**Figure 2–source data 1: Summary statistics for Figure 2**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Fig. 2b: MEMBRANE CAPACITANCE (pF)** |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean** **± SD** | 121.6 ± 36.6 | 402.3 ± 105.4 | 462.4± 147.0 | 109.6, 2, 20 | <0.0001 | -130.8 ± 38.1 | 205.0 ± 69.4 | 340.0 ± 142.5 | 295.9 ± 109.2 | 37.77, 3, 127 | <0.0001 |
| **No. of animals** | 10 | 5 | 8 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 1-7 | 4-8 | 1-7 |   |   | 3-8 | 4-8 | 1-8 | 1-5 |   |   |
| **Total cells** | 43 | 25 | 35 |   |   | 48 | 28 | 28 | 27 |   |   |
| **Fig. 2c: MEMBRANE CONDUCTANCE (nS)** |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean** **± SD** | 10.6 ± 3.8 | 20.1± 5.3 | 23.7 ± 7.4 | 33.58, 2, 18 | <0.0001 | 8.0 ± 2.4 | 13.1 ± 4.5 | 18.6 ± 9.2 | 13.8 ± 4.1 | 17.83, 3, 25 | <0.0001 |
| **No. of animals** | 8 | 5 | 8 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 3-6 | 4-8 | 1-6 |   |   | 3-8 | 4-8 | 1-8 | 1-5 |   |   |
| **Total cells** | 34 | 24 | 30 |   |   | 47 | 27 | 27 | 28 |   |   |
| **Fig. 2d: RMP (mV)** |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean** **± SD** | -74.5± 3.8 | -69.0 ± 5.1 | -64.9 ± 5.3 | 24.20, 2, 20 | <0.0001 | -69.7 ± 5.0 | -70.0 ± 5.3 | -67.6 ± 5.3 | -64.7 ± 7.3 | 4.292, 3, 25 | 0.0142 |
| **No. of animals** | 10 | 5 | 8 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 1-7 | 4-8 | 1-7 |   |   | 3-8 | 4-8 | 1-8 | 1-5 |   |   |
| **Total cells** | 42 | 25 | 36 |   |   | 52 | 28 | 28 | 31 |   |   |
| **Fig. 2f: AP INPUT-OUTPUT CURVE** |  |  |  |  |  |
|  | **No. of neurons** | **Mixed-effects ANOVA** |  | **No. of neurons** | **Mixed-effects ANOVA** |
|  | Fixed effects (type III) | **F (DFn, DFd)** | **p-value** |  | Fixed effects (type III) | **F (DFn, DFd)** | **p-value** |
| **Control** | 36 | **Current step** | F (20, 1657) = 263.4 | <0.0001 | **Control** | 50 | **Current step** | F (20, 2579) = 600.7 | <0.0001 |
| ***RhebY35L*** | 22 | **Group** | F (2, 83) = 39.60 | <0.0001 | ***Depdc5KO*** | 27 | **Group** | F (3, 129) = 22.84 | <0.0001 |
| ***MTORS2215Y*** | 28 | **Current step x group** | F (40, 1657) = 22.84 | <0.0001 | ***PtenKO*** | 28 | **Current step x group** | F (60, 2579) = 14.66 | <0.0001 |
|  |  |  |  |   |   | ***Tsc1KO*** | 28 |  |  |  |  |
| **Fig. 2g: RHEBOBASE (pA)** |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean****± SD** | 544.4 ± 225.3 | 1356.0 ± 634.6 | 1422.0 ± 624.2 | 26.74, 2, 77 | <0.0001 | 484.4 ± 180.1 | 737.3 ± 315.2 | 1071.0 ± 503.5 | 935.7 ± 444.4 | 16.84, 3, 25 | <0.0001 |
| **No. of animals** | 7 | 5 | 6 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 4-6 | 4-7 | 1-7 |   |   | 3-7 | 4-7 | 1-8 | 1-5 |   |   |
| **Total cells** | 32 | 24 | 24 |   |   | 50 | 27 | 28 | 28 |   |   |
| **Fig. 2h: 1st ISI (ms)** |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |  |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean****± SD** | 15.9 ± 6.4 | 28.7 ± 9.8 | 24.3 ± 10.8 | 9.386, 2, 18 | 0.0016 | 15.5 ± 7.0 | 22.7 ± 9.1 | 12.2 ± 8.0 | 23.1 ± 10.3 | 8.105, 3, 24 | 0.0007 |
| **No. of animals** | 9 | 5 | 7 |   |   | 10 | 5 | 5 | 8 |   |   |
| **No. cells/ animal** | 1-6 | 1-4 | 1-7 |   |   | 3-7 | 4-7 | 2-5 | 2-4 |   |   |
| **Total cells** | 36 | 16 | 25 |   |   | 50 | 27 | 18 | 23 |   |   |
| **Fig. 2i: AP THRESHOLD (mV)**  |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean** **± SD** | -39.2 ± 6.3 | -44.0± 12.3 | -41.0 ± 8.4 | 1.937, 2, 78 | 0.1511 | -35.8± 9.4 | -39.0 ± 7.9 | -40.0 ± 14.4 | -37.9 ± 11.1 | 0.6587, 3, 25 | 0.5852 |
| **No. of animals** | 7 | 5 | 6 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 4-6 | 3-7 | 1-6 |   |   | 2-6 | 3-7 | 1-7 | 1-5 |   |   |
| **Total cells** | 33 | 23 | 25 |   |   | 42 | 25 | 26 | 25 |   |   |
| **Fig. 2j: AP PEAK AMPLITUDE (mV)**  |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean** **± SD** | 138.1 ± 10.1 | 139.0 ± 11.6 | 120.3 ± 8.9 | 20.51, 2, 15 | <0.0001 | 135.0 ± 9.1 | 136.1± 12.9 | 138.3 ± 21.1 | 126.1 ± 7.8 | 4.239, 3, 116 | 0.0070 |
| **No. of animals** | 7 | 5 | 6 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 4-6 | 3-7 | 1-6 |   |   | 2-6 | 3-7 | 1-7 | 1-5 |   |   |
| **Total cells** | 33 | 23 | 25 |   |   | 43 | 25 | 27 | 25 |   |   |
| **Fig. 2k: AP HALF-WIDTH (ms)** |  |  |  |  |  |
|  |  |  |  | **Nested one-way ANOVAa** |  |  |  |  | **Nested one-way ANOVAa** |
|  | **Control** | ***RhebY35L*** | ***MTORS2215Y*** | **F, DFn, DFd** | **p-value** | **Control** | ***Depdc5KO*** | ***PtenKO*** | ***Tsc1KO*** | **F, DFn, DFd** | **p-value** |
| **Mean** **± SD** | 1.79 ± 0.30 | 1.50 ± 0.18 | 1.44 ± 0.14 | 18.45, 2, 15 | <0.0001 | 1.76 ± 0.43 | 1.65± 0.23 | 1.44 ± 0.24 | 1.52 ± 0.23 | 3.679, 3, 25 | 0.0254 |
| **No. of animals** | 7 | 5 | 6 |   |   | 10 | 5 | 6 | 8 |   |   |
| **No. cells/ animal** | 4-6 | 3-7 | 1-6 |   |   | 2-6 | 3-7 | 1-7 | 1-5 |   |   |
| **Total cells** | 33 | 23 | 25 |   |   | 43 | 25 | 27 | 25 |   |   |

aThe nested one-way ANOVA fits a mixed-effects model wherein the main factor is treated as a fixed factor and the nested factor is treated as a random factor.

bPost-hoc analyses were performed using Holm-Šídák multiple comparison test. Significant post-hoc results (p<0.05) are denoted with symbols (\*, #, Ɏ) on the graphs, with the number of symbols 1-4 denoting the significant levels p<0.05, <0.01, <0.001, and <0.0001, respectively. For all two-way repeated measured and mixed-effects model ANOVA, all significant results (p<0.05) are denoted with one symbol regardless of the significant level for clearness on the graphs.