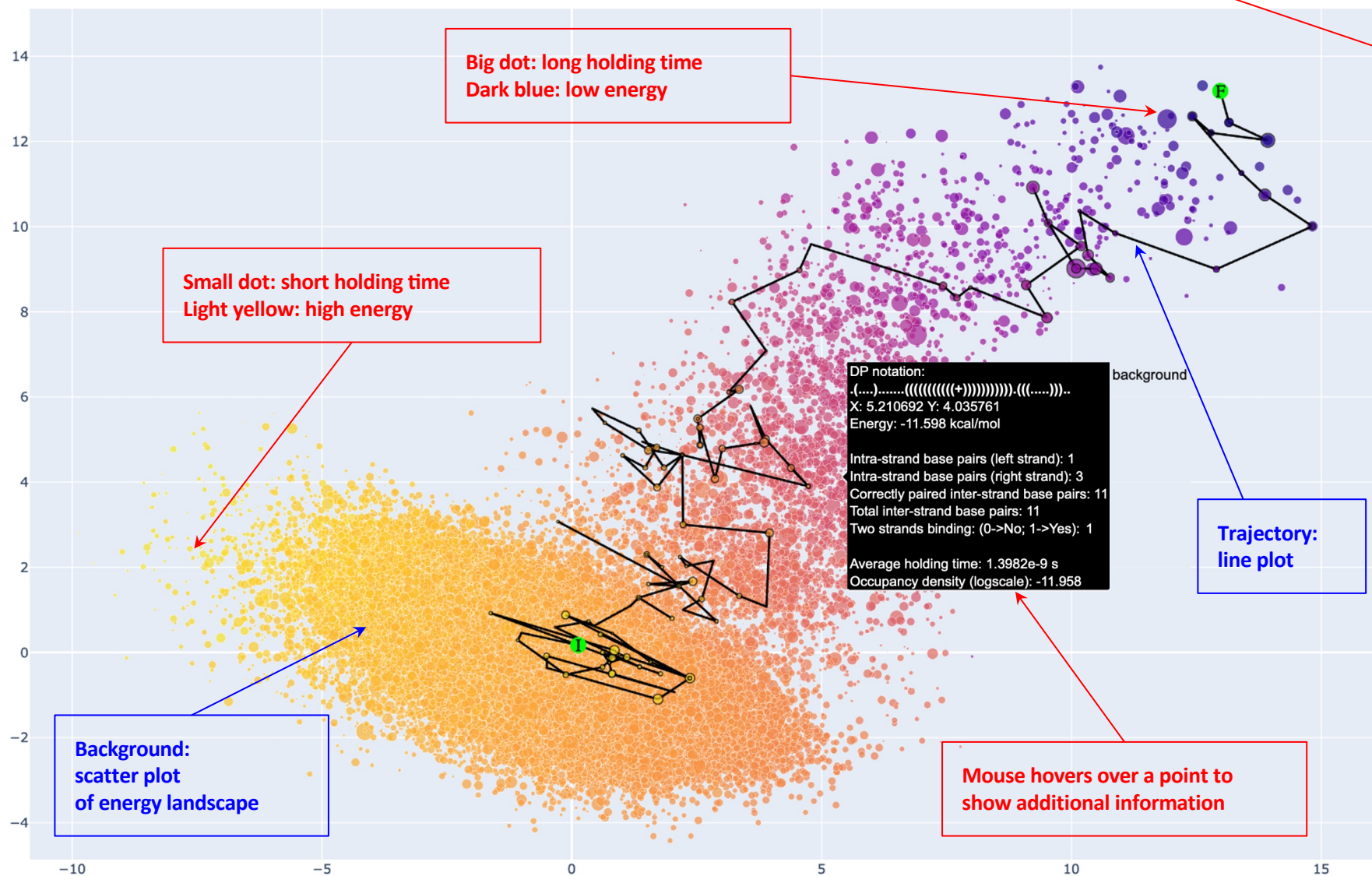
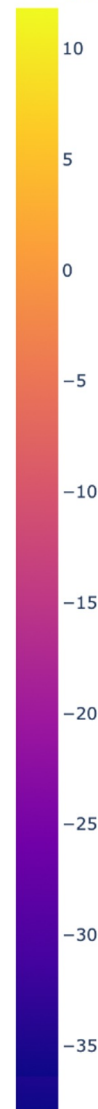


Free energy (kcal/mol)



DP notation:
.(....).....((((((((((+)))))))))).((((.....)))..
X: 5.210692 Y: 4.035761
Energy: -11.598 kcal/mol

Intra-strand base pairs (left strand): 1
Intra-strand base pairs (right strand): 3
Correctly paired inter-strand base pairs: 11
Total inter-strand base pairs: 11
Two strands binding: (0->No; 1->Yes): 1

Average holding time: 1.3982e-9 s
Occupancy density (logscale): -11.958

Click to show
the trajectory

Save
plot



Free
scaling



Select area



Zoom
in/out



Reset



Single Trajectory

- background
- trace 1
- trace 2
- trace 3
- trace 4
- trace 5
- trace 6
- trace 7
- trace 8
- trace 9
- trace 10
- trace 11
- trace 12
- trace 13
- trace 14
- trace 15
- trace 16
- trace 17
- trace 18
- trace 19
- trace 20
- trace 21
- trace 22
- trace 23
- trace 24
- trace 25
- trace 26
- trace 27
- trace 28
- trace 29
- trace 30
- trace 31
- trace 32
- trace 33
- trace 34
- trace 35
- trace 36
- trace 37
- trace 38
- trace 39