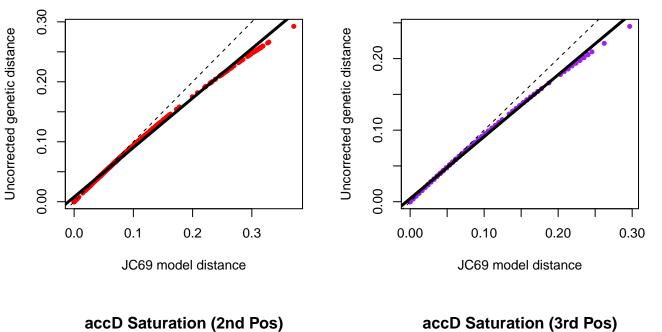
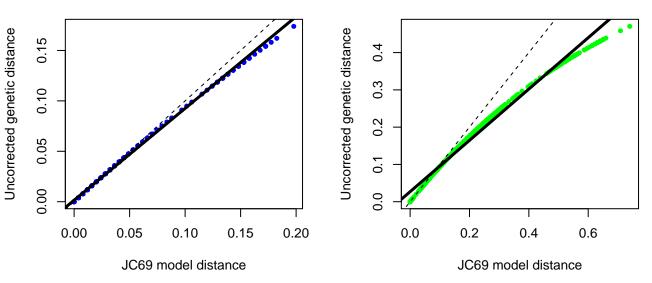


accD Saturation (All Bases)

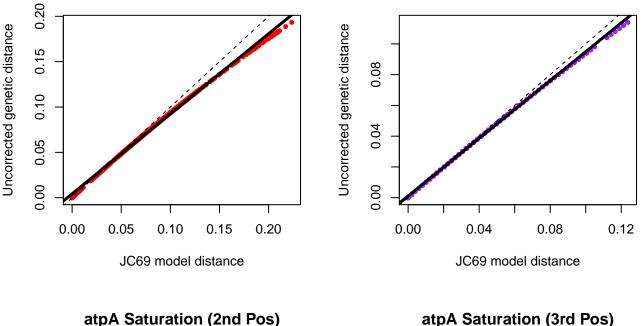


accD Saturation (3rd Pos)

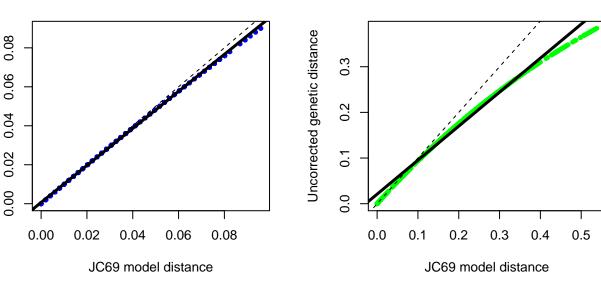




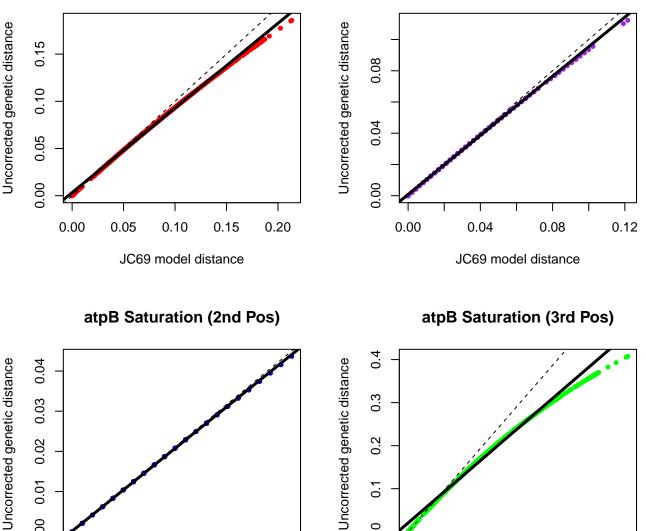
Uncorrected genetic distance

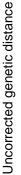


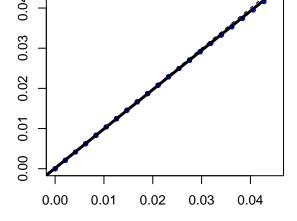


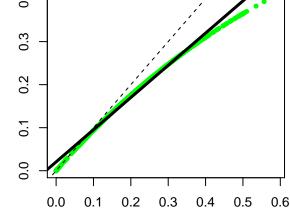




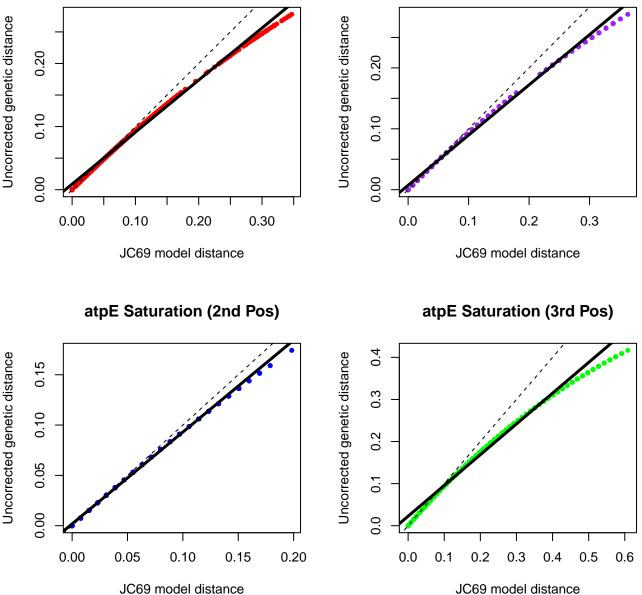




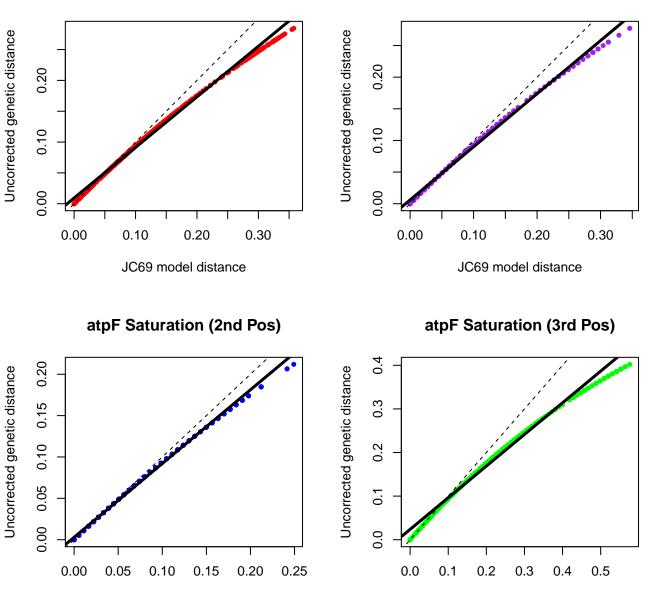










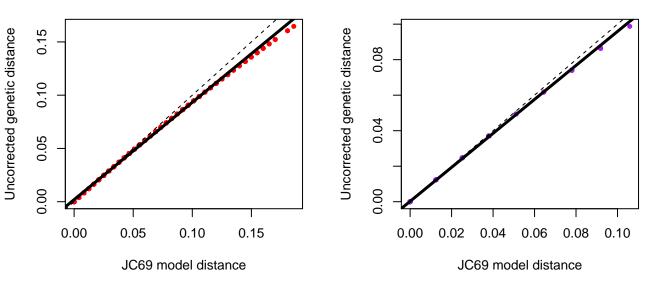


JC69 model distance

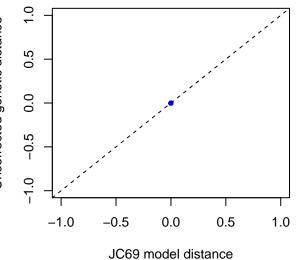
JC69 model distance



atpH Saturation (All Bases)

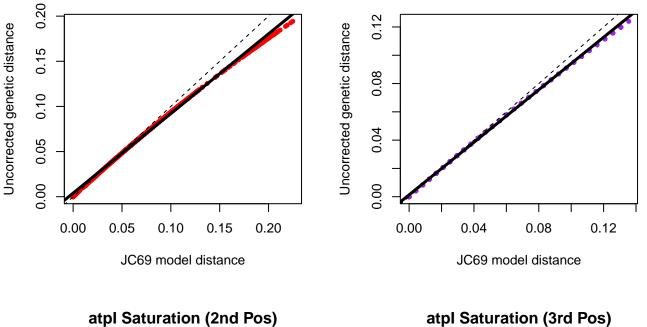


atpH Saturation (2nd Pos)

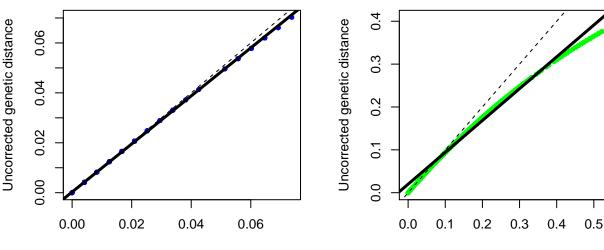


Uncorrected genetic distance





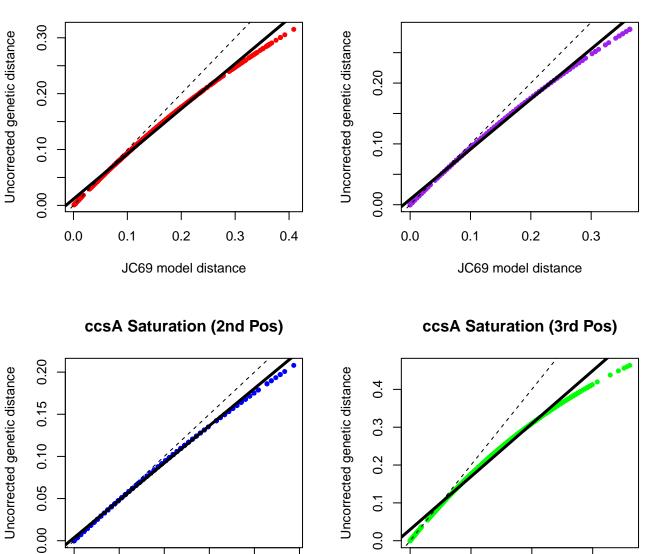




JC69 model distance



ccsA Saturation (All Bases)



JC69 model distance

0.10

0.15

0.20

0.25

0.0

0.00

0.05

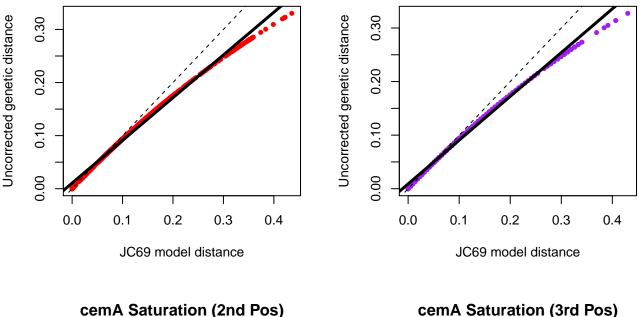
JC69 model distance

0.4

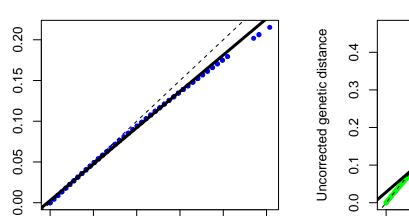
0.6

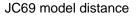


cemA Saturation (All Bases)









0.4

0.6

0.2

JC69 model distance

0.15

0.20

0.25

0.0

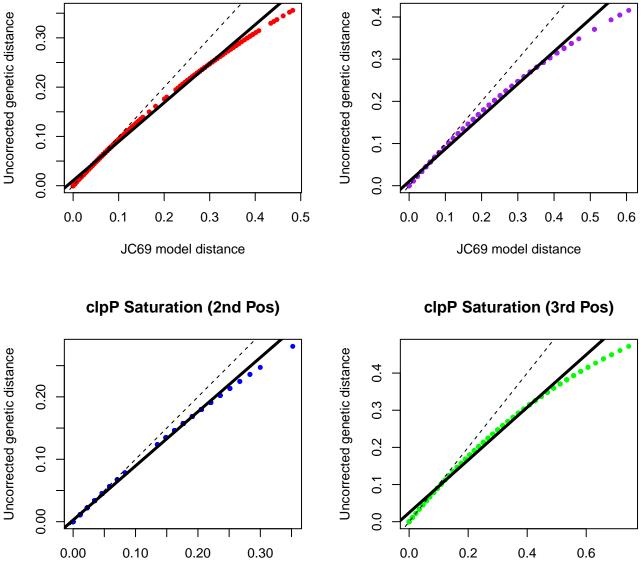
0.10

Uncorrected genetic distance

0.00



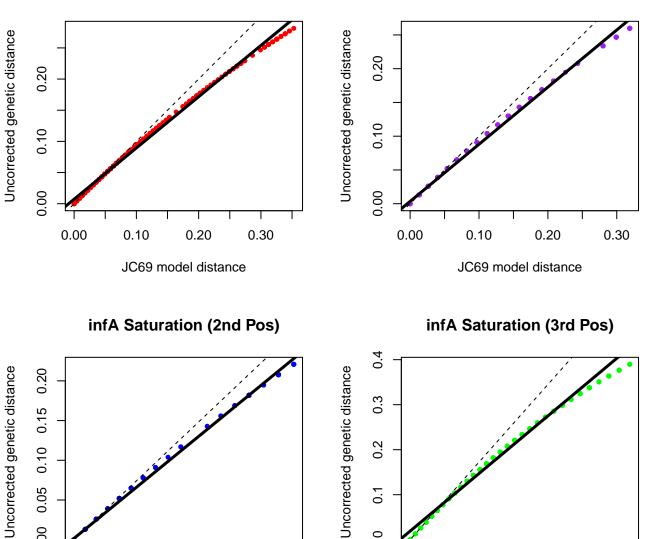
clpP Saturation (All Bases)



JC69 model distance



infA Saturation (All Bases)



0.0

0.0

0.1

JC69 model distance

0.15

0.20

0.25

0.10

0.00

0.00

0.05

JC69 model distance

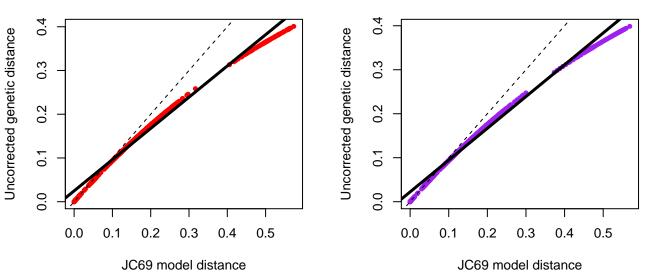
0.3

0.4

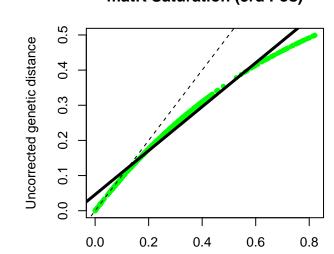
0.5



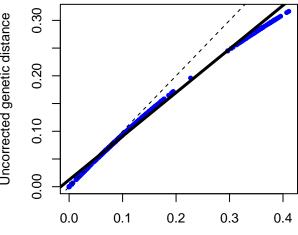
matK Saturation (All Bases)









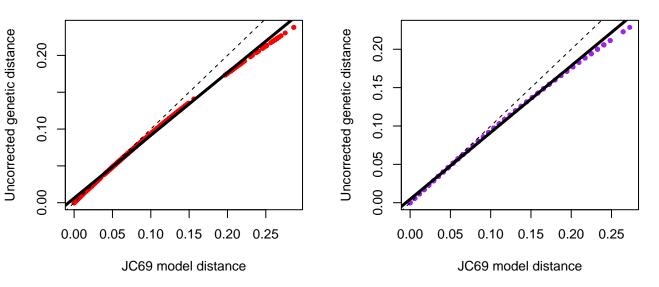


JC69 model distance

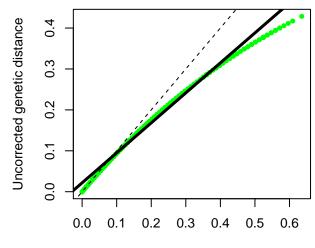
JC69 model distance



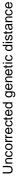
ndhA Saturation (All Bases)

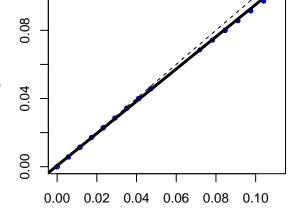


ndhA Saturation (3rd Pos)





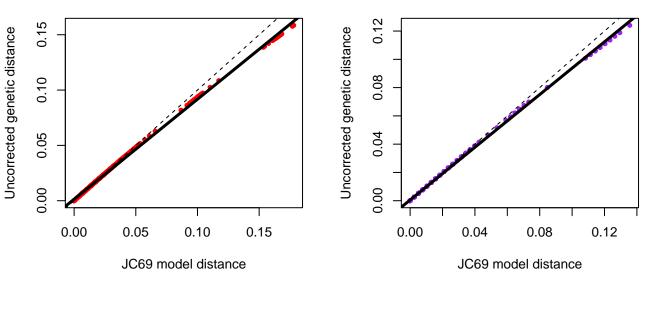




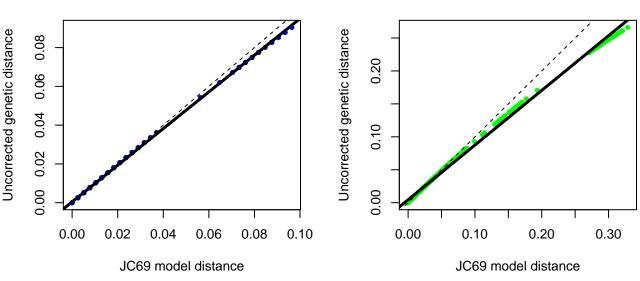
JC69 model distance



ndhB Saturation (2nd Pos)

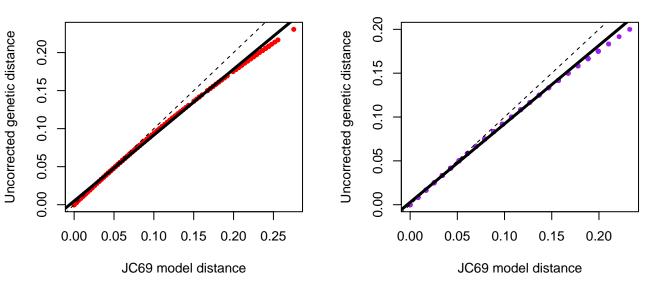


ndhB Saturation (3rd Pos)

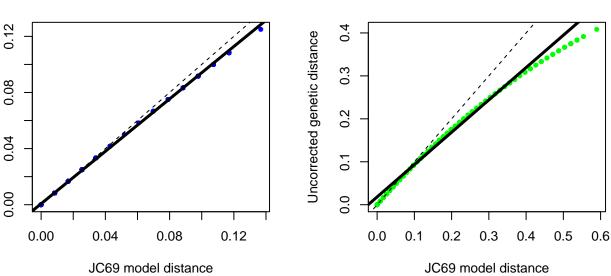




ndhC Saturation (2nd Pos)

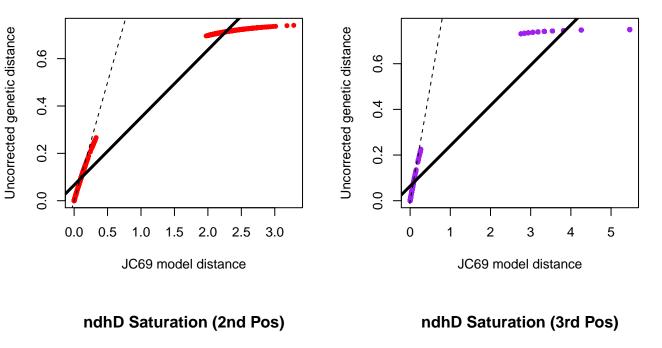


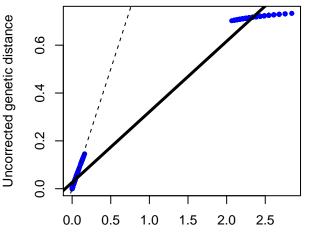


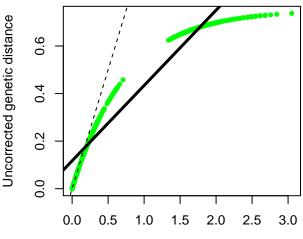




ndhD Saturation (All Bases)

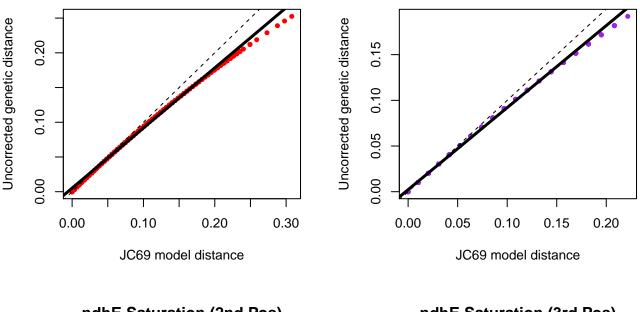




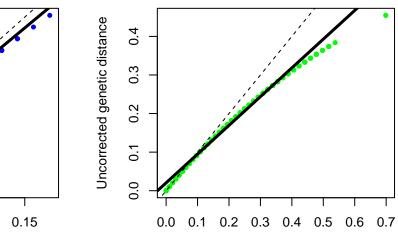


JC69 model distance



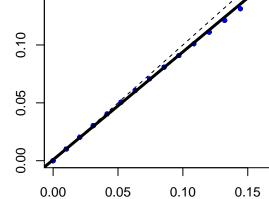










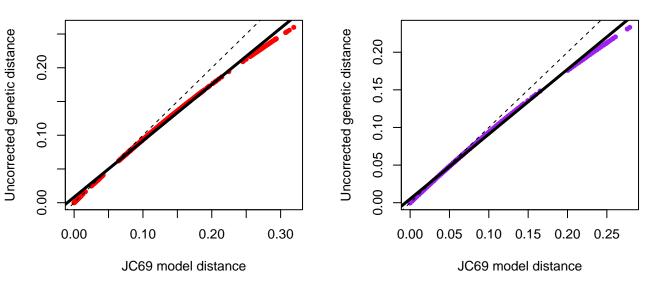


JC69 model distance

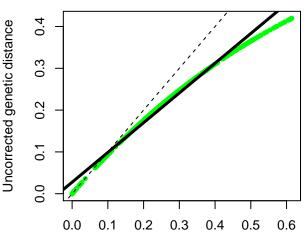
JC69 model distance



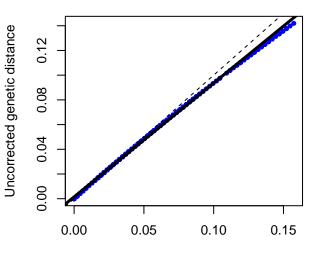
ndhF Saturation (All Bases)



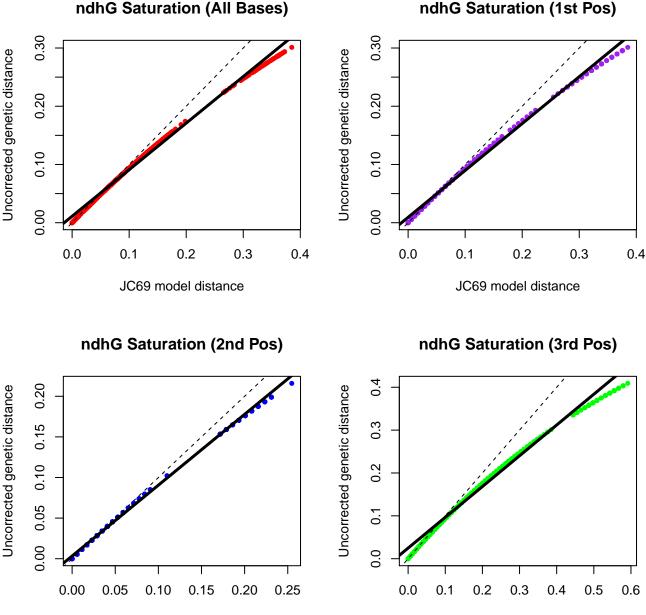
ndhF Saturation (3rd Pos)







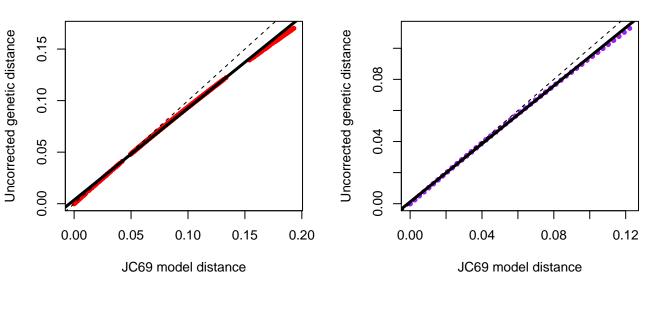
JC69 model distance



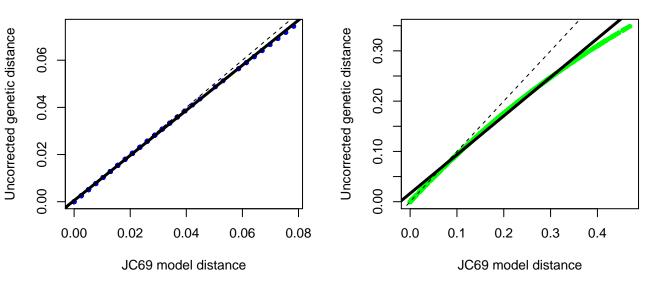
JC69 model distance

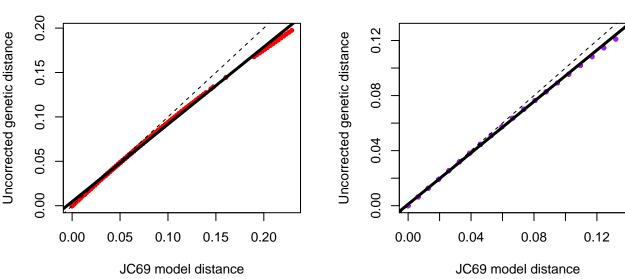


ndhH Saturation (2nd Pos)



ndhH Saturation (3rd Pos)

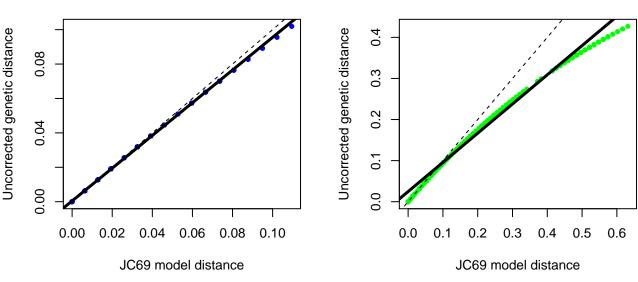




ndhl Saturation (2nd Pos)

ndhl Saturation (3rd Pos)

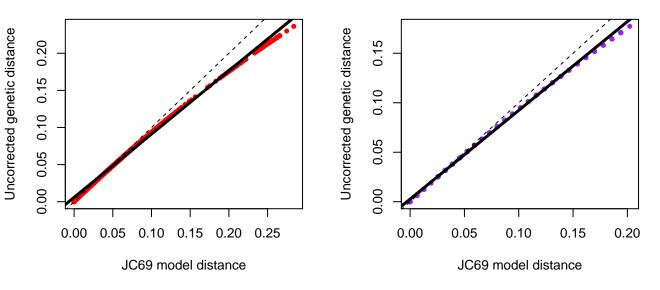
ndhl Saturation (1st Pos)



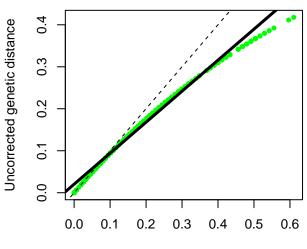
ndhl Saturation (All Bases)



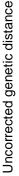
ndhJ Saturation (All Bases)

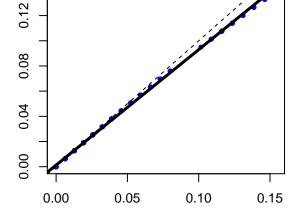


ndhJ Saturation (3rd Pos)



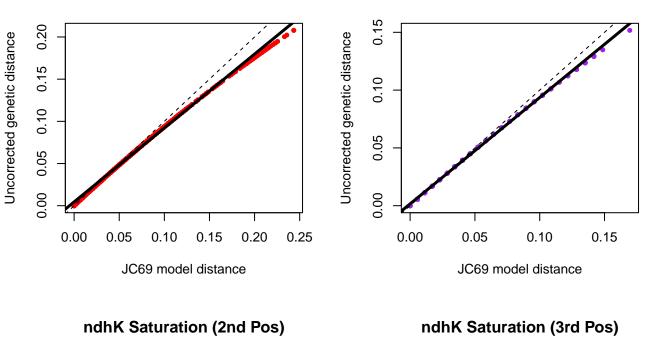
ndhJ Saturation (2nd Pos)





JC69 model distance



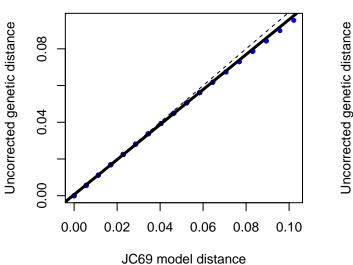


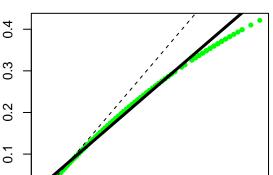
0.0

0.0

0.1

0.2



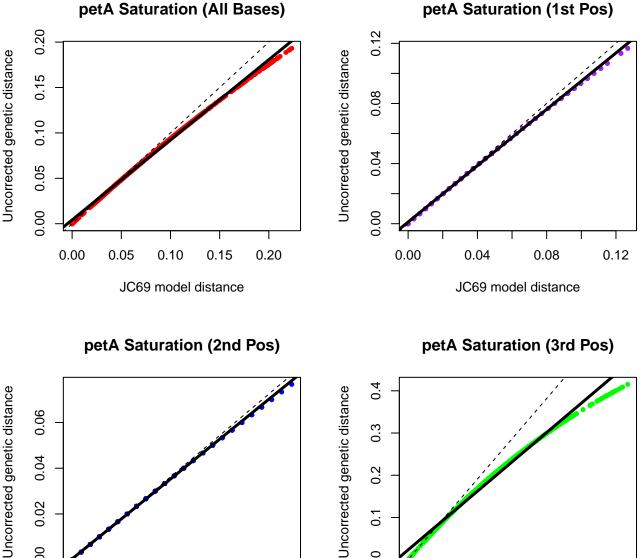


JC69 model distance

0.3

0.4

0.5



0.0

0.0

0.1

0.2



0.02 0.00

0.00

JC69 model distance

0.04

0.06

0.08

0.02

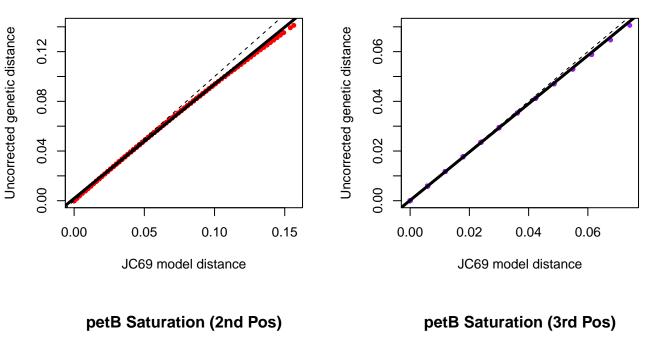
JC69 model distance

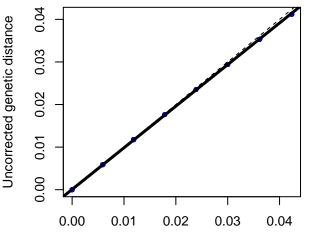
0.3

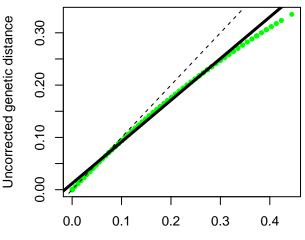
0.4

0.5





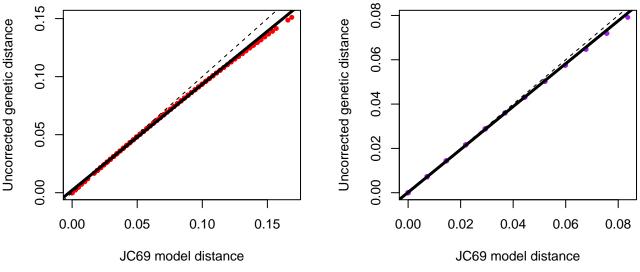




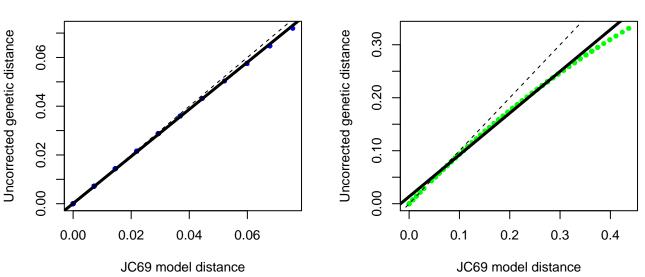


petD Saturation (All Bases)

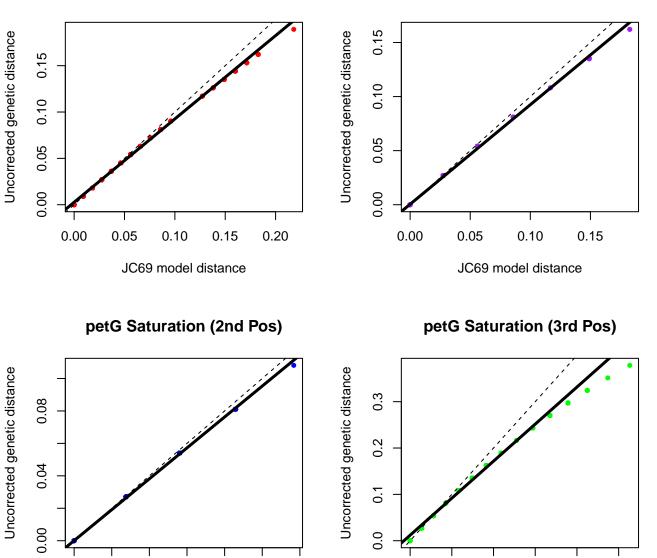
petD Saturation (2nd Pos)



petD Saturation (3rd Pos)







0.06

0.08

0.10

0.12

0.00

0.02

0.04

JC69 model distance

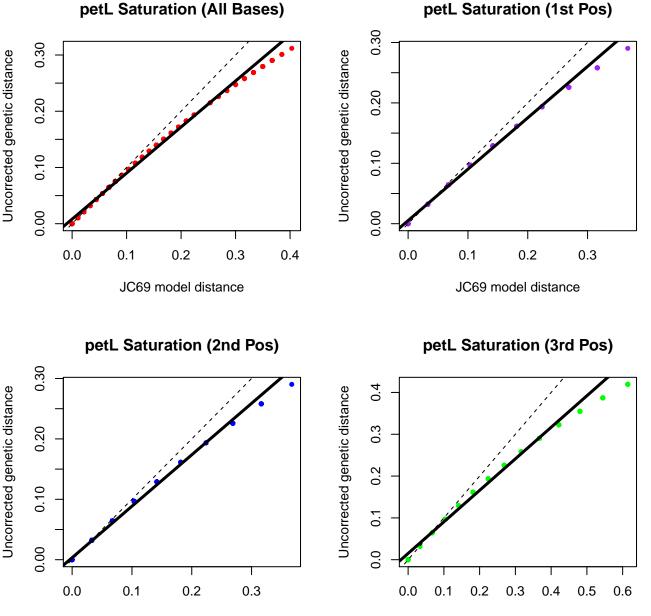
0.3

0.4

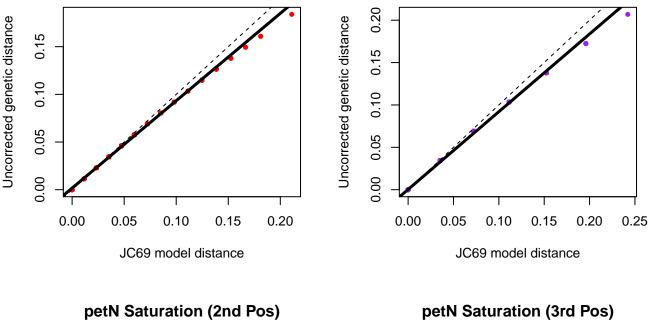
0.5

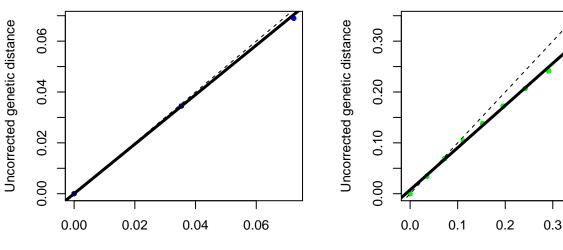
0.2

0.0







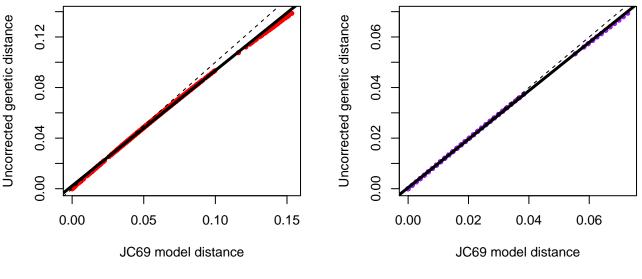


JC69 model distance

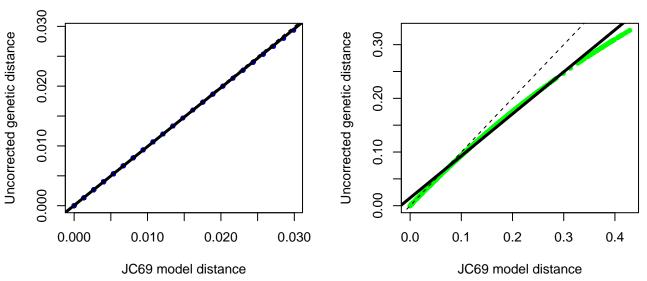


psaA Saturation (All Bases)

psaA Saturation (2nd Pos)

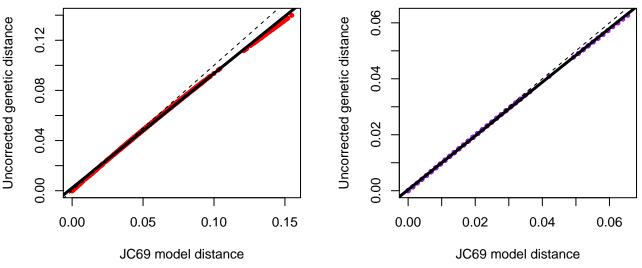




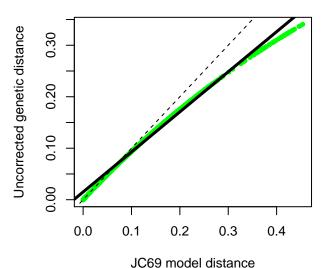




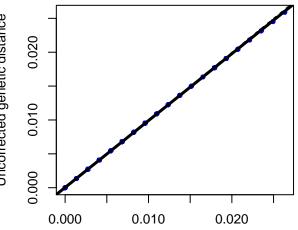
psaB Saturation (All Bases)



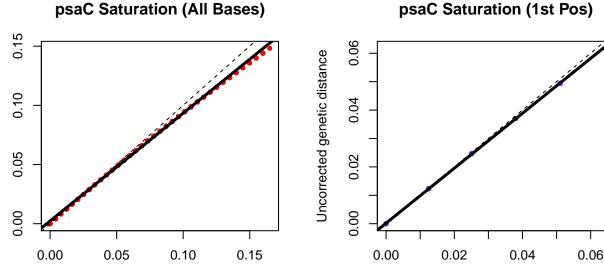
psaB Saturation (3rd Pos)







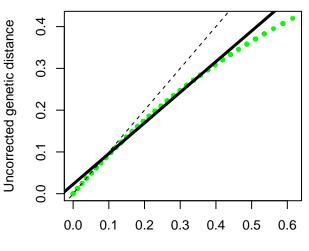
JC69 model distance

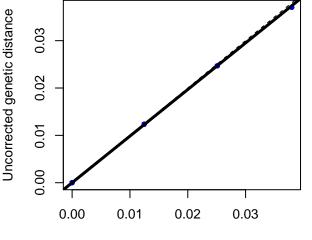


Uncorrected genetic distance

psaC Saturation (3rd Pos)

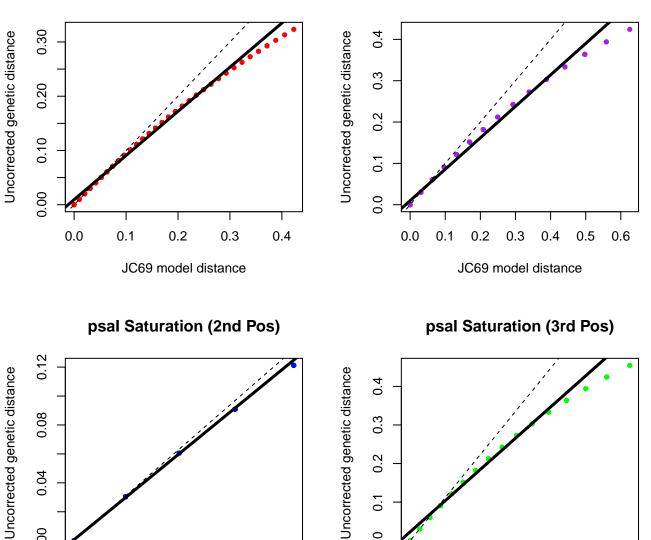






JC69 model distance





0.1

0.0

0.0

0.1

0.2

JC69 model distance

0.08

0.12

0.04

0.00

0.00

JC69 model distance

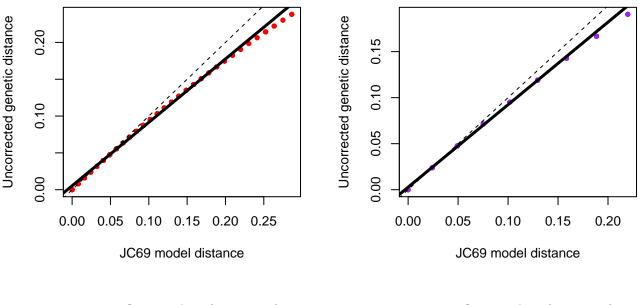
0.4

0.5

0.6

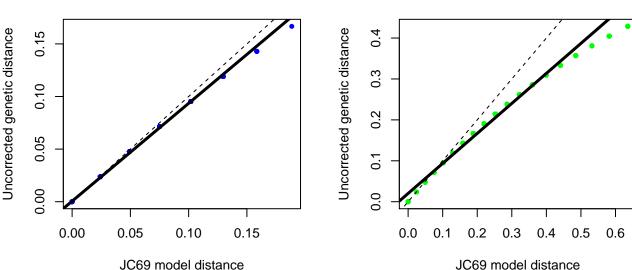
0.7





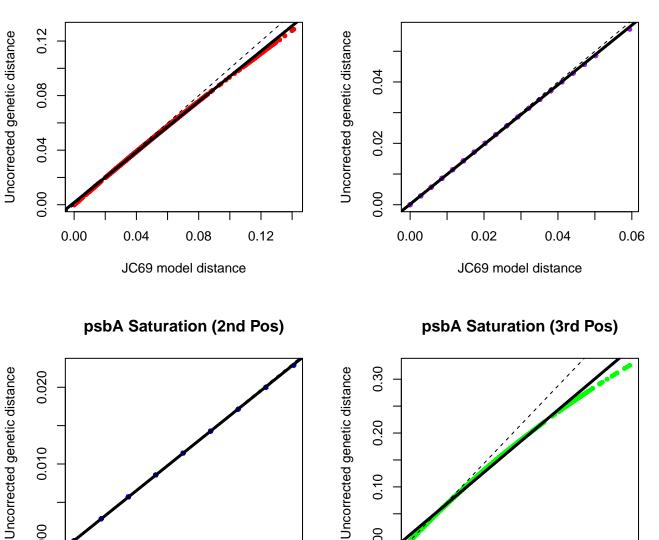








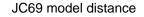
psbA Saturation (All Bases)



0.00

0.0

0.1



0.015

0.020

0.010

0.000

0.000

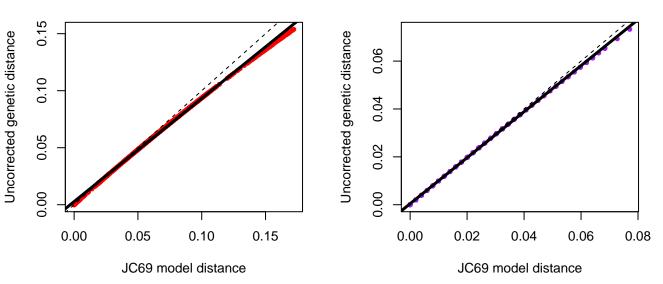
0.005



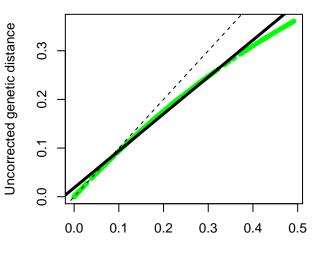
0.3

0.4



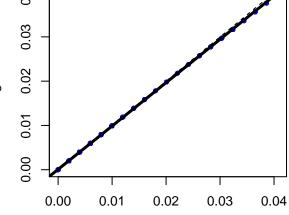


psbB Saturation (3rd Pos)





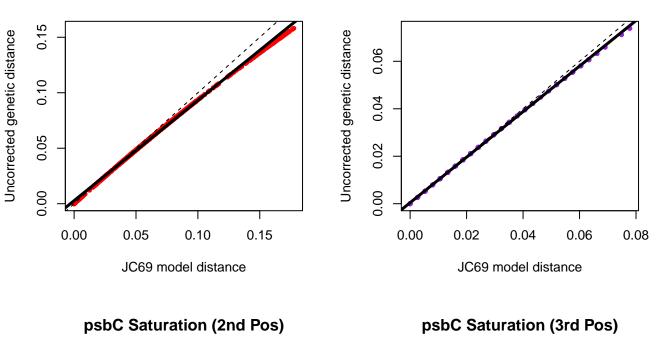


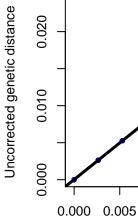


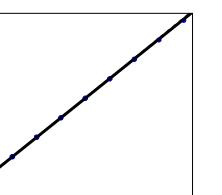
JC69 model distance

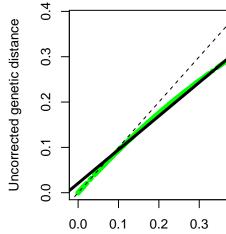


psbC Saturation (All Bases)









0.015

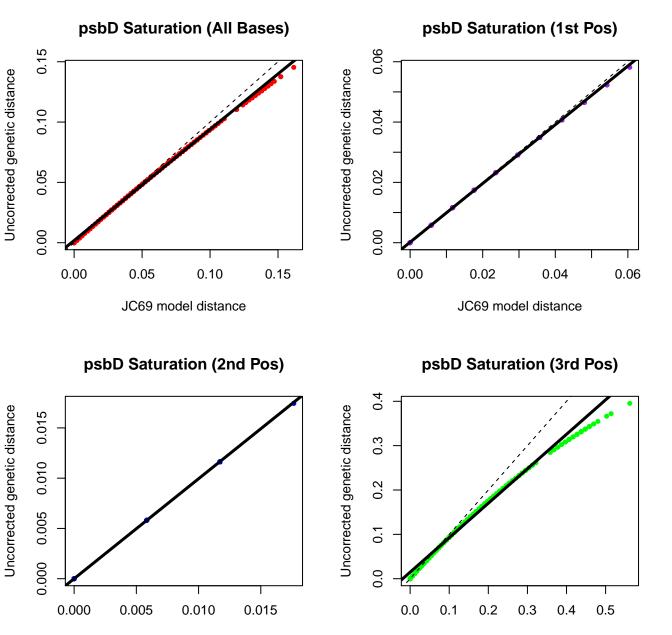
0.020

0.025

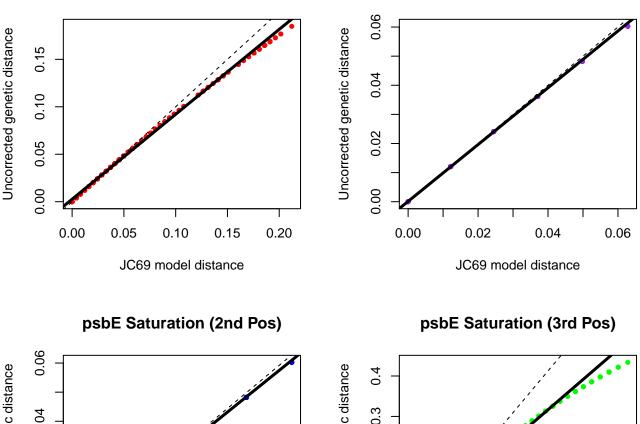
0.010

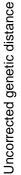
JC69 model distance

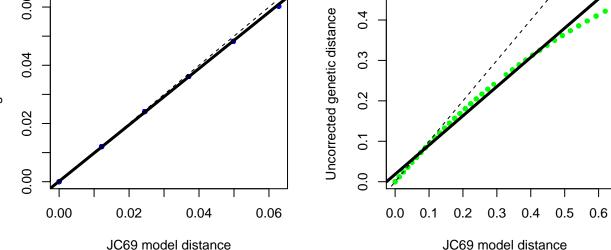
0.4

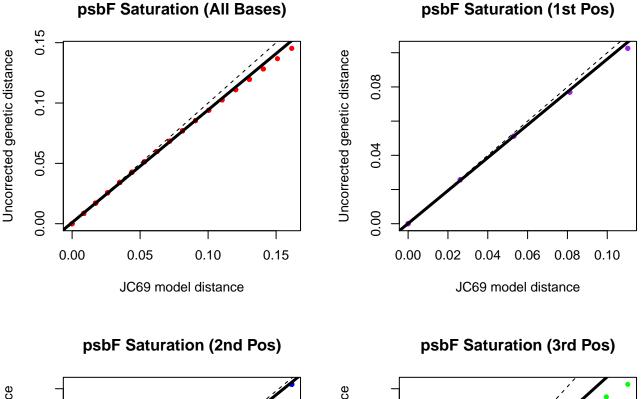








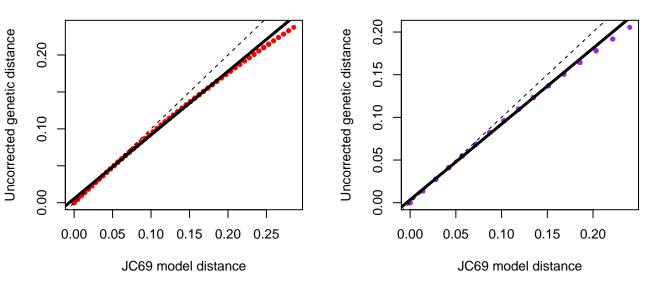




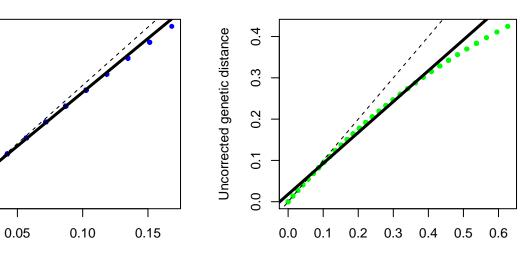


Uncorrected genetic distance 0.30 0.04 0.20 0.02 0.10 0.00 0.00 0.2 0.00 0.01 0.02 0.03 0.04 0.05 0.0 0.1 0.3 0.4 0.5 JC69 model distance JC69 model distance









psbH Saturation (2nd Pos)

0.15

0.10

0.05

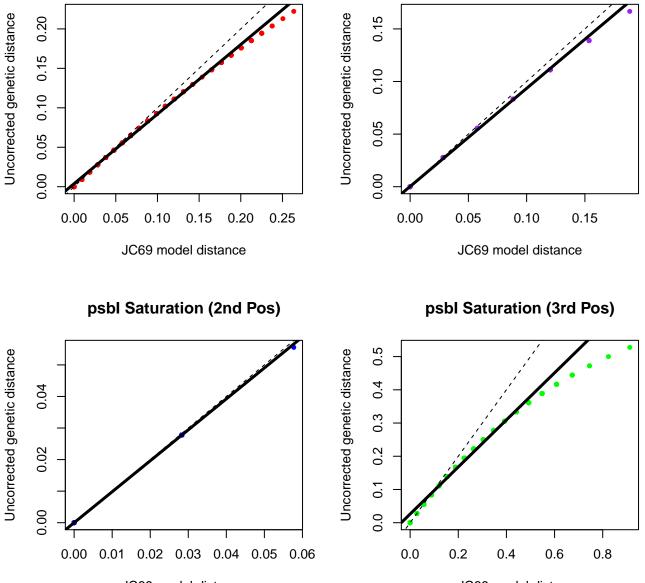
0.00

0.00

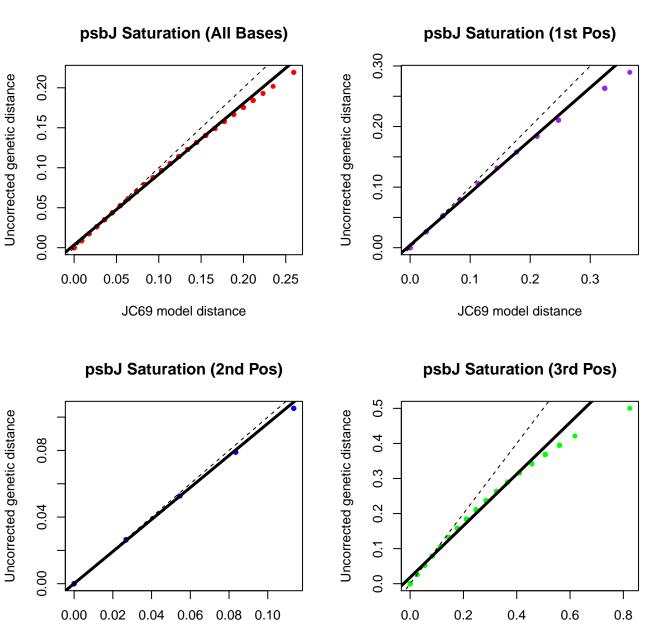
JC69 model distance

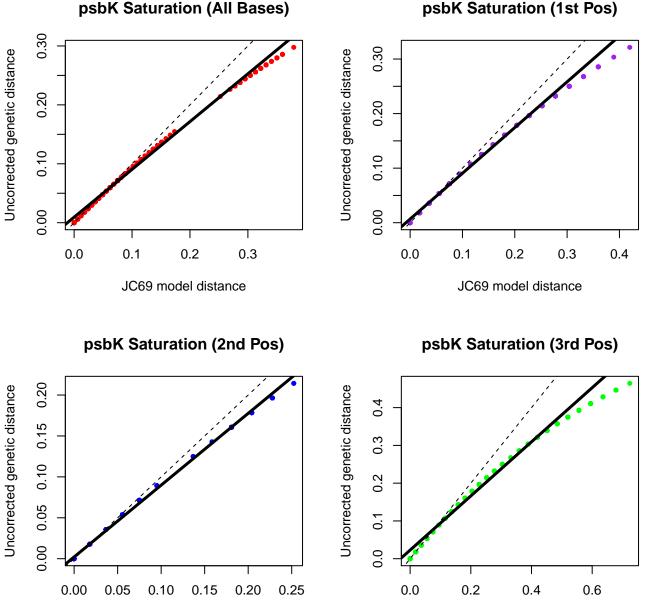


psbl Saturation (All Bases)



JC69 model distance



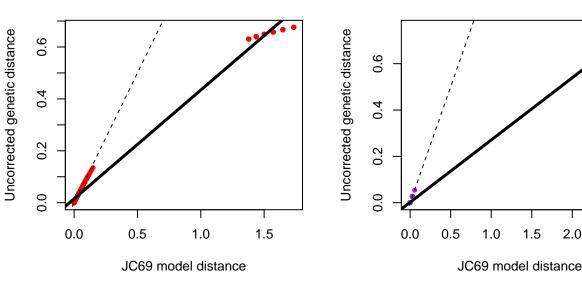


psbK Saturation (All Bases)

JC69 model distance

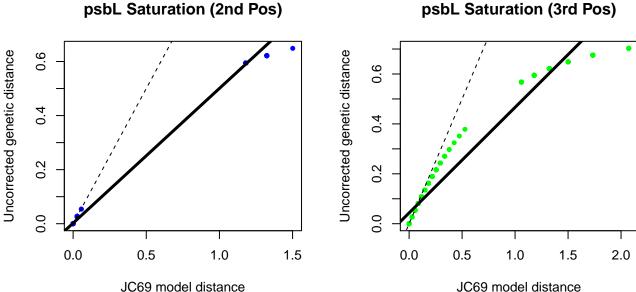
psbL Saturation (All Bases)

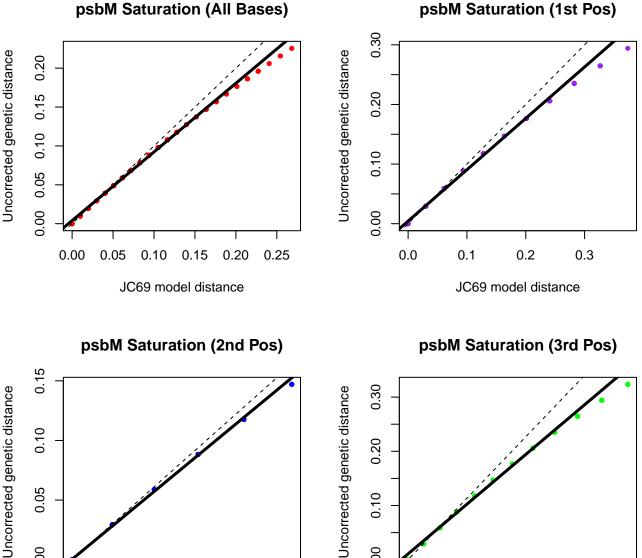
psbL Saturation (1st Pos)





2.5





0.00

0.0

0.1



0.05 0.00

0.00

JC69 model distance

0.10

0.15

0.05

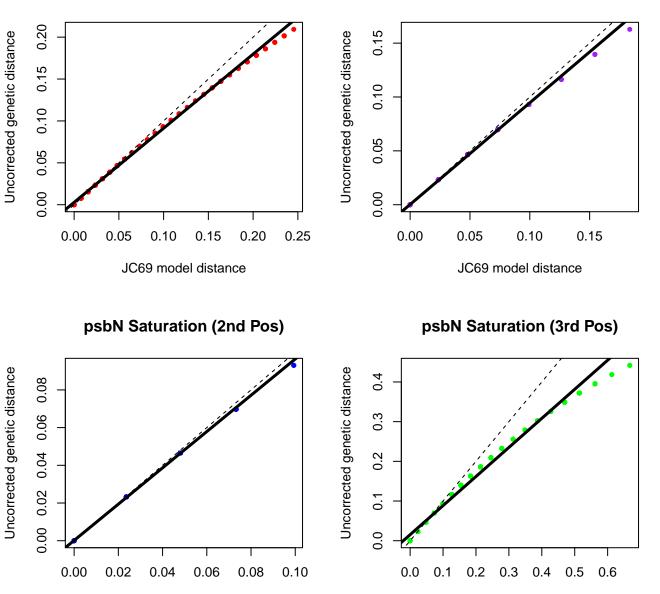
JC69 model distance

0.2

0.3

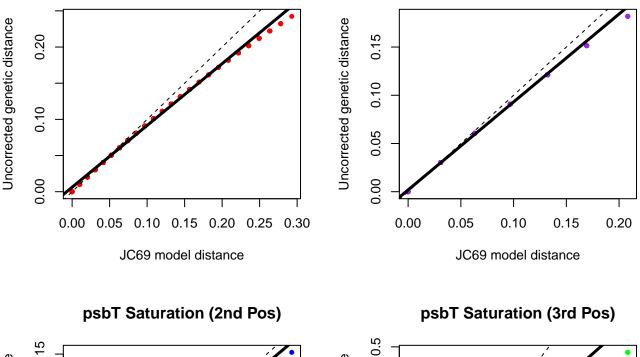


psbN Saturation (All Bases)

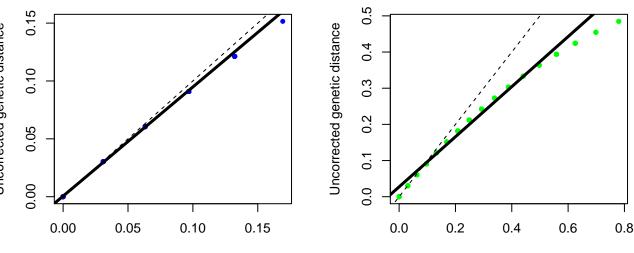


JC69 model distance





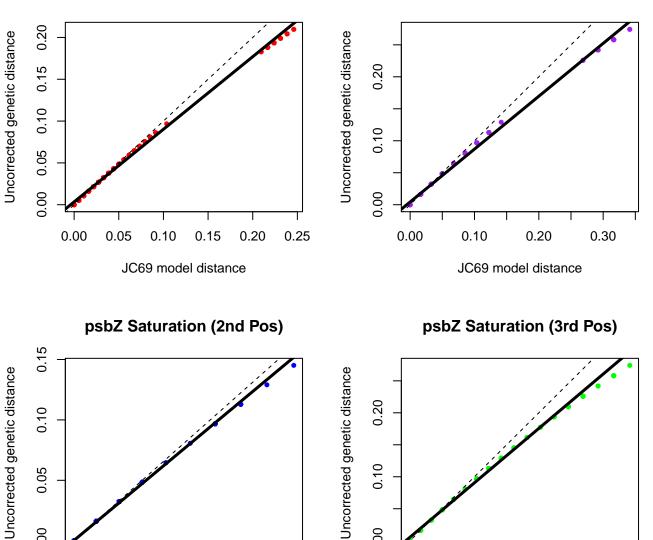




JC69 model distance



psbZ Saturation (All Bases)



0.00

0.00

0.10

0.15

0.05

0.00

0.00

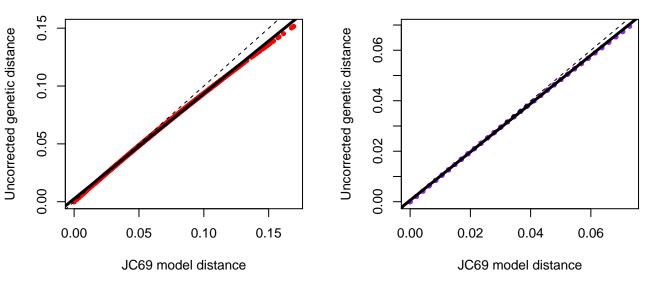


0.20

0.30

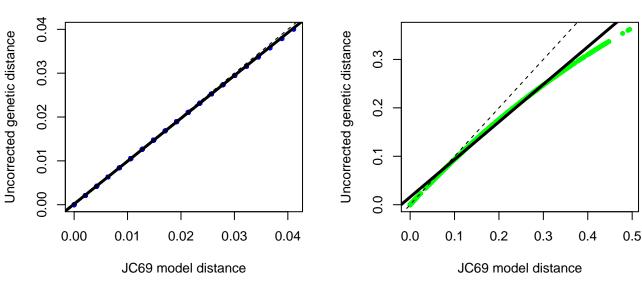






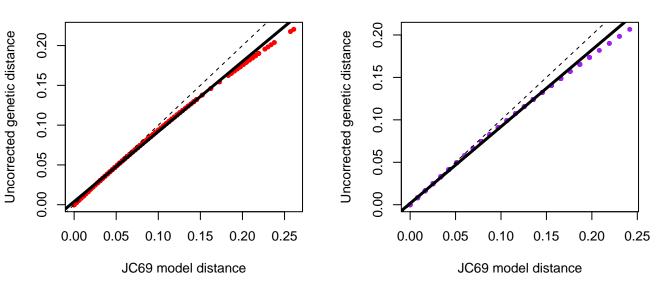




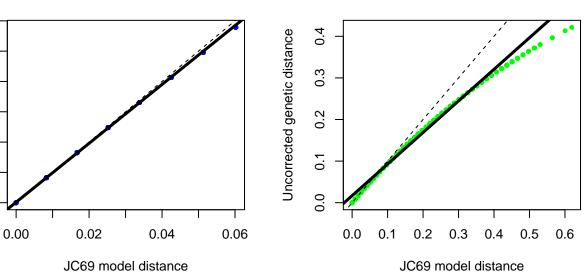




rpl14 Saturation (2nd Pos)



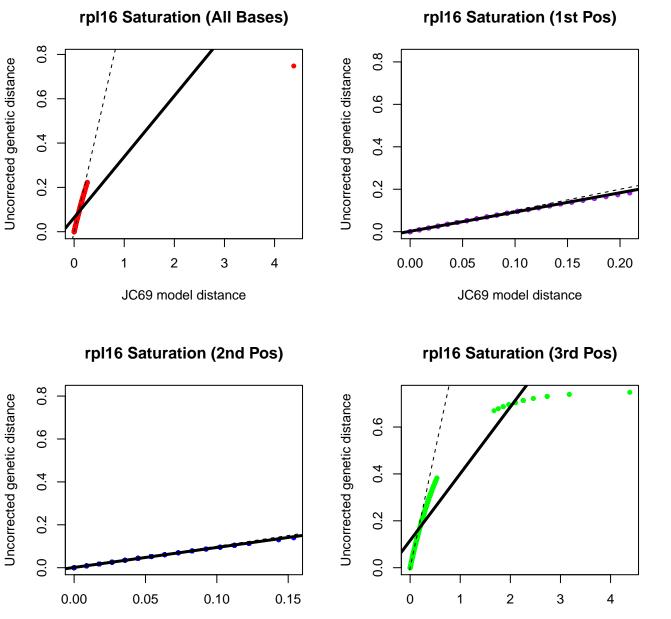
rpl14 Saturation (3rd Pos)



0.06

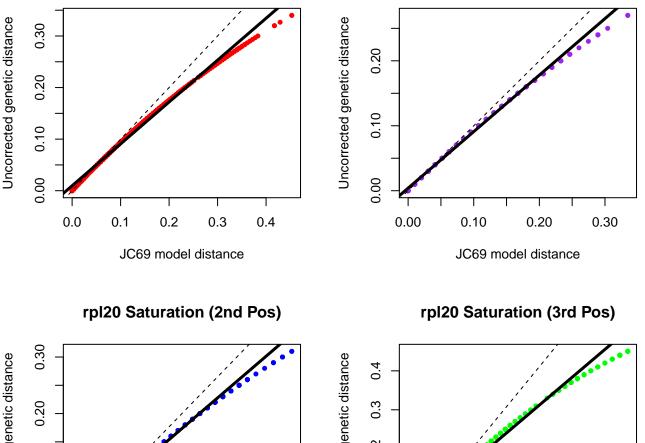
0.04

0.02





rpl20 Saturation (All Bases)



Uncorrected genetic distance

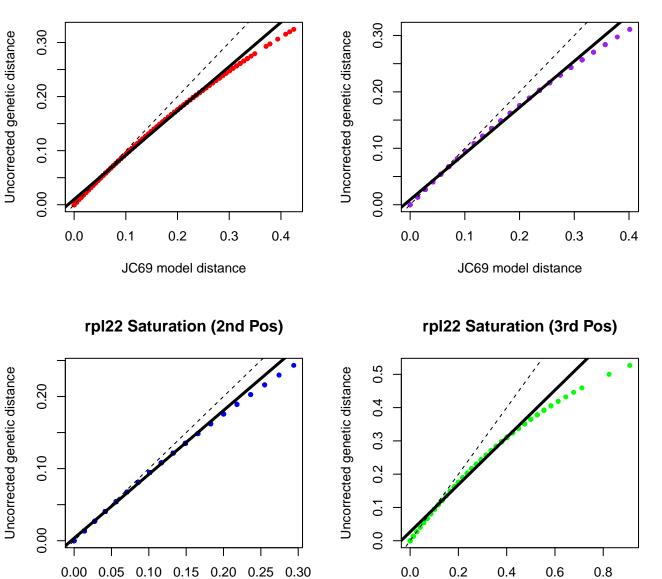
Uncorrected genetic distance 0.10 0.00 0.0 0.1 0.2 0.3 0.4

JC69 model distance

0.2 0.1 0.0 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7



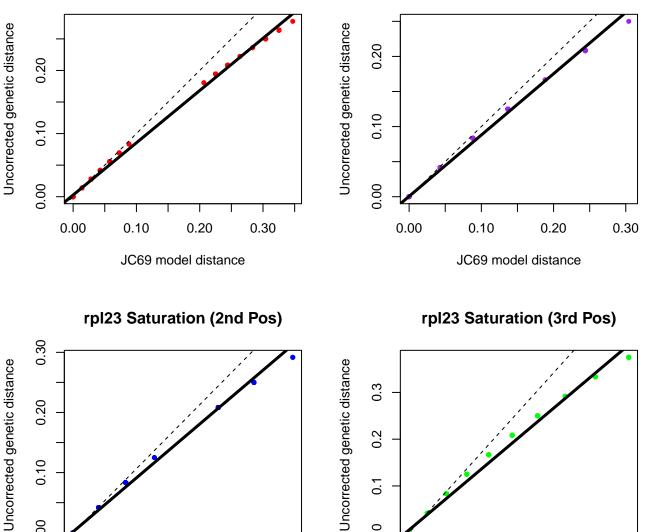
rpl22 Saturation (All Bases)



JC69 model distance



rpl23 Saturation (All Bases)



0.0

0.0

0.1



0.00

0.0

JC69 model distance

0.1

0.2

0.3

JC69 model distance

0.3

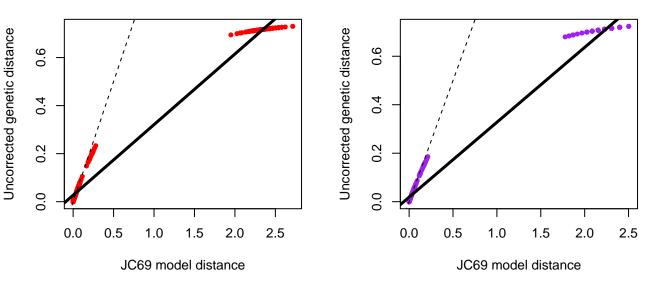
0.4

0.5

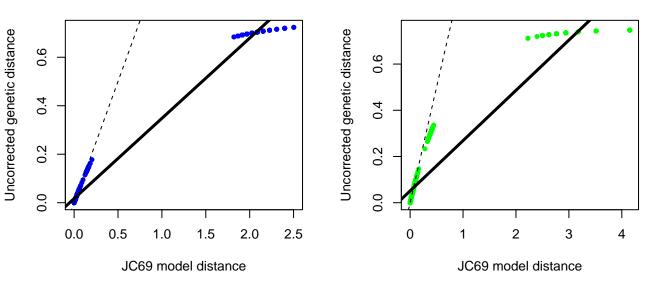


rpl2 Saturation (2nd Pos)

rpl2 Saturation (1st Pos)

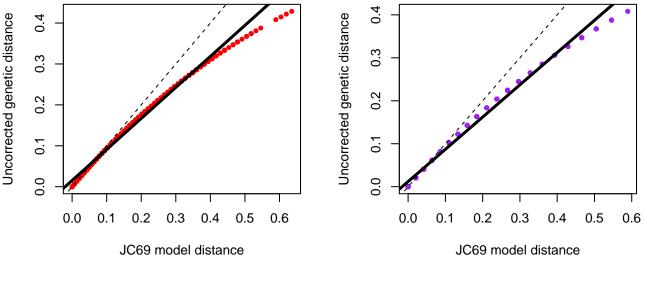




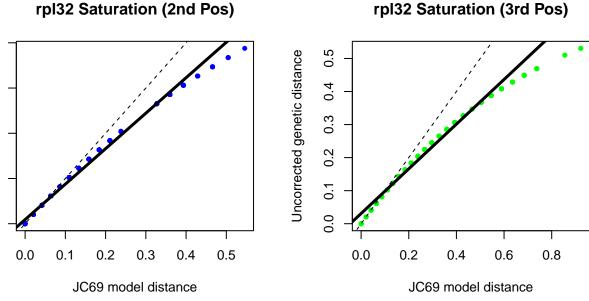




rpl32 Saturation (All Bases)



rpl32 Saturation (3rd Pos)



0.4

0.3

0.2

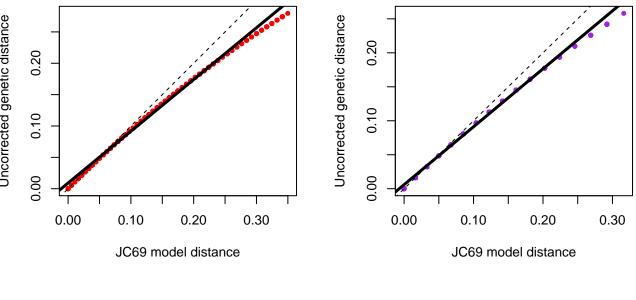
0.1

0.0

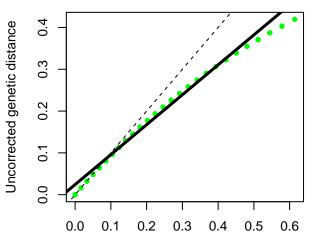
Uncorrected genetic distance



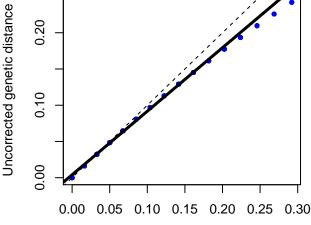
rpl33 Saturation (All Bases)



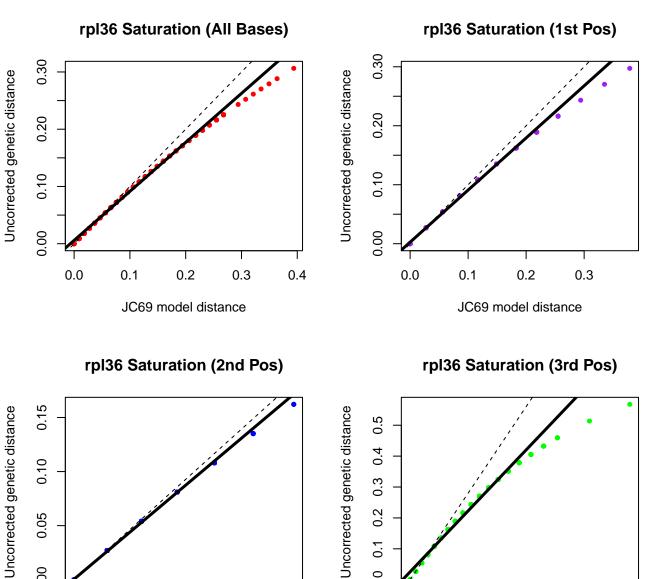








JC69 model distance



0.0

0.0

0.2



0.05

0.10

0.15

0.00

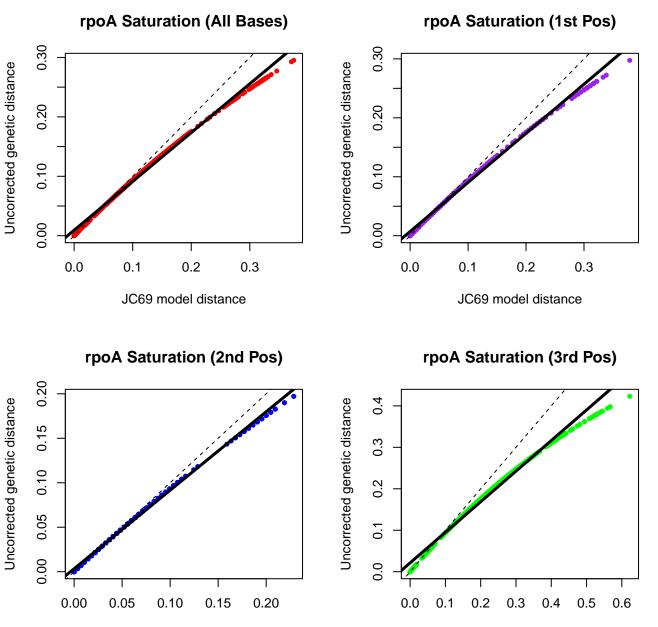
0.00

JC69 model distance

0.6

0.8

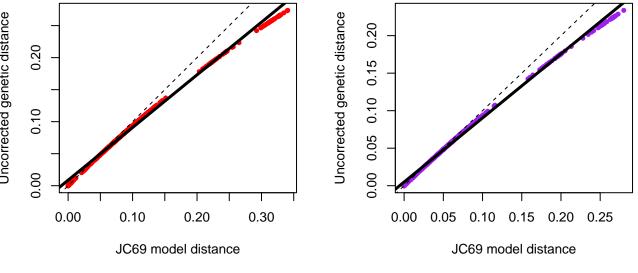
1.0



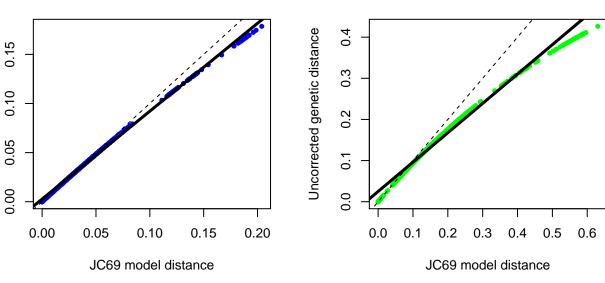


rpoB Saturation (All Bases)

rpoB Saturation (2nd Pos)



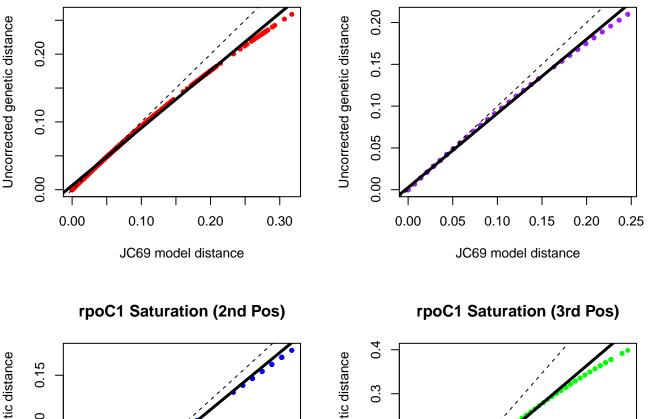
rpoB Saturation (3rd Pos)



Uncorrected genetic distance



rpoC1 Saturation (All Bases)



Uncorrected genetic distance

0.00 0.05 0.10 0.15 0.20 JC69 model distance

JC69 model distance

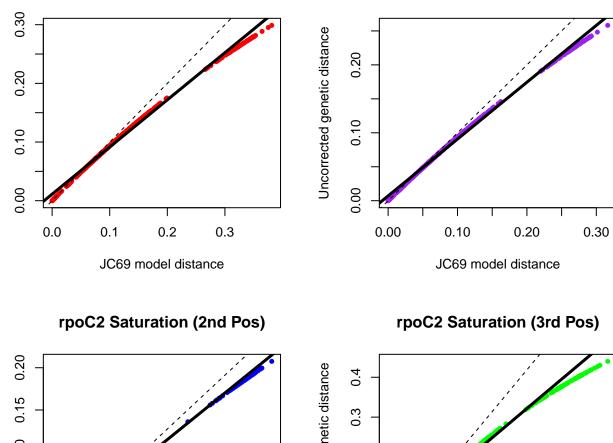
0.3

0.2

0.0

0.1

0.4



rpoC2 Saturation (All Bases)

rpoC2 Saturation (1st Pos)

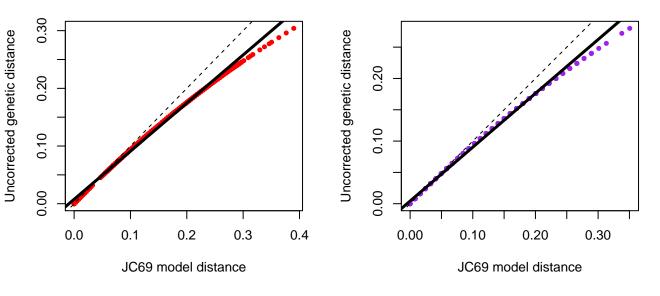
Uncorrected genetic distance

Uncorrected genetic distance 0.10 0.2 0.05 0.1 0.00 0.0 0.2 0.00 0.05 0.10 0.15 0.20 0.25 0.0 0.1 0.3 0.4 0.5 0.6 JC69 model distance JC69 model distance

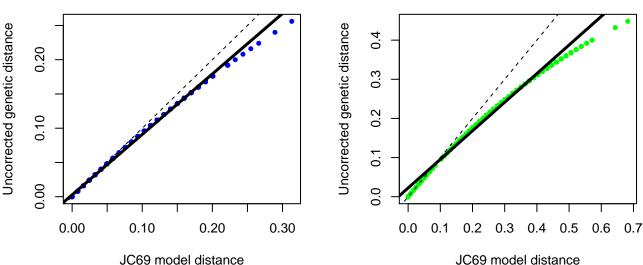


rps11 Saturation (2nd Pos)



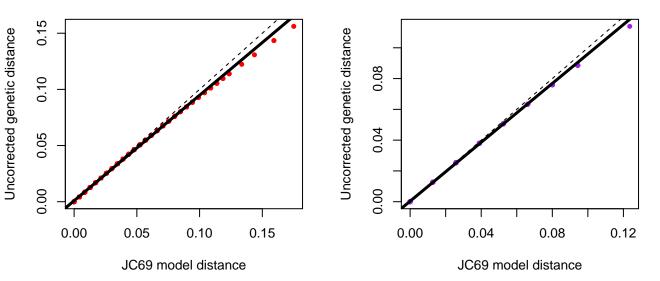




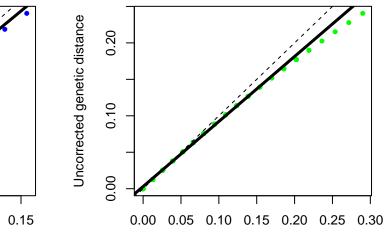




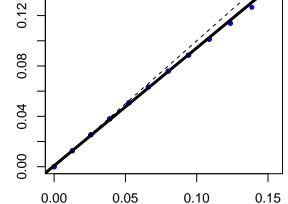




rps12 Saturation (3rd Pos)

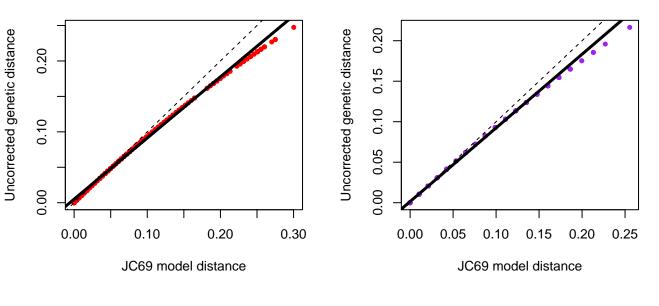


rps12 Saturation (2nd Pos)

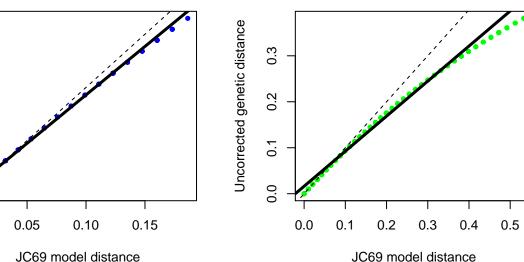


JC69 model distance







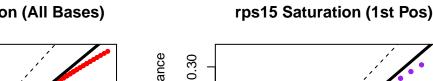


rps14 Saturation (2nd Pos)

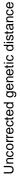
0.10

0.05

0.00







Uncorrected genetic distance 0.3 0.20 0.2 0.10 0.1 0.00 0.0 0.0 0.2 0.1 0.2 0.3 0.4 0.5 0.0 0.1

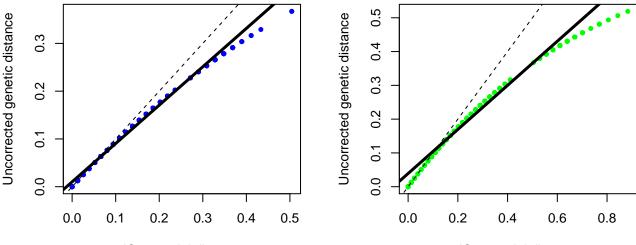
JC69 model distance

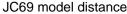


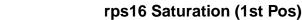
JC69 model distance

0.3

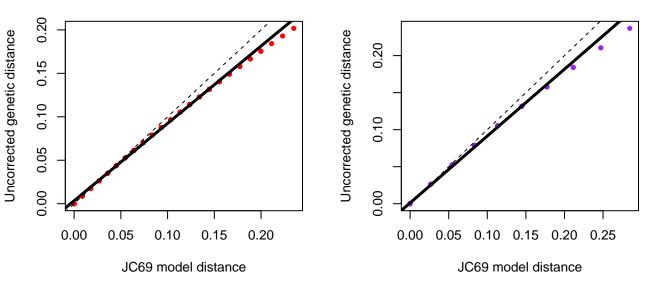




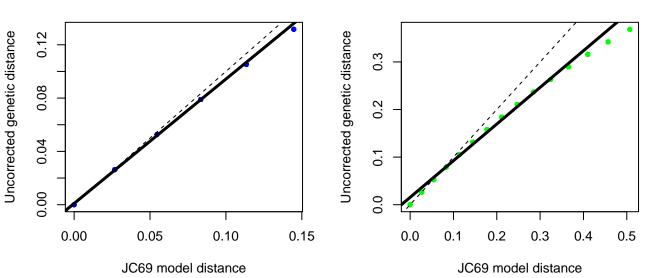




rps16 Saturation (2nd Pos)



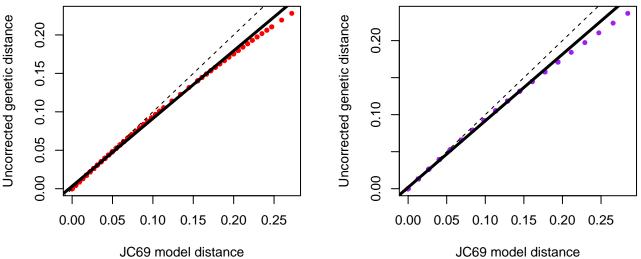




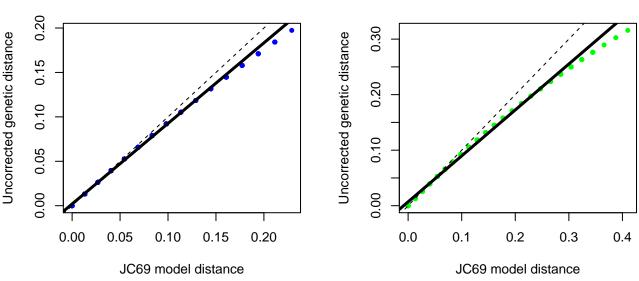


rps18 Saturation (All Bases)

rps18 Saturation (2nd Pos)

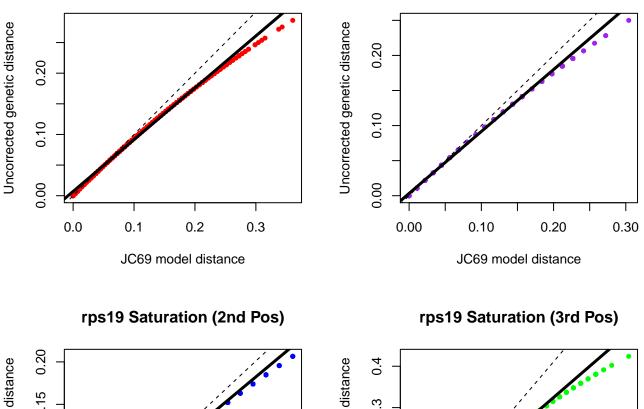


rps18 Saturation (3rd Pos)

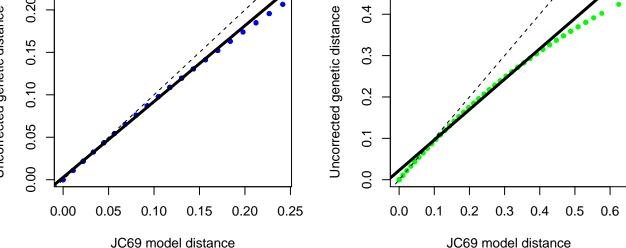




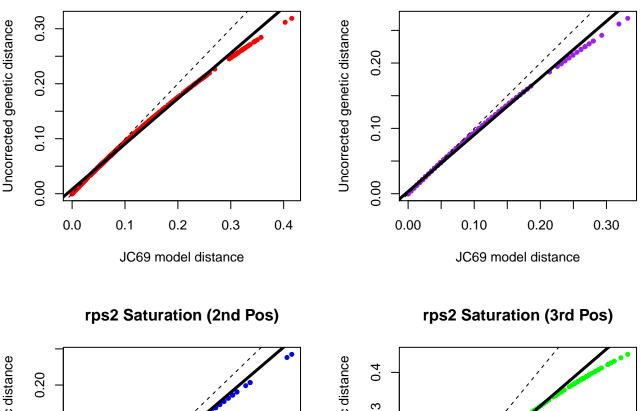
rps19 Saturation (All Bases)



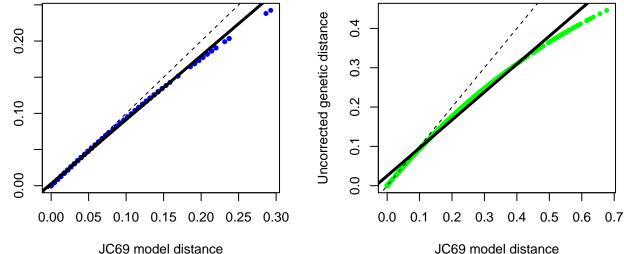
Uncorrected genetic distance

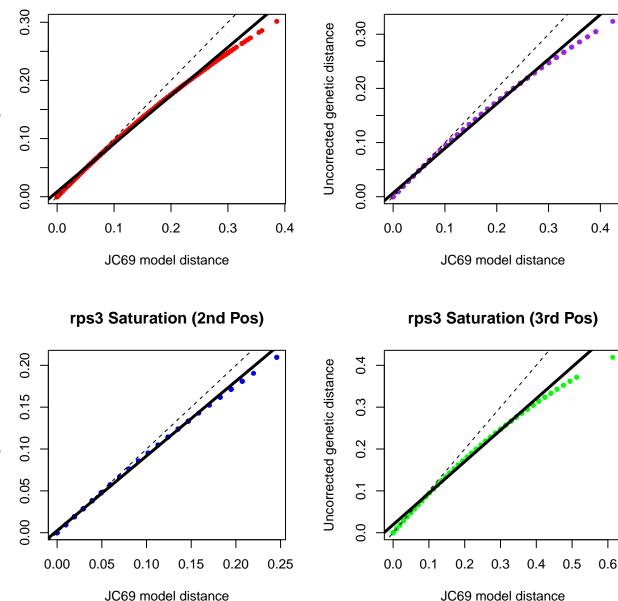










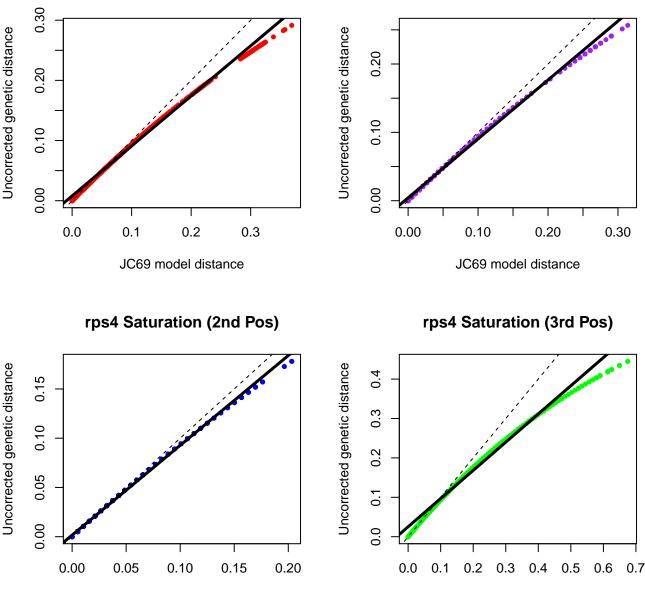


rps3 Saturation (1st Pos)

rps3 Saturation (All Bases)

Uncorrected genetic distance

Uncorrected genetic distance

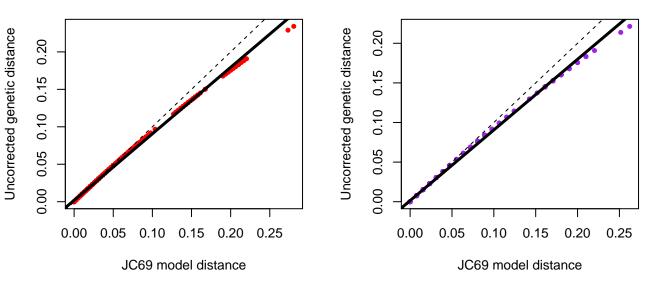


rps4 Saturation (All Bases)

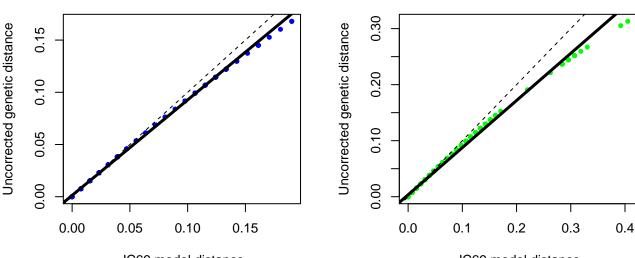
JC69 model distance

rps4 Saturation (1st Pos)





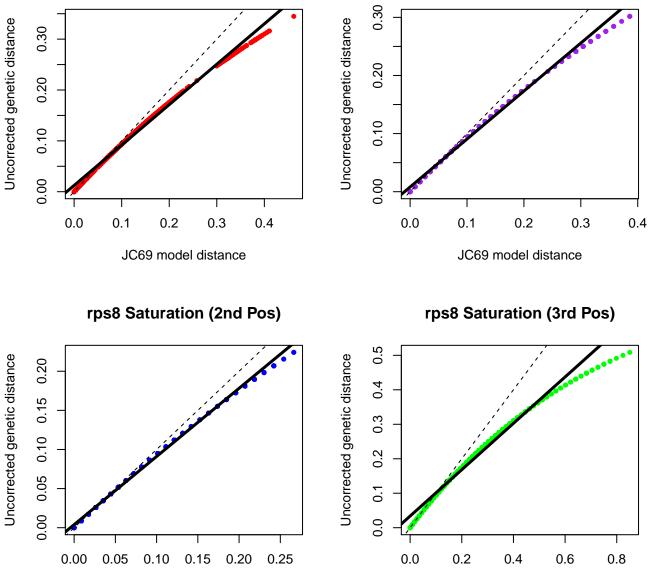
rps7 Saturation (3rd Pos)

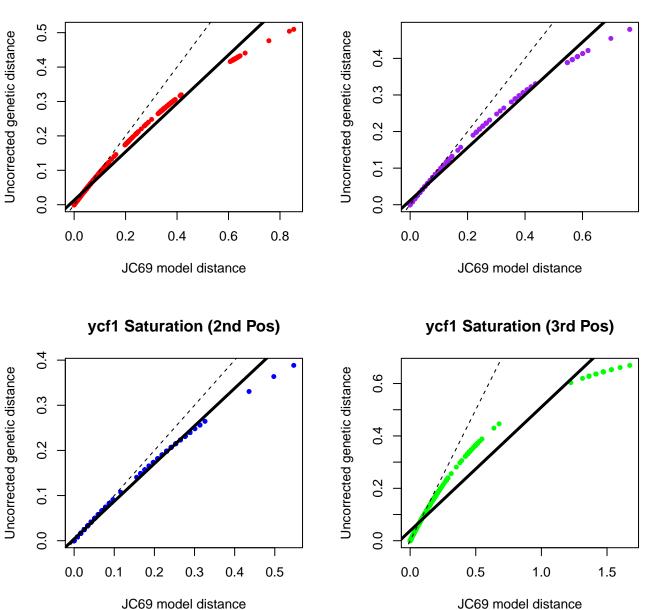


rps7 Saturation (2nd Pos)

JC69 model distance





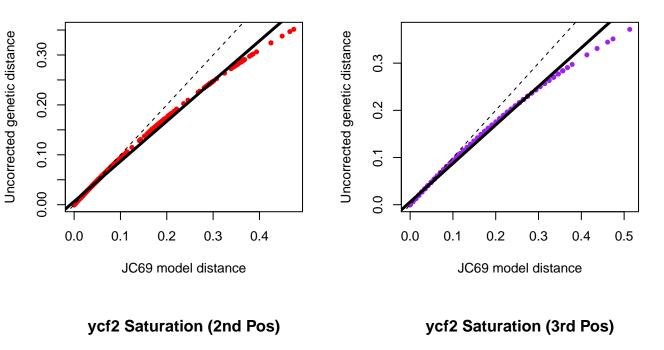


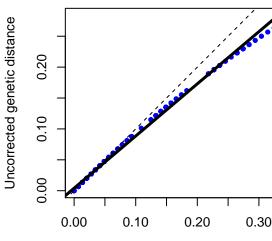
ycf1 Saturation (All Bases)

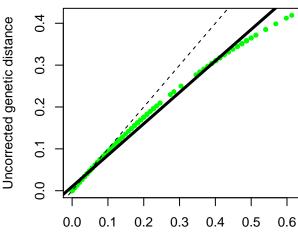
ycf1 Saturation (1st Pos)



ycf2 Saturation (All Bases)

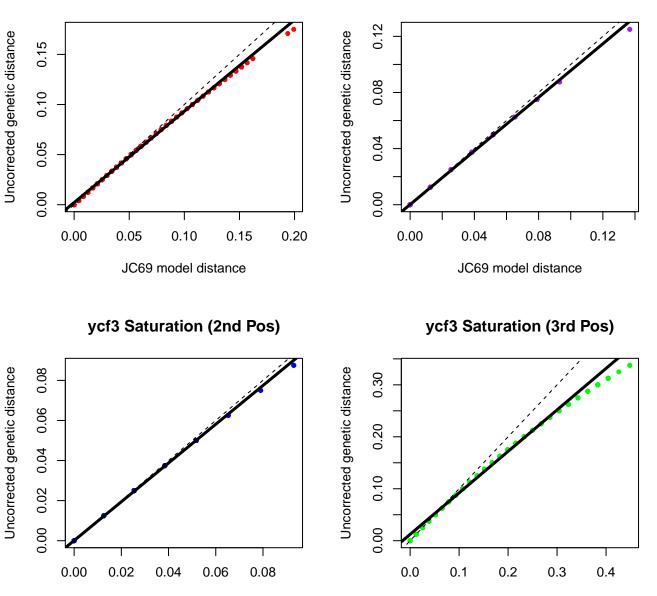






JC69 model distance







ycf4 Saturation (All Bases)

