

Figure S5 Behavioral Analyses of the dat-1(vt21) and dat-1(vt22) strains. A. Vt21 and vt22 mimic the dat-1 Swip phenotype as measured by automated thrashing analysis. Individual animals were recorded using a video capture system and then analyzed with customed-designed Thrasher software that assigns multiple linear elements projecting from the worm centroid. The position of these linear elements are tracked and converted off-line to movement frequency as a function of time. Batch conversions are generated, providing mean values and SEM along moving averages. Error bars are not shown in these plots for simplicity. Dat-1(ok157), vt21 and vt22 were found to be significantly different from N2 using two-way ANOVA with Bonferroni posttests of mutants to N2, with each mutant possessing a P<0.001 after the one minute mark. B. Mutation of the postsynaptic receptor DOP-3 fully rescues the paralysis phenotype of vt21 and vt22. Analyses were performed as described in B, where vt21 and vt22 were both found to be significantly different from the double mutants vt21;dop-3(vs106) and vt22;dop-3(vs106) with P< 0.001 after 1 minute. C. Heat map representations of dat-1(vt21) and dat-1(vt22) swimming traces. Analyses were performed as described in Figure 4. D. vt21 and vt22 display enhanced sensitivity to exogenous DA when tested on solid medium, as compared to N2, but are indistinguishable from dat-1. For these assays, 10 L4 stage worms were placed on plates containing increasing concentrations of exogenous DA, incubated for 20 min and then scored for 10 sec as paralyzed or moving. Dose-response curves were compared using two-way ANOVA with Bonferroni posttests comparing mutants to N2, in which dat-1, vt21 + vt22 were all found to be significantly different from N2, with a P<0.001 at 15 and 20 mM DA. Data derive from at least 4 tests per strain per DA concentration. Error bars represent SEM. Exogenous DA dose response profiles and data analysis were performed as described in the Methods and in Figure 3.