

Figures and figure supplements

Quantification of gait parameters in freely walking wild type and sensory deprived Drosophila melanogaster

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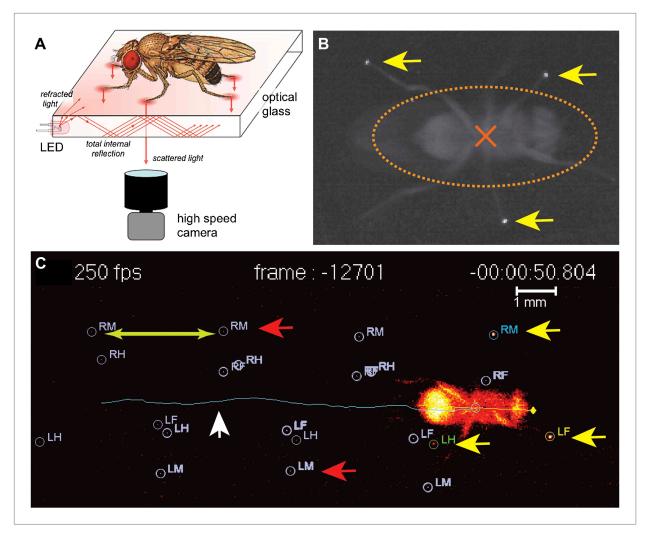


Figure 1. fTIR apparatus and FlyWalker software. (A). Schematic of the fTIR optical effect. LED light sources are located at the edges of an optical glass and light propagates within the glass via internal reflection. Tarsal contacts lead to light scattering detected by a high-speed camera. See

Figure 1—figure supplement 1 for more details. (B). Single frame of a fTIR video. The fTIR effect can be seen for three legs in stance phase (yellow arrows). Background light partially illuminates the fly's body (orange dashed ellipse; the center of the body is indicated by an orange cross). (C). Image generated by the FlyWalker software. The fly's footprints and body center are tracked throughout the video. Present footprints are identified and labeled (yellow arrows). The fly body and trajectory are visualized by a blue line (white arrow). Past footprints can also be recorded (red arrows). A scale bar can be introduced. Step length is defined as the distance between two consecutive footprints (green arrows).

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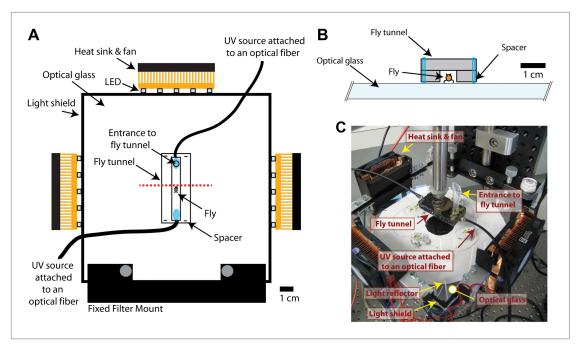


Figure 1—figure supplement 1. Additional information on fTIR apparatus. (A). Detailed schematic of the fTIR apparatus. Red dashed line shows position of the cross-section image in panel (B). (B) Cross-section of schematic at red dashed line in panel (A). (C) Photograph of the fTIR apparatus. DOI: 10.7554/eLife.00231.004

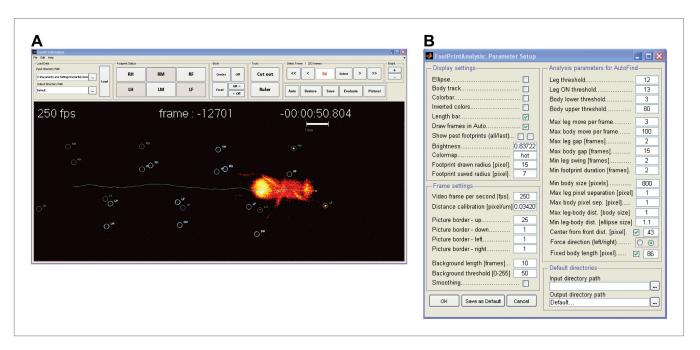


Figure 1—figure supplement 2. FlyWalker program. **(A)**. Screenshot of the FlyWalker graphical user interface used to edit fTIR videos. **(B)**. Screenshot of the FlyWalker parameters window.



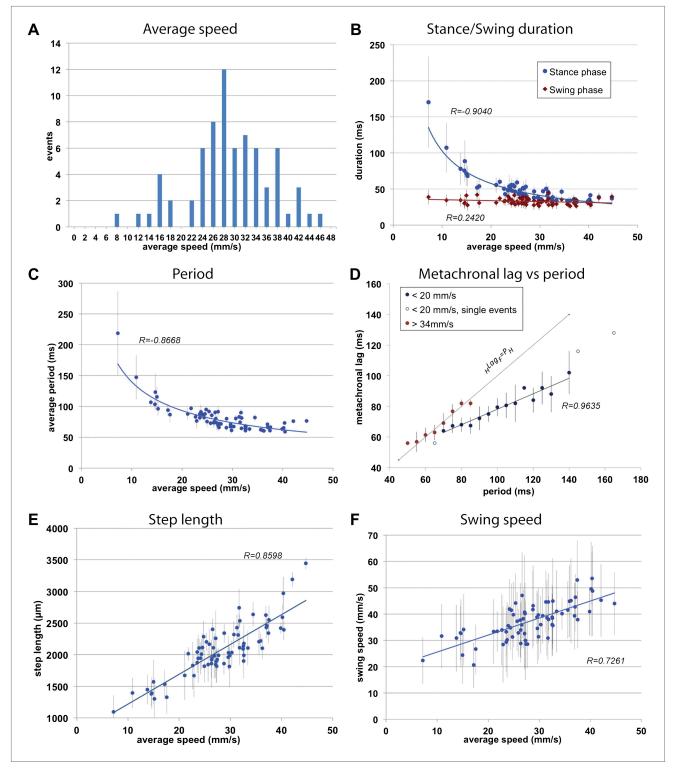


Figure 2. General walking parameters. (A). Speed histogram of 71 videos recorded for wild type flies, with 2 mm/s bins. Average speeds vary between 7.2 and 44.7 mm/s, with 28 mm/s the most represented speed. Speeds at ~20 mm/s are underrepresented in this dataset. (B). Durations of stance (blue) and swing (red) phases as a function of speed. Swing phase duration remains mostly constant while stance phase duration is inversely proportional to speed. Graphical fits for swing and stance phases versus speed are represented in red and blue lines, respectively. (C). Step period is inversely proportional to average speed. The blue line is a graphical fit. (D). Duration of the metachronal lag as a function of the hindleg period for slow (blue) and fast (red) flies. metachronal lag closely matches hindleg period for fast flies ($_{\rm HLag_F} \approx {\rm Period}$). Regression line for the slow walking flies: $_{\rm HLag_F} = {\rm Period} \times 0.505 + 27.627$. Figure 2. Continued on next page



Figure 2. Continued

A total of 220 metachronal waves were used, error bars correspond the standard error of the mean. (E),(F). Step length (E) and swing speed (F) increase linearly with speed. A graphical fit is shown by the blue lines. Error bars in (B–F) correspond to standard error of the mean. DOI: 10.7554/eLife.00231.008

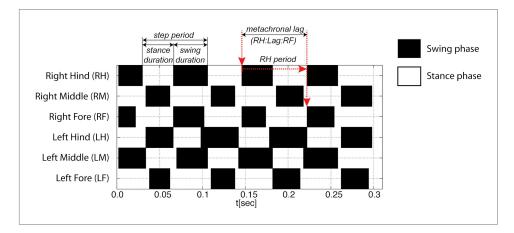


Figure 2—figure supplement 1. Gait definitions. For each leg, swing phases are represented in black and stance phases are represented in white (from top to bottom: right hind (RH); right mid (RM); right front (RF); left hind (LH); left middle (LM); left front (LF)). Step period is the time for one complete cycle (initiation of consecutive stance phases) and is divided into stance duration and swing duration. The metachronal lag is defined as the time between the start of sequential swing phases for the hind and forelegs on the ipsilateral side (red arrows). DOI: 10.7554/eLife.00231.009



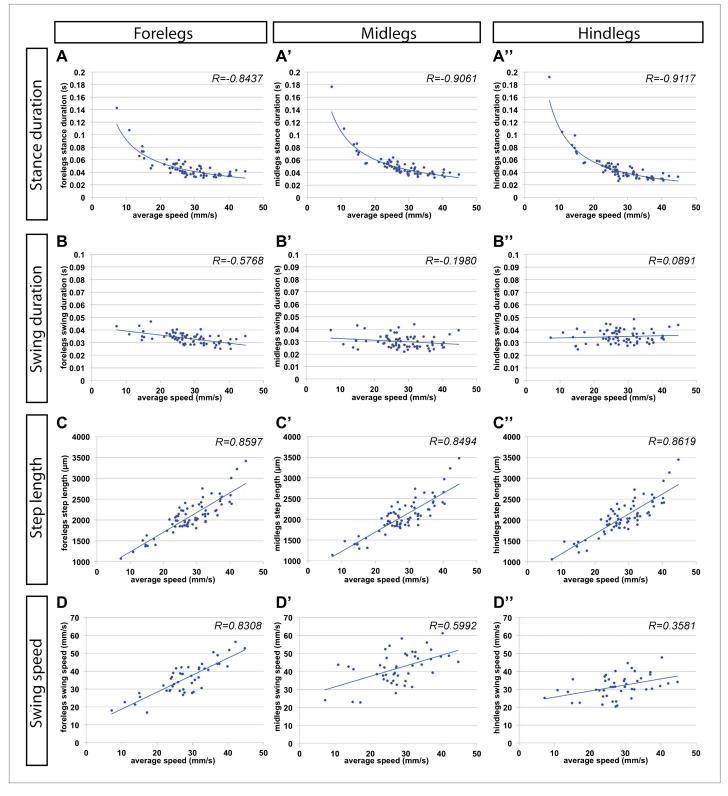


Figure 2—figure supplement 2. Gait parameters by segment. **(A)–(D)**. Each column corresponds to the legs in a particular segment. Graphical fits are represented by blue lines. **(A)** *Stance duration*. **(B)** *Swing duration*. **(C)** *Step length*. **(D)** *Swing speed*. DOI: 10.7554/eLife.00231.010



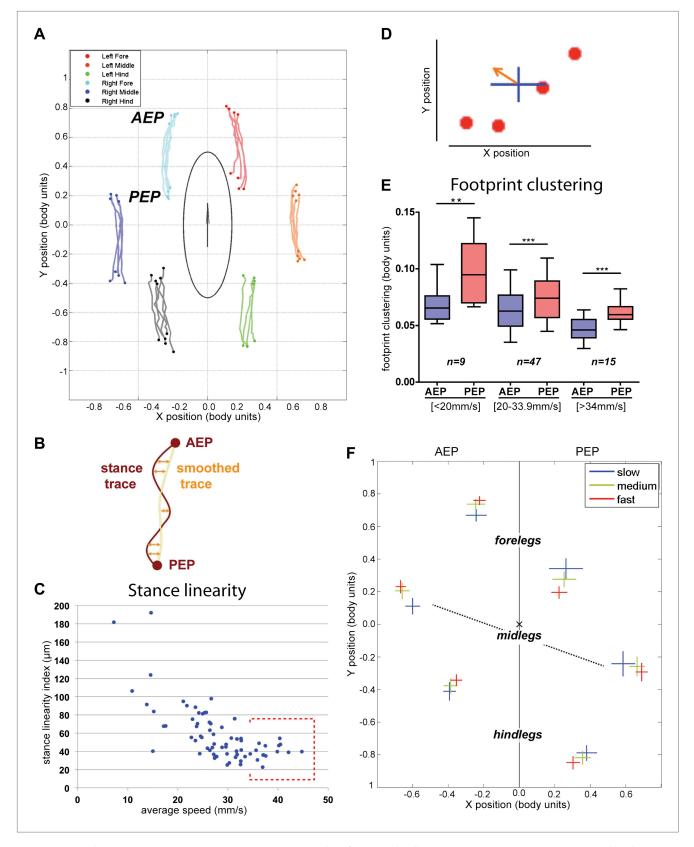


Figure 3. Spatial parameters. **(A)**. Stance traces. Representative plot of an animal walking at 28.82 mm/s. Traces are generated by the position of the stance phase footprints relative to the body center (set at 0.0,0.0). For each leg, stance onset corresponds to the Anterior Extreme Position (AEP) while Figure 3. Continued on next page



Figure 3. Continued

stance offset is termed *Posterior Extreme Position* (*PEP*). (**B**). Method to quantify the *stance linearity index*. For each stance trace (brown), a smoothed trace is generated (using data from every five frames; yellow trace), and the average of the difference between these two lines (orange arrows) corresponds to the *stance linearity index*. (**C**). *Stance linearity* as a function of speed. Each data point corresponds to the average of all traces for all six legs for a single video greater than ~34 mm/s, *stance linearity* becomes constant (red box). (**D**). Method to quantify *footprint clustering*. For each set of *AEP/PEP* footprints (red circles), an average ±STD xy point is created (blue cross). The *footprint clustering* value is calculated as the vector sum of the two STD values (orange arrow). (**E**). Quantification of *footprint clustering*. Data were grouped into slow (<20 mm/s), medium (between 20 and 34 mm/s) and fast (>34 mm/s) speeds. Boxplots represent the median as the middle line, with the lower and upper edges of the boxes representing the 25% and 75% quartiles, respectively; the whiskers represent the range of the full data set. Values are normalized for body size. Asterisks indicate the significance of the decrease in *footprint clustering* between the *AEP* and the *PEP* (using the paired parametric t test in the case of the slow speed group and the Wilcoxon non-parametric test for the remaining groups, **p<0.005, ***p<0.001). A comparison between the three speed groups also displays statistical significance (Kruskal–Wallis-ANOVA, p values of 0.0015 and 0.0014 for *AEP* and *PEP*, respectively). Dunn's *post hoc* significance tests show: slow *AEP* vs medium *AEP*, not significant (NS); medium *AEP* vs fast *AEP*, **; slow *PEP* vs medium *PEP*, NS; medium *PEP* vs fast *PEP*, NS; slow *PEP* vs fast *PEP*, ***. (**F**). Footprint position relative to the body center. Data were pooled as in the previous panel. *AEP* and *PEP* values for each leg are represented on the left and right section

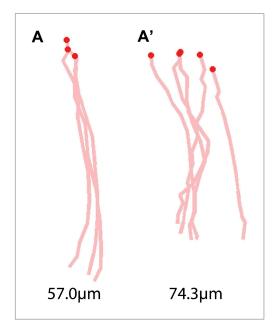


Figure 3—figure supplement 1. Representative examples of stance traces and corresponding stance linearity values. A less linear stance trace (e.g., (A') compared to (A)) corresponds a higher stance linearity value.



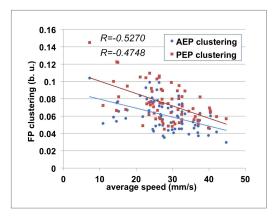


Figure 3—figure supplement 2. AEP and PEP clustering values for all segments as a function of speed. Linear trend lines for AEP and PEP versus average speed are shown as blue and red lines, respectively.

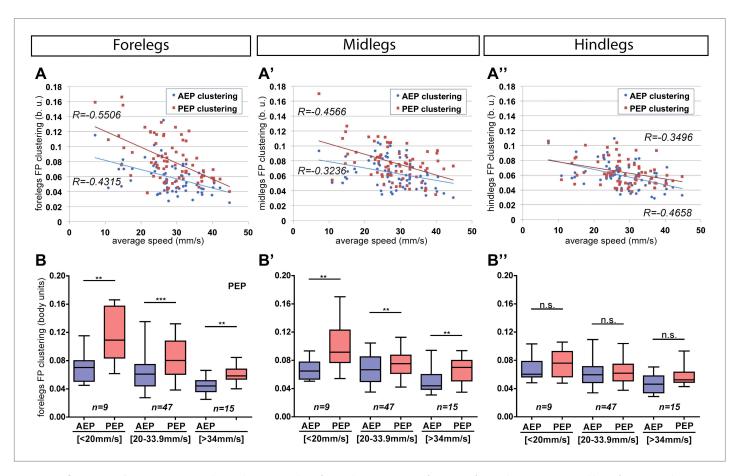


Figure 3—figure supplement 3. AEP and PEP clustering values for each segment as a function of speed. (A). Linear trend lines for AEP and PEP are represented as blue and red lines, respectively. (A) forelegs; (A') midlegs; (B). Boxplots of the data presented in (A). Data were grouped into slow (<20 mm/s), medium (between 20 and 34 mm/s) and fast (>34 mm/s) animals. Boxplots represent the median as the middle line, with the lower and upper edges of the boxes representing the 25% and 75% quartiles, respectively, and the whiskers representing the range of the full data set. Values are normalized for body size. Asterisks indicate statistical significance between AEP and PEP. (Data analyzed by the Wilcoxon or paired-t test the, **p<0.005, ***p<0.001). Values are normalized for body size.



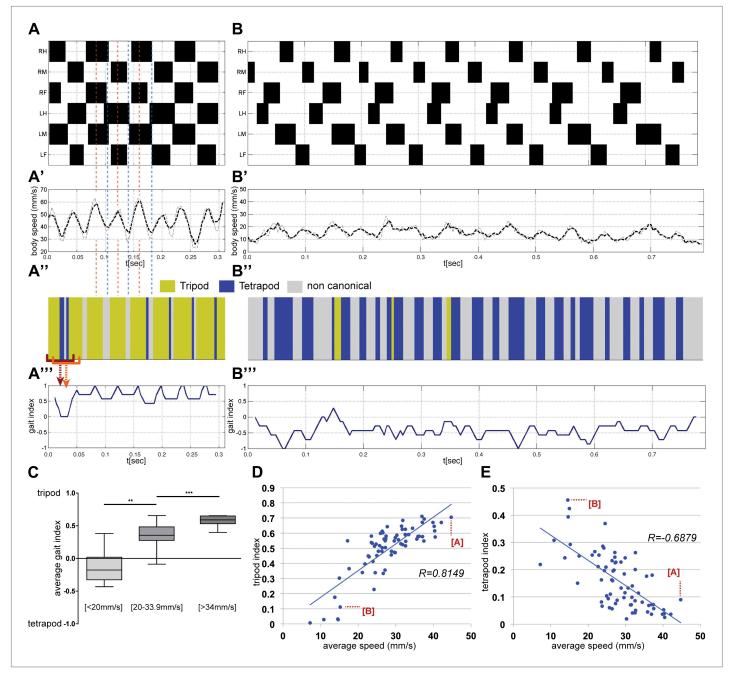


Figure 4. Gait parameters. (A),(B). Upper panels show the step pattern for representative videos of animals walking at 44.7 (A) and at 14.6 mm/s (B). For each leg swing phases are represented in black (from top to bottom: right hind (RH); right mid (RM); right front (RF); left hind (LH); left middle (LM); left front (LF)). (A'),(B') show the *instantaneous speed* for the same video. Thick and thin lines correspond to integration times of 25 or 12.5 ms, respectively. In faster animals (A), peak speeds are observed halfway through the stance phase (red dashed line), while minimum speeds are observed during gait transitions (blue dashed line). (A"),(B"), for each frame the corresponding gait was color coded as follows: green (tripod), blue (tetrapod), and gray (non-canonical). Red brackets indicate the eight frame windows used to generate the gait index plots (A"'),(B"'). Tripod = +1; tetrapod = -1; and noncanonical = 0. (C). Quantification of the average gait index for three speed groups. For each video the average gait index was calculated for all frames (p<0.0001 for Kruskal–Wallis-ANOVA test. Dunn's post hoc significance test: **p<0.005, ***p<0.001. (D),(E). Tripod (D) and tetrapod (E) indexes as a function of speed. Graphical fits are represented in blue. Data for points labeled (A) and (B) are shown in panels (A) and (B), respectively.



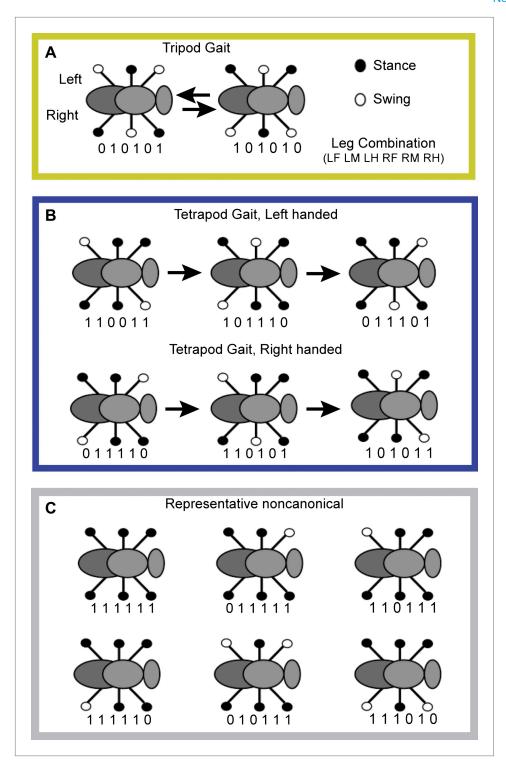


Figure 4—figure supplement 1. Gait features. (**A**)–(**C**). Walking gaits and leg combinations. Each leg can be either in a swing or stance phase, represented by white or black circles, respectively. Underneath each combination the corresponding numerical code ('1' and '0' correspond to stance or swing, respectively. Digit order: left foreleg; left midleg; left hindleg; right foreleg; right midleg; right hindleg. (**A**) Tripod gait combinations. (**B**) Tetrapod gait combinations. Tetrapod gait can be either right or left handed depending on which side is swinging more anteriorly. (**C**) Representative noncanonical combinations, which are not represented by tripod or tetrapod configurations.



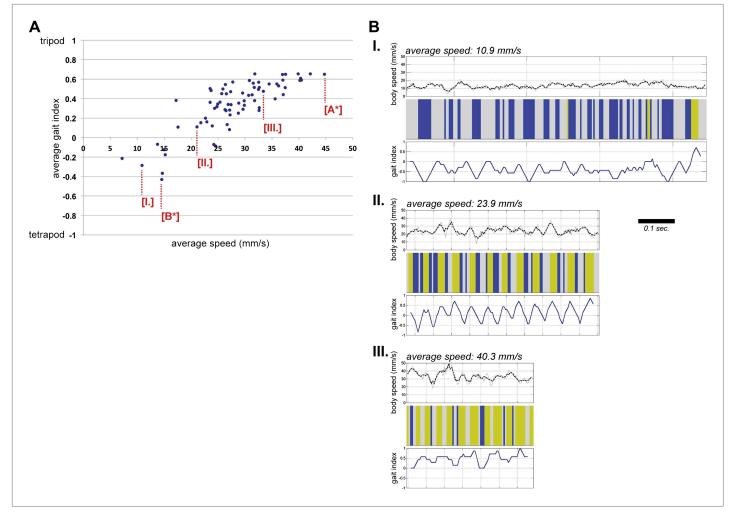


Figure 4—figure supplement 2. Gait index plot. (A). Average gait index as a function of speed. Each point corresponds to the average of all frames in a video (Figure 4A",B"). (B). Instantaneous speed, gait map and gait index plots for data points labeled in (A) by I., II., and III. are shown. Data for points labeled A* and B* are shown in Figure 4A,B, respectively.

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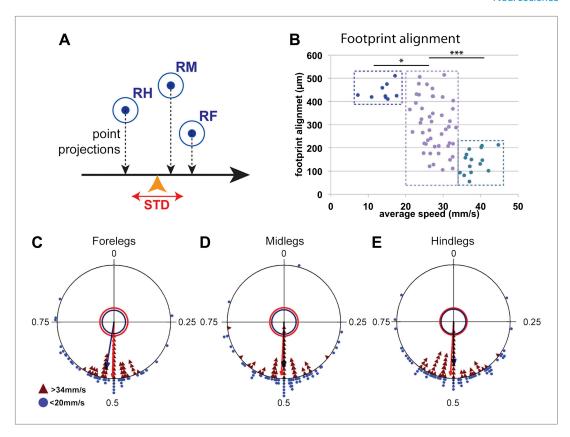


Figure 5. Coordination parameters. **(A)**. Method to calculate *footprint alignment*. For each set of footprints, the projection points along the displacement axis (horizontal black arrow) are calculated. *Footprint alignment* corresponds to the standard deviation (STD) of the average point (orange arrowhead). **(B)**. Quantification of *footprint alignment* versus speed. Values are color coded according to speed group: <20 mm/s in dark blue; between 20 and 34 mm/s in purple and >34 mm/s in turquoise. Values cluster into three main groups indicated by the dotted boxes (p<0.0001 for Kruskal–Wallis-ANOVA test. Dunn's post hoc significance test: *p<0.005, ****p<0.001). **(C)**–(**E**). Radial plots of contralateral leg phases comparing slow (blue triangles) and fast (red triangles) walking animals. Mean r vectors for slow and fast animals are represented by blue and red arrows, respectively. Inner circles indicate a Rayleigh p value of 0.05. DOI: 10.7554/eLife.00231.019

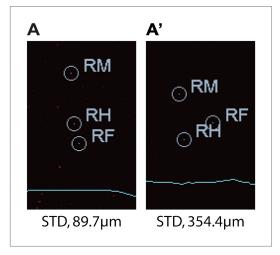


Figure 5—figure supplement 1. Representative images of footprints with different footprint alignment values.



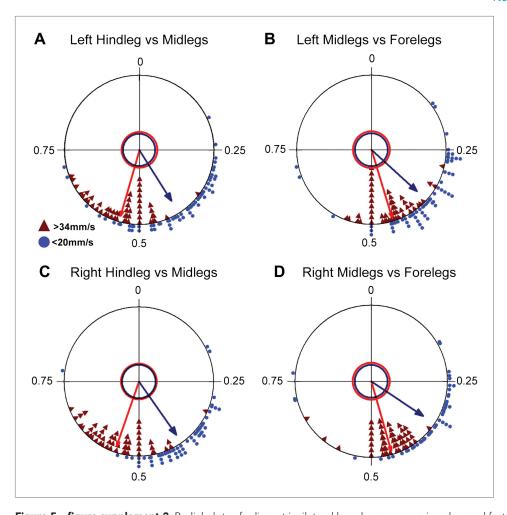


Figure 5—figure supplement 2. Radial plots of adjacent ipsilateral leg phases comparing slow and fast walking animals. Individual data points for slow and fast flies are represented by blue circles and red triangles, respectively. Mean r vectors for slow and fast animals are represented by blue and red arrows, respectively. Inner circles indicate a Rayleigh p value of 0.05.



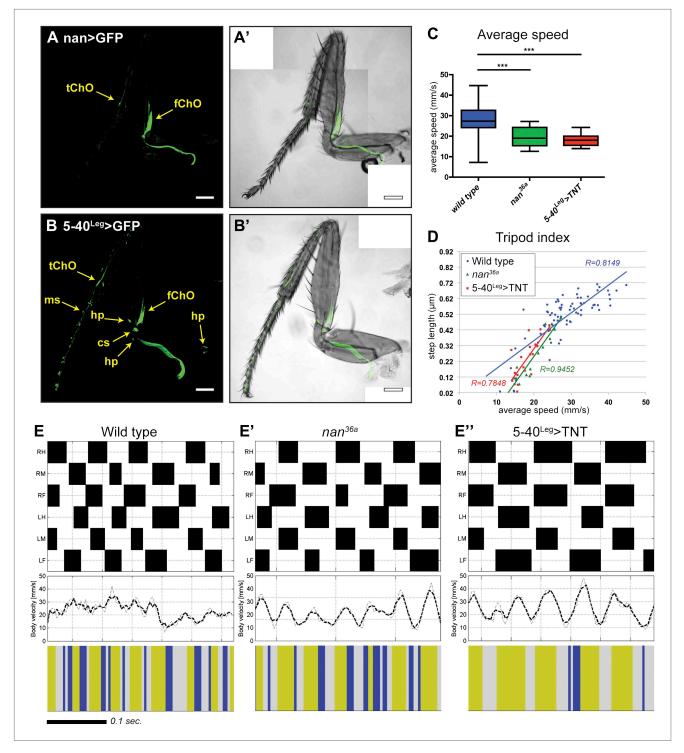


Figure 6. Effects of sensory deprivation on walking. (**A**). Expression pattern driven by *nanchung*-Gal4. Genotype: *F-Gal4*, *UAS-GFP*. (**B**). GFP expression under combinatorial control of *5-40-Gal4* and *dac*^{RE}-flp. Genotype: *5-40-Gal4*, *dac*^{RE}-flp, *UAS-FRT*-stop-FRP-GFP. All classes of sensory neurons in the leg express GFP. tChO, tibia chordotonal organ; fChO, femur chordotonal organ; hp, hair plates; cs, campaniform sensilla; ms, mechanosensory brisles. Bar, 100 μm. (**C**). Average speed. Boxplots with the median as the middle line and the lower and upper edges of the boxes representing the 25% and 75% quartiles, respectively; the whiskers represent the range of the full data set. Statistical analysis with one-way-ANOVA (p<0.0001) followed by Tukey's *post hoc* test, ***p<0.001. (**D**). *Tripod index*. Lines represent graphical fits. See *Table 2* for statistical analysis. (**E**). Gait patterns, instantaneous speeds, and gait maps for representative wild type and sensory deprived animals walking at similar speeds. See *Figure 2—figure supplement 1* for details. Tripod gait properties remain unchanged.



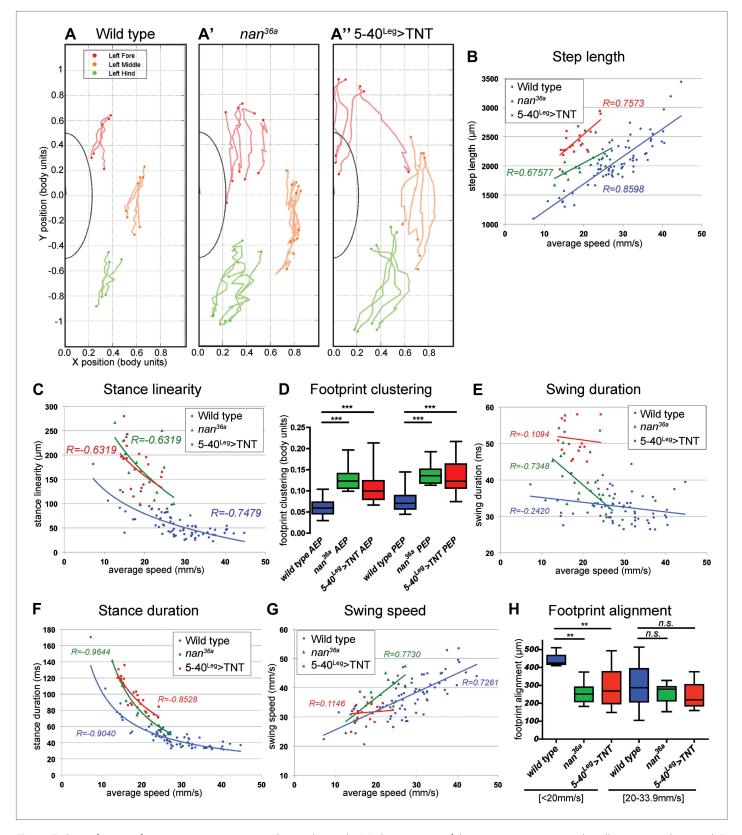


Figure 7. Quantification of gait parameters in sensory deprived animals. (A). Stance traces of three representative animals walking at a similar speed. For simplicity, only left stance traces are shown. Traces for sensory deprived genotypes display a longer step length; higher jitter and a more variable AEP and PEP. (See panels (B)–(D) and (H) for quantification). In (B),(C) and (E)–(G), colored lines represent graphical fits. See Table 2 for statistical analysis of Figure 7. Continued on next page



Figure 7. Continued

(B), (C), (E), (F) and (G). (B). Step length. Sensory deprived animals display an increased step length. (C). Stance linearity. Sensory deprived animals display a more jittery movement. (D). AEP and PEP footprint clustering. Wild type data correspond to the speed range of nan³⁶⁹ and 5-40^{1-eg}>TNT (12.6 to 27.7 mm/s). In (D), (H), boxplots represent the median as the middle line and the lower and upper edges of the boxes representing the 25% and 75% quartiles, respectively; the whiskers represent the range of the full data set. Statistical analysis with Kruskal–Wallis-ANOVA (p<0.0001) followed by Dunn's post hoc test, ***p<0.001. (E). Swing duration. (F). Stance duration. (G). Swing speed. (H). Footprint alignment. Slow-walking sensory deprived animals display more aligned footprints compared to wild type flies. Data were grouped into slow (<20 mm/s) and medium speeds (between 20 and 34 mm/s). Kruskal–Wallis-ANOVA test: p=0.0003 and NS for the slow and medium speed groups, respectively. Asterisks indicate the significance of the decrease in footprint clustering between genotypes. (Data analyzed by the Dunn's post hoc significance test, **p<0.005).