360118
[3]	chr12	[9381129, 9386803]	-		A2MP1	0.1070964
[4]	chr8	[18027971, 18081198]	+		NAT1	0.4463454
[5]	chr8	[18248755, 18258723]	+		NAT2	0.2569173
[6]	chr14	[95058395, 95090390]	+		SERPINA3	0.3149317
...
[23053]	chr2	[38177477, 38263484]	-		RMDN2-AS1	0.8379452
[23054]	chr4	[98288077, 98411315]	+		STPG2-AS1	0.6557142
[23055]	chr9	[137711261, 137764464]	-		LOC101448202	0.2690768
[23056]	chr17	[4607525, 4608824]	+		LOC101559451	0.9891573

seqlengths:

chr1	chr2	...	chrUn_g1000249
249250621	243199373	...	38502