Mouse, chr.2, high-GC (24-36Mb) region, B4A - SINE/B4

GC = 48%, max. div. = 27, $\mu > 2.5$, no power-law
Mouse, chr.2, high_GC (160-174.5 Mb) region, B4A-SINE/B4

GC = 47%, max. div. = 23, E=0.9, μ=0.73
Mouse, chr.2, low-GC (39-56.5 Mb) region, B4A - SINE/B4

GC = 39%, max. div. = 27, E=1.5, μ=0.55
Mouse chr.2, low-GC (77.5-90 Mb) region, B4A-SINE/B4

GC = 39%, div.max. = 29, E=1.80, μ=0.85
Mouse chr.2, high-GC (160-174.5 Mb) region, RSINE1-SINE/B4

GC = 47%, max. div. = 31, E=0.7, μ=2.16
Mouse chr.2, low-GC (39-56.5 Mb) region, RSINE1 - SINE/B4

GC = 39\%, \text{max. div.} = 27, \ E=1.36, \ \mu=0.87
Mouse chr.2, low-GC (77.5-90 Mb) region, RSINE1-SINE/B4

$GC = 39\%, \; div.\; max. = 27, \; E = 1.95, \; \mu = 0.86$
Mouse, Chr.4, GC-rich region (123-143 Mb), B3-SINE/B2

max. deviation = 24%, E = 1.15, μ = 1.78, r² = 0.99
Mouse, Chr. 4, GC-poor region (64-84 Mb), B3-SINE/B2

max. deviation = 24%, $E = 1.48$, $\mu = 1$, $r^2 = 0.99$
Mouse, Chr.5, GC-rich region (110-125 Mb), B4A-SINE/B4_d28

max. divergence = 28%, no power-law
Mouse, Chr.5, GC-poor region (45-60Mb), B4A-SINE/B4

max. divergence = 28%, E=1.26, μ=0.8, r²=0.99
Mouse, Chr.5, GC-rich region (110-125 Mb), B1_Mus1-SINE/Alu

max. divergence = 11%, E=1.03, μ=1.83
Mouse, Chr.5, GC-rich region (45-60 Mb), B1_Mus1-SINE/Alu

divergence = 11\%, E=0.8, \mu=1.35
Mouse, Chr.5, GC-rich region (110-125 Mb), B2_Mm2-SINE/B2

max. divergence = 5%, $E=0.9$, $\mu=0.82$
Mouse, Chr.5, GC-poor region (45-60 Mb), B2_Mm2-SINE/B2

max. divergence = 7%, $E=2.40$, $\mu=0.40$
Mouse, Chr.15, GC-rich region (72-90Mb), B1_Mus1-SINE/Alu

max. divergence = 11%, E=1.36, μ=1.15, r²=0.99
Mouse, Chr.15, GC-poor region (4-24 Mb), B1_Mus1-SINE/Alu

max. divergence = 11%, no power-law
Mouse, Chr.15, GC-rich region (72-90 Mb), B2_Mm2-SINE/B2

max. divergence = 7\%, E=1.15, \mu=1.32, r^2=0.99
Mouse, Chr.15, 1st GC-poor region (40-58 Mb), B2_Mm2-SINE/B2

max. divergence = 7%, $E=1.48$, $\mu=0.33$, $r^2=0.98$
Mouse, Chr.15, 2nd GC-poor region (4-24 Mb), B2_Mm2-SINE/B2

max. divergence = 7%  E=1.92  μ=0.65
Mouse, Chr.15, GC-rich region (72-90 Mb), B3-SINE/B2

max. divergence = 23%, E=0.70, μ=2.35
Mouse, Chr.15, GC-poor region (40Mb-58Mb), B3-SINE/B2

max. divergence = 23%, E=1.36, \( \mu=0.93 \), \( r^2 = 0.99 \)
Mouse, Chr.15, GC-rich region (72-90 Mb), RSINE1-SINE/B4

max. divergence = 26%, no power-law
Mouse, Chr.15, GC-poor region (40-58 Mb), RSINE1-SINE/B4

max. divergence = 26%, E=1.48, μ=0.45, r²=0.97
Mouse, Chr.2, GC-rich region (24-36 Mb), B1_Mus1-SINE/Alu

max. divergence = 11%, E=0.80, μ=1.10
Mouse, Chr.2, GC-poor region (39-56.5 Mb), B1_Mus1- SINE/Alu

max. divergence = 14%, no power-law
Dog, Chr.1, GC-rich region (108.5-118 Mb), MIR-SINE/MIR

max. divergence = 31%, E=0.81, μ=2.28
Dog, Chr.1, GC-poor region (60-70 Mb), MIR-SINE/MIR

max. divergence = 33%, E<0.6, μ>>2.5, no power-law
Dog, Chr.1, GC-rich region (108.5-118 Mb), MIR3-SINE/MIR

max. divergence = 37%, $E=0.70$, $\mu=1.00$
Dog, Chr.1, GC-poor region (60-70 Mb), MIR3-SINE/MIR

max. divergence = 37%, no power-law
Dog, Chr.10, GC-rich region (20-31 Mb), MIR-SINE/MIR
max. deviation = 32%, E=0.9, μ=1.5, r²=0.99
Dog, Chr.10, GC-poor region (53-63 Mb), MIR-SINE/MIR

max. divergence = 32%, no power-law
Dog, Chr.10, GC-rich region (20-31 Mb), MIRb-SINE/MIR

max. divergence = 31%, E=1.48, μ=1.45
Dog, Chr.10, GC-poor region (53-63 Mb), MIRb-SINE/MIR

max. divergence = 31%, $E=0.80$, $\mu=2.15$
Dog, Chr.12, GC-rich region (3.5-16 Mb), MIR-SINE/MIR

max. divergence = 33%, $E=0.80$, $\mu > 2.5$ no power-law
Dog, Chr.12, GC-poor region (25-45 Mb), MIR-SINE/MIR

max. divergence = 35%, no power-law
Dog, Chr.12, GC-rich region (3.5-16 Mb), MIRb-SINE/MIR

max. divergence = 28%, no power-law
Dog, Chr.12, GC-poor region (25-45 Mb), MIRb-SINE/MIR

max. divergence = 28%, no power-law
Dog, Chr.X, GC-rich region (0.4-10.4 Mb), SINEC_a1-SINE/Lys

max. divergence = 8%, E=1.15, μ=0.67
Dog, Chr.X, GC-poor region (64-76 Mb), SINEC_a1-SINE/Lys

max. divergence = 9%, E=1.13, μ=0.35
Dog, Chr.X, GC-rich region (0.4-10.4 Mb), SINEC_c2-SINE/Lys

max. divergence = 18%, E=0.90, μ=1.60
Dog, Chr.X, GC-poor region (64-76 Mb), SINEC_c2-SINE/Lys
max. divergence = 18%, no power-law