

First list the sessions in the database.

```
warning('off')
T = make_sessions_table
```

T = 35x18 table

...

| | sess_date | rat | sessid | mat_file_name | probe_serial | days_implanted |
|----|-------------|--------|--------|------------------------|---------------|----------------|
| 1 | 27-May-2019 | "K265" | 699605 | ""X:\RATTER\PhysDat... | "17131311562" | 5 |
| 2 | 30-May-2019 | "A242" | 700519 | "X:\RATTER\PhysData... | "17131311621" | 10 |
| 3 | 03-Jun-2019 | "A242" | 701531 | "X:\RATTER\PhysData... | "17131311621" | 14 |
| 4 | 06-Jun-2019 | "A242" | 702016 | "X:\RATTER\PhysData... | "17131311621" | 17 |
| 5 | 07-Jun-2019 | "A242" | 702361 | "X:\RATTER\PhysData... | "17131311621" | 18 |
| 6 | 10-Jun-2019 | "A242" | 703121 | "X:\RATTER\PhysData... | "17131311621" | 21 |
| 7 | 12-Jun-2019 | "A242" | 703771 | "X:\RATTER\PhysData... | "17131311621" | 23 |
| 8 | 20-Jun-2019 | "A242" | 706158 | "X:\RATTER\PhysData... | "17131311621" | 31 |
| 9 | 28-Jun-2019 | "A242" | 708534 | "X:\RATTER\PhysData... | "17131311621" | 39 |
| 10 | 15-Jul-2019 | "A230" | 713429 | "X:\RATTER\PhysData... | "17131308571" | 13 |
| 11 | 26-Sep-2019 | "A241" | 734656 | "X:\RATTER\PhysData... | "18194823302" | 15 |
| 12 | 26-Sep-2019 | "A243" | 734648 | "X:\RATTER\PhysData... | "18194824132" | 13 |
| 13 | 30-Sep-2019 | "X046" | 735284 | "X:\RATTER\PhysData... | "18194823122" | 15 |
| 14 | 09-Oct-2019 | "A241" | 737985 | "X:\RATTER\PhysData... | "18194823631" | 28 |
| 15 | 10-Oct-2019 | "A243" | 738225 | "X:\RATTER\PhysData... | "18194824132" | 27 |
| 16 | 29-Oct-2019 | "A243" | 743494 | "X:\RATTER\PhysData... | "18194824132" | 46 |
| 17 | 01-Nov-2019 | "A243" | 744398 | "X:\RATTER\PhysData... | "18194824132" | 49 |
| 18 | 11-Dec-2019 | "A241" | 753483 | "X:\RATTER\PhysData... | "18194823631" | 91 |
| 19 | 14-Jan-2020 | "A241" | 760297 | "X:\RATTER\PhysData... | "18194823631" | 125 |
| 20 | 22-Jan-2020 | "A241" | 762805 | "X:\RATTER\PhysData... | "18194823631" | 133 |
| 21 | 23-Jan-2020 | "A241" | 763087 | "X:\RATTER\PhysData... | "18194823631" | 134 |
| 22 | 11-Feb-2020 | "A249" | 768938 | "X:\RATTER\PhysData... | "18194819132" | 7 |
| 23 | 04-Sep-2020 | "A249" | 784371 | "X:\RATTER\PhysData... | "18194819132" | 213 |
| 24 | 07-Sep-2020 | "A249" | 784741 | "X:\RATTER\PhysData... | "18194819132" | 216 |
| 25 | 07-Sep-2020 | "A256" | 784742 | "X:\RATTER\PhysData... | "18194819321" | 6 |
| 26 | 09-Sep-2020 | "A249" | 785000 | "X:\RATTER\PhysData... | "18194819132" | 218 |
| 27 | 11-Sep-2020 | "A249" | 785272 | "X:\RATTER\PhysData... | "18194819132" | 220 |
| 28 | 11-Sep-2020 | "A256" | 785275 | "X:\RATTER\PhysData... | "18194819321" | 10 |

| | sess_date | rat | sessid | mat_file_name | probe_serial | days_implanted |
|----|-------------|--------|--------|-------------------------|---------------|----------------|
| 29 | 23-Sep-2020 | "A249" | 786975 | "X:\RATTER\PhysData..." | "18194819132" | 232 |
| 30 | 23-Sep-2020 | "A256" | 786978 | "X:\RATTER\PhysData..." | "18194819321" | 22 |
| 31 | 25-Sep-2020 | "A249" | 787269 | "X:\RATTER\PhysData..." | "18194819132" | 234 |
| 32 | 18-Sep-2020 | "A249" | 786254 | "X:\RATTER\PhysData..." | "18194819132" | 227 |
| 33 | 18-Sep-2020 | "A256" | 786258 | "X:\RATTER\PhysData..." | "18194819321" | 17 |
| 34 | 21-Sep-2020 | "A256" | 786682 | "X:\RATTER\PhysData..." | "18194819321" | 20 |
| 35 | 28-Sep-2020 | "A256" | 787674 | "X:\RATTER\PhysData..." | "18194819321" | 27 |

Load one of them (the last one, just for examples).

```
Cells = load(T.mat_file_name{end})
```

Cells = *struct with fields:*

```

    rec: [1x1 struct]
    jrc_file: 'E:\Adrian\A256\A256_2020_09_28\A256_2020_09_28_g0\spikesort_2020_11_23_09_09_42_ks2jrc\A256...'
    mat_file_name: 'E:\Adrian\A256\A256_2020_09_28\A256_2020_09_28_g0\spikesort_2020_11_23_09_09_42_ks2jrc\A256...'
    made_by: 'make_Cells_from_JRC'
    params: 'time_s_from_last_ts_to_include_spike'
    Trials: [1x1 struct]
    nTrials: 479
    sessid: 787674
    last_modified: 25-Nov-2020 14:47:22
    rat: 'A256'
    sess_date: '2020-09-28'
    bank: [368x1 double]
    electrode: [368x1 double]
    unitCount: [368x1 double]
    unitISIRatio: [1x368 double]
    unitLRatio: [1x368 double]
    unitIsoDist: [1x368 double]
    unitVppRaw: [368x1 single]
    meanWfGlobalRaw: [61x306x368 single]
    clusterNotes: {368x1 cell}
    waveformSim: [368x368 double]
    raw_spike_time_s: {368x1 cell}
    firing_rate: [1x368 double]
    spike_time_s: [1x1 struct]
    kSpikeWindowS: [1x1 struct]
    ks_good: [1x368 logical]
    frac_isi_violation: [368x1 double]
    waveform: [1x368 struct]
    recorded: {368x1 cell}
    distance_from_tip: [368x1 double]
    penetration: [1x1 struct]
    hemisphere: 'right'
    probe_serial: '18194819321'
    DV: [368x1 double]
    ML: [368x1 double]
    AP: [368x1 double]
    regions: [1x368 double]

```

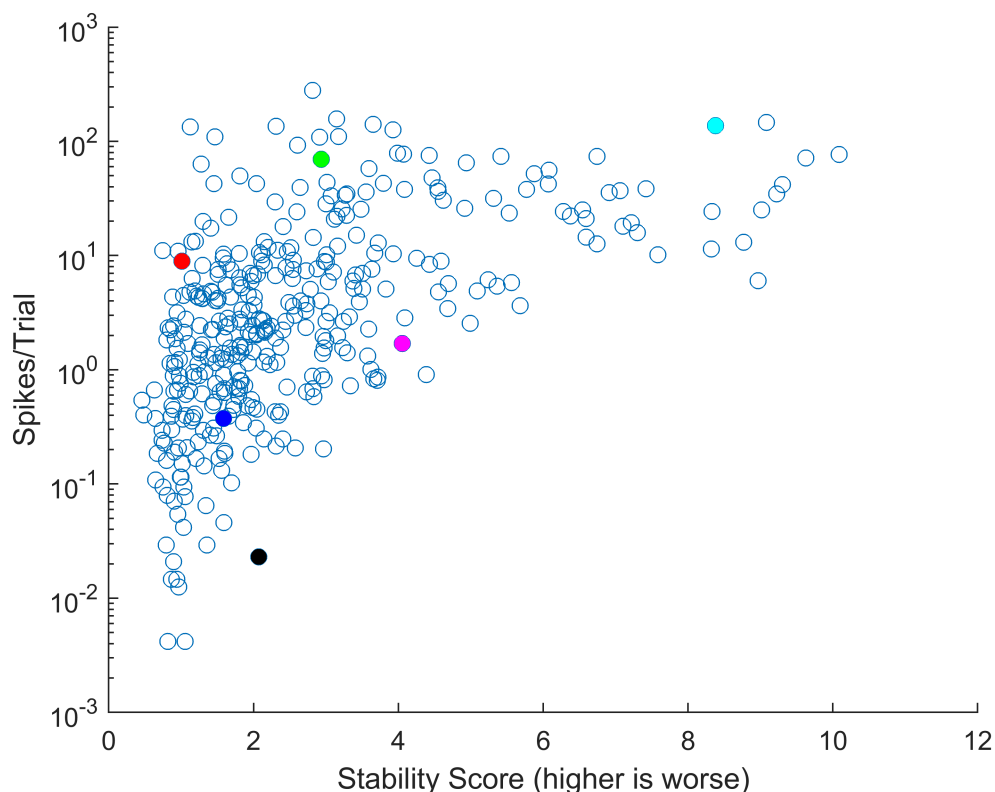
Use the new utility function "calculate_unit_stability" to get the stability metric and number of spikes per trial for each cell. The stability metric is the ratio of sd's of two distributions of trial-averaged firing rates A and B. A is the distribution obtained by repeatedly sampling groups of contiguous trials. B is the distribution obtained

by repeatedly sampling random groups of trials. If the ratio is large, it means sampling groups of trials that are contiguous in time gives a broader distribution of rates than you'd expect -- i.e. the variability in firing rate across time is systematic. For a neuron whose firing rate statistics do not change across the session, the stability metric would be exactly 1.

```
[stability,presence,nspks] = calculate_unit_stability(Cells)
```

```
stability = 1×368
    8.7693    2.5784    4.2533    4.0718    3.2662    1.8006    1.6544    2.4046 ...
presence = 1×368
    0.9875    0.1065    0.9958    1.0000    1.0000    0.8706    0.1795    0.5699 ...
nspks = 1×368
    13.1127    0.2067    9.4593    77.3779    33.7244    10.4280    0.3925    2.2296 ...
```

```
cla;scatter(stability,nspks);hold on
examples = [156 267 234 246 361 37];
colors ={'r','g','c','m','k','b'};
for i=1:6
    scatter(stability(examples(i)),nspks(examples(i)),colors{i},'filled');hold on;
end
xlabel('Stability Score (higher is worse)')
ylabel('Spikes/Trial')
set(gca,'yscale','log')
```

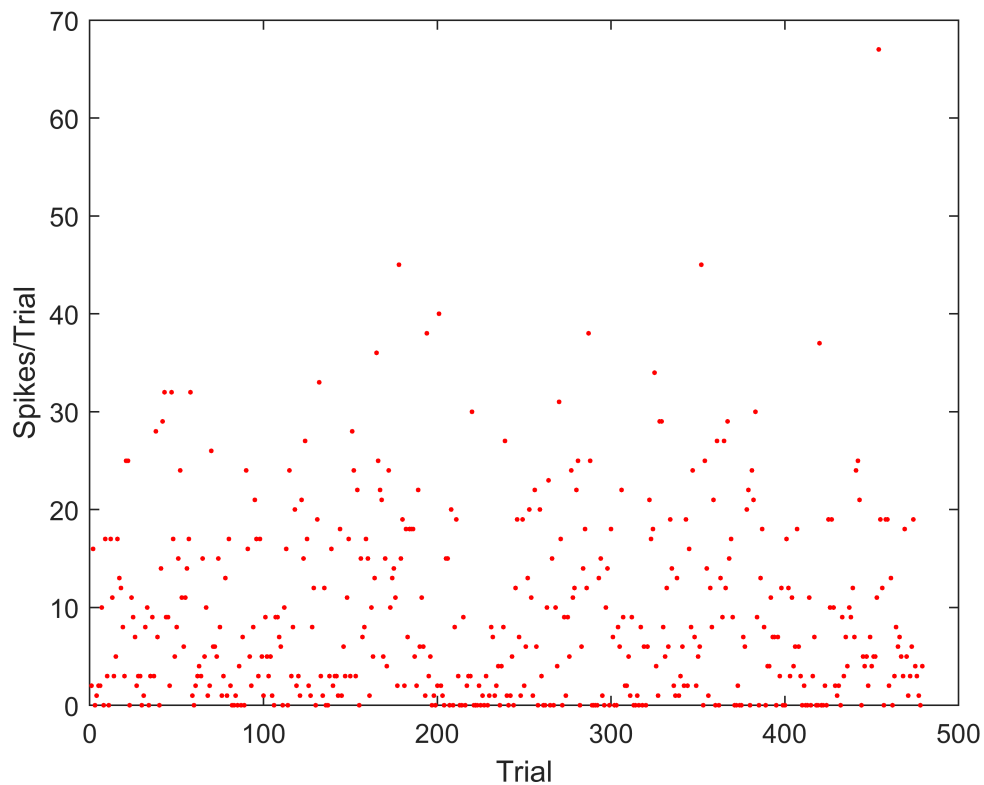


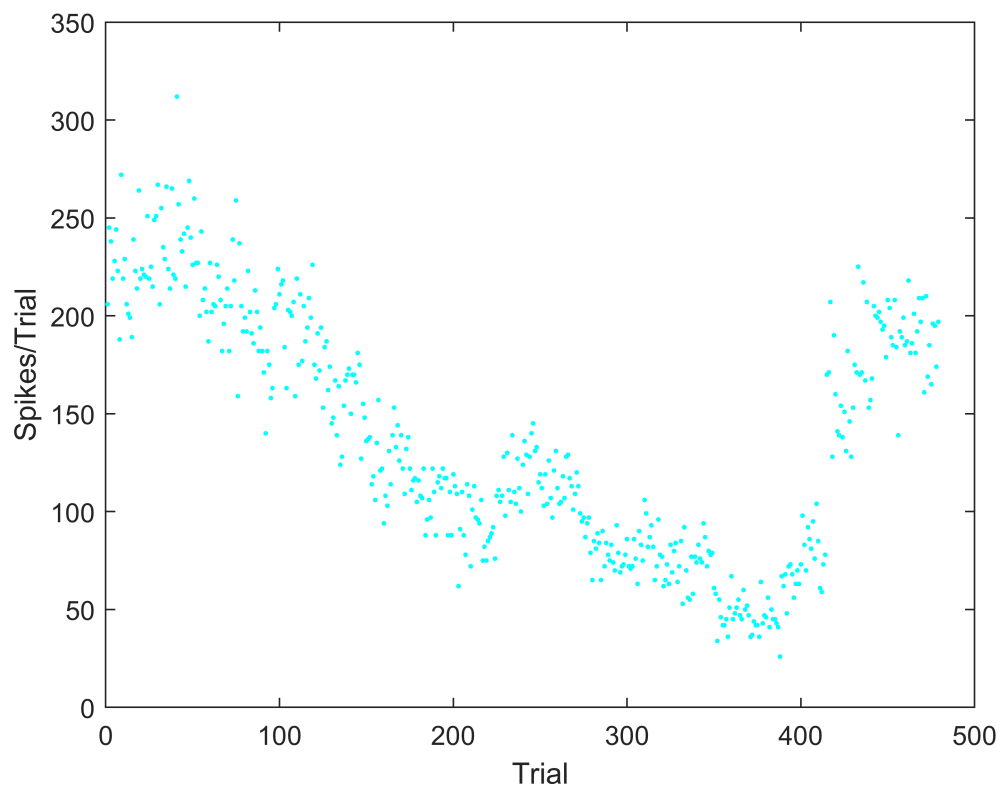
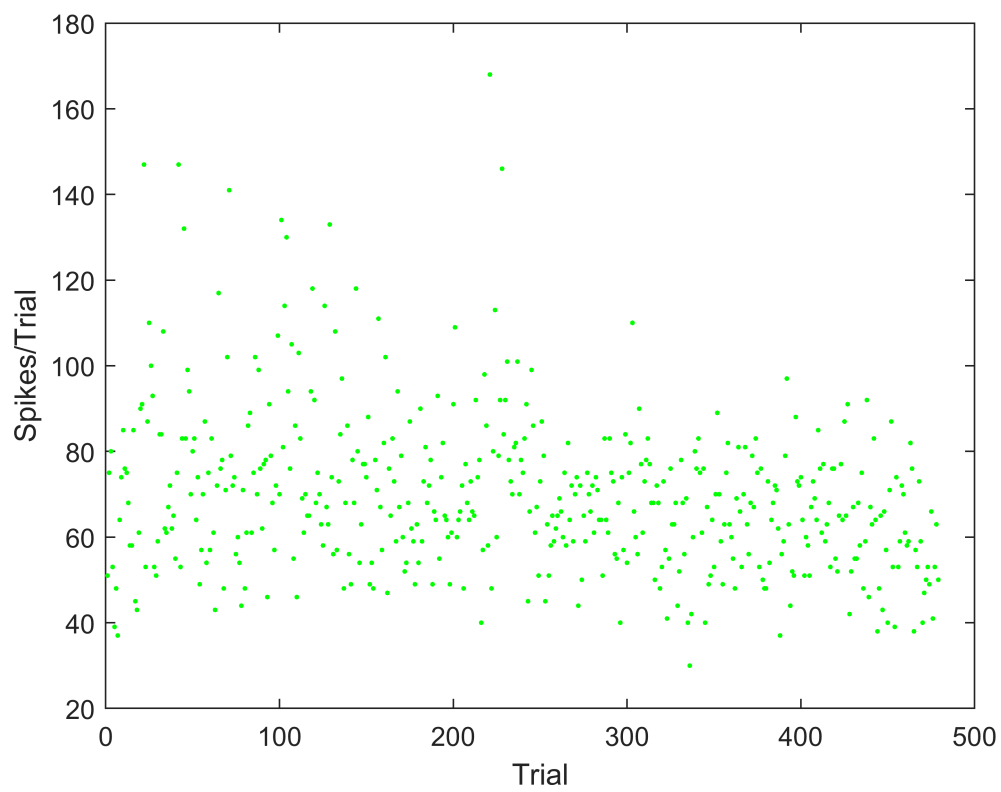
What you can see is that the majority of cells form a cluster with stability slightly above 1 (i.e. some excess variability across time but not much). Then there is a long tail of cells with stability values much greater than 1.

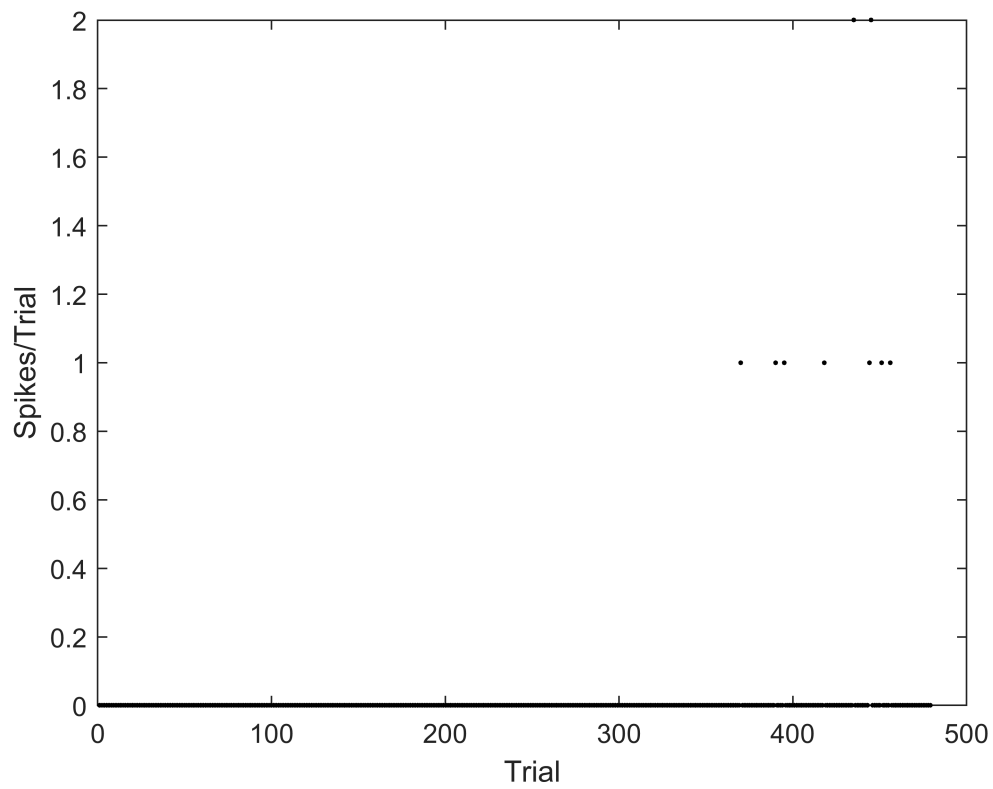
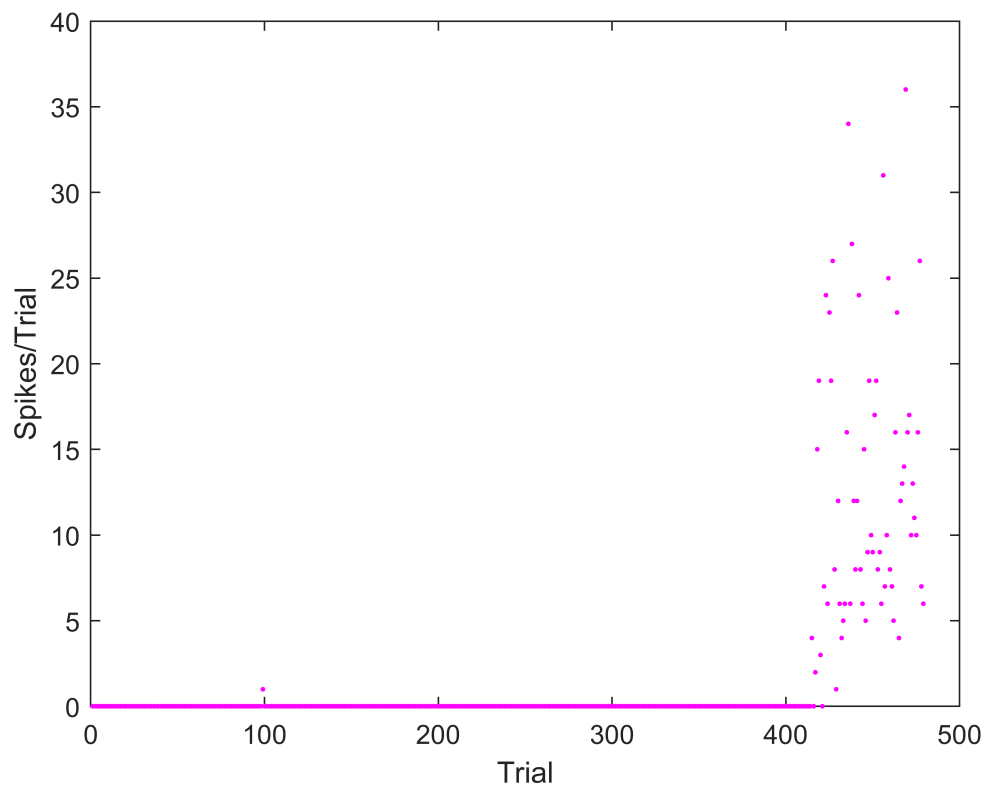
Cells with very low firing rates do not show large stability scores because Poisson variability (which shows up in both the A and B distribution) swamps variability across time. You probably want to exclude these cells anyway.

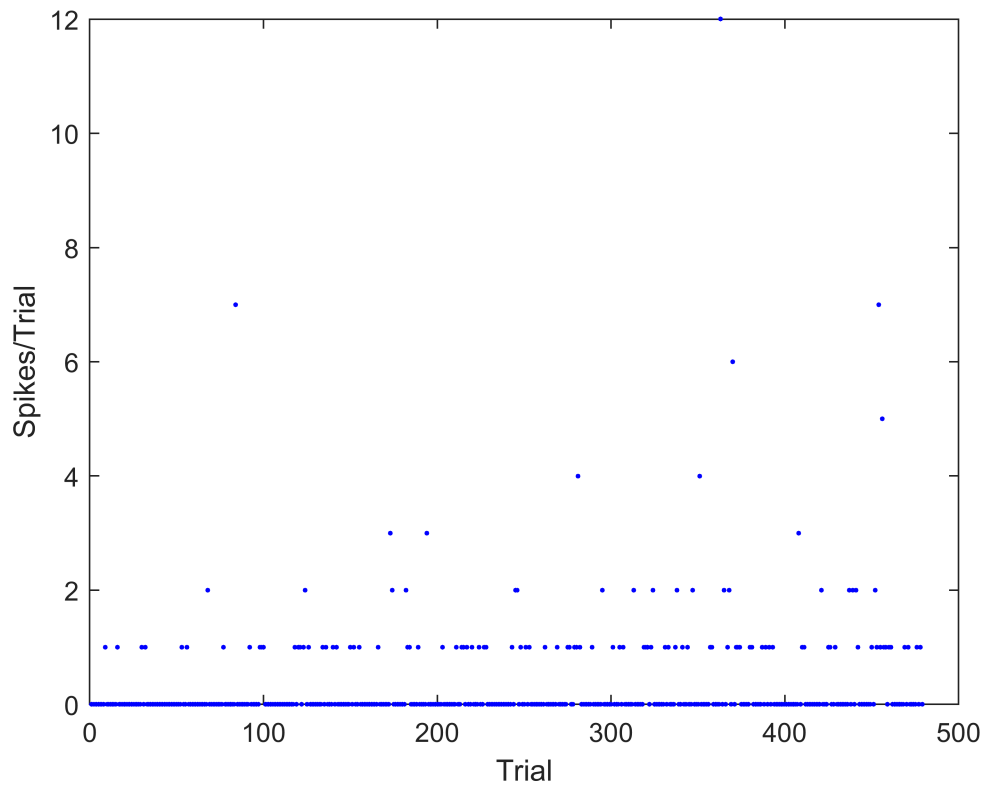
To get a sense of what the firing rate of cells in the above distribution look like, I've flagged 6 example cells. I've plotted spikes/trial across the session for each cell. A threshold of about 4 or 5 for the stability score seems reasonable to exclude cells which are truly anomalously unstable. This would exclude 10-15% of cells for this session.

```
for i=1:6
    figure;plot(cellfun(@numel,Cells.spike_time_s.cpoke_in{examples(i)}),sprintf('.%s',colors{i}
end
```









The function "calculate_unit_stability" also returns a value called "presence." This is the fraction of trials for which a unit spiked. Excluding cells for which that value is less than 0.5 (i.e. fired on less than half of trials) seems also reasonable and excludes the low firing rate cells (about 1/3 of cells).

```
figure
scatter(stability,nspsks)
hold on
scatter(stability(presence<0.5),nspsks(presence<0.5),'k','filled')
set(gca,'yscale','log')
xlabel('Stability Score (higher is worse)')
ylabel('Spikes/Trial')
```

