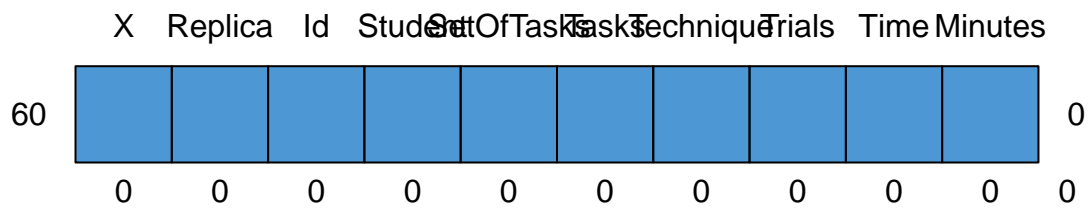


```
ccwo <- read.csv("datasetatoms.csv")
ccwo$Minutes[ccwo$Minutes < 0] <- NA
```

```
md.pattern(ccwo)
```

```
## /\      /\
## {  `---'  }
## {  0    0  }
## ==> V <== No need for mice. This data set is completely observed.
## \  \|/  /
## `-----'
```



```
## X Replica Id Student SetOfTasks Tasks Technique Trials Time Minutes
## 60 1 1 1 1 1 1 1 1 1 1 0
## 0 0 0 0 0 0 0 0 0 0 0 0
```

```
##
## iter imp variable
## 1 1
## 1 2
## 1 3
## 1 4
## 1 5
## 2 1
```

##	2	2
##	2	3
##	2	4
##	2	5
##	3	1
##	3	2
##	3	3
##	3	4
##	3	5
##	4	1
##	4	2
##	4	3
##	4	4
##	4	5
##	5	1
##	5	2
##	5	3
##	5	4
##	5	5
##	6	1
##	6	2
##	6	3
##	6	4
##	6	5
##	7	1
##	7	2
##	7	3
##	7	4
##	7	5
##	8	1
##	8	2
##	8	3
##	8	4
##	8	5
##	9	1
##	9	2
##	9	3
##	9	4
##	9	5
##	10	1
##	10	2
##	10	3
##	10	4
##	10	5
##	11	1
##	11	2
##	11	3
##	11	4
##	11	5
##	12	1
##	12	2
##	12	3
##	12	4
##	12	5

##	13	1
##	13	2
##	13	3
##	13	4
##	13	5
##	14	1
##	14	2
##	14	3
##	14	4
##	14	5
##	15	1
##	15	2
##	15	3
##	15	4
##	15	5
##	16	1
##	16	2
##	16	3
##	16	4
##	16	5
##	17	1
##	17	2
##	17	3
##	17	4
##	17	5
##	18	1
##	18	2
##	18	3
##	18	4
##	18	5
##	19	1
##	19	2
##	19	3
##	19	4
##	19	5
##	20	1
##	20	2
##	20	3
##	20	4
##	20	5
##	21	1
##	21	2
##	21	3
##	21	4
##	21	5
##	22	1
##	22	2
##	22	3
##	22	4
##	22	5
##	23	1
##	23	2
##	23	3
##	23	4

##	23	5
##	24	1
##	24	2
##	24	3
##	24	4
##	24	5
##	25	1
##	25	2
##	25	3
##	25	4
##	25	5
##	26	1
##	26	2
##	26	3
##	26	4
##	26	5
##	27	1
##	27	2
##	27	3
##	27	4
##	27	5
##	28	1
##	28	2
##	28	3
##	28	4
##	28	5
##	29	1
##	29	2
##	29	3
##	29	4
##	29	5
##	30	1
##	30	2
##	30	3
##	30	4
##	30	5
##	31	1
##	31	2
##	31	3
##	31	4
##	31	5
##	32	1
##	32	2
##	32	3
##	32	4
##	32	5
##	33	1
##	33	2
##	33	3
##	33	4
##	33	5
##	34	1
##	34	2
##	34	3

##	34	4
##	34	5
##	35	1
##	35	2
##	35	3
##	35	4
##	35	5
##	36	1
##	36	2
##	36	3
##	36	4
##	36	5
##	37	1
##	37	2
##	37	3
##	37	4
##	37	5
##	38	1
##	38	2
##	38	3
##	38	4
##	38	5
##	39	1
##	39	2
##	39	3
##	39	4
##	39	5
##	40	1
##	40	2
##	40	3
##	40	4
##	40	5
##	41	1
##	41	2
##	41	3
##	41	4
##	41	5
##	42	1
##	42	2
##	42	3
##	42	4
##	42	5
##	43	1
##	43	2
##	43	3
##	43	4
##	43	5
##	44	1
##	44	2
##	44	3
##	44	4
##	44	5
##	45	1
##	45	2

```

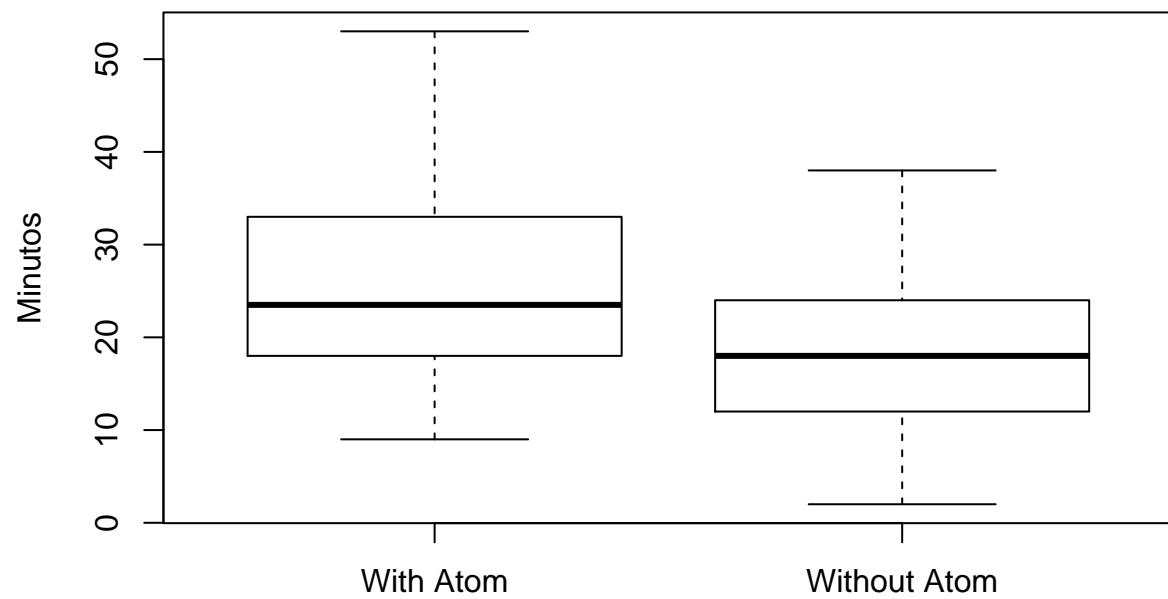
## 45 3
## 45 4
## 45 5
## 46 1
## 46 2
## 46 3
## 46 4
## 46 5
## 47 1
## 47 2
## 47 3
## 47 4
## 47 5
## 48 1
## 48 2
## 48 3
## 48 4
## 48 5
## 49 1
## 49 2
## 49 3
## 49 4
## 49 5
## 50 1
## 50 2
## 50 3
## 50 4
## 50 5

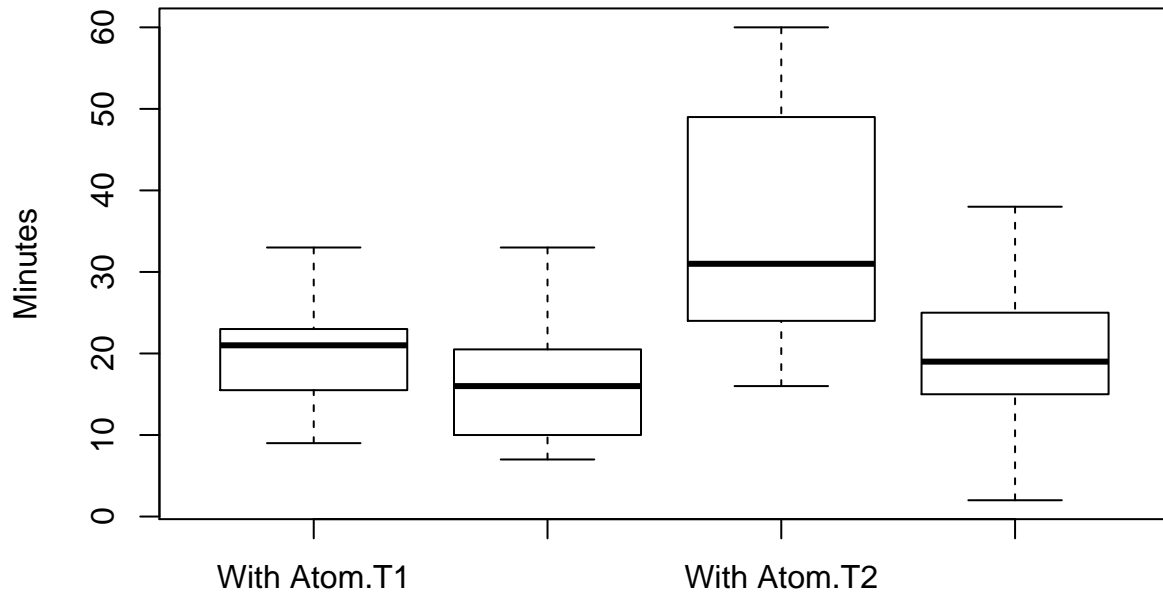
```

Warning: Number of logged events: 1

##	Replica	Id	SetOfTasks	Technique	Trials	Time
## 1	41	1	ST1	Without Atom	7	13
## 2	41	1	ST2	With Atom	3	16
## 3	41	2	ST1	With Atom	5	21
## 4	41	2	ST2	Without Atom	3	15
## 5	42	1	ST1	Without Atom	5	14
## 6	42	1	ST2	With Atom	12	39
## 7	42	2	ST1	With Atom	7	16
## 8	42	2	ST2	Without Atom	12	22
## 9	43	1	ST1	Without Atom	5	18
## 10	43	1	ST2	With Atom	9	53
## 11	43	2	ST1	With Atom	9	21
## 12	43	2	ST2	Without Atom	5	15
## 13	44	1	ST1	Without Atom	4	9
## 14	44	1	ST2	With Atom	4	24
## 15	44	2	ST1	With Atom	6	17
## 16	44	2	ST2	Without Atom	6	23
## 17	45	1	ST1	With Atom	9	25
## 18	45	1	ST2	Without Atom	7	28
## 19	45	2	ST1	Without Atom	5	7
## 20	45	2	ST2	With Atom	8	18
## 21	46	1	ST1	With Atom	7	32
## 22	46	1	ST2	Without Atom	9	32

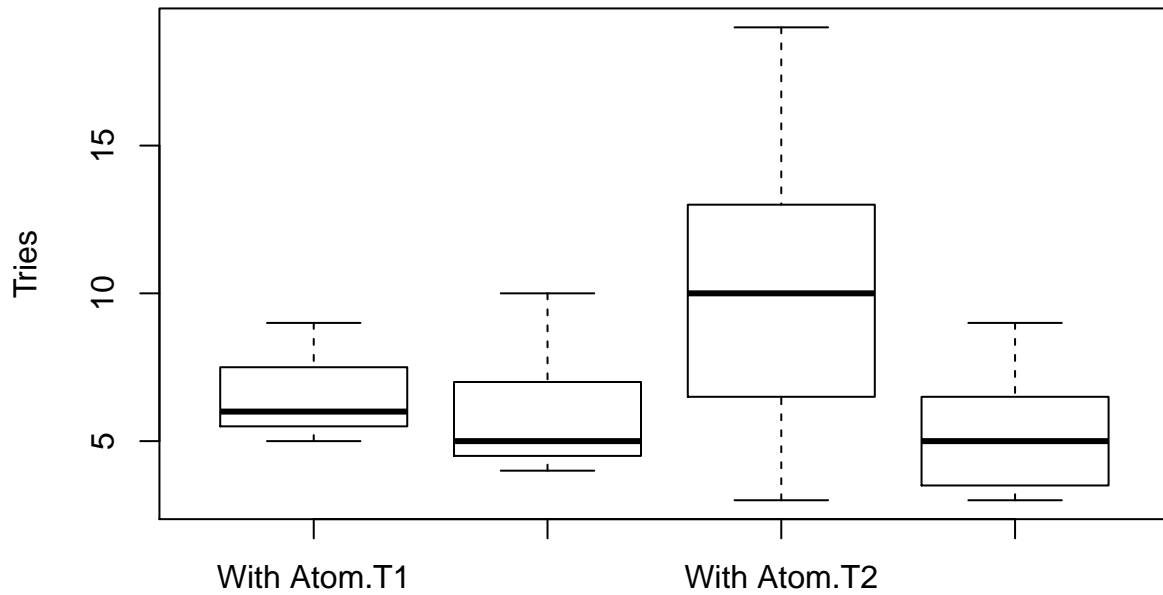
## 23	46	2	ST1 Without Atom	4	10
## 24	46	2	ST2 With Atom	12	31
## 25	47	1	ST1 Without Atom	4	18
## 26	47	1	ST2 With Atom	5	53
## 27	47	2	ST1 With Atom	5	21
## 28	47	2	ST2 Without Atom	4	24
## 29	48	1	ST1 With Atom	7	15
## 30	48	1	ST2 Without Atom	3	12
## 31	48	2	ST1 Without Atom	10	25
## 32	48	2	ST2 With Atom	29	60
## 33	49	1	ST1 Without Atom	8	33
## 34	49	1	ST2 With Atom	10	23
## 35	49	2	ST1 With Atom	6	9
## 36	49	2	ST2 Without Atom	3	2
## 37	50	1	ST1 With Atom	8	21
## 38	50	1	ST2 Without Atom	5	19
## 39	50	2	ST1 Without Atom	7	19
## 40	50	2	ST2 With Atom	10	29
## 41	51	1	ST1 Without Atom	5	22
## 42	51	1	ST2 With Atom	6	24
## 43	51	2	ST1 With Atom	6	9
## 44	51	2	ST2 Without Atom	5	15
## 45	52	1	ST1 With Atom	9	23
## 46	52	1	ST2 Without Atom	24	38
## 47	52	2	ST1 Without Atom	7	16
## 48	52	2	ST2 With Atom	14	52
## 49	53	1	ST1 With Atom	5	11
## 50	53	1	ST2 Without Atom	4	8
## 51	53	2	ST1 Without Atom	5	9
## 52	53	2	ST2 With Atom	15	31
## 53	54	1	ST1 Without Atom	4	27
## 54	54	1	ST2 With Atom	7	34
## 55	54	2	ST1 With Atom	6	33
## 56	54	2	ST2 Without Atom	3	26
## 57	55	1	ST1 With Atom	5	23
## 58	55	1	ST2 Without Atom	4	19
## 59	55	2	ST1 Without Atom	5	10
## 60	55	2	ST2 With Atom	19	46





##	Replica	Tasks	Technique	Trials	Time
## 1	41	T1	With Atom	5	21
## 2	41	T1	Without Atom	7	13
## 3	41	T2	With Atom	3	16
## 4	41	T2	Without Atom	3	15
## 5	42	T1	With Atom	7	16
## 6	42	T1	Without Atom	5	14
## 7	42	T2	With Atom	12	39
## 8	42	T2	Without Atom	12	22
## 9	43	T1	With Atom	9	21
## 10	43	T1	Without Atom	5	18
## 11	43	T2	With Atom	9	53
## 12	43	T2	Without Atom	5	15
## 13	44	T1	With Atom	6	17
## 14	44	T1	Without Atom	4	9
## 15	44	T2	With Atom	4	24
## 16	44	T2	Without Atom	6	23
## 17	45	T1	With Atom	9	25
## 18	45	T1	Without Atom	5	7
## 19	45	T2	With Atom	8	18
## 20	45	T2	Without Atom	7	28
## 21	46	T1	With Atom	7	32
## 22	46	T1	Without Atom	4	10
## 23	46	T2	With Atom	12	31
## 24	46	T2	Without Atom	9	32
## 25	47	T1	With Atom	5	21

## 26	47	T1 Without Atom	4	18
## 27	47	T2 With Atom	5	53
## 28	47	T2 Without Atom	4	24
## 29	48	T1 With Atom	7	15
## 30	48	T1 Without Atom	10	25
## 31	48	T2 With Atom	29	60
## 32	48	T2 Without Atom	3	12
## 33	49	T1 With Atom	6	9
## 34	49	T1 Without Atom	8	33
## 35	49	T2 With Atom	10	23
## 36	49	T2 Without Atom	3	2
## 37	50	T1 With Atom	8	21
## 38	50	T1 Without Atom	7	19
## 39	50	T2 With Atom	10	29
## 40	50	T2 Without Atom	5	19
## 41	51	T1 With Atom	6	9
## 42	51	T1 Without Atom	5	22
## 43	51	T2 With Atom	6	24
## 44	51	T2 Without Atom	5	15
## 45	52	T1 With Atom	9	23
## 46	52	T1 Without Atom	7	16
## 47	52	T2 With Atom	14	52
## 48	52	T2 Without Atom	24	38
## 49	53	T1 With Atom	5	11
## 50	53	T1 Without Atom	5	9
## 51	53	T2 With Atom	15	31
## 52	53	T2 Without Atom	4	8
## 53	54	T1 With Atom	6	33
## 54	54	T1 Without Atom	4	27
## 55	54	T2 With Atom	7	34
## 56	54	T2 Without Atom	3	26
## 57	55	T1 With Atom	5	23
## 58	55	T1 Without Atom	5	10
## 59	55	T2 With Atom	19	46
## 60	55	T2 Without Atom	4	19



##	X	Replica	Id	Student	SetOfTasks	Tasks
## 1	1	41	1	paulo	ST1	AV1.2:C01.2:DE1.2
## 2	1	41	1	paulo	ST2	AV2.1:C02.1:DE2.1
## 3	1	41	2	Romário	ST1	AV1.1:C01.1:DE1.1
## 4	1	41	2	Romário	ST2	AV2.2:C02.2:DE2.2
## 5	2	42	1	Igor	ST1	AV1.2:C01.2:DE1.2
## 6	2	42	1	Igor	ST2	AV2.1:C02.1:DE2.1
## 7	2	42	2	Hyago	ST1	AV1.1:C01.1:DE1.1
## 8	2	42	2	Hyago	ST2	AV2.2:C02.2:DE2.2
## 9	3	43	1		ST1	AV1.2:C01.2:DE1.2
## 10	3	43	1		ST2	AV2.1:C02.1:DE2.1
## 11	3	43	2		ST1	AV1.1:C01.1:DE1.1
## 12	3	43	2		ST2	AV2.2:C02.2:DE2.2
## 13	4	44	1		ST1	AV1.2:C01.2:DE1.2
## 14	4	44	1		ST2	AV2.1:C02.1:DE2.1
## 15	4	44	2		ST1	AV1.1:C01.1:DE1.1
## 16	4	44	2		ST2	AV2.2:C02.2:DE2.2
## 17	5	45	1	Matheus Costa	ST1	AV1.1:C01.1:DE1.1
## 18	5	45	1	Matheus Costa	ST2	AV2.2:C02.2:DE2.2
## 19	5	45	2	Davi Jose	ST1	AV1.2:C01.2:DE1.2
## 20	5	45	2	Davi Jose	ST2	AV2.1:C02.1:DE2.1
## 21	6	46	1	Marlon Lúcio	ST1	AV1.1:C01.1:DE1.1
## 22	6	46	1	Marlon Lúcio	ST2	AV2.2:C02.2:DE2.2
## 23	6	46	2	Jackson Barbosa da Silva	ST1	AV1.2:C01.2:DE1.2
## 24	6	46	2	Jackson Barbosa da Silva	ST2	AV2.1:C02.1:DE2.1
## 25	7	47	1	Allan	ST1	AV1.2:C01.2:DE1.2

## 26	7	47	1	Allan	ST2 AV2.1:C02.1:DE2.1
## 27	7	47	2	Dielson Sales de Carvalho	ST1 AV1.1:C01.1:DE1.1
## 28	7	47	2	Dielson Sales de Carvalho	ST2 AV2.2:C02.2:DE2.2
## 29	8	48	1	Thiago	ST1 AV1.1:C01.1:DE1.1
## 30	8	48	1	Thiago	ST2 AV2.2:C02.2:DE2.2
## 31	8	48	2		ST1 AV1.2:C01.2:DE1.2
## 32	8	48	2		ST2 AV2.1:C02.1:DE2.1
## 33	9	49	1	Rodrigo Lima	ST1 AV1.2:C01.2:DE1.2
## 34	9	49	1	Rodrigo Lima	ST2 AV2.1:C02.1:DE2.1
## 35	9	49	2	Jardel Costa	ST1 AV1.1:C01.1:DE1.1
## 36	9	49	2	Jardel Costa	ST2 AV2.2:C02.2:DE2.2
## 37	10	50	1	Felipe Pontes	ST1 AV1.1:C01.1:DE1.1
## 38	10	50	1	Felipe Pontes	ST2 AV2.2:C02.2:DE2.2
## 39	10	50	2	Jairo Souza	ST1 AV1.2:C01.2:DE1.2
## 40	10	50	2	Jairo Souza	ST2 AV2.1:C02.1:DE2.1
## 41	11	51	1	julios	ST1 AV1.2:C01.2:DE1.2
## 42	11	51	1	julios	ST2 AV2.1:C02.1:DE2.1
## 43	11	51	2	Romero Malaquias	ST1 AV1.1:C01.1:DE1.1
## 44	11	51	2	Romero Malaquias	ST2 AV2.2:C02.2:DE2.2
## 45	12	53	1	Bruno Georgevich Ferreira	ST1 AV1.1:C01.1:DE1.1
## 46	12	53	1	Bruno Georgevich Ferreira	ST2 AV2.2:C02.2:DE2.2
## 47	12	53	2	jadson	ST1 AV1.2:C01.2:DE1.2
## 48	12	53	2	jadson	ST2 AV2.1:C02.1:DE2.1
## 49	13	55	1		ST1 AV1.1:C01.1:DE1.1
## 50	13	55	1		ST2 AV2.2:C02.2:DE2.2
## 51	13	55	2	Arthur	ST1 AV1.2:C01.2:DE1.2
## 52	13	55	2	Arthur	ST2 AV2.1:C02.1:DE2.1
## 53	14	52	1		ST1 AV1.1:C01.1:DE1.1
## 54	14	52	1		ST2 AV2.2:C02.2:DE2.2
## 55	14	52	2	vinicius lopes	ST1 AV1.2:C01.2:DE1.2
## 56	14	52	2	vinicius lopes	ST2 AV2.1:C02.1:DE2.1
## 57	15	54	1	Thiago Tenorio	ST1 AV1.2:C01.2:DE1.2
## 58	15	54	1	Thiago Tenorio	ST2 AV2.1:C02.1:DE2.1
## 59	15	54	2	Joao Victor Ribeiro Ferro	ST1 AV1.1:C01.1:DE1.1
## 60	15	54	2	Joao Victor Ribeiro Ferro	ST2 AV2.2:C02.2:DE2.2
##				Technique Trials Time Minutes	
## 1	Without Atom	7	13	13	
## 2	With Atom	3	16	16	
## 3	With Atom	5	21	21	
## 4	Without Atom	3	15	15	
## 5	Without Atom	5	14	14	
## 6	With Atom	12	39	39	
## 7	With Atom	7	16	16	
## 8	Without Atom	12	22	22	
## 9	Without Atom	5	18	18	
## 10	With Atom	9	53	53	
## 11	With Atom	9	21	21	
## 12	Without Atom	5	15	15	
## 13	Without Atom	4	9	9	
## 14	With Atom	4	24	24	
## 15	With Atom	6	17	17	
## 16	Without Atom	6	23	23	
## 17	With Atom	9	25	25	
## 18	Without Atom	7	28	28	

## 19	Without Atom	5	7	7
## 20	With Atom	8	18	18
## 21	With Atom	7	32	32
## 22	Without Atom	9	32	32
## 23	Without Atom	4	10	10
## 24	With Atom	12	31	31
## 25	Without Atom	4	18	18
## 26	With Atom	5	53	53
## 27	With Atom	5	21	21
## 28	Without Atom	4	24	24
## 29	With Atom	7	15	15
## 30	Without Atom	3	12	12
## 31	Without Atom	10	25	25
## 32	With Atom	29	60	60
## 33	Without Atom	8	33	33
## 34	With Atom	10	23	23
## 35	With Atom	6	9	9
## 36	Without Atom	3	2	2
## 37	With Atom	8	21	21
## 38	Without Atom	5	19	19
## 39	Without Atom	7	19	19
## 40	With Atom	10	29	29
## 41	Without Atom	5	22	22
## 42	With Atom	6	24	24
## 43	With Atom	6	9	9
## 44	Without Atom	5	15	15
## 45	With Atom	5	11	11
## 46	Without Atom	4	8	8
## 47	Without Atom	5	9	9
## 48	With Atom	15	31	31
## 49	With Atom	5	23	23
## 50	Without Atom	4	19	19
## 51	Without Atom	5	10	10
## 52	With Atom	19	46	46
## 53	With Atom	9	23	23
## 54	Without Atom	24	38	38
## 55	Without Atom	7	16	16
## 56	With Atom	14	52	52
## 57	Without Atom	4	27	27
## 58	With Atom	7	34	34
## 59	With Atom	6	33	33
## 60	Without Atom	3	26	26

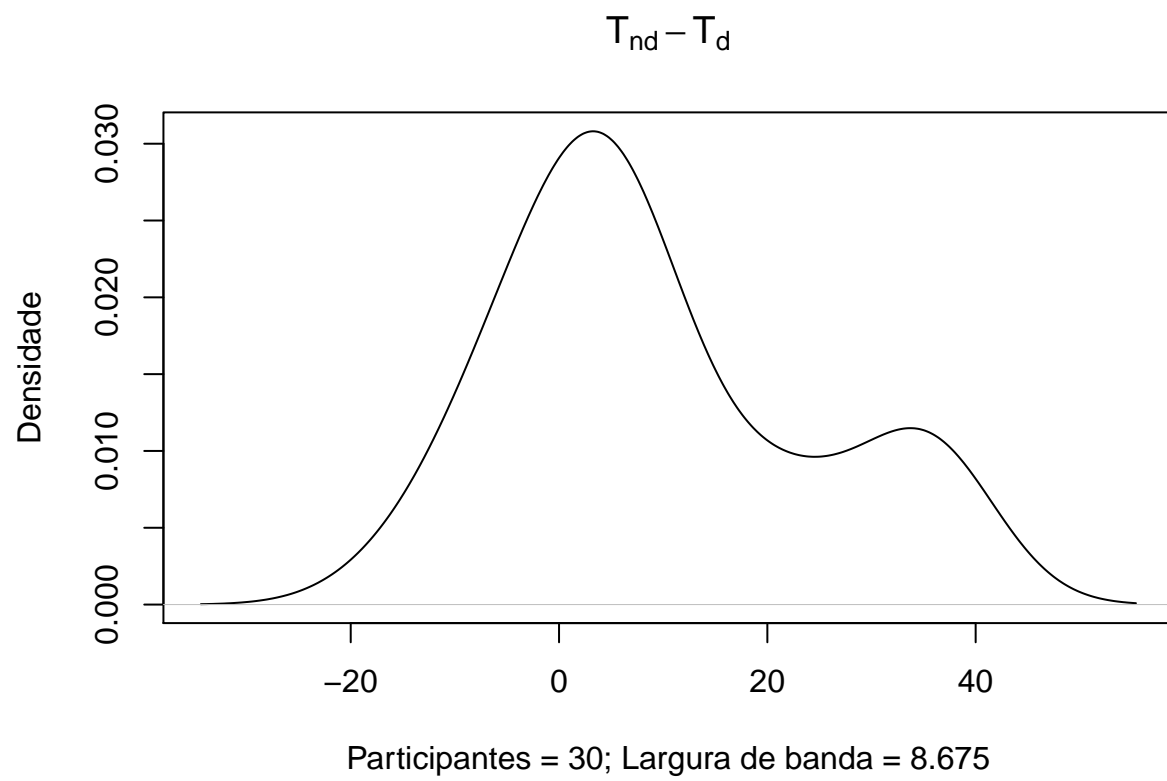
##	Replica	Tasks	Minutes	Time
## 1	41 AV1.2:CO1.2:DE1.2		13	13
## 2	41 AV1.1:CO1.1:DE1.1		21	21
## 3	42 AV1.2:CO1.2:DE1.2		14	14
## 4	42 AV1.1:CO1.1:DE1.1		16	16
## 5	43 AV1.2:CO1.2:DE1.2		18	18
## 6	43 AV1.1:CO1.1:DE1.1		21	21
## 7	44 AV1.2:CO1.2:DE1.2		9	9
## 8	44 AV1.1:CO1.1:DE1.1		17	17
## 9	45 AV1.1:CO1.1:DE1.1		25	25
## 10	45 AV1.2:CO1.2:DE1.2		7	7

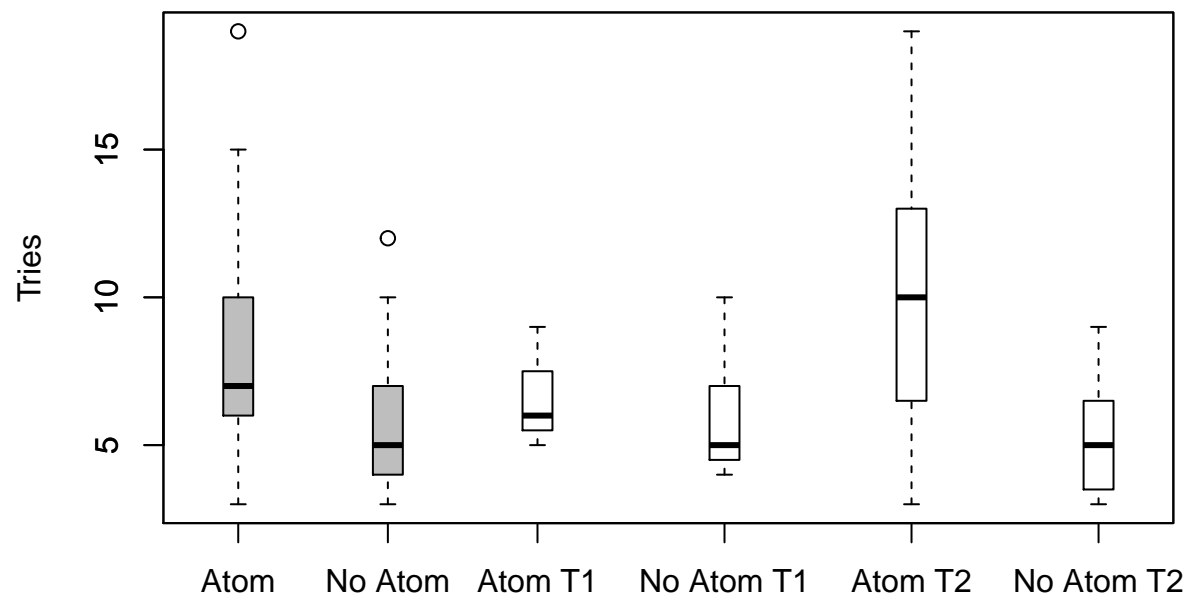
## 11	46	AV1.1:C01.1:DE1.1	32	32
## 12	46	AV1.2:C01.2:DE1.2	10	10
## 13	47	AV1.2:C01.2:DE1.2	18	18
## 14	47	AV1.1:C01.1:DE1.1	21	21
## 15	48	AV1.1:C01.1:DE1.1	15	15
## 16	48	AV1.2:C01.2:DE1.2	25	25
## 17	49	AV1.2:C01.2:DE1.2	33	33
## 18	49	AV1.1:C01.1:DE1.1	9	9
## 19	50	AV1.1:C01.1:DE1.1	21	21
## 20	50	AV1.2:C01.2:DE1.2	19	19
## 21	51	AV1.2:C01.2:DE1.2	22	22
## 22	51	AV1.1:C01.1:DE1.1	9	9
## 23	53	AV1.1:C01.1:DE1.1	11	11
## 24	53	AV1.2:C01.2:DE1.2	9	9
## 25	55	AV1.1:C01.1:DE1.1	23	23
## 26	55	AV1.2:C01.2:DE1.2	10	10
## 27	52	AV1.1:C01.1:DE1.1	23	23
## 28	52	AV1.2:C01.2:DE1.2	16	16
## 29	54	AV1.2:C01.2:DE1.2	27	27
## 30	54	AV1.1:C01.1:DE1.1	33	33

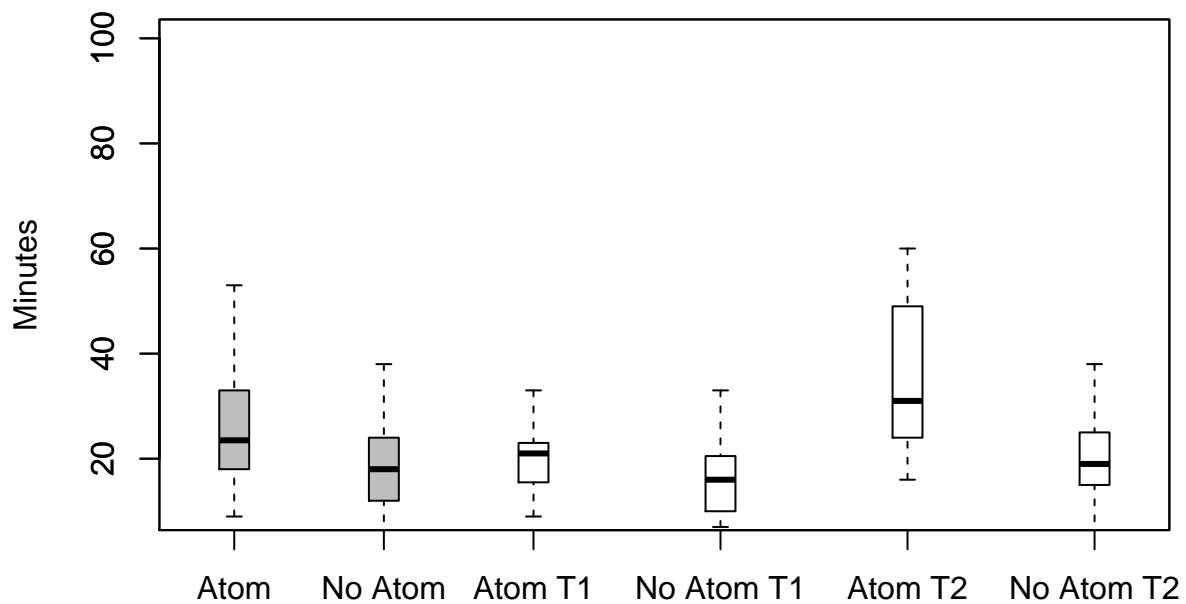
##	Replica	Tasks	Minutes	Time
## 1	41	AV2.1:C02.1:DE2.1	16	16
## 2	41	AV2.2:C02.2:DE2.2	15	15
## 3	42	AV2.1:C02.1:DE2.1	39	39
## 4	42	AV2.2:C02.2:DE2.2	22	22
## 5	43	AV2.1:C02.1:DE2.1	53	53
## 6	43	AV2.2:C02.2:DE2.2	15	15
## 7	44	AV2.1:C02.1:DE2.1	24	24
## 8	44	AV2.2:C02.2:DE2.2	23	23
## 9	45	AV2.2:C02.2:DE2.2	28	28
## 10	45	AV2.1:C02.1:DE2.1	18	18
## 11	46	AV2.2:C02.2:DE2.2	32	32
## 12	46	AV2.1:C02.1:DE2.1	31	31
## 13	47	AV2.1:C02.1:DE2.1	53	53
## 14	47	AV2.2:C02.2:DE2.2	24	24
## 15	48	AV2.2:C02.2:DE2.2	12	12
## 16	48	AV2.1:C02.1:DE2.1	60	60
## 17	49	AV2.1:C02.1:DE2.1	23	23
## 18	49	AV2.2:C02.2:DE2.2	2	2
## 19	50	AV2.2:C02.2:DE2.2	19	19
## 20	50	AV2.1:C02.1:DE2.1	29	29
## 21	51	AV2.1:C02.1:DE2.1	24	24
## 22	51	AV2.2:C02.2:DE2.2	15	15
## 23	53	AV2.2:C02.2:DE2.2	8	8
## 24	53	AV2.1:C02.1:DE2.1	31	31
## 25	55	AV2.2:C02.2:DE2.2	19	19
## 26	55	AV2.1:C02.1:DE2.1	46	46
## 27	52	AV2.2:C02.2:DE2.2	38	38
## 28	52	AV2.1:C02.1:DE2.1	52	52
## 29	54	AV2.1:C02.1:DE2.1	34	34
## 30	54	AV2.2:C02.2:DE2.2	26	26

[1] 18.23333

[1] 27.7







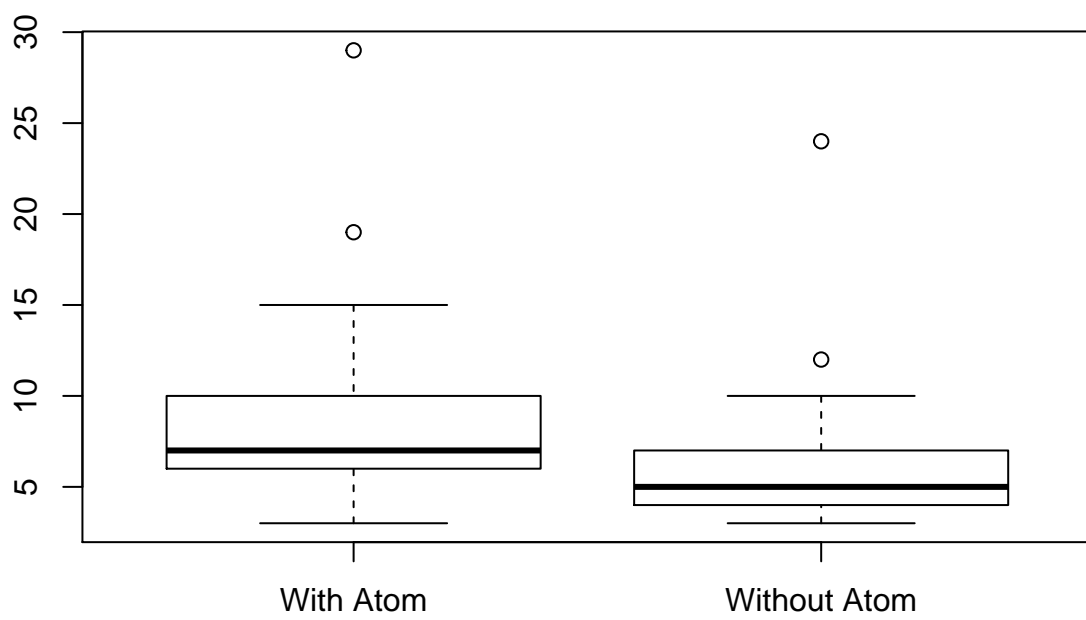
```
totalTrials <- sqldf("select Replica, Id, SetOfTasks,
                      Technique, sum(Trials) as Trials
                      from ccwood
                      group by Replica, Id,SetOfTasks, Technique")
```

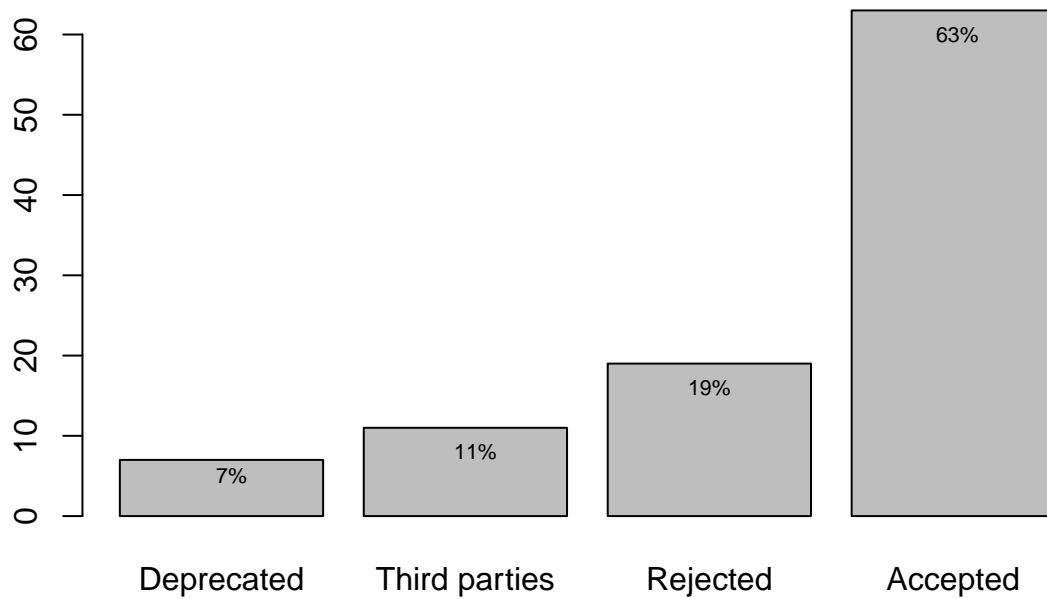
totalTrials

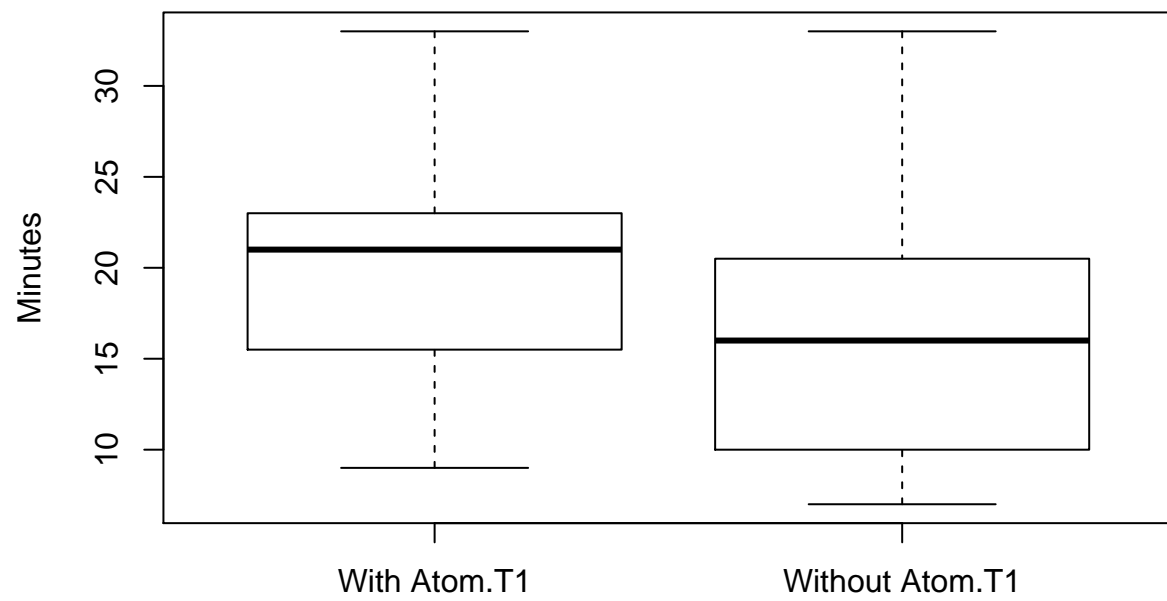
##	Replica	Id	SetOfTasks	Technique	Trials
## 1	41	1	ST1	Without Atom	7
## 2	41	1	ST2	With Atom	3
## 3	41	2	ST1	With Atom	5
## 4	41	2	ST2	Without Atom	3
## 5	42	1	ST1	Without Atom	5
## 6	42	1	ST2	With Atom	12
## 7	42	2	ST1	With Atom	7
## 8	42	2	ST2	Without Atom	12
## 9	43	1	ST1	Without Atom	5
## 10	43	1	ST2	With Atom	9
## 11	43	2	ST1	With Atom	9
## 12	43	2	ST2	Without Atom	5
## 13	44	1	ST1	Without Atom	4
## 14	44	1	ST2	With Atom	4
## 15	44	2	ST1	With Atom	6
## 16	44	2	ST2	Without Atom	6
## 17	45	1	ST1	With Atom	9
## 18	45	1	ST2	Without Atom	7

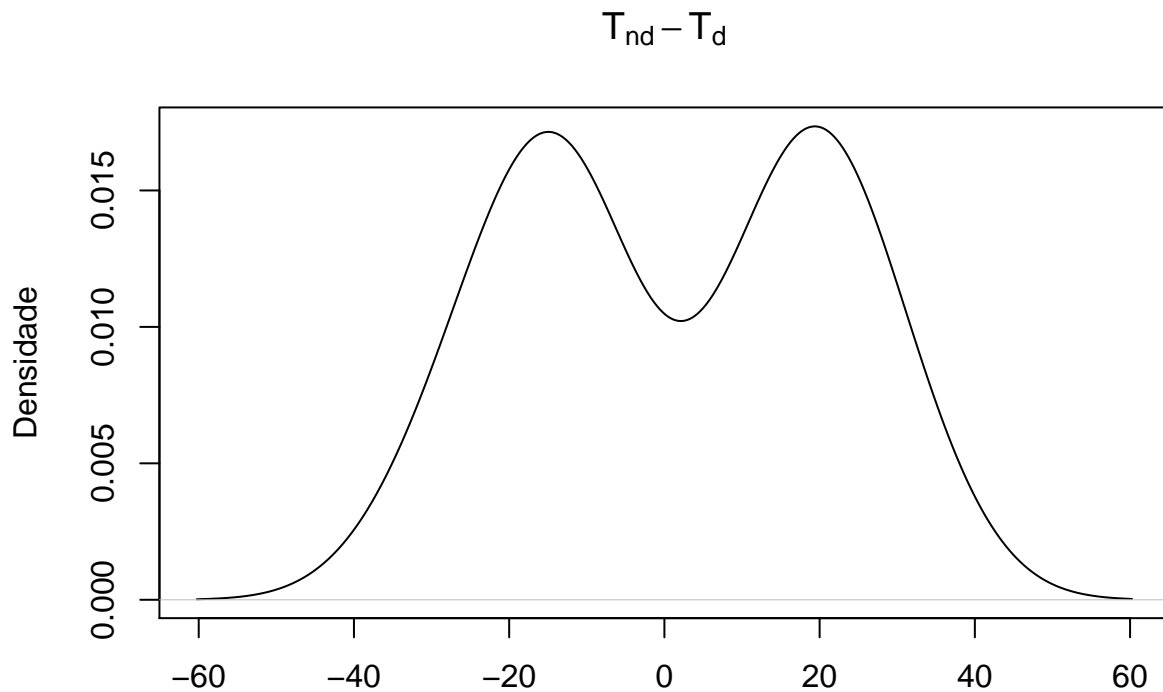
## 19	45	2	ST1 Without Atom	5
## 20	45	2	ST2 With Atom	8
## 21	46	1	ST1 With Atom	7
## 22	46	1	ST2 Without Atom	9
## 23	46	2	ST1 Without Atom	4
## 24	46	2	ST2 With Atom	12
## 25	47	1	ST1 Without Atom	4
## 26	47	1	ST2 With Atom	5
## 27	47	2	ST1 With Atom	5
## 28	47	2	ST2 Without Atom	4
## 29	48	1	ST1 With Atom	7
## 30	48	1	ST2 Without Atom	3
## 31	48	2	ST1 Without Atom	10
## 32	48	2	ST2 With Atom	29
## 33	49	1	ST1 Without Atom	8
## 34	49	1	ST2 With Atom	10
## 35	49	2	ST1 With Atom	6
## 36	49	2	ST2 Without Atom	3
## 37	50	1	ST1 With Atom	8
## 38	50	1	ST2 Without Atom	5
## 39	50	2	ST1 Without Atom	7
## 40	50	2	ST2 With Atom	10
## 41	51	1	ST1 Without Atom	5
## 42	51	1	ST2 With Atom	6
## 43	51	2	ST1 With Atom	6
## 44	51	2	ST2 Without Atom	5
## 45	52	1	ST1 With Atom	9
## 46	52	1	ST2 Without Atom	24
## 47	52	2	ST1 Without Atom	7
## 48	52	2	ST2 With Atom	14
## 49	53	1	ST1 With Atom	5
## 50	53	1	ST2 Without Atom	4
## 51	53	2	ST1 Without Atom	5
## 52	53	2	ST2 With Atom	15
## 53	54	1	ST1 Without Atom	4
## 54	54	1	ST2 With Atom	7
## 55	54	2	ST1 With Atom	6
## 56	54	2	ST2 Without Atom	3
## 57	55	1	ST1 With Atom	5
## 58	55	1	ST2 Without Atom	4
## 59	55	2	ST1 Without Atom	5
## 60	55	2	ST2 With Atom	19

```
boxplot(totalTrials$Trials~totalTrials$Technique)
```









Participantes = 30; Largura de banda = 4.082

```
totalTime <- sqldf("select Replica, Id, SetOfTasks,
                    Technique, sum(Trials) as Trials, sum(Minutes) as Time
                    from ccwood where Tasks = 'AV1.1:C01.1:DE1.1' or Tasks = 'AV1.2:C01.2:DE1.2'
                    group by Replica, Id,SetOfTasks, Technique")
```

```
totalTime$Time <- with(totalTime, log2(Time))
```

```
totalTime$Replica = as.factor(totalTime$Replica)
totalTime$Id = as.factor(totalTime$Id)
totalTime$SetOfTasks = as.factor(totalTime$SetOfTasks)
totalTime$Technique = as.factor(totalTime$Technique)
```

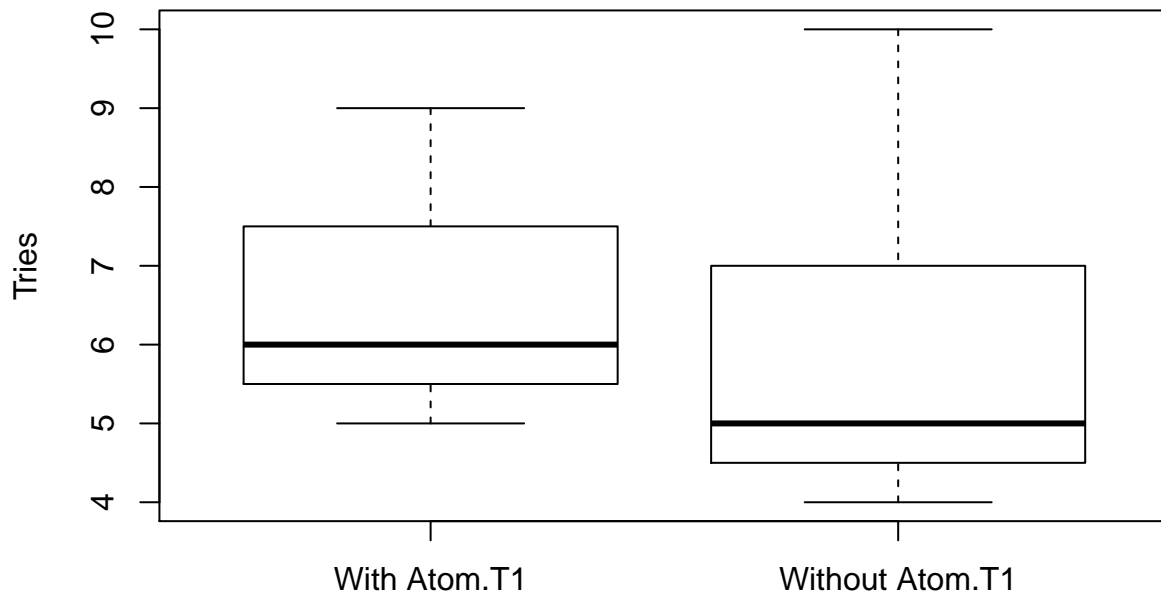
```
totalTime.gvlma = gvlma(lm(Time ~ Technique, data=totalTime))
summary(totalTime.gvlma)
```

```
##
## Call:
## lm(formula = Time ~ Technique, data = totalTime)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.1119 -0.5229  0.1858  0.3257  1.1251
##
## Coefficients:
```

```
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      4.2066     0.1609  26.147  <2e-16 ***
## TechniqueWithout Atom -0.2873     0.2275  -1.263    0.217
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.6231 on 28 degrees of freedom
## Multiple R-squared:  0.05387,    Adjusted R-squared:  0.02008
## F-statistic: 1.594 on 1 and 28 DF,  p-value: 0.2171
##
##
## ASSESSMENT OF THE LINEAR MODEL ASSUMPTIONS
## USING THE GLOBAL TEST ON 4 DEGREES-OF-FREEDOM:
## Level of Significance = 0.05
##
## Call:
## gvlma(x = lm(Time ~ Technique, data = totalTime))
##
##               Value p-value          Decision
## Global Stat      2.611e+00  0.6249 Assumptions acceptable.
## Skewness         2.499e-01  0.6171 Assumptions acceptable.
## Kurtosis         9.031e-01  0.3419 Assumptions acceptable.
## Link Function    1.105e-15  1.0000 Assumptions acceptable.
## Heteroscedasticity 1.458e+00  0.2272 Assumptions acceptable.
```

```
summary(aov(lm(Time ~ Technique, data=totalTime)))
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Technique   1  0.619  0.6189   1.594  0.217
## Residuals  28 10.870  0.3882
```



```
totalTrials <- sqldf("select Replica, Id, SetOfTasks,
                      Technique, sum(Trials) as Trials
                      from ccwood where Tasks = 'AV1.1:C01.1:DE1.1' or Tasks = 'AV1.2:C01.2:DE1.2'
                      group by Replica, Id,SetOfTasks, Technique")
totalTrials$Trials <- with(totalTrials, Trials + 1)
totalTrials$Trials <- with(totalTrials, log2(Trials))
```

```
totalTrials$Replica = as.factor(totalTrials$Replica)
totalTrials$Id = as.factor(totalTrials$Id)
totalTrials$SetOfTasks = as.factor(totalTrials$SetOfTasks:totalTrials$Technique)
totalTrials$Technique = as.factor(totalTrials$Technique)
```

```
summary(aov(Trials ~ Replica + Replica:Id + SetOfTasks + Technique, data=totalTrials))
```

```
##           Df Sum Sq Mean Sq
## Replica    14  1.6643   0.1189
## SetOfTasks    1  0.3582   0.3582
## Replica:Id   14  1.1195   0.0800
```

```
totalTrials.gvlma = gvlma(lm(Trials ~ Technique, data=totalTrials))
summary(totalTrials.gvlma)
```

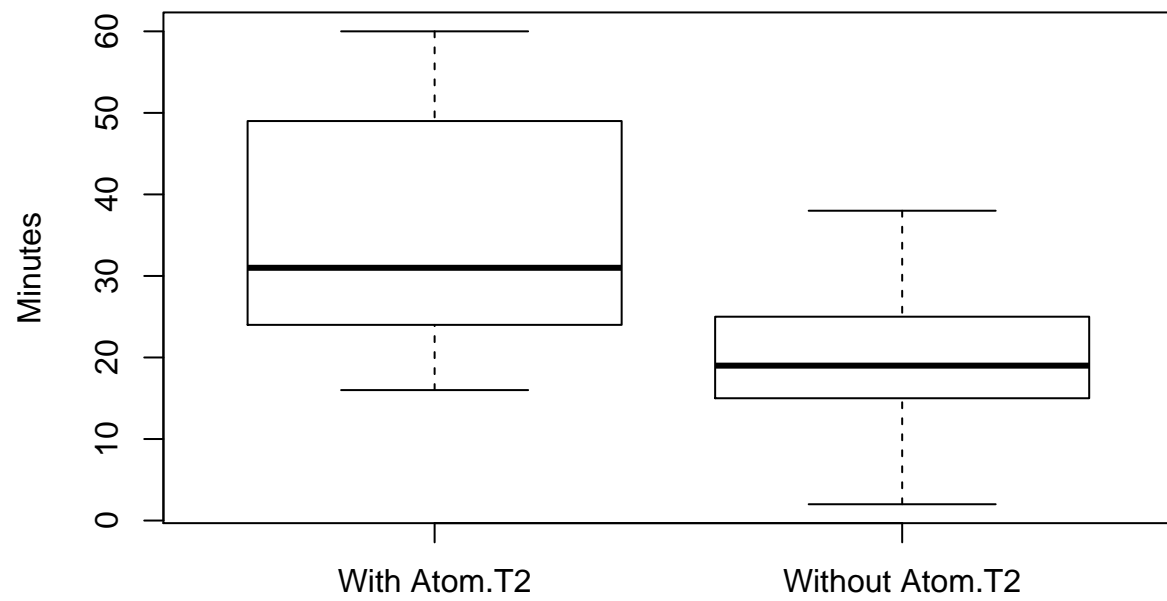
```
##
## Call:
```

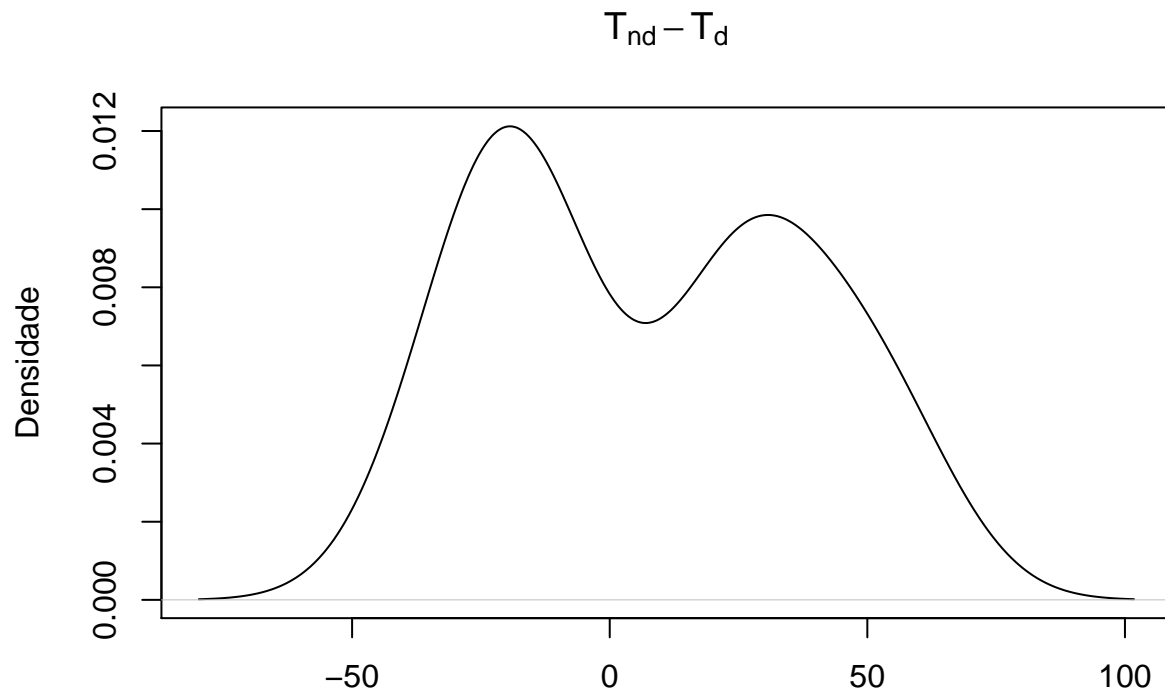


```
## lm(formula = Trials ~ Technique, data = totalTrials)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.3732 -0.2741 -0.1063  0.2927  0.7643
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      2.91367    0.08141  35.789  <2e-16 ***
## TechniqueWithout Atom -0.21854    0.11514  -1.898   0.068 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3153 on 28 degrees of freedom
## Multiple R-squared:  0.114, Adjusted R-squared:  0.08236
## F-statistic: 3.603 on 1 and 28 DF,  p-value: 0.06803
##
##
## ASSESSMENT OF THE LINEAR MODEL ASSUMPTIONS
## USING THE GLOBAL TEST ON 4 DEGREES-OF-FREEDOM:
## Level of Significance = 0.05
##
## Call:
## gvlma(x = lm(Trials ~ Technique, data = totalTrials))
##
##              Value p-value              Decision
## Global Stat      -3.799e+01  1.0000 Assumptions acceptable.
## Skewness          1.729e+00  0.1886 Assumptions acceptable.
## Kurtosis          3.856e-01  0.5346 Assumptions acceptable.
## Link Function     -4.011e+01  1.0000 Assumptions acceptable.
## Heteroscedasticity 9.333e-04  0.9756 Assumptions acceptable.
```

```
summary(aov(Trials ~ Technique, data=totalTrials))
```

```
##              Df Sum Sq Mean Sq F value Pr(>F)
## Technique      1  0.3582   0.3582   3.603  0.068 .
## Residuals     28  2.7838   0.0994
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```





Participantes = 30; Largura de banda = 5.528

```
totalTime <- sqldf("select Replica, Id, SetOfTasks,
                    Technique, sum(Trials) as Trials, sum(Minutes) as Time
                    from ccwood where Tasks = 'AV2.1:C02.1:DE2.1' or Tasks = 'AV2.2:C02.2:DE2.2'
                    group by Replica, Id,SetOfTasks, Technique")
```

```
totalTime$Time <- with(totalTime, log2(Time))
```

```
totalTime$Replica = as.factor(totalTime$Replica)
totalTime$Id = as.factor(totalTime$Id)
totalTime$SetOfTasks = as.factor(totalTime$SetOfTasks)
totalTime$Technique = as.factor(totalTime$Technique)
```

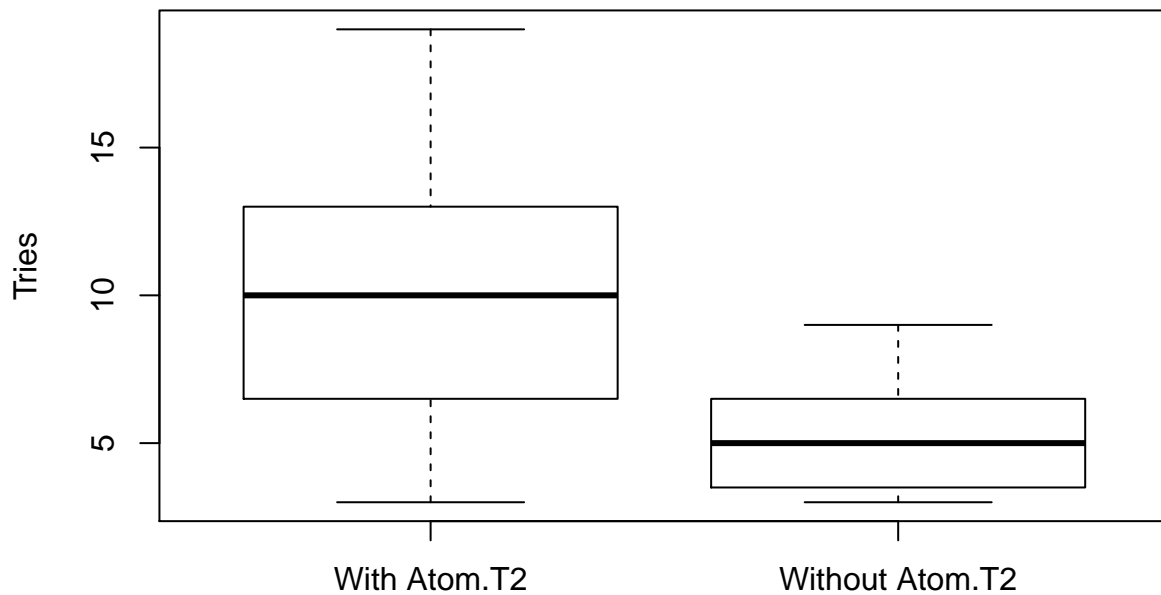
```
totalTime.gvlma = gvlma(lm(Time ~ Technique, data=totalTime))
summary(totalTime.gvlma)
```

```
##
## Call:
## lm(formula = Time ~ Technique, data = totalTime)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.0750 -0.3861  0.1105  0.5966  1.1729
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
```

```
## (Intercept)          5.0393      0.2169  23.232 < 2e-16 ***
## TechniqueWithout Atom -0.9643      0.3068  -3.143  0.00393 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.8401 on 28 degrees of freedom
## Multiple R-squared:  0.2608, Adjusted R-squared:  0.2344
## F-statistic: 9.881 on 1 and 28 DF,  p-value: 0.003928
##
##
## ASSESSMENT OF THE LINEAR MODEL ASSUMPTIONS
## USING THE GLOBAL TEST ON 4 DEGREES-OF-FREEDOM:
## Level of Significance = 0.05
##
## Call:
## gvlma(x = lm(Time ~ Technique, data = totalTime))
##
##              Value    p-value              Decision
## Global Stat      3.981e+01 4.734e-08 Assumptions NOT satisfied!
## Skewness         1.471e+01 1.257e-04 Assumptions NOT satisfied!
## Kurtosis         2.491e+01 6.006e-07 Assumptions NOT satisfied!
## Link Function    -1.813e-14 1.000e+00 Assumptions acceptable.
## Heteroscedasticity 1.963e-01 6.577e-01 Assumptions acceptable.
```

```
summary(aov(lm(Time ~ Technique, data=totalTime)))
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## Technique      1  6.974    6.974    9.881 0.00393 **
## Residuals     28 19.762    0.706
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## Kruskal-Wallis rank sum test
##
## data:  Time by Technique
## Kruskal-Wallis chi-squared = 9.1927, df = 1, p-value = 0.00243
```



```
totalTrials <- sqldf("select Replica, Id, SetOfTasks,
                      Technique, sum(Trials) as Trials
                      from ccwood where Tasks = 'AV2.1:C02.1:DE2.1' or Tasks = 'AV2.2:C02.2:DE2.2'
                      group by Replica, Id,SetOfTasks, Technique")
```

```
totalTrials$Trials <- ifelse(totalTrials$Trials == 0, 1, totalTrials$Trials)
totalTrials$Trials <- with(totalTrials, log2(Trials))
```

```
totalTrials$Replica = as.factor(totalTrials$Replica)
totalTrials$Id = as.factor(totalTrials$Id)
totalTrials$SetOfTasks = as.factor(totalTrials$SetOfTasks:totalTime$Id)
totalTrials$Technique = as.factor(totalTrials$Technique)
```

```
summary(aov(Trials ~ Technique, data=totalTrials))
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
## Technique  1  4.852   4.852   6.608 0.0158 *
## Residuals 28 20.560   0.734
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
totalTrials.gvlma = gvlma(lm(Trials ~ Technique, data=totalTrials))
summary(totalTrials.gvlma)
```

```
##
## Call:
## lm(formula = Trials ~ Technique, data = totalTrials)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.62191 -0.56706 -0.08059  0.39815  2.18244
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      3.2069      0.2213  14.494 1.53e-14 ***
## TechniqueWithout Atom -0.8044      0.3129  -2.571  0.0158 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.8569 on 28 degrees of freedom
## Multiple R-squared:  0.1909, Adjusted R-squared:  0.162
## F-statistic: 6.608 on 1 and 28 DF,  p-value: 0.01576
##
##
## ASSESSMENT OF THE LINEAR MODEL ASSUMPTIONS
## USING THE GLOBAL TEST ON 4 DEGREES-OF-FREEDOM:
## Level of Significance = 0.05
##
## Call:
## gvlma(x = lm(Trials ~ Technique, data = totalTrials))
##
##              Value p-value      Decision
## Global Stat      1.636e+00  0.8023 Assumptions acceptable.
## Skewness         1.544e+00  0.2140 Assumptions acceptable.
## Kurtosis         7.535e-02  0.7837 Assumptions acceptable.
## Link Function    7.228e-15  1.0000 Assumptions acceptable.
## Heteroscedasticity 1.623e-02  0.8986 Assumptions acceptable.
```

```
summary(aov(Trials ~ Technique, data=totalTrials))
```

```
##              Df Sum Sq Mean Sq F value Pr(>F)
## Technique      1  4.852   4.852   6.608 0.0158 *
## Residuals     28 20.560   0.734
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
ss = summary(aov(Trials ~ Technique, data=totalTrials))[[1]]$"Sum Sq"
eta.sq = ss[1]/(ss[1] + ss[2])
print(paste0("The eta-squared is ",toString(round(eta.sq,3))))
```

```
## [1] "The eta-squared is 0.191"
```

```
q <- TukeyHSD(aov(Trials~Technique, data=totalTrials))
q
```

```
## Tukey multiple comparisons of means
```

```
##      95% family-wise confidence level
##
## Fit: aov(formula = Trials ~ Technique, data = totalTrials)
##
## $Technique
##               diff      lwr      upr      p adj
## Without Atom-With Atom -0.8043511 -1.445296 -0.1634066 0.0157578
```

```
slices <- c(67, 33)
lbls <- c("Perfectiva", "Não Perfectiva")
pct <- round(slices/sum(slices)*100)
lbls <- paste(lbls, pct) # add percents to labels
lbls <- paste(lbls,"%",sep="") # ad % to labels
pie(slices,labels = lbls, col=rainbow(length(lbls))
)
```

