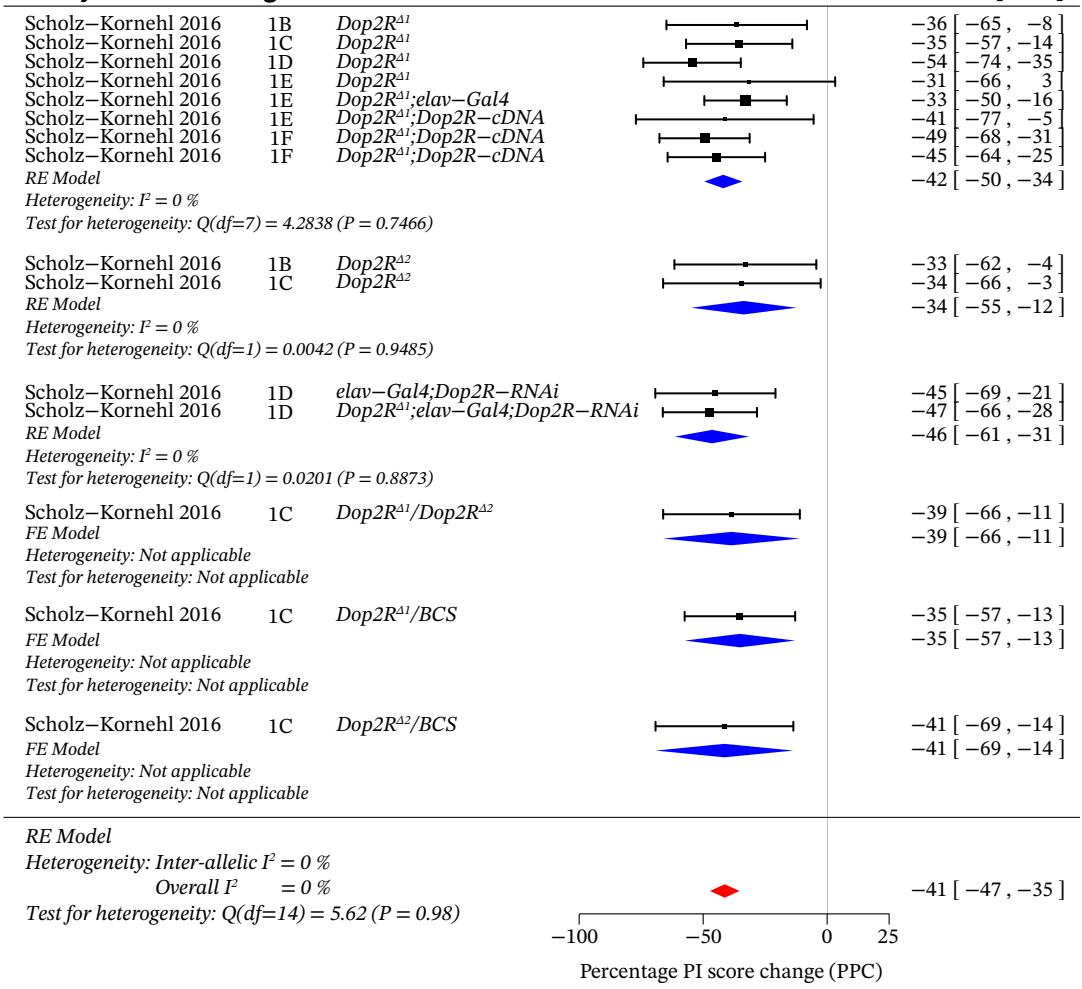
Study
Figure
Allele
%PPC [95CI]
Scholz−Kornehl 2016 1B Dop2R^Δ1 −36 [−65 , −8]
Scholz−Kornehl 2016 1C Dop2R^Δ1 −35 [−57 , −14]
Scholz−Kornehl 2016 1D Dop2R^Δ1 −54 [−74 , −35]
Scholz−Kornehl 2016 1E Dop2R^Δ1 −31 [−66 , 3]
Scholz−Kornehl 2016 1E Dop2R^Δ1;elav−Gal4 −33 [−50 , −16]
Scholz−Kornehl 2016 1E Dop2R^Δ1;Dop2R−cDNA −41 [−77 , −5]
Scholz−Kornehl 2016 1F Dop2R^Δ1;Dop2R−cDNA −49 [−68 , −31]
Scholz−Kornehl 2016 1F Dop2R^Δ1;Dop2R−cDNA −45 [−64 , −25]
RE Model −42 [−50 , −34]
Heterogeneity: I² = 0 %
Test for heterogeneity: Q(df=7) = 4.2838 (P = 0.7466)
Scholz−Kornehl 2016 1B Dop2R^Δ2 −33 [−62 , −4]
Scholz−Kornehl 2016 1C Dop2R^Δ2 −34 [−66 , −3]
RE Model −34 [−55 , −12]
Heterogeneity: I² = 0 %
Test for heterogeneity: Q(df=1) = 0.0042 (P = 0.9485)
Scholz−Kornehl 2016 1D elav−Gal4;Dop2R−RNAi −45 [−69 , −21]
Scholz−Kornehl 2016 1D Dop2R^Δ1;elav−Gal4;Dop2R−RNAi −47 [−66 , −28]
RE Model −46 [−61 , −31]
Heterogeneity: I² = 0 %
Test for heterogeneity: Q(df=1) = 0.0201 (P = 0.8873)
Scholz−Kornehl 2016 1C Dop2R^Δ1/Dop2R^Δ2 −39 [−66 , −11]
FE Model −39 [−66 , −11]
Heterogeneity: Not applicable
Test for heterogeneity: Not applicable
Scholz−Kornehl 2016 1C Dop2R^Δ1/BCS −35 [−57 , −13]
FE Model −35 [−57 , −13]
Heterogeneity: Not applicable
Test for heterogeneity: Not applicable
Scholz−Kornehl 2016 1C Dop2R^Δ2/BCS −41 [−69 , −14]
FE Model −41 [−69 , −14]
Heterogeneity: Not applicable
Test for heterogeneity: Not applicable
RE Model
Heterogeneity: Inter-allelic I² = 0 %
Overall I² = 0 %
Test for heterogeneity: Q(df=14) = 5.62 (P = 0.98)
−41 [−47 , −35]
−100
−50
0
25
Percentage PI score change (PPC)