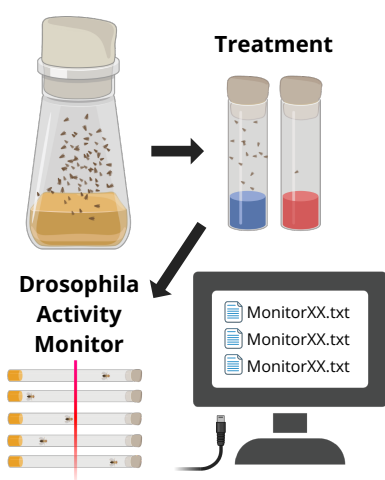
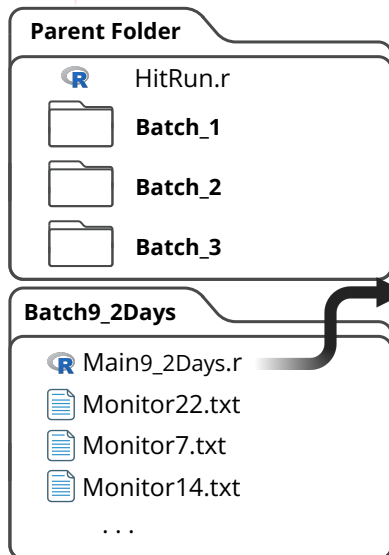


A



B



C

D

```

1 Title = "Batch9_2Days"
2 # -----
3 info <- data.table::data.table(
4
5   file = rep(c("Monitor22.txt", "Monitor7.txt",
6               "Monitor14.txt", "Monitor3.txt",
7               "Monitor15.txt", "Monitor39.txt",
8               "Monitor53.txt", "Monitor27.txt",
9               "Monitor52.txt", "Monitor17.txt",
10              "Monitor21.txt", "Monitor50.txt"), each = 32),
11
12   region_id = 1:32,
13   status = "OK",
14   start_datetime = "2019-08-17 10:00:00",
15   stop_datetime = "2019-08-19 10:20:00",
16   Sex = "M",
17   Genotype = rep(c("SIP-L2-2", "SIP-L2-2",
18                   "SIP-S2-8", "SIP-S2-8",
19                   "SIP-S2-1", "SIP-S2-1",
20                   "CS", "CS",
21                   "SIP-S2-10", "SIP-S2-10",
22                   "SIP-L2-1", "SIP-L2-1"), each = 32),
23
24   Temperature = "21.5C",
25   Treatment = rep(c("Iso", "Grp",
26                   "Iso", "Grp",
27                   "Iso", "Grp",
28                   "Iso", "Grp",
29                   "Iso", "Grp"), each = 32),
30
31   Environment = "2D",
32   Light = "12:12"
33 )

```