

Behaviour tracking platforms



A

Metadata

id	machine_name	date	...	condition	sex	...	p
xxx...xx x	machine_001	2016-09-01	...	A	M	...	p ₁
xxx...xx y	machine_001	2016-09-01	...	B	M	...	p ₂
xxx...xx z	machine_002	2016-09-03	...	A	F	...	p ₃
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
n	machine _n	date _n	...	condition _n	sex _n	...	p _n

Platform fields

Experiment fields

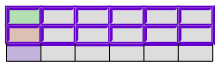
Data

id	t	activity	...	q
xxx...xx x	1	1	⋮	q _{1,1}
xxx...xx x	2	0	⋮	q _{1,2}
xxx...xx x	3	0	⋮	q _{1,3}
xxx...xx x	⋮	⋮	⋮	⋮
xxx...xx y	⋮	⋮	⋮	⋮
xxx...xx z	1	0	⋮	q _{3,1}
xxx...xx z	2	2	⋮	q _{3,2}
xxx...xx z	3	0	⋮	q _{3,3}
xxx...xx z	⋮	⋮	⋮	⋮
⋮	⋮	⋮	⋮	⋮
n	⋮	⋮	⋮	q _{n,k_n}

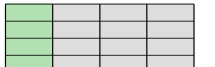
+

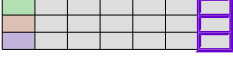
B

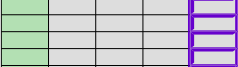
Metadata

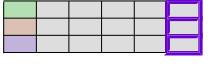
Select	dt[CRITERIA, meta = TRUE]  <pre>> male_meta <- dt[sex == "M", meta = TRUE]</pre>
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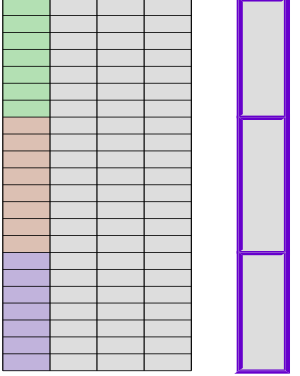
Data

dt[CRITERIA]  <pre>> late_dt <- dt[t > 5]</pre> <p>Note: metadata is updated when selection removes all data from one id.</p>
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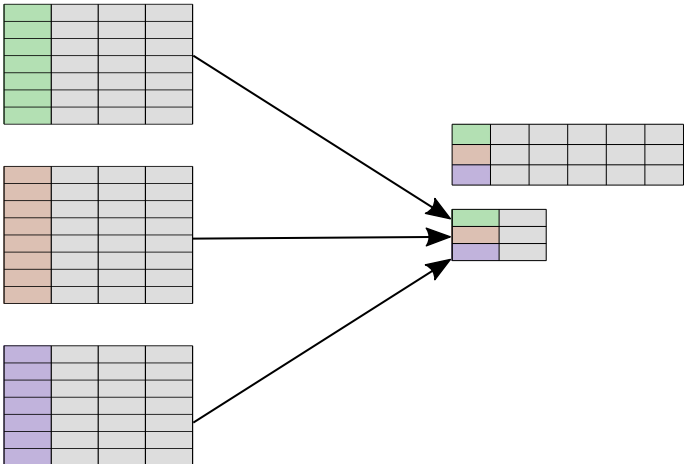
Alter, create & delete (meta)variables	dt[, X := value, meta = TRUE]  <pre>> dt[, genotype := "wt", meta = TRUE] > dt[, sex := NULL, meta = TRUE] # delete</pre>
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dt[, Y := value]  <pre>> dt[, t_2 := t-1] > dt[, t := NULL] # delete t</pre> <p>Note: update data in place. No copy of dt in memory.</p>
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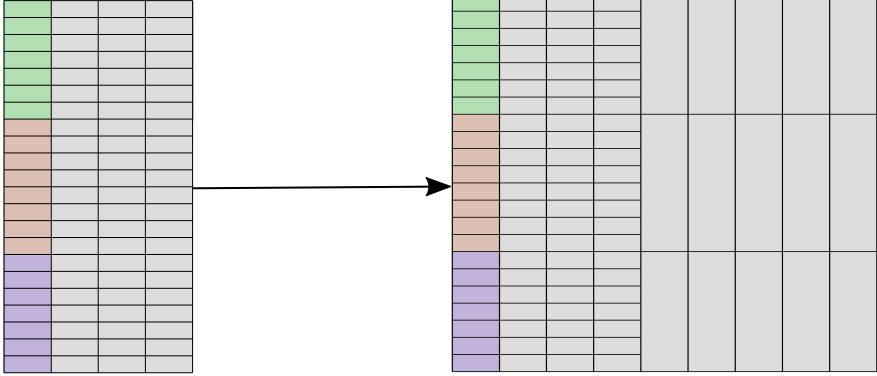
Expand metavariables as variables	dt[xmv(X)]  <pre>> dt <- dt[xmv(sex) == "M"] # select data with sex > dt[, s := xmv(sex)] # copy metavariable as variable</pre>
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Aggregate & summary	dt[, OPERATION, by = id] <pre>> # mean activity, per individual > dt <- dt[,.(mean_act = mean(activity)), by = id] > dt[, .N, by = id] # count reads per id</pre>
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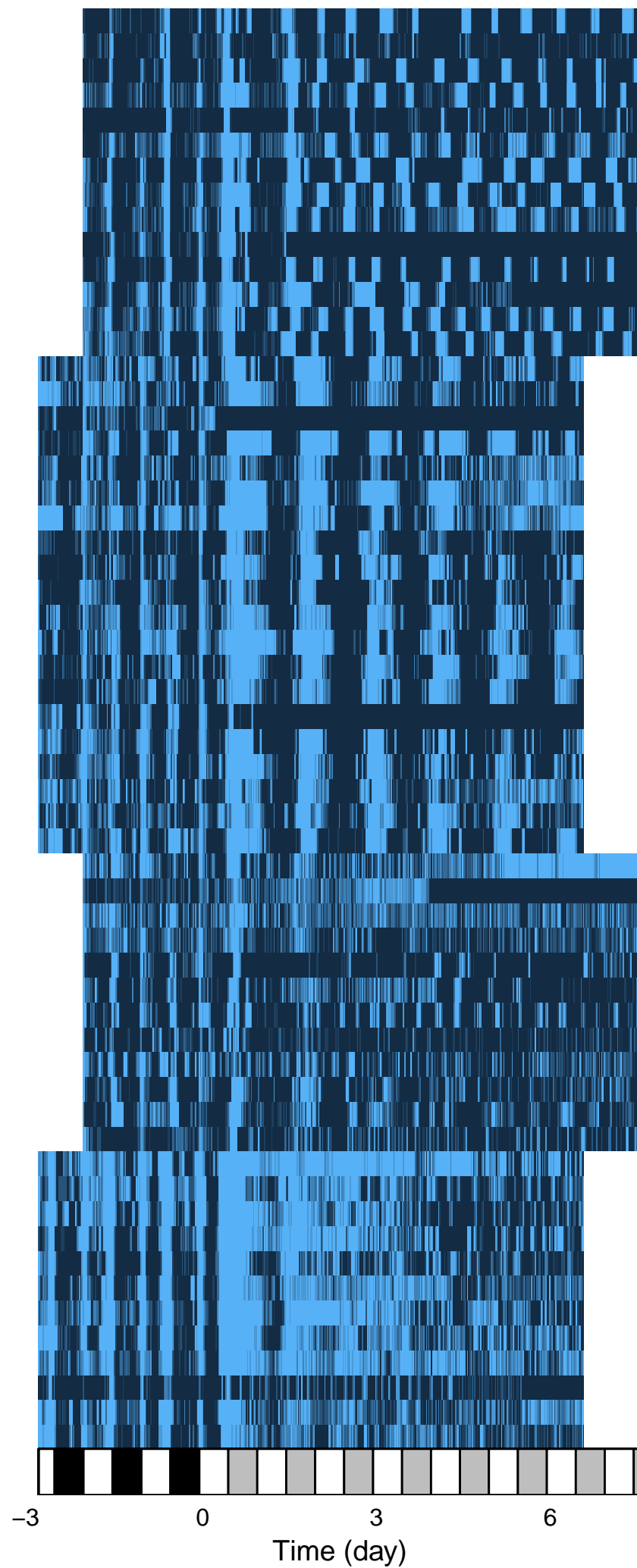
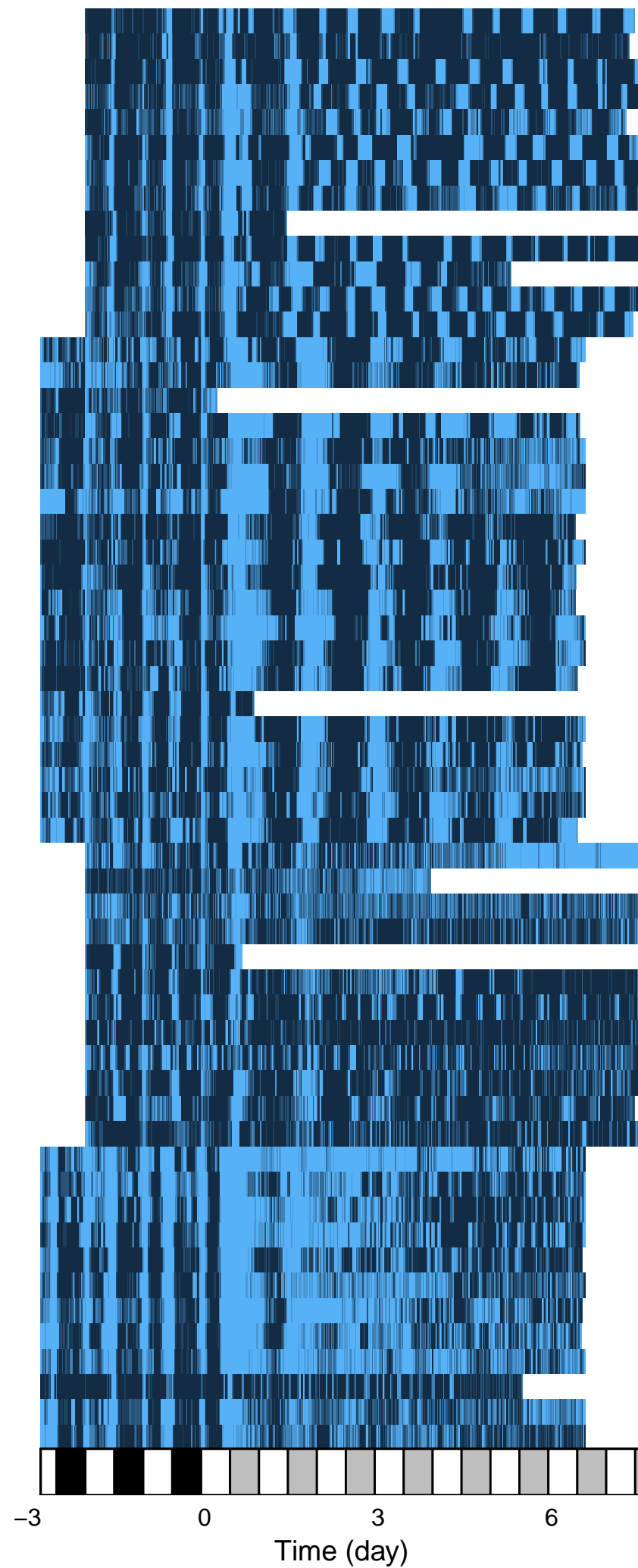
OPERATION 
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Join data & metadata	rejoin(dt) <pre>> full_table <- rejoin(dt)</pre> <p>Note: used mostly after aggregation or preprocessing</p>
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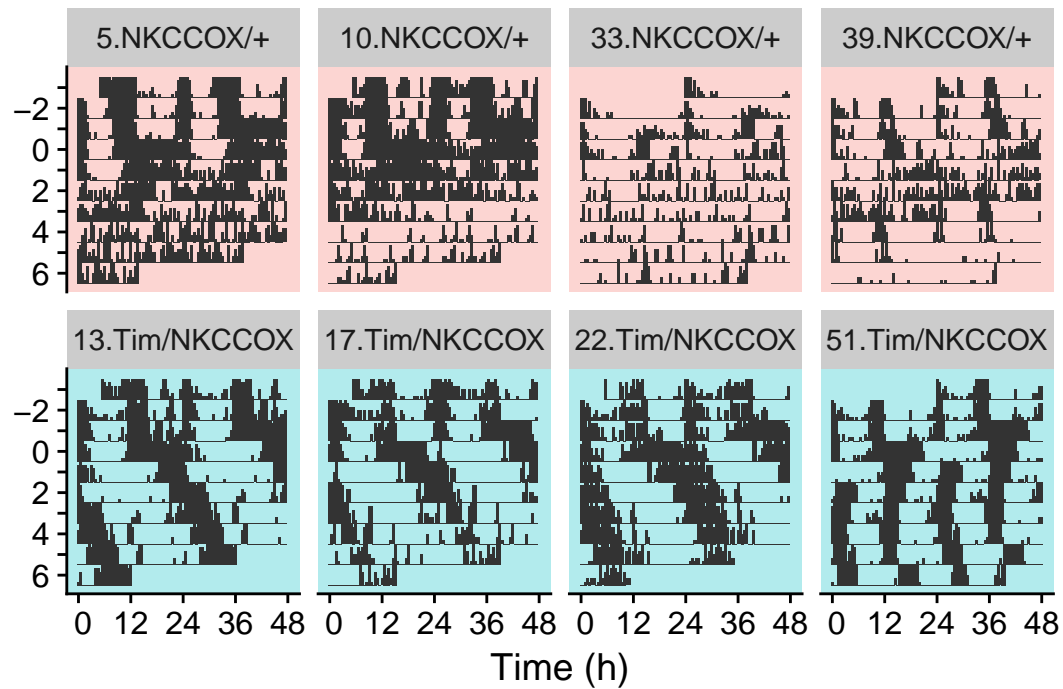
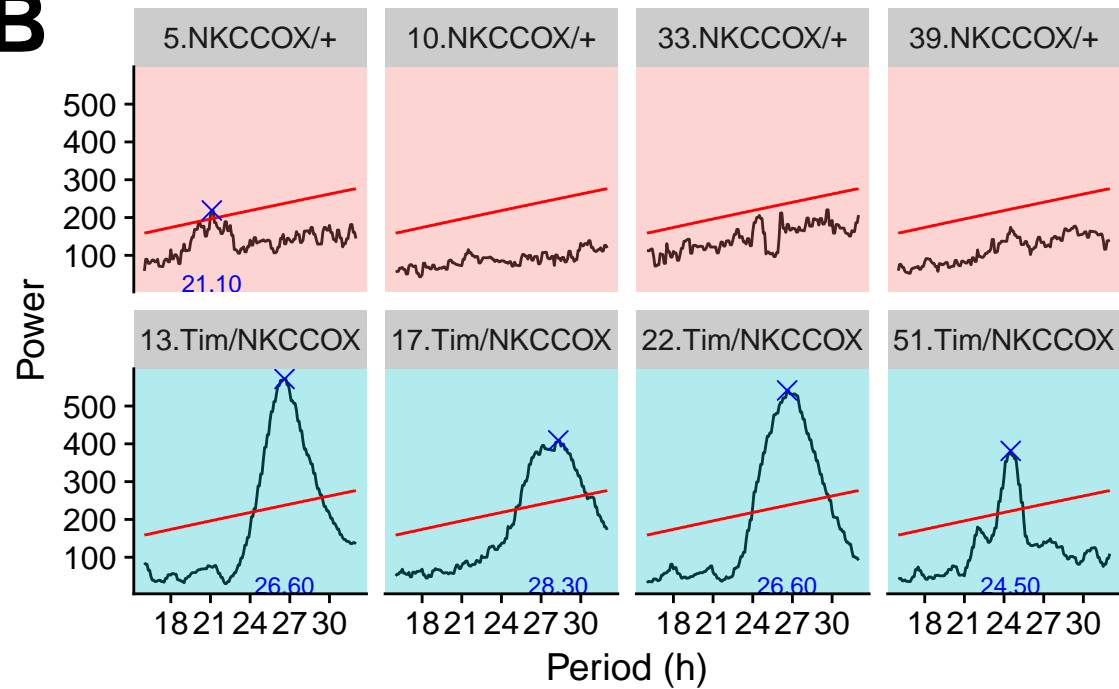
REJOIN 

A

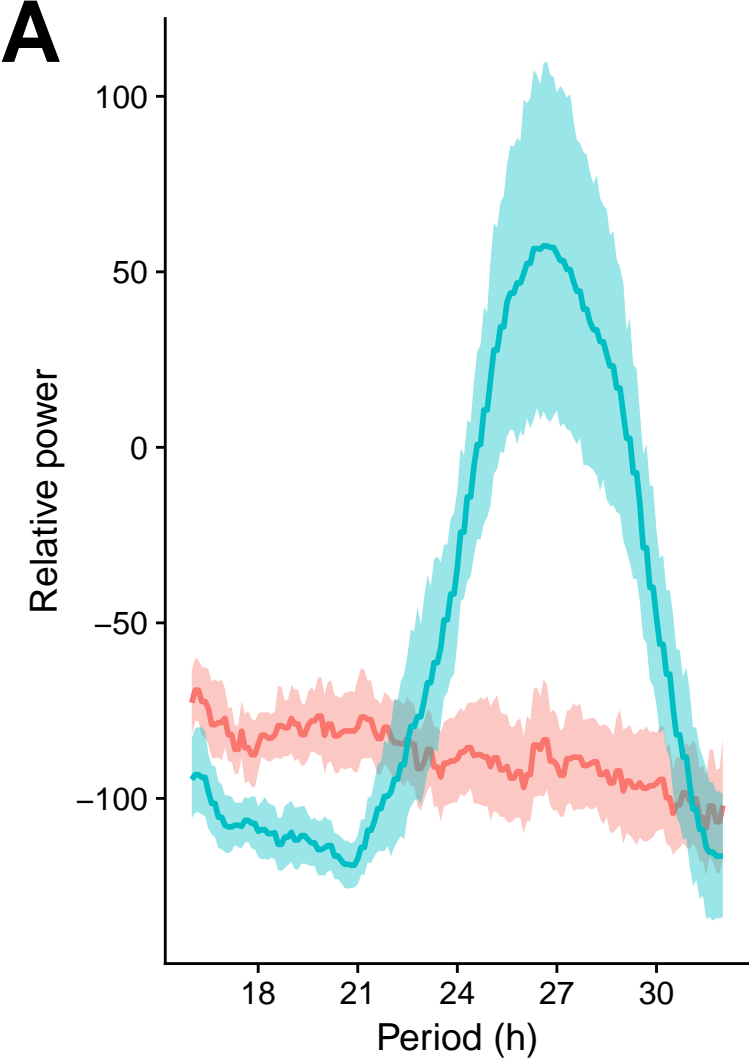
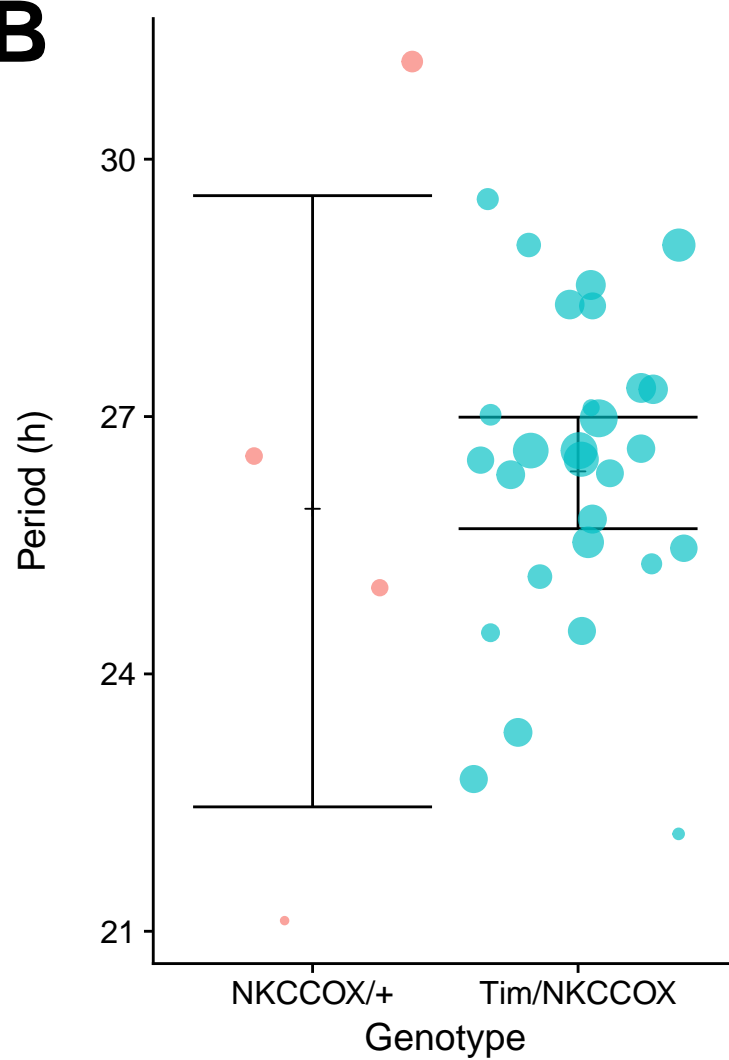
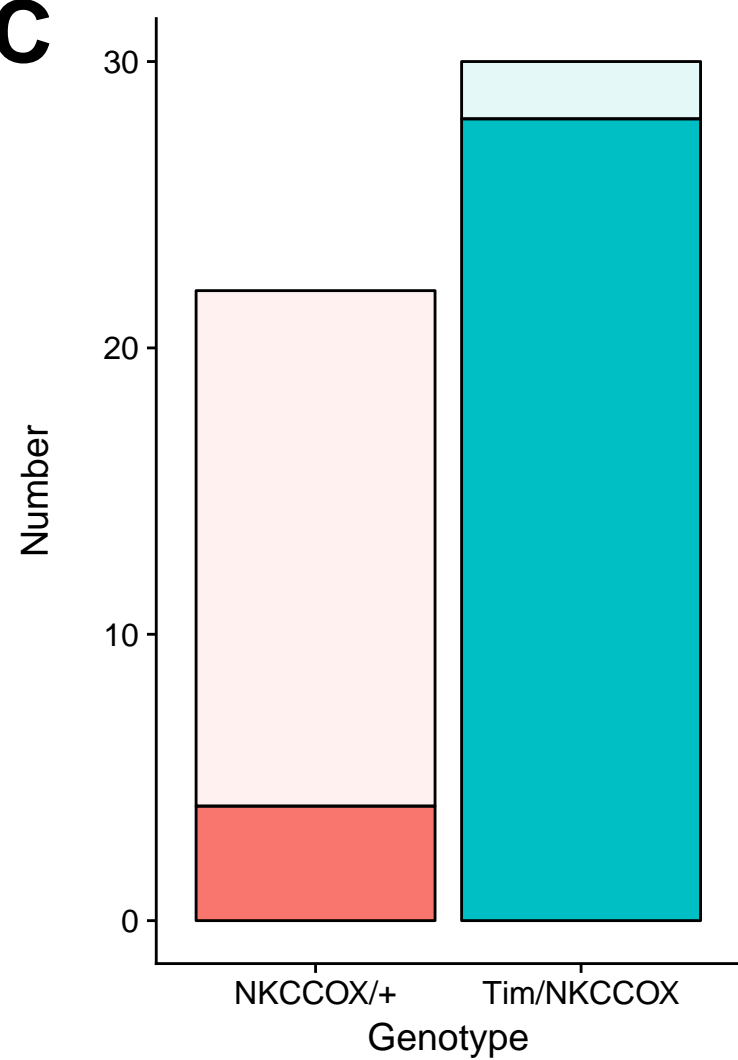
58.Tim/NKCCOX
57.Tim/NKCCOX
56.Tim/NKCCOX
55.Tim/NKCCOX
54.Tim/NKCCOX
53.Tim/NKCCOX
52.Tim/NKCCOX
51.Tim/NKCCOX
50.Tim/NKCCOX
49.Tim/NKCCOX
48.Tim/NKCCOX
47.Tim/NKCCOX
46.Tim/NKCCOX
45.Tim/NKCCOX
32.Tim/NKCCOX
31.Tim/NKCCOX
30.Tim/NKCCOX
29.Tim/NKCCOX
28.Tim/NKCCOX
27.Tim/NKCCOX
26.Tim/NKCCOX
25.Tim/NKCCOX
24.Tim/NKCCOX
23.Tim/NKCCOX
22.Tim/NKCCOX
21.Tim/NKCCOX
20.Tim/NKCCOX
19.Tim/NKCCOX
18.Tim/NKCCOX
17.Tim/NKCCOX
16.Tim/NKCCOX
15.Tim/NKCCOX
14.Tim/NKCCOX
13.Tim/NKCCOX
44.NKCCOX/+
43.NKCCOX/+
42.NKCCOX/+
41.NKCCOX/+
40.NKCCOX/+
39.NKCCOX/+
38.NKCCOX/+
37.NKCCOX/+
36.NKCCOX/+
35.NKCCOX/+
34.NKCCOX/+
33.NKCCOX/+
12.NKCCOX/+
11.NKCCOX/+
10.NKCCOX/+
9.NKCCOX/+
8.NKCCOX/+
7.NKCCOX/+
6.NKCCOX/+
5.NKCCOX/+
4.NKCCOX/+
3.NKCCOX/+
2.NKCCOX/+
1.NKCCOX/+

**B**

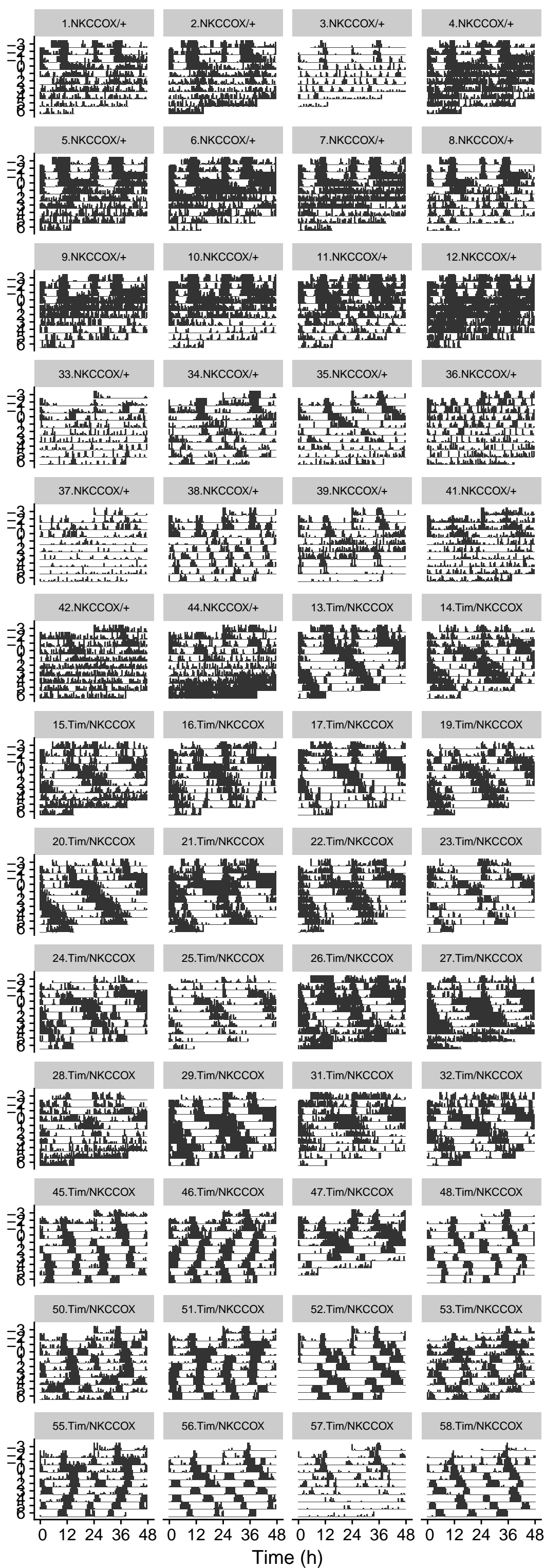
Moving 0% 50% 100%

A**B**

Genotype ■ NKCCOX/+ ■ Tim/NKCCOX

A**B****C**

Genotype ■ NKCCOX/+ ■ Tim/NKCCOX Peak power ● 300 ● 400 ● 500 ■ Arrhythmic ■ Rhythmic

A**B**