



**Figure S1. Resting membrane potential of 12 PD neurons does not change after 8 hr. of impalement with two microelectrodes, Related to Figure 2.** Each line shows the membrane potential measurement of a single neuron at the start of the experiment in TTX (t=0) and then after 8 hr. of two-electrode voltage clamp (t=8). *P* value generated using a two-tailed paired t test.

<u>Activity-Dependent Correlations</u>	<u>p values</u>			<u>Pearson's <i>r</i></u>		
	Control	Silent	TEVC	Control	Silent	TEVC
<i>Same slope (Figure 3A)</i>						
<i>IH vs. BKCa</i>	0.003	0.280	0.001	0.805	0.358	0.836
<i>IRK vs. CaV2</i>	0.022	0.717	0.005	0.678	-0.132	0.746
<i>NALCN vs. CaV2</i>	0.006	0.804	0.002	0.737	0.090	0.801
<i>IH vs. SKCa</i>	0.033	0.142	0.013	0.643	0.473	0.691
<i>Shaw1 vs. Shaker</i>	<0.0001	0.301	<0.0001	0.879	0.344	0.933
<i>Shaw2 vs. CaV2</i>	0.001	0.610	0.041	0.849	-0.198	0.596
<i>IRK vs. NALCN</i>	0.015	0.153	0.045	0.705	-0.462	0.586
<i>Shaw2 vs. IRK</i>	0.001	0.630	0.041	0.864	-0.174	0.596
<i>Shaw2 vs. Shaw1</i>	0.019	0.478	0.015	0.691	0.254	0.680
<i>SKCa vs. Shaker</i>	0.003	0.068	0.001	0.772	0.569	0.822
<i>Shal vs. IH</i>	0.007	0.070	0.001	0.790	0.566	0.846
<i>Shab vs. NALCN</i>	0.036	0.255	0.012	0.607	0.375	0.696
<i>Shaw2 vs. Shaker</i>	0.041	0.315	0.003	0.622	-0.354	0.780
<i>BKCa vs. SKCa</i>	0.033	0.127	0.003	0.615	0.490	0.801
<i>Different slope (Figure 3B)</i>						
<i>NALCN vs. Shaker</i>	<0.0001	0.059	0.009	0.871	0.585	0.712
<i>NALCN vs. SKCa</i>	0.001	0.182	0.021	0.838	0.434	0.654
<i>Shaker vs. IRK</i>	0.019	0.942	0.003	0.691	-0.025	0.769
<i>Shaw1 vs. IRK</i>	0.007	0.456	0.001	0.758	0.251	0.821
<i>Shaw1 vs. CaV2</i>	0.015	0.417	<0.0001	0.855	0.289	0.927
<i>NALCN vs. Shaw1</i>	<0.0001	0.831	0.003	0.843	-0.073	0.772
<i>Shaker vs. CaV2</i>	0.018	0.102	<0.0001	0.666	0.547	0.918

**Table S1. Correlated patterns of gene expression that arise due to voltage activity of the membrane, Related to Figure 2 and Figure 3.**

<u>Feedback-Independent Correlations (Figure 2C<sub>2</sub>)</u>	<i>p</i> values			Pearson's <i>r</i>		
	Control	Silent	TEVC	Control	Silent	TEVC
<i>BKKCa vs. NaV</i>	0.028	0.033	0.041	0.686	0.643	0.652
<i>Shab vs. CaV2</i>	0.011	0.026	<0.0001	0.701	0.695	0.932
<i>NaV vs. Shal</i>	0.014	0.010	0.023	0.775	0.735	0.673
<i>Shab vs. SKKCa</i>	0.021	0.010	0.002	0.653	0.737	0.786
<i>Shaw1 vs. SKKCa</i>	0.004	0.027	0.002	0.756	0.660	0.792
<i>Shaw1 vs. Shab</i>	0.001	0.006	<0.0001	0.820	0.764	0.923
<i>Shal vs. BKKCa</i>	0.024	0.024	0.007	0.672	0.671	0.757
<i>Shab vs. Shaker</i>	0.001	0.030	<0.0001	0.814	0.650	0.922

  

<u>Activity-Independent Correlations</u>	<i>p</i> values			Pearson's <i>r</i>		
	Control	Silent	TEVC	Control	Silent	TEVC
Same Trend as Figure 2C <sub>3</sub>						
<i>CaV2 vs. CaV1</i>	<0.0001	0.376	0.234	0.857	0.315	0.391
<i>IRK vs. CaV1</i>	0.004	0.793	0.310	0.783	-0.090	0.338
<i>Shaw1 vs CaV1</i>	0.015	0.071	0.549	0.680	0.564	0.203
<i>Shaw2 vs. CaV1</i>	<0.0001	0.840	0.995	0.945	0.074	0.003

  

<u>Activity-Restricted Correlations</u>	<i>p</i> values			Pearson's <i>r</i>		
	Control	Silent	TEVC	Control	Silent	TEVC
Same Trend as Figure 2C <sub>4</sub>						
<i>SKKCa vs. CaV1</i>	0.418	<0.0001	0.816	0.258	0.894	-0.080
<i>NaV vs. CaV1</i>	0.186	0.013	0.696	0.455	0.716	-0.142
<i>Shab vs CaV1</i>	0.177	0.036	0.555	0.417	0.635	0.200
<i>NaV vs. IH</i>	0.378	0.014	0.302	0.335	0.710	0.342
<i>Shal vs. CaV1</i>	0.765	0.041	0.286	-0.102	0.622	-0.354

**Table S2. Correlated patterns of gene expression that arise as a result of factors other than the naturalistic waveform activity of PD, Related to Figure 2.** *Feedback-Independent Correlations:* Correlated patterns of gene expression that arise independent of voltage activity of the membrane and chemical feedback. *Activity-Independent Correlations:* Correlated relationships that are present in the control group, but eliminated in both TTX (silent) and TTX+Activity (TEVC) groups. *Activity-Restricted Correlations:* Correlated relationships that are present only when activity was suppressed.

<u>Never Correlated</u>	<u>Became correlated with TTX and TTX + TEVC</u>	<u>Only correlated with TTX+ TEVC</u>
<i>BKKCa vs. CaV1</i>	<i>Shab vs. BKKCa</i>	<i>CaV2 vs. BKKCa</i>
<i>BKKCa vs. Shaw 2</i>	<i>NaV vs. CaV2</i>	<i>IRK vs. BKKCa</i>
<i>IH vs. CaV1</i>	<i>Shal vs. NALCN</i>	<i>NALCN vs. BKKCa</i>
<i>NALCN vs. CaV1</i>	<i>SKKCa vs. NALCN</i>	<i>Shaker vs. BKKCa</i>
<i>Shaker vs. CaV1</i>	<i>SKKCa vs. Shal</i>	<i>Shaw1 vs. BKKCa</i>
<i>IH vs. CaV2</i>	<i>Shaker vs. Shal</i>	<i>SKKCa vs. CaV2</i>
<i>Shal vs. CaV2</i>	<i>NaV vs. SKKCa</i>	<i>Shab vs. IRK</i>
<i>IH vs. IRK</i>	<i>Shab vs. NaV</i>	<i>NaV vs. NALCN</i>
<i>IH vs. NALCN</i>	<i>Shaker vs. NaV</i>	<i>Shaw1 vs. Shal</i>
<i>IH vs. Shab</i>		<i>Shaw2 vs. Shal</i>
<i>IH vs. Shaker</i>		<i>Shaw1 vs. NaV</i>
<i>IH vs. Shaw1</i>		<i>Shaw2 vs. NaV</i>
<i>IH vs. Shaw2</i>		<i>Shaw2 vs. Shab</i>
<i>Shal vs. IRK</i>		
<i>SKKCa vs. IRK</i>		
<i>NaV vs. IRK</i>		
<i>Shaw2 vs. BKKCa</i>		
<i>Shab vs. Shal</i>		
<i>Shaw2 vs. SKKCa</i>		

**Table S3. Channel pairs that were never correlated, became with either TTX or TTX+TEVC, or were only correlated with TTX+TEVC, Related to Figure 2.**

<u>Became correlated with TTX and TTX + TEVC</u>	<u>Significant difference between slopes?</u>	<u>ANCOVA p value</u>
<i>Shab vs. BKCa</i>	No	0.299
<i>NaV vs. CaV2</i>	Yes	0.044
<i>Shal vs. NALCN</i>	No	0.560
<i>SKKCa vs. Shal</i>	Yes	0.021
<i>Shaker vs. Shal</i>	No	0.109
<i>NaV vs. SKKCa</i>	No	0.225
<i>Shab vs. NaV</i>	No	0.227
<i>Shaker vs. NaV</i>	No	0.146

**Table S4. ANCOVA results for correlations that emerged after loss of inputs with or without activity, Related to Figure 2.**

Target	Forward Primer	Reverse Primer	Probe	Labeling	Efficiency
<i>NALCN</i>	TCGCTTCCACGGTGTACATTC	GCGGTGCCTTTGTTCTCAG	TCTTCGTCTTCCTTGGCTGCATGA	FAM-BHQ1	97.1
<i>IRK</i>	TACAGTGGCGTTGGACTCTAC	TCCACCACACCAAGGCAAATAG	TCGTGTTGCGTATGTCATTATCAGC	CAL Fluor Red 610-BHQ2	93.9
<i>IH</i>	TCGGTGCCACTAGACTACATC	GACCCGCGTGGAGAATCTG	TCCTCATCTTCAACCAGGACTTCAGC	Quasar 670-BHQ2	95.9
<i>SKKCa</i>	GCATCGGAGCATTGAACAGAA	GCCCGGACAGATAGTCATCAG	CAACTTCAACACTCGGTTTGCCTCAA	Quasar 705-BHQ2	96.1
<i>Shaw1</i>	CGCGTCACTCCTCAGGACTT	CCCAGCACCAGGAAGAACAC	TGATACAGACTTTCGGTGCATCCGC	Quasar 705-BHQ2	87.4
<i>Shaw2</i>	GAACGCCATCAAGCACTATCATC	ATGGCGCCCGACAGCTTAG	TGGCTTGAAGGACGGTCTCACA	Quasar 670-BHQ2	91.5
<i>CbNaV</i>	TCAACGGGAGGTACCATAAGTG	TCGCTGTTCACCAAGAGTAG	CGGAGGGATTGAAGCTCAACGCA	CAL Fluor Red 610-BHQ2	81
<i>Shaker</i>	GAGGCTCAGAAGACCAGTCAAC	TGGCGATATCACCGAGCTCAT	CACTCGATGTCTTCGCGGAGGAGAT	CAL Fluor Gold 540-BHQ1	90.7
<i>Shab</i>	GAGCCGGACAGACAGGAAC	TGCGCCTCCTTCTGTAGTC	AAGAACCACGAACACCACATGGGTC	FAM-BHQ1	93.5
<i>CaV1</i>	CCAGGCCTTCTACTGGCTCATT	GCTGGCGATAGTGCTCACTG	TGTGCTCGTCTTCCTCAACACGG	FAM-BHQ1	96
<i>CaV2</i>	ATCCGGCGGACAGTAAAGC	GTTCCGGCAGCAACACAAC	TGGTTCTACTGGTTCGTCATCATACTTGT	CAL Fluor Gold 540-BHQ1	97.9
<i>Shal</i>	GACACCACCTTACCTCCATTC	GAACCATGTCCCGTATCCTA	CGGCGTTTTGGTACACCATTGTCAC	Quasar 670-BHQ2	110.7
<i>BKKCa</i>	GCTCAAACCTCGGCTTCATTG	CTGCGTGTCTGGAGAAGTTT	AGAATCCCGCGCTAAACATGACT	Quasar 705-BHQ2	96
<i>vGluT</i>	GCGTTCGTGGACCTTCTAC	TCAGCCACCCTGTAATGGAA	ATCACAGCCAACCTACTTCAGCGAG	CAL Fluor Gold 540-BHQ1	95.3
<i>ChAT</i>	GGACCGCCTGGCTAAGTAC	TCGCGGAGTCCATAAGG	AGGCGGCGCTCAAGCTTCAGAC	CAL Fluor Red 610-BHQ2	93.1
<i>vAChT</i>	GCGTCAGCTGCTTCTTCT	CAGCAGTGCCGTGTCTATGAG	TTCGCCAGCAACTACTGGGTGTT	Quasar 670-BHQ2	92.8
<i>ACHE</i>	GGGCAACATGGGCATGTAC	GGTCACCACCGAAGAATCAATG	AGGCGCTGGCCATCAAGTGGATAC	Quasar 705-BHQ2	94.2

**Table S5. Primers and probes used for multiplex qRT-PCR, Related to STAR Methods.**  
Each box in the table represents a group of genes that were multiplexed in a single PCR run.