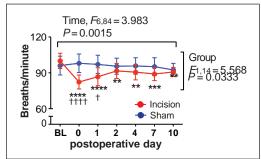
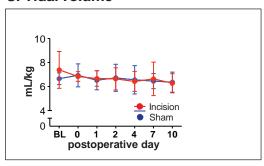
A. Minute ventilation

Time, F6,84 = 3.207; P = 0.0069 900 600 F6,84 = 2.633 P = 0.0218 P = 0.0218

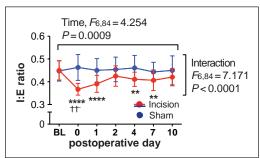
B. Breathing frequency



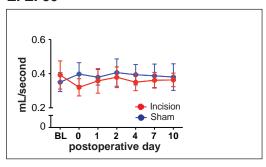
C. Tidal volume



D. I:E ratio



E. EF50



F. Body weight

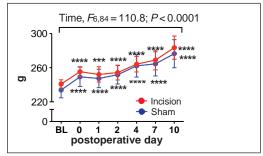


Figure S2. Raw (non-normalized) data for the Protocol C, the effect of upper abdominal incision vs. sham incision on the ventilatory parameters. (A) Minute ventilation of the Incision group after the surgery, on POD 0 through POD 10, was significantly lower compared with baseline (POD 0: P = 0.0001; POD 1: P = 0.0006; POD 2: P = 0.0070; POD 4: P = 0.0010; POD 7: P = 0.0015; POD 10: P = 0.00150.0003). On the other hand, there was no statistical difference in minute ventilation of the Sham group throughout the testing period. (B) Breathing frequency of the Incision group on POD 0 though POD 10 was significantly lower compared to baseline (POD 0: P < 0.0001; POD 1: P < 0.0001; POD 2: P =0.0072; POD 4: P = 0.0019; POD 7: P = 0.0002; POD 10: P = 0.0022). The mean breathing frequency values of the Sham group were not different throughout the testing period. (C) For tidal volume, there was no significant main effect or interaction. (D) Inspiratory-to-expiratory time ratio (I:E ratio) of the Incision group was lower compared with baseline on POD 0 through POD 7 (POD 0: P < 0.0001; POD 1: P < 0.0001; POD 4: P = 0.0054; POD 7: P = 0.0017); I:E ratio of the Sham group was not significantly different throughout the testing period. (E) For expiratory flow at 50% expired volume (EF50), there was no significant main effect or interaction. (F) Body weight on POD 0 through POD 10 were significantly greater compared with baseline in both the Incision and the Sham groups. Data are presented as mean \pm SD. N = 8 per group. **P < 0.01; ***P < 0.001; ****P < 0.0001 vs. Baseline, and $\dagger P < 0.05$; $\dagger \dagger \dagger P < 0.001$; $\dagger \dagger \dagger \dagger P < 0.0001$ vs. Sham by two-way ANOVA with repeated measured in one factor, followed by Sidak's multiple comparison tests. BL = baseline.