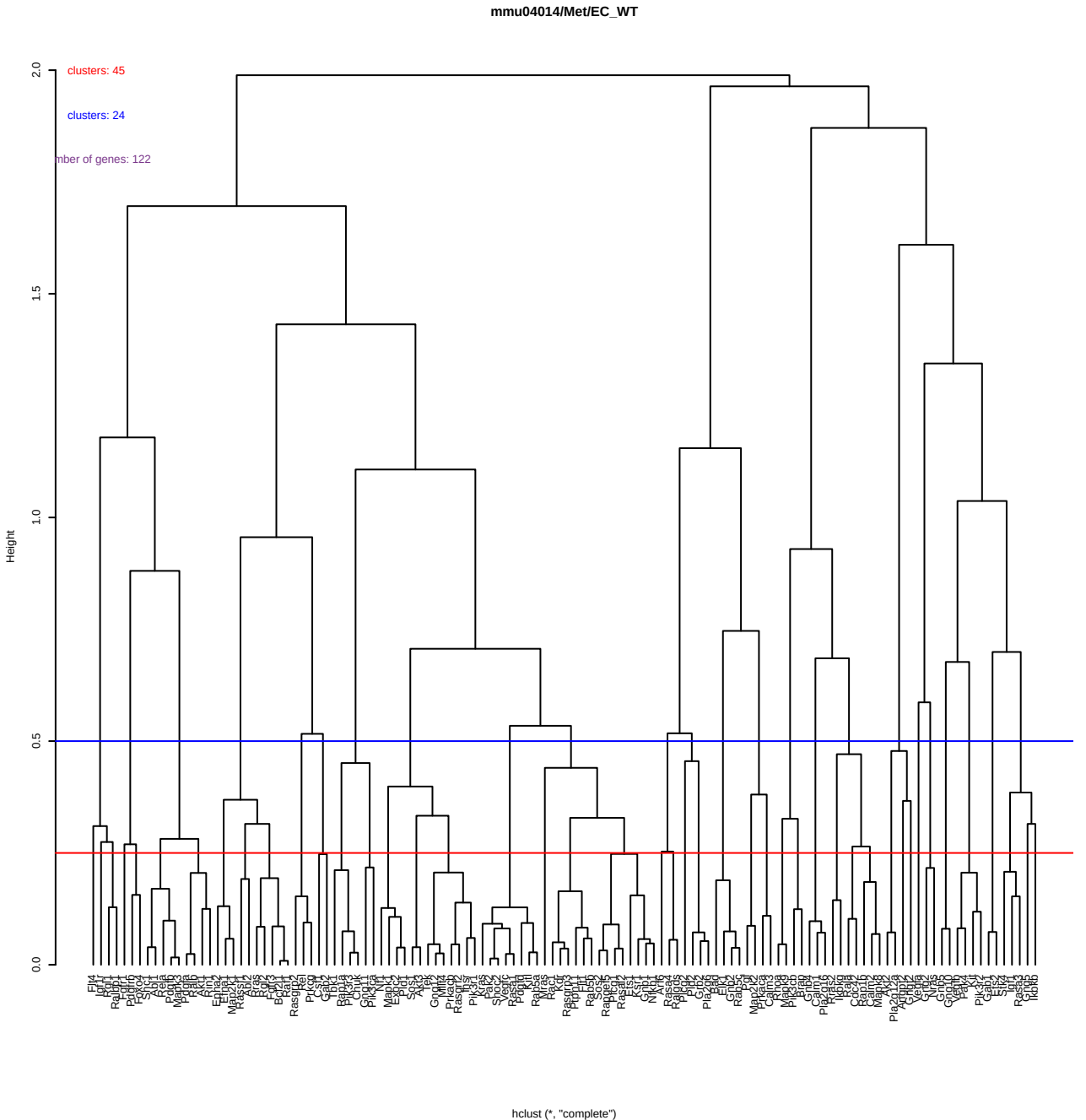
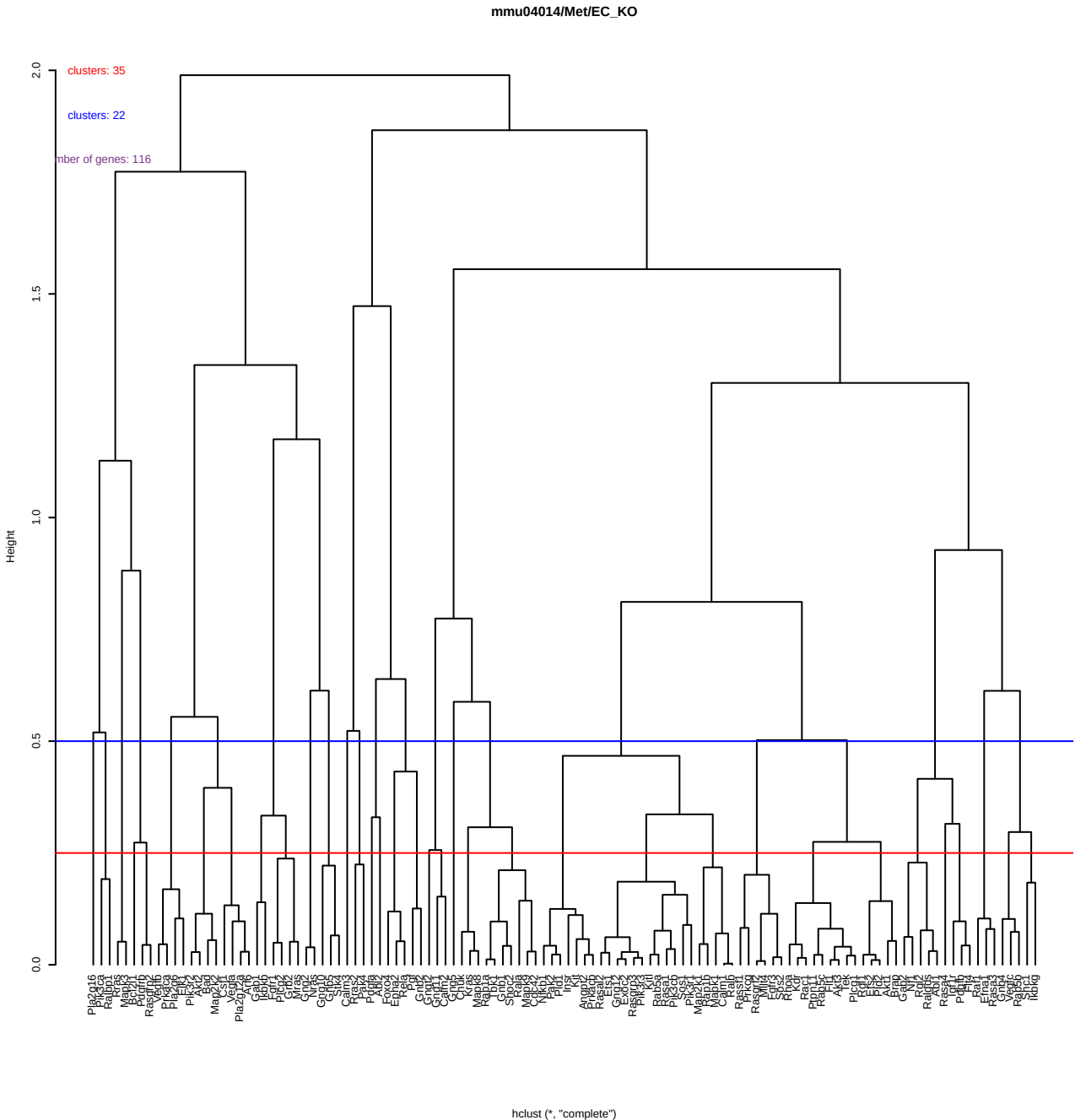
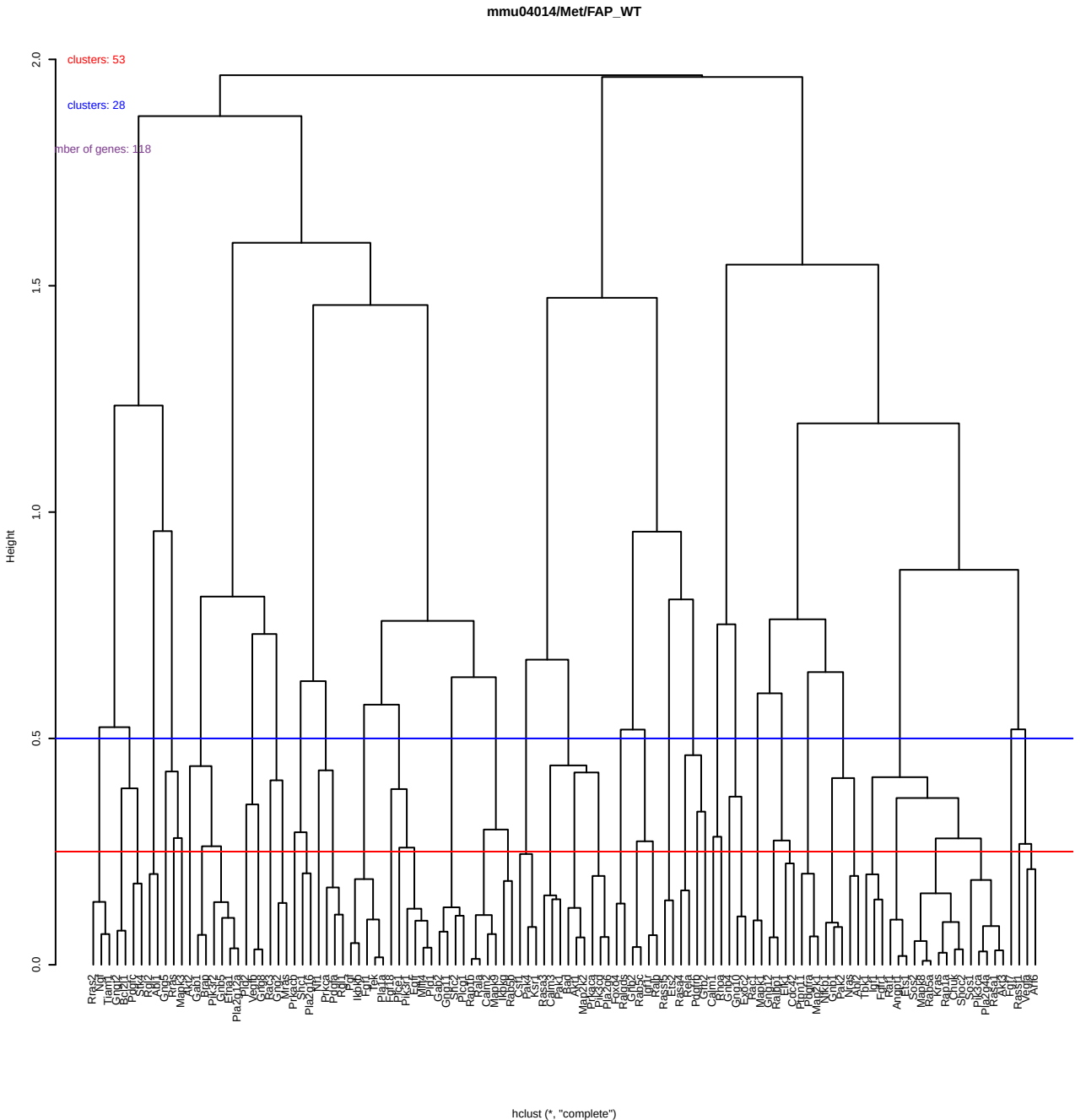
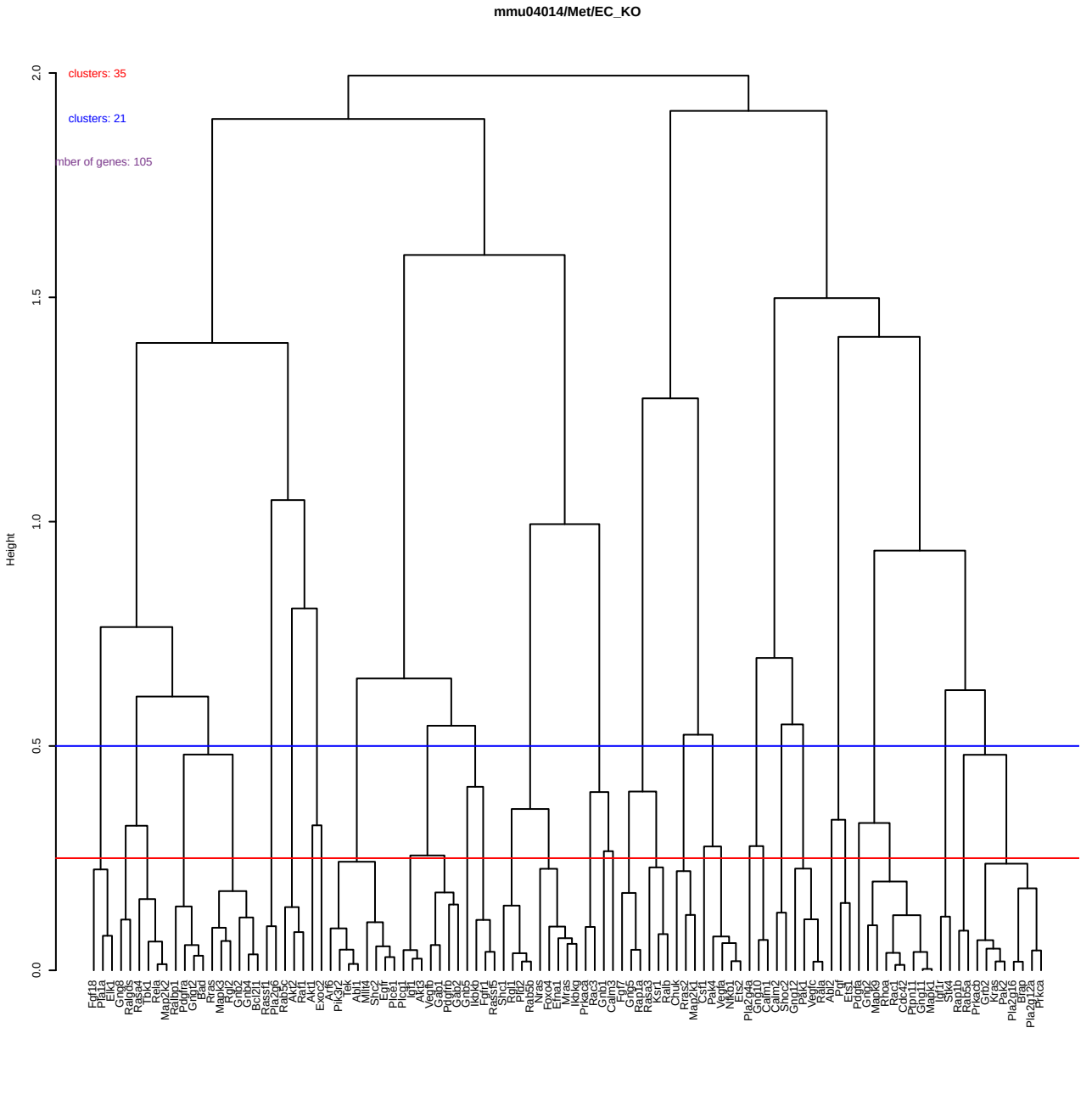
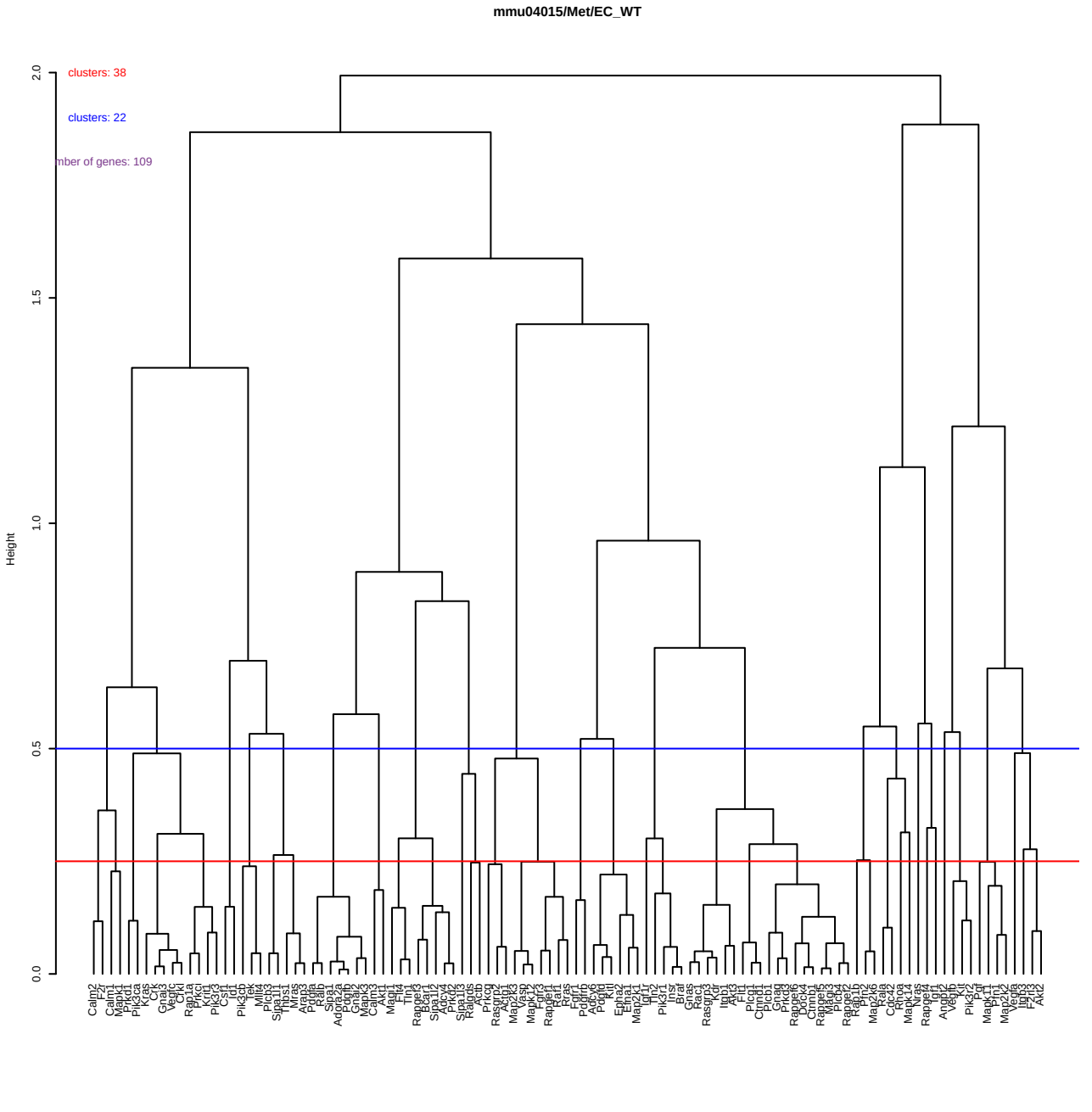

mmu04014/Met/EC_WT
clusters: 45
clusters: 24
mber of genes: 122
Height
0.0
0.5
1.0
1.5
2.0
hclust (*, "complete")


mmu04014/Met/EC_KO
clusters: 35
clusters: 22
mber of genes: 116
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

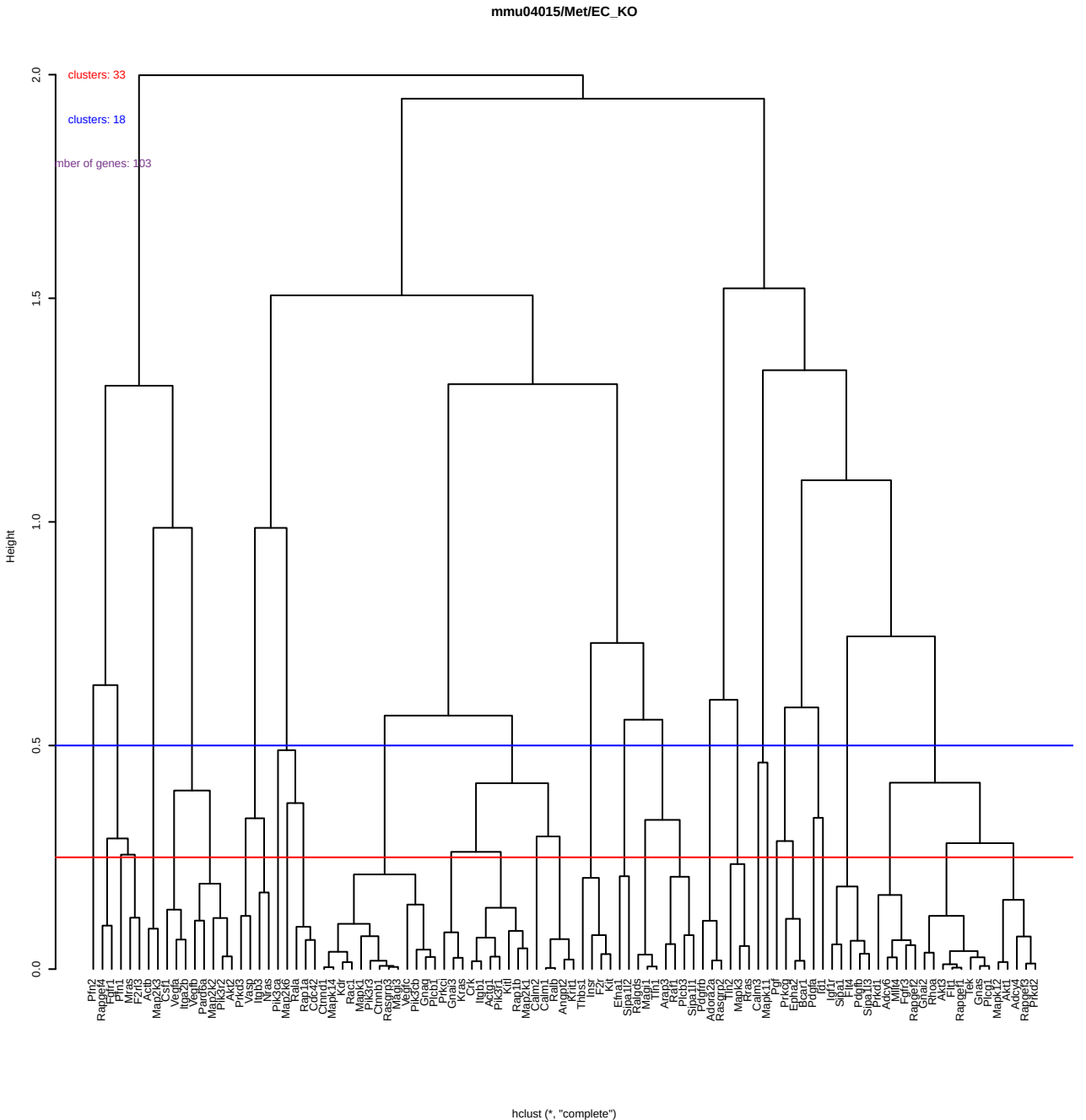

mmu04014/Met/FAP_WT
clusters: 53
clusters: 28
mber of genes: 118
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

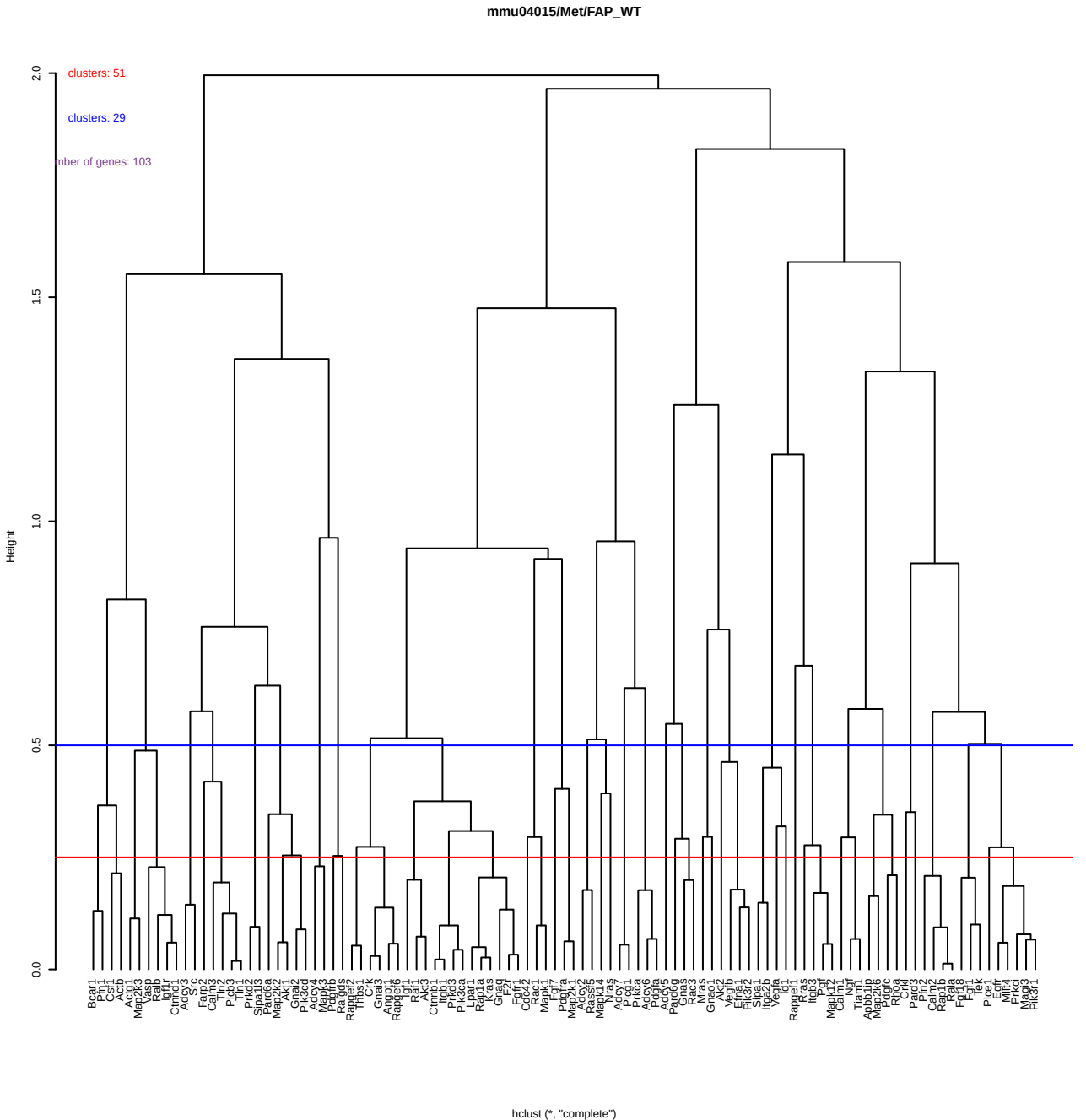


hclust (*, "complete")

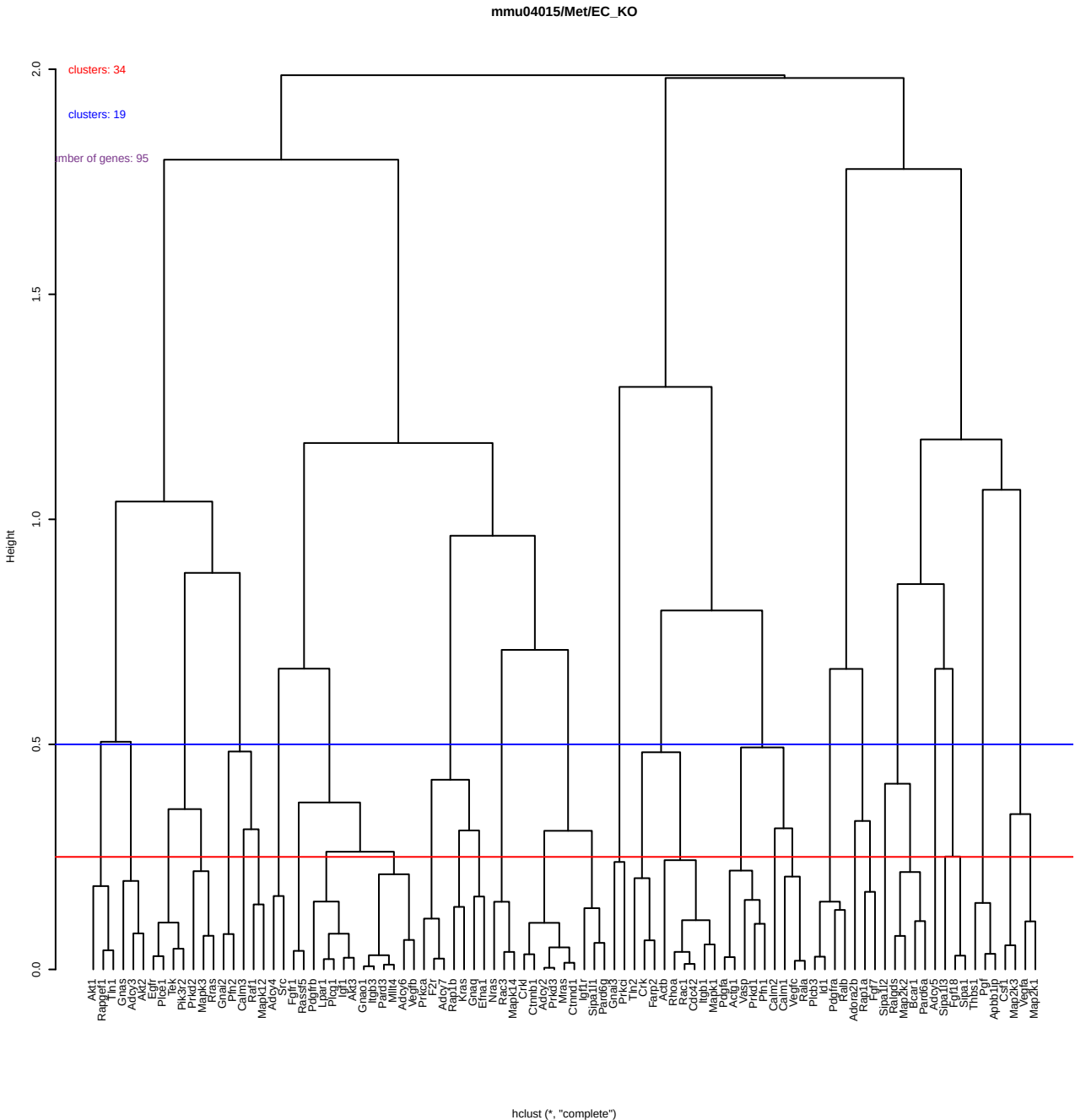


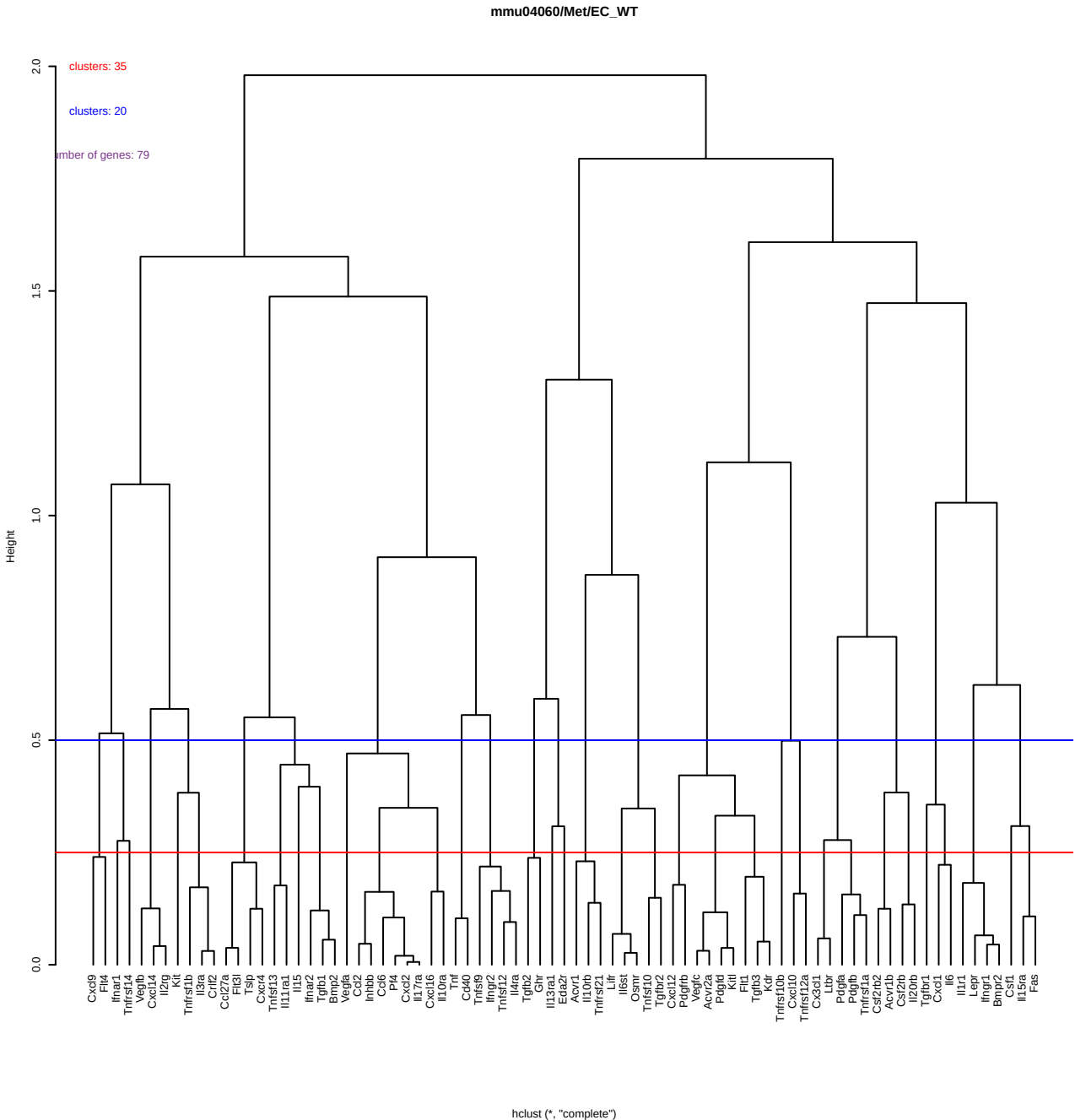
hclust (*, "complete")

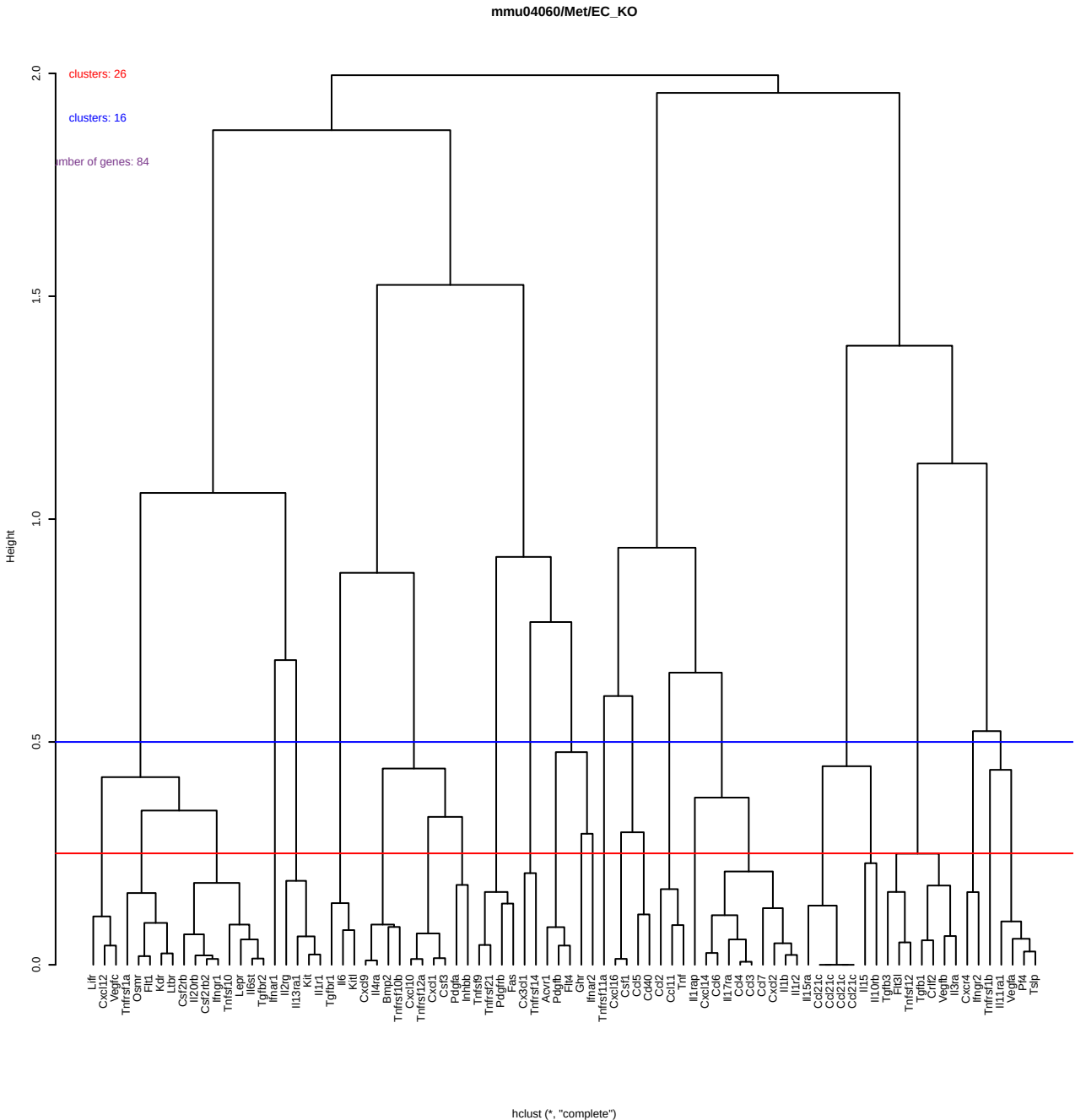

mmu04015/Met/EC_KO
clusters: 33
clusters: 18
mber of genes: 103
Height
2.0
1.5
1.0
0.5
0.0
Pfn2
Rapgef4
Flrt1
Pfn1
Mras
F2rl3
Actb
Map2k3
Csf1
Vegfa
Itga2b
Vegfb
Pardba
Map2k7
Pik3r2
Akt2
Prkd3
Vasp
Ivns1abp
Nras
Pik3ca
Map2k6
Rcla
Rapgef1a
Cdc42
Ctnnd1
Mapk14
Kdr
Rac1
Mapk1
Pik3r3
Ctnnb1
Rasgrp3
Magi3
Vegfc
Pik3cb
Cbl
Plcb1
Prkci
Gnai3
Kras
Crk
Itgb1
Acig1
Pik3r1
Xiap
Ran
Map2k1
Calm2
Calm1
Cam
Angpt2
Klf1
Thbs1
Hsr
Fzd7
Kit
Efna1
Sipa1l2
Rapgef5
Magi1
Tln1
Arap3
Rac3
Pik3r5
Sipa1l1
Pdgfrb
Adora2a
Rasgrp2
Gnb1
Mapk3
Rras
Calm3
Mapk11
Pgf
Prkcq
Epha2
Ecar1
Pdgfra
Igf1
Igf1r
Sipa1
Flt4
Pdgfb
Sipa1l3
Prkd1
Acvr2
Nfkb1
Fgfr3
Rapgef2
Gnai2
Rhoa
Akt3
Flt1
Rapgef1
Tek
Gnas
Plcg1
Mapk12
Akt1
Adcy4
Rapgef3
Prkd2
hclust (*, "complete")

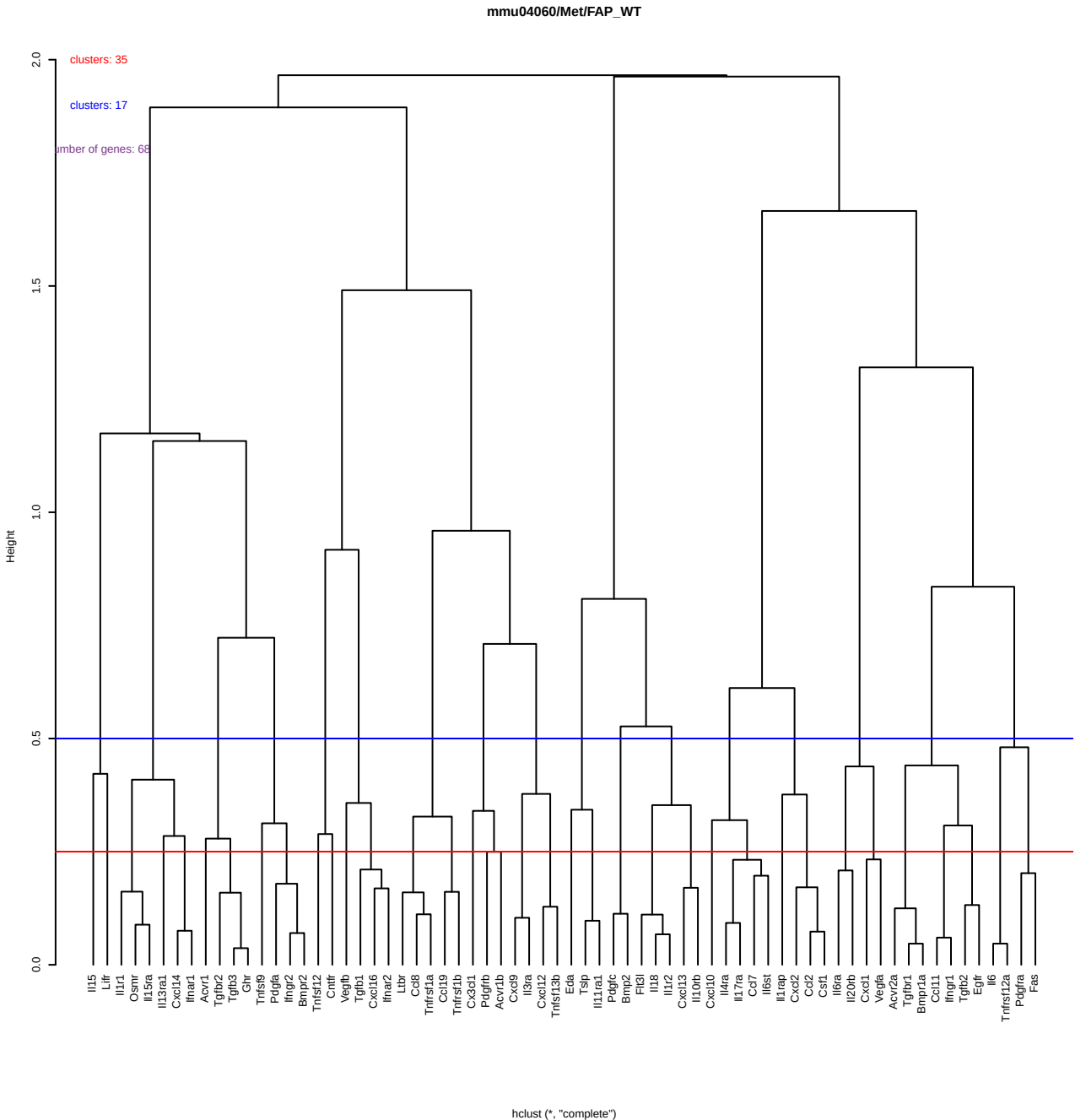


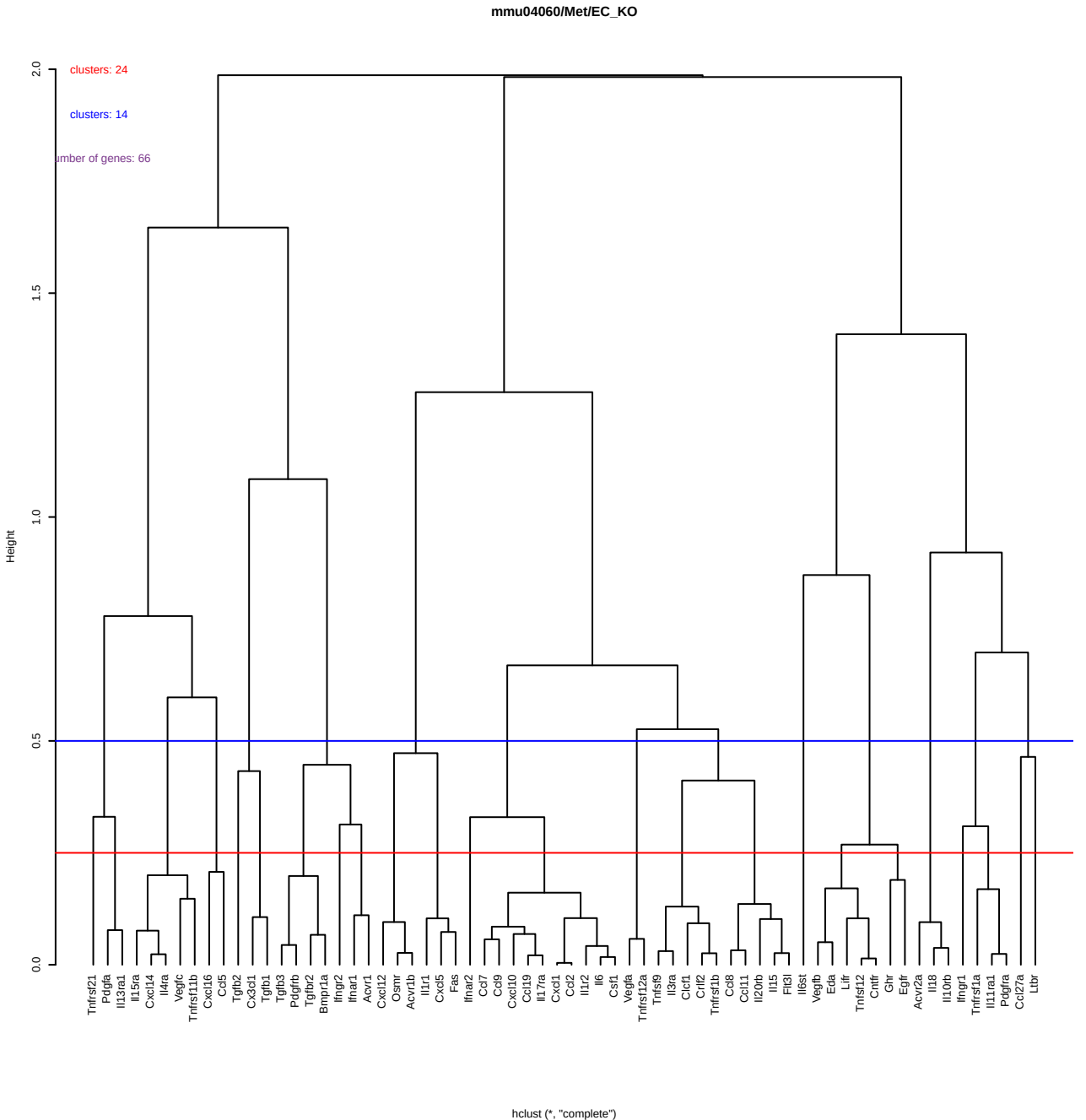
hclust (*, "complete")

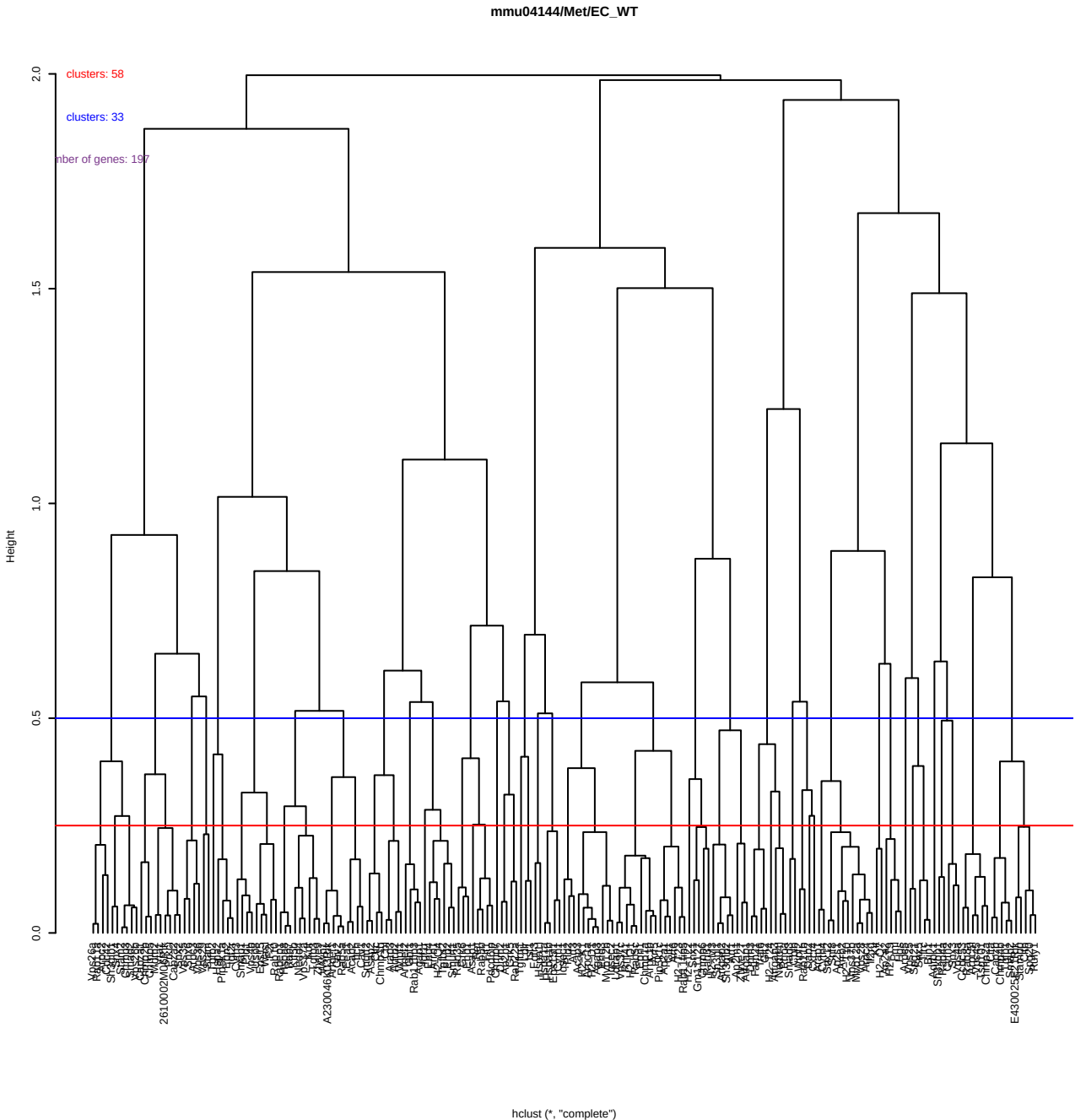

mmu04015/Met/EC_KO
clusters: 34
clusters: 19
mber of genes: 95
Height
0.0
0.5
1.0
1.5
2.0
Akt1
Rapgef1
Tln1
Gnas
Adcy3
Akt2
Esrrb
Plcb1
Tek
Pik3r2
Prkd2
Mapk3
Kras
Gnai2
Pfn2
Calm3
Rat1
Mapk12
Adcy4
Sos1
Fgf1
Rasgrf5
Pdgfrb
Lpar1
Plcg1
Ap1
Akt3
Gnao1
Itgb3
Pard3
Mll4
Adcy6
Vegfb
Pik3ca
F2r
Adcy7
Rap1b
Kras
Gnaq
Ehmt1
Nras
Rac3
Mapk14
Crk
Ctnnb1
Adcy2
Pik3cb
Mras
Ctnnd1
Igf1r
Sipa1l1
Pardbp
Gnai3
Prkcd
Tln2
Crk
Farp2
Actb
Rhoa
Rac1
Cdc42
Itgb1
Mapk1
Pdgfa
Acrg1
Yes1
Pik3r1
Pfn1
Calm2
Calm1
Vegfc
Ralia
Prkcd3
Id1
Pdgfra
Ralb
Adora2b
Rap1a
Egfr
Sipa1l2
Ralgds
Map2k2
Bcar1
Pardba
Adcy5
Sos1
Fgf18
Slpa1
Thbs1
Pgf
Apbb1p
Crk
Mapk3
Vegfa
Map2k1
hclust (*, "complete")

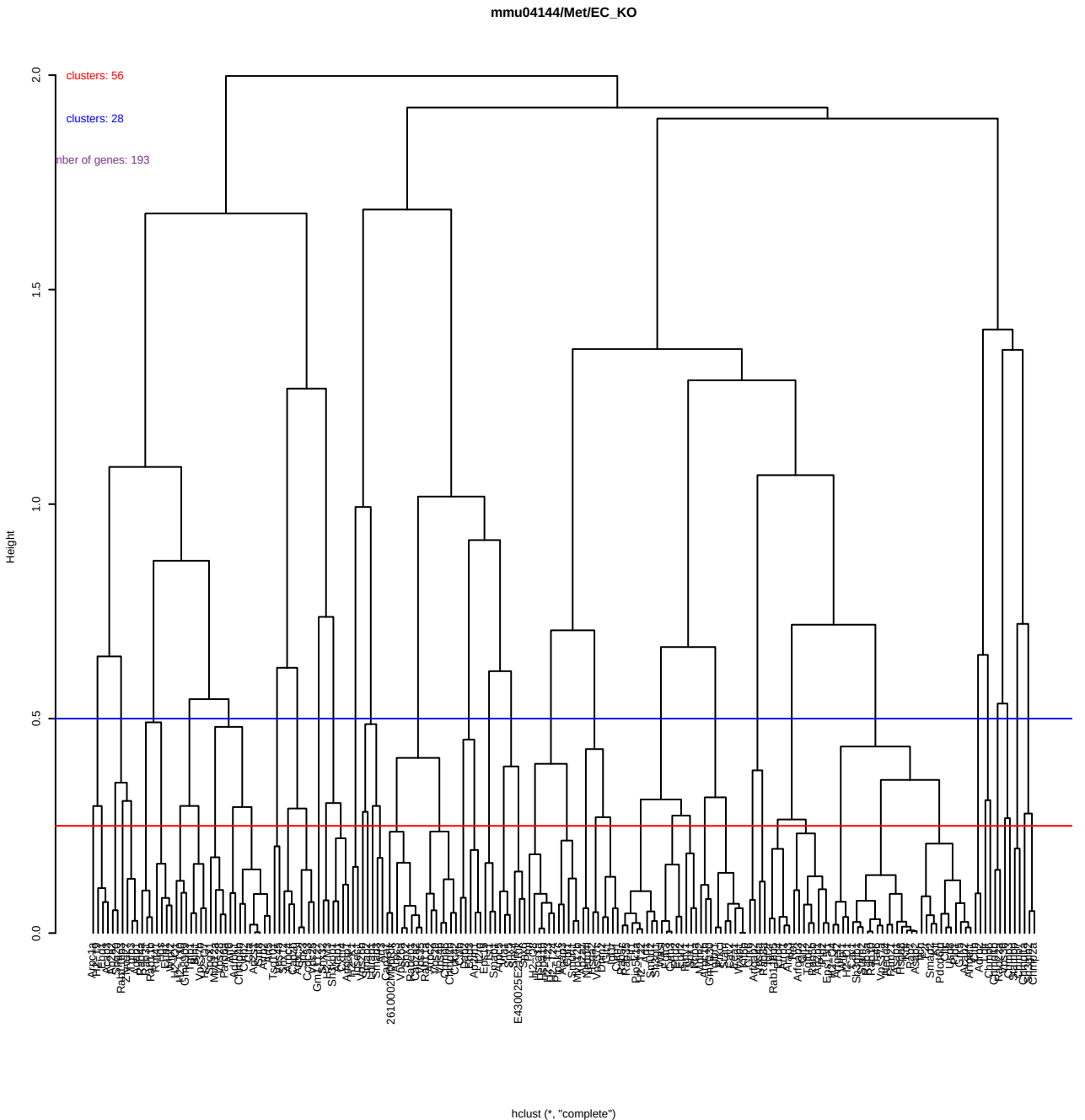

mmu04060/Met/EC_WT
clusters: 35
clusters: 20
mber of genes: 79
Height
2.0
1.5
1.0
0.5
0.0
Cxcl9
Flt4
Ifnar1
Tnfrsf14
Vegfb
Cxcl14
Il2rg
Kit
Tnfrsf1b
Il3ra
Crlf2
Ccl27a
Flt3l
Tslp
Cxcr4
Tnfsf13
Il11ra1
Il15
Ifnar2
Tgfb1
Bmp2
Vegfa
Ccl2
Inhbb
Ccl6
Pf4
Cxcl2
Il17ra
Cxcl16
Il10ra
Tnf
Cd40
Tnfsf9
Ifngr2
Tnfsf12
Il4ra
Tgfb2
Ghr
Il13ra1
Eda2r
Acvr1
Il10rb
Tnfrsf21
Lifr
Il6st
Osmr
Tnfsf10
Tgfbr2
Cxcl12
Pdgfrb
Vegfc
Acvr2a
Pdgfd
Kitl
Flt1
Tgfb3
Kdr
Tnfrsf10b
Cxcl10
Tnfrsf12a
Cx3cl1
Ltbr
Pdgfa
Pdgfb
Tnfrsf1a
Csf2rb2
Acvr1b
Csf2rb
Il20rb
Tgfbr1
Cxcl1
Il6
Il1r1
Lepr
Ifngr1
Bmpr2
Csf1
Il15ra
Fas
hclust (*, "complete")

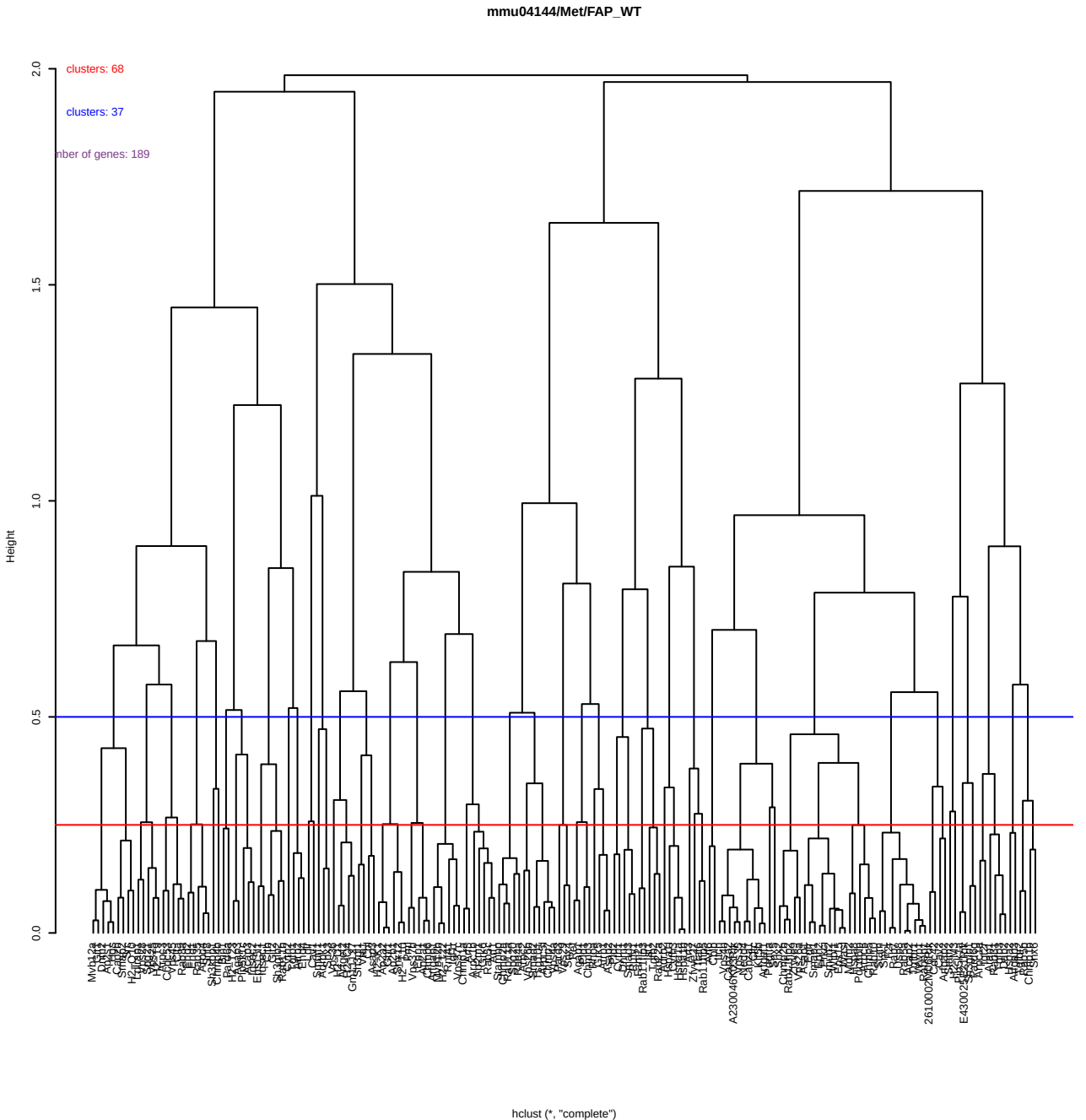

mmu04060/Met/EC_KO
clusters: 26
clusters: 16
mber of genes: 84
Height
0.0
0.5
1.0
1.5
2.0
Lifr
Cxcl12
Vegfc
Tnfrsf1a
Osmr
Flt1
Kdr
Ltbr
Csf2rb
Il20rb
Csf2rb2
Ifngr1
Tnfsf10
Lepr
Il6st
Tgfbr2
Ifnar1
Il2rg
Il13ra1
Kit
Il1r1
Il1r11
Tgfbr1
Il6
Kitl
Cxcl9
Il4ra
Bmp2
Tnfrsf10b
Cxcl10
Tnfrsf12a
Cxcl1
Csf3
Pdgfa
Inhbb
Tnfsf9
Tnfrsf21
Pdgfrb
Fas
Cx3cl1
Tnfrsf14
Acvr1
Pdgfb
Flt4
Ghr
Ifnar2
Tnfrsf11a
Cxcl16
Csf1
Ccr5
Cd40
Cd2
Ccl11
Tnf
Il1rap
Cxcl14
Ccl6
Il17ra
Ccl4
Ccl3
Ccl7
Cxcl2
Il1b
Il1r2
Il15ra
Ccl21c
Ccl21c
Ccl21c
Ccl21c
Ccl21c
Il15
Il10rb
Tgfb3
Flt3l
Tnfrsf12
Tgfb1
Ccrl2
Vegfb
Il3ra
Cxcr4
Ifngr2
Tnfrsf1b
Il11ra1
Vegfa
Pf4
Tslp
hclust (*, "complete")

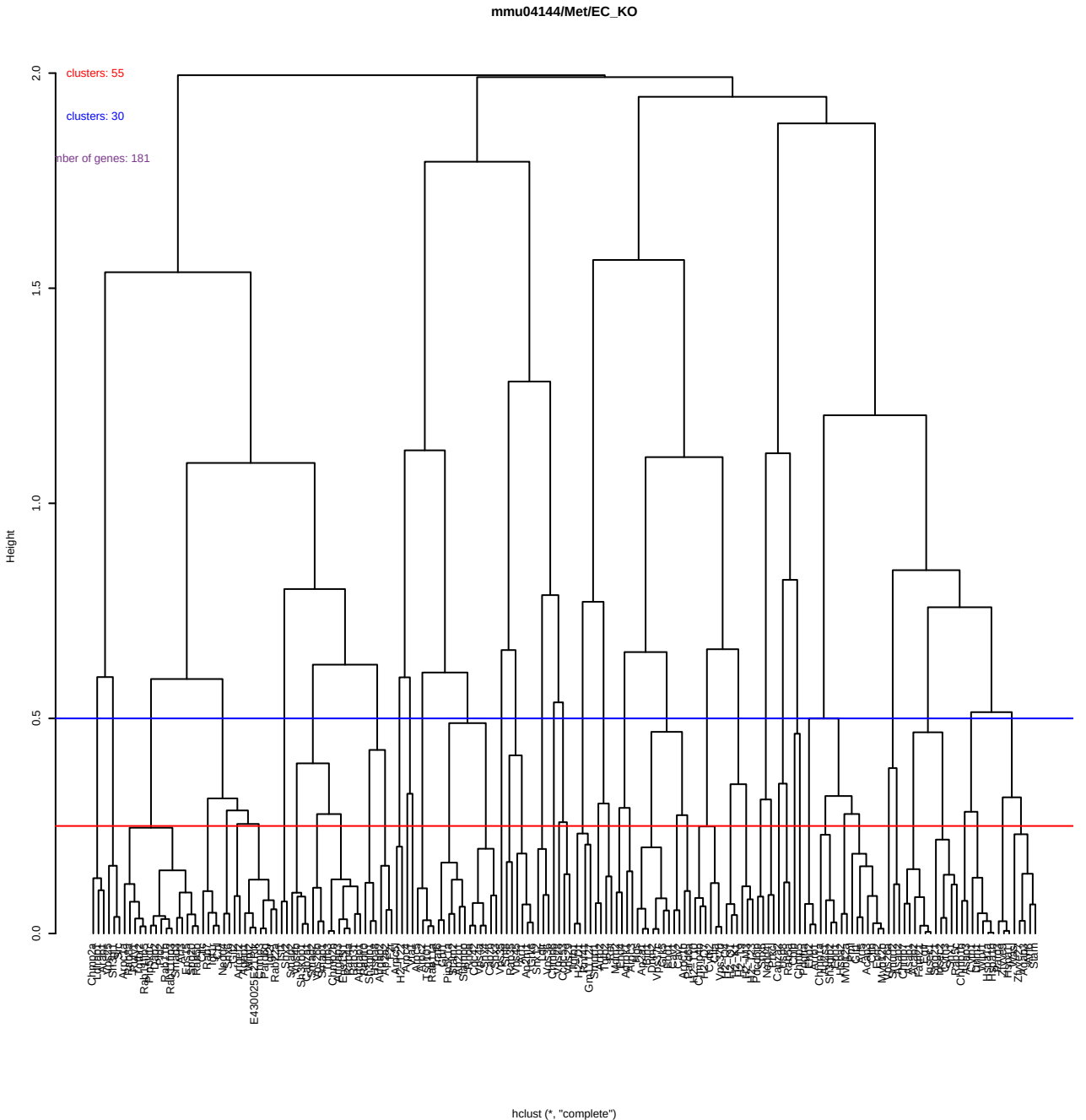

mmu04060/Met/FAP_WT
clusters: 35
clusters: 17
umber of genes: 68
Height
2.0
1.5
1.0
0.5
0.0
Il15
Lif
Il1r1
Osmr
Il15ra
Il13ra1
Cxcl14
Ifnar1
Acvr1
Tgfbr2
Tgfb3
Ghr
Tnfsf9
Pdgfa
Ifngr2
Bmpr2
Tnfsf12
Cntfr
Vegfb
Tgfb1
Cxcl16
Ifnar2
Ltbr
Ccl8
Tnfrsf1a
Ccl19
Tnfrsf1b
Cx3cl1
Pdgfrb
Acvr1b
Cxcl9
Il3ra
Cxcl12
Tnfsf13b
Eda
Tslp
Il11ra1
Pdgfc
Bmp2
Flt3l
Il18
Il1r2
Cxcl13
Il10rb
Cxcl10
Il4ra
Il17ra
Ccl7
Il6st
Il1rap
Cxcl2
Ccl2
Csf1
Il6ra
Il20rb
Cxcl1
Vegfa
Acvr2a
Tgfbr1
Bmpr1a
Ccl11
Ifngr1
Tgfb2
Egfr
Il6
Tnfrsf12a
Pdgfra
Fas
hclust (*, "complete")

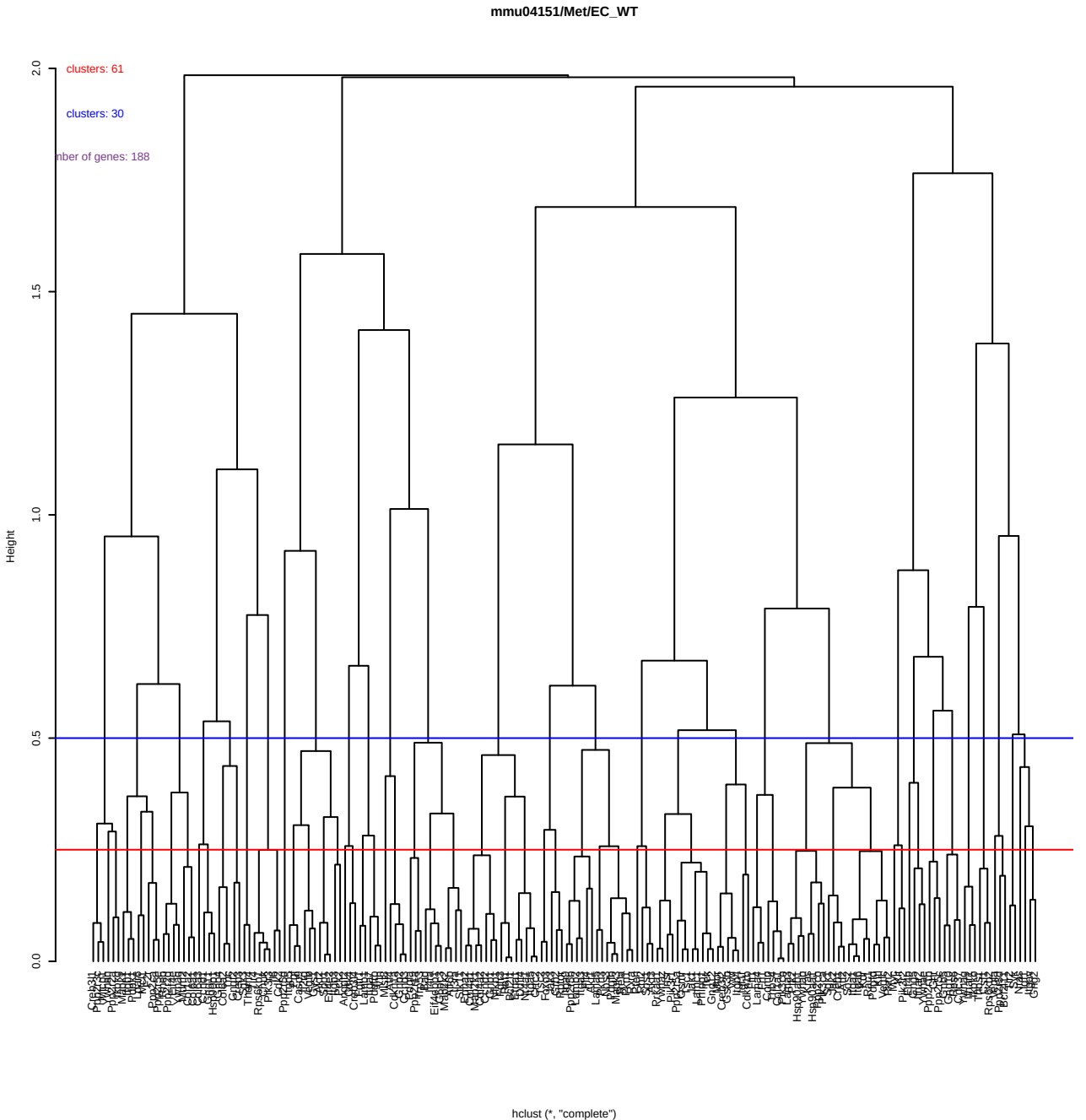

mmu04060/Met/EC_KO
clusters: 24
clusters: 14
umber of genes: 66
Height
2.0
1.5
1.0
0.5
0.0
Tnfrsf21
Pdgfa
Il13ra1
Il15ra
Cxcl14
Il4ra
Vegfc
Tnfrsf11b
Cxcl16
Ccl5
Tgfb2
Cx3cl1
Tgfb1
Tgfb3
Pdgfrb
Tgfbr2
Bmpr1a
Ifngr2
Ifnar1
Acvr1
Cxcl12
Osmr
Acvr1b
Il1r1
Cxcl5
Fas
Ifnar2
Ccl7
Ccl9
Cxcl10
Ccl19
Il17ra
Cxcl1
Ccl2
Il1r2
Il6
Csf1
Vegfa
Tnfrsf12a
Tnfsf9
Il3ra
Clcf1
Crlf2
Tnfrsf1b
Ccl8
Ccl11
Il20rb
Il15
Flt3l
Il6st
Vegfb
Eda
Lifr
Tnfsf12
Cntfr
Ghr
Egfr
Acvr2a
Il18
Il10rb
Ifngr1
Tnfrsf1a
Il11ra1
Pdgfra
Ccl27a
Ltbr
hclust (*, "complete")

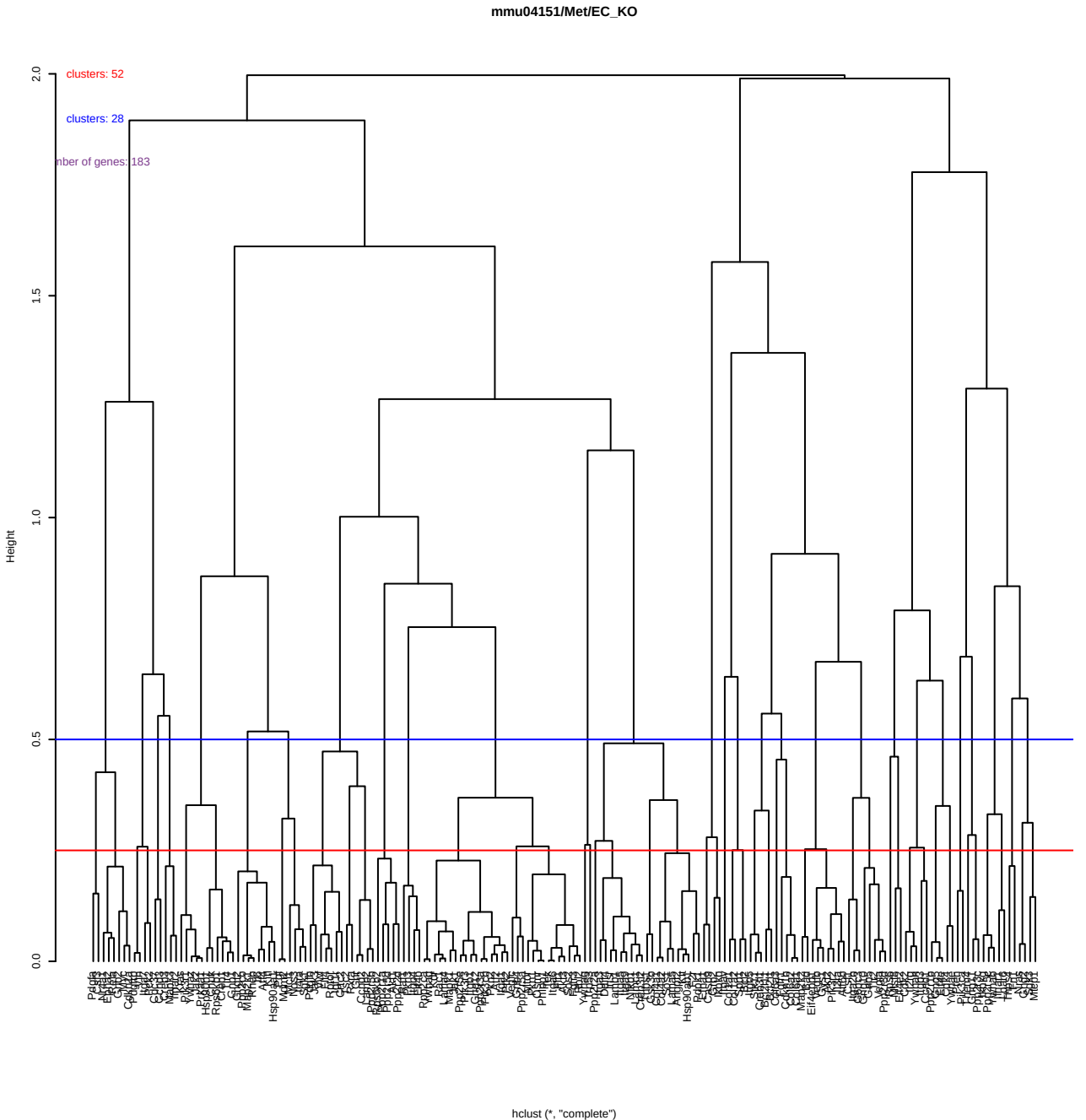

mmu04144/Met/EC_WT
clusters: 58
clusters: 33
mber of genes: 197
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

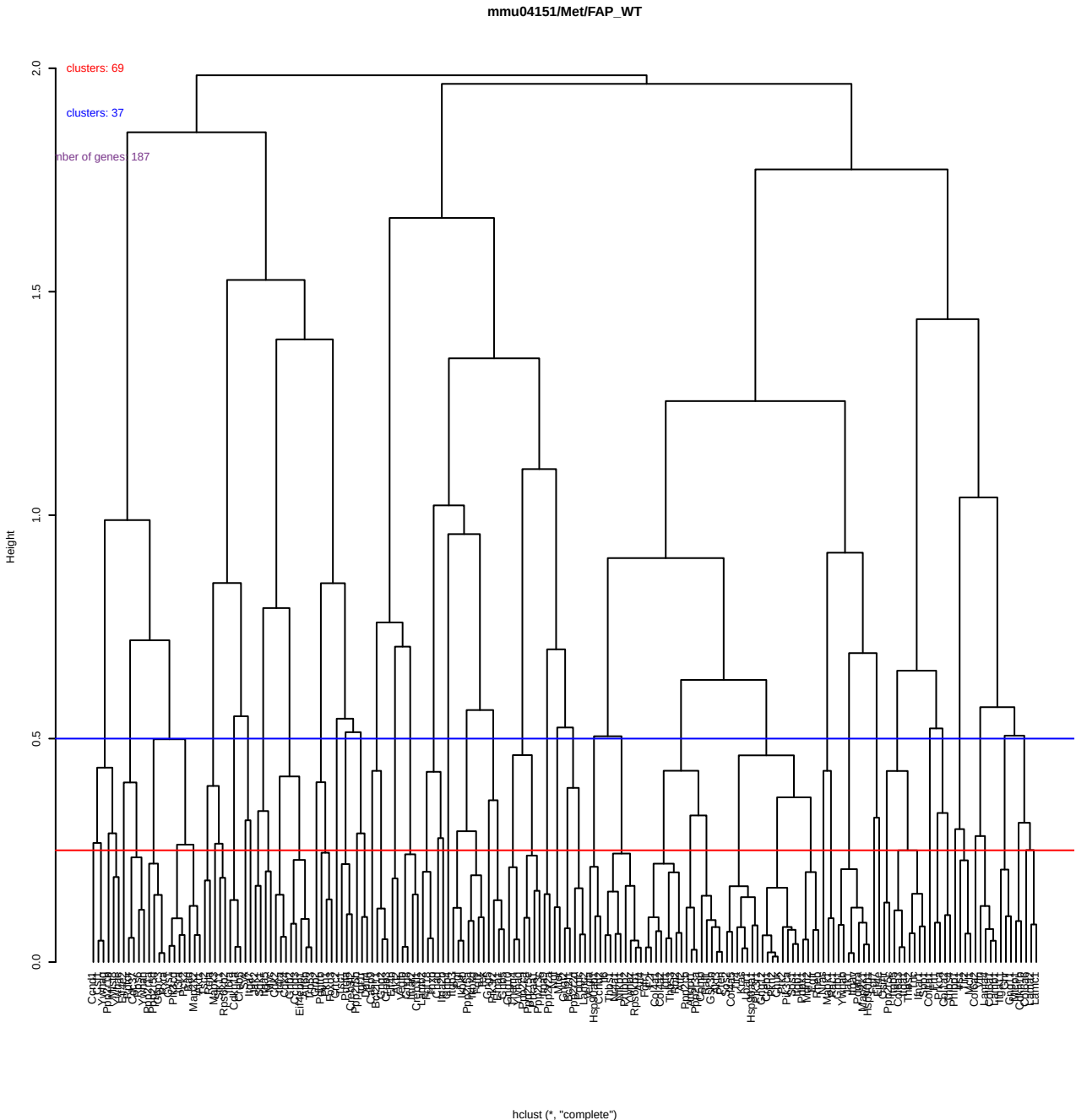

mmu04144/Met/EC_KO
clusters: 56
clusters: 28
mber of genes: 193
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

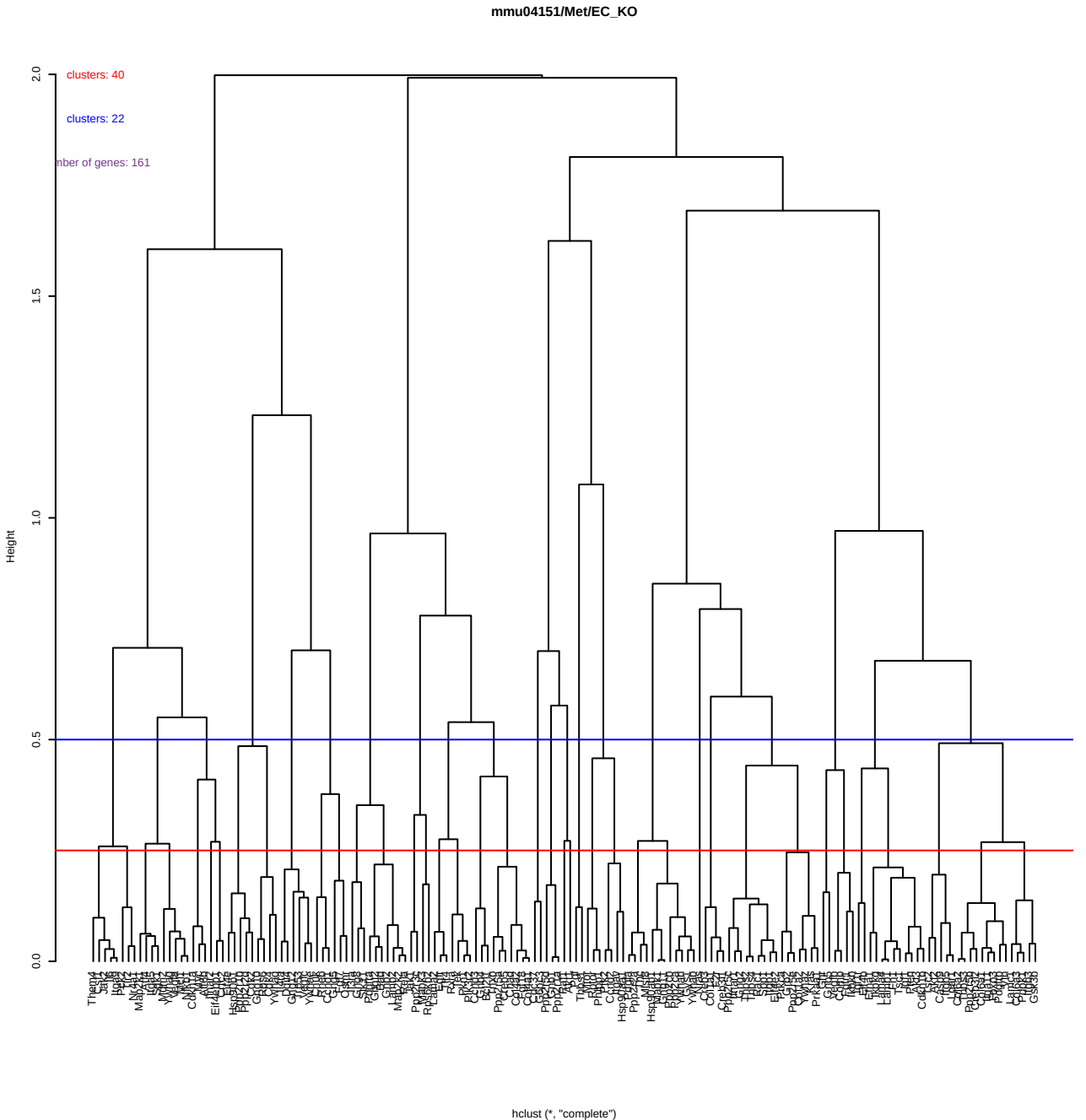

mmu04144/Met/FAP_WT
clusters: 68
clusters: 37
mber of genes: 189
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

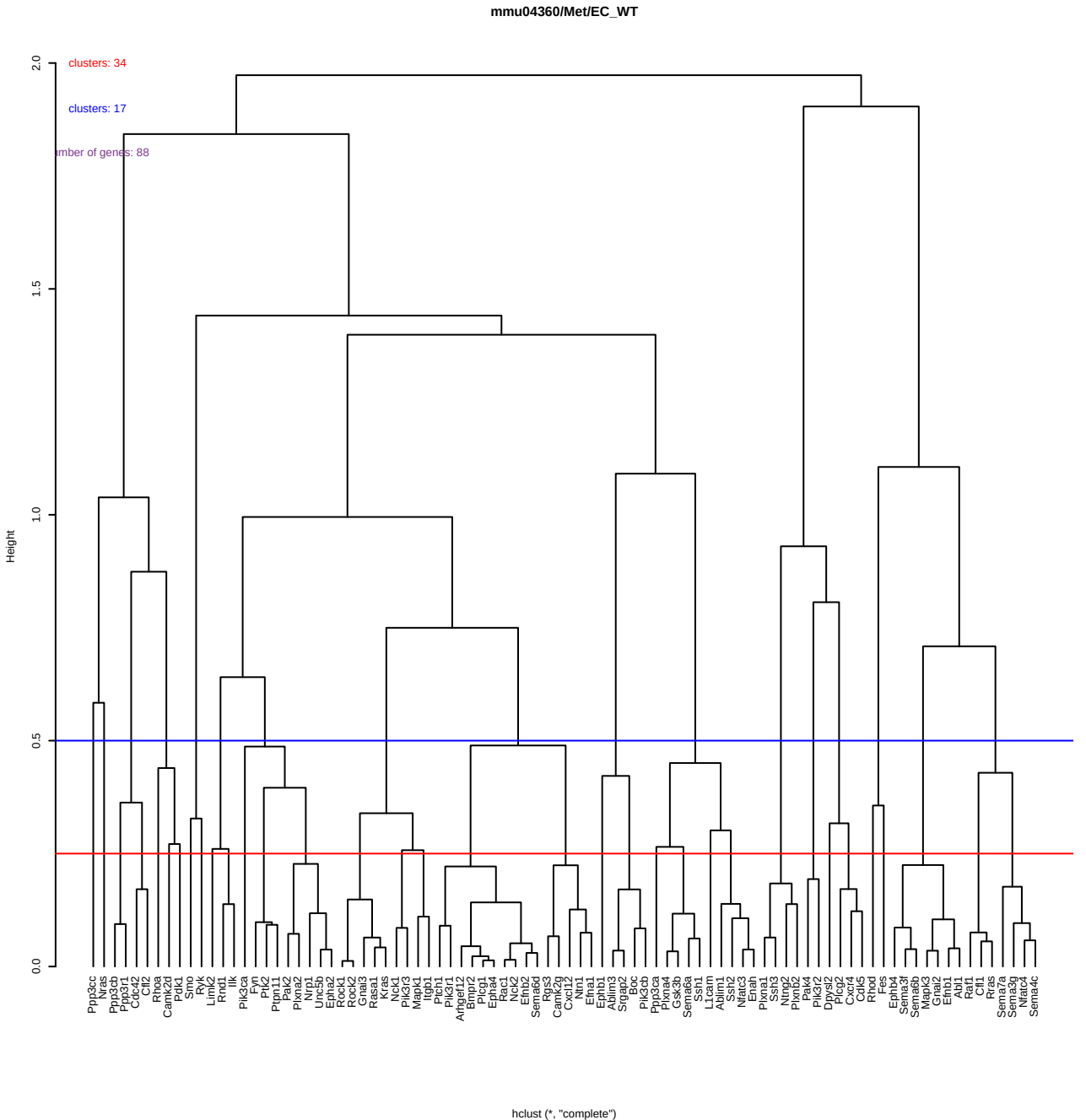

mmu04144/Met/EC_KO
clusters: 55
clusters: 30
mber of genes: 181
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

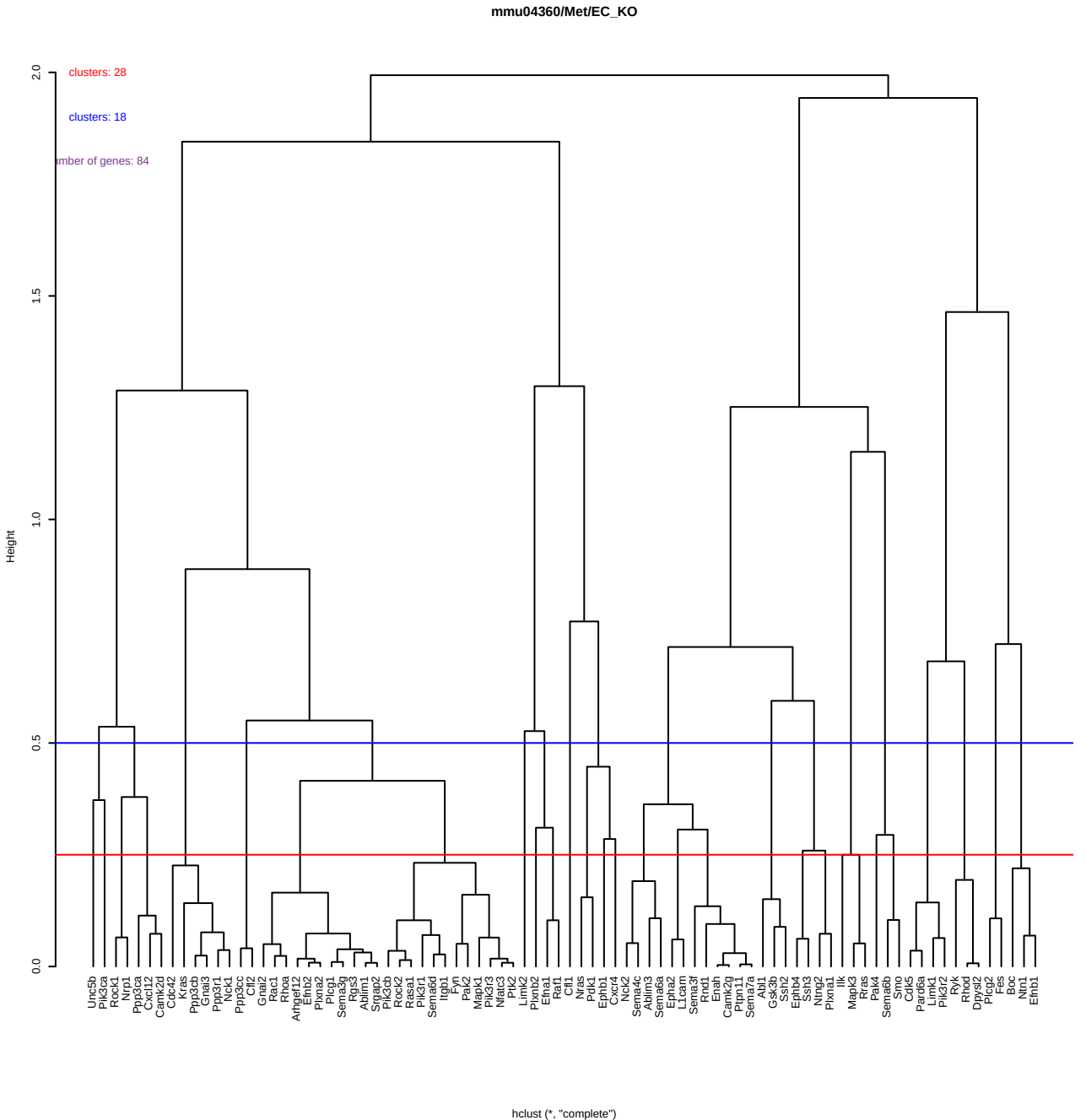

mmu04151/Met/EC_WT
clusters: 61
clusters: 30
mber of genes: 188
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

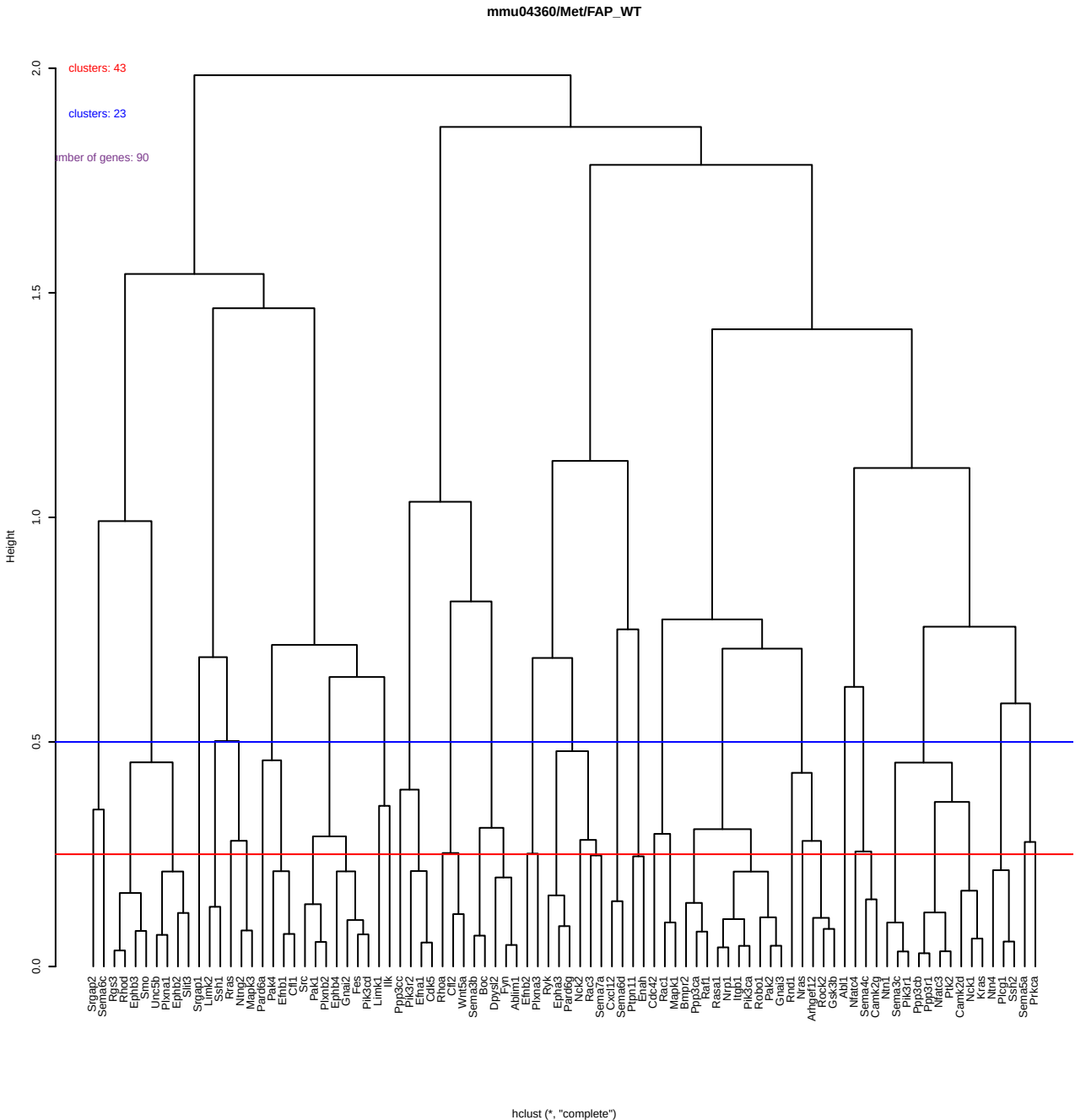

mmu04151/Met/EC_KO
clusters: 52
clusters: 28
nber of genes: 183
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")


mmu04151/Met/FAP_WT
clusters: 69
clusters: 37
mber of genes: 187
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

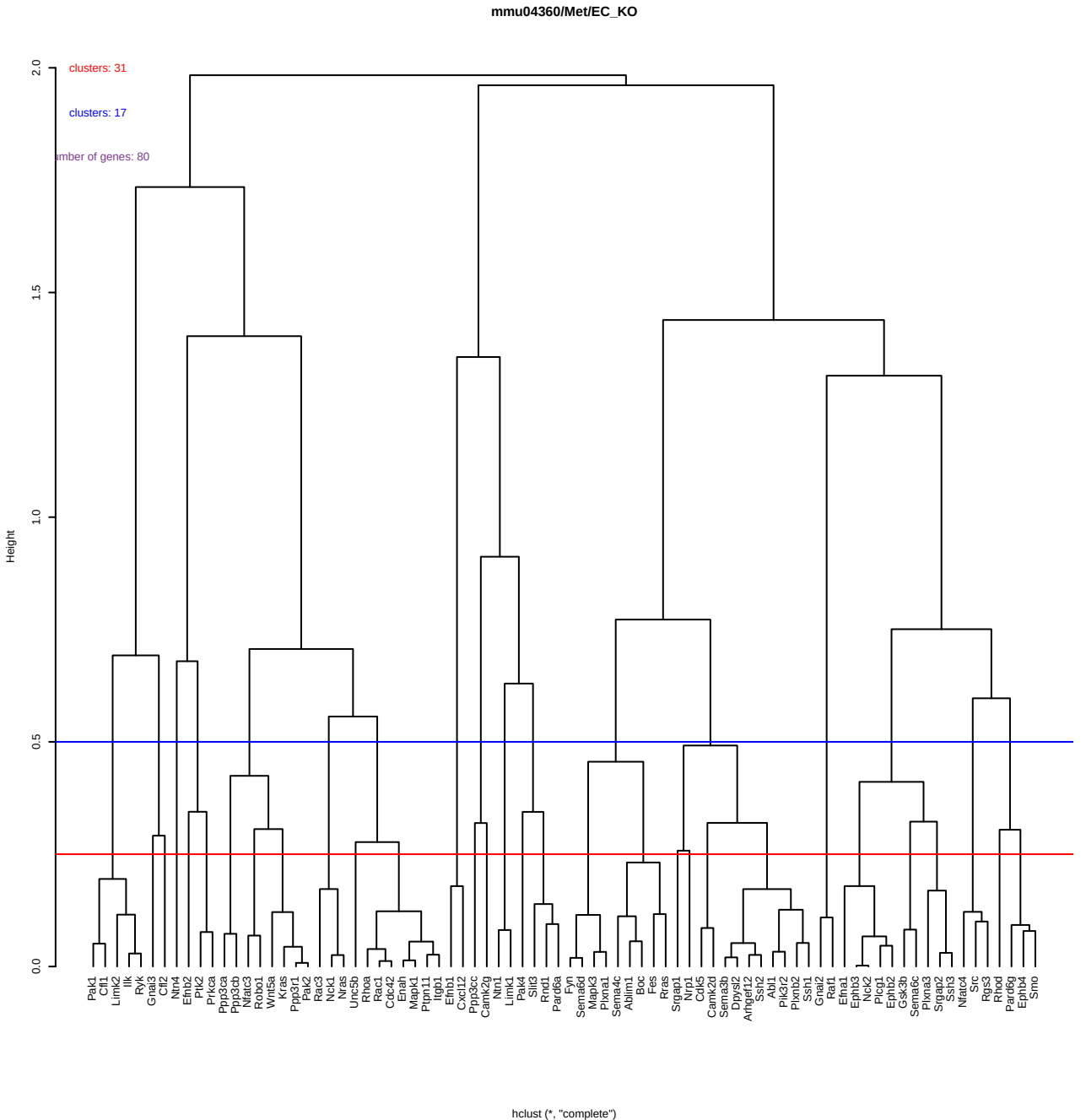

mmu04151/Met/EC_KO
clusters: 40
clusters: 22
nber of genes: 161
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

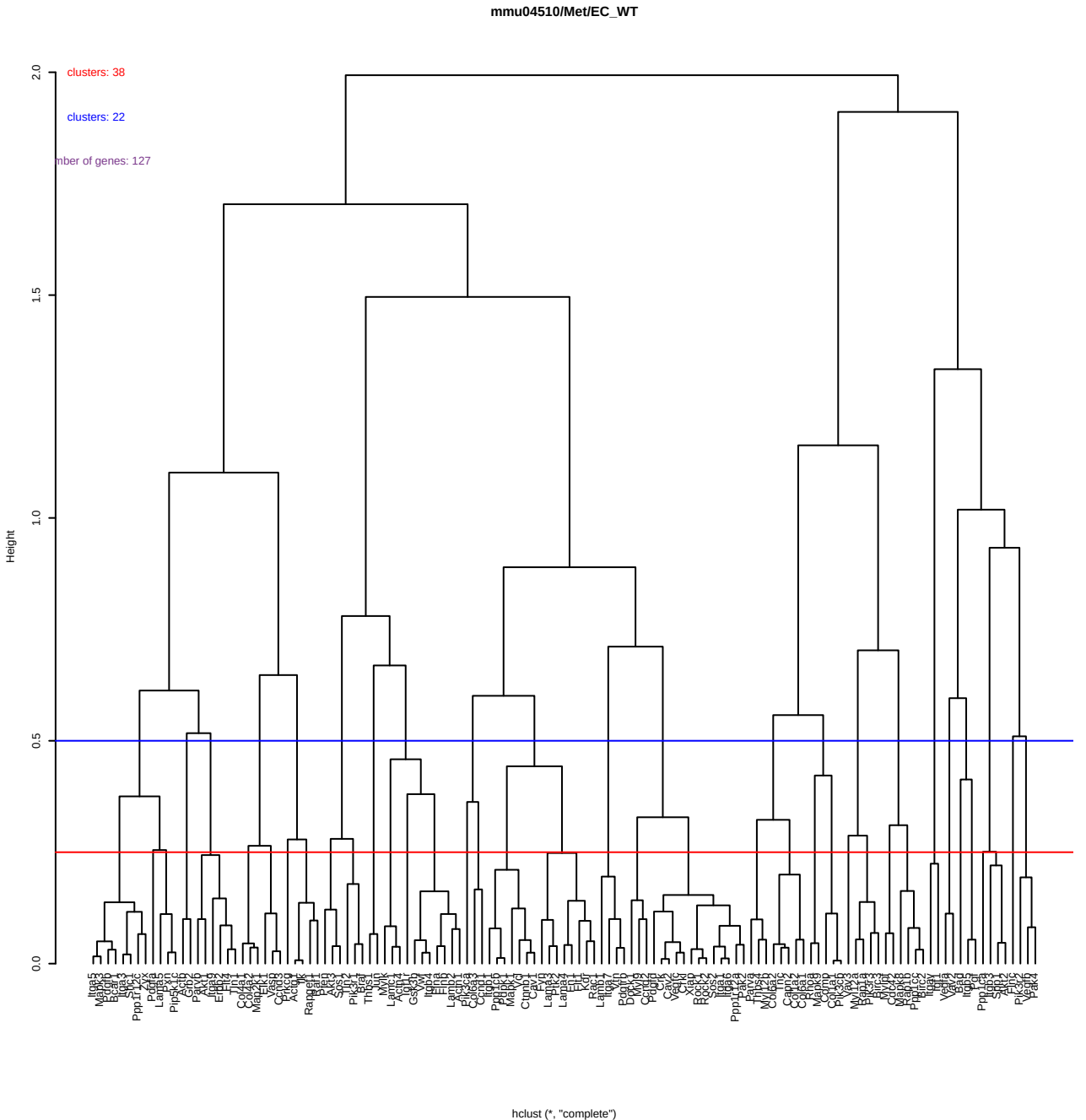

mmu04360/Met/EC_WT
clusters: 34
clusters: 17
mber of genes: 88
Height
2.0
1.5
1.0
0.5
0.0
Ppp3cc
Nras
Ppp3cb
Ppp3r1
Cdc42
Cfl2
Rhoa
Camk2d
Pdk1
Smo
Prkx
Limk2
Rnd1
Ilk
Pik3ca
Fyn
Pik2
Ptpn11
Pak2
Plxna2
Nrp1
Unc5b
Epha2
Rock1
Rock2
Gnai3
Rasa1
Kras
Nck1
Pik3r3
Mapk1
Itgb1
Plch1
Pik3r1
Arhgef12
Ptpn2
Plcb1
Epha4
Rac1
Nck2
Efnb2
Sema6d
Rgs3
Camk2g
Cxcl12
Nrn1
Efna1
Epha7
Ablim3
Srgap2
Boc
Pik3cb
Ppp3ca
Plxna4
Gsk3b
Sema6a
Ssh1
L1cam
Abhd17a
Slit2
Nfatc3
Enah
Plxna1
Ssh3
Ntng2
Plxnb2
Pak4
Pik3r2
Dpysl2
Plcg2
Cxcl4
Cxcl5
Rhod
Fes
Ephb4
Sema3f
Sema6b
Mapk3
Gnai2
Efnb1
Abl1
Rac1
Rac3
Rras
Sema7a
Sema3g
Nfatc4
Sema4c
hclust (*, "complete")

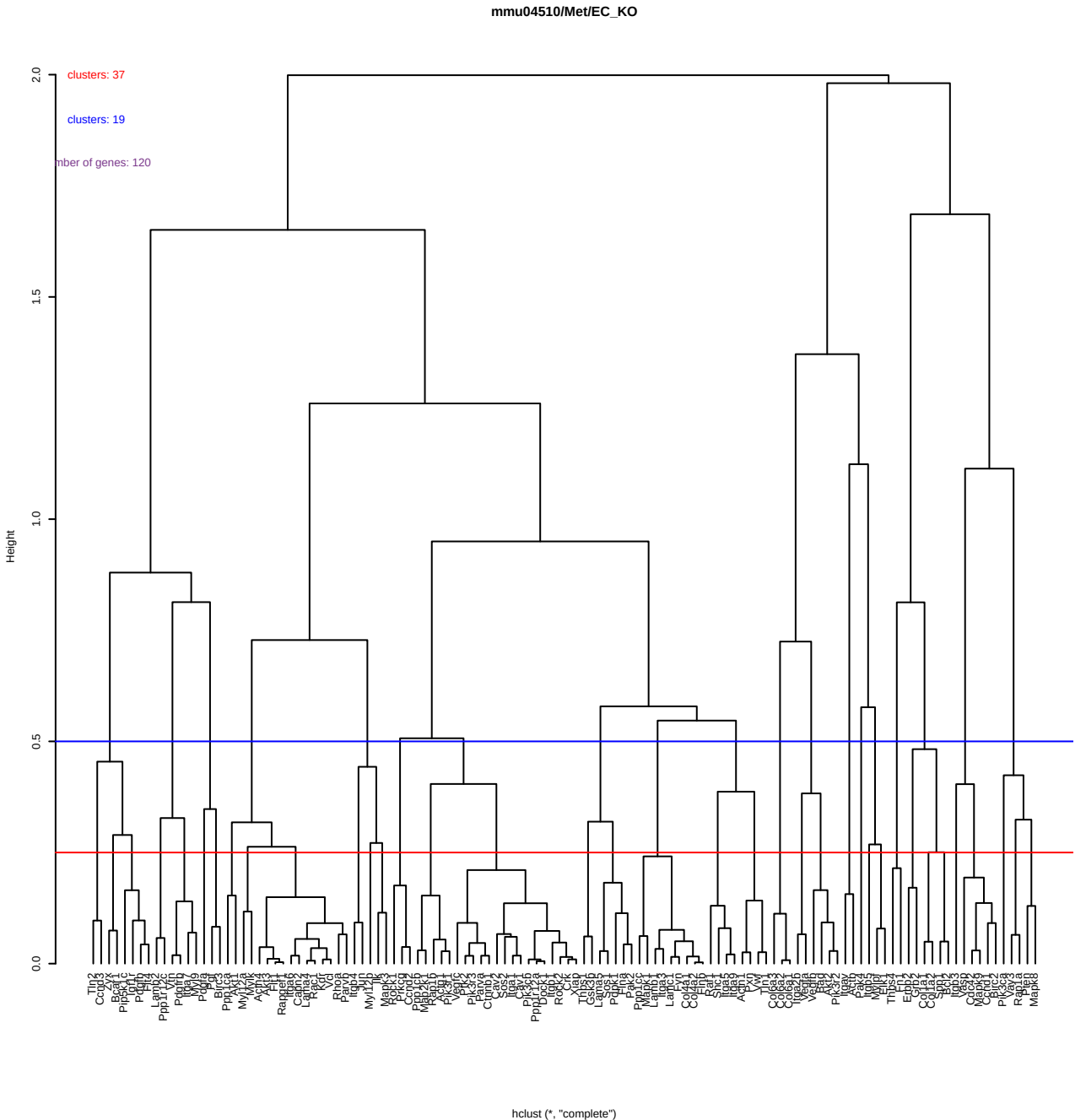

mmu04360/Met/EC_KO
clusters: 28
clusters: 18
mber of genes: 84
Height
2.0
1.5
1.0
0.5
0.0
Unc5b
Pik3ca
Rock1
Nrp1
Ppp3ca
Cxcl12
Camk2d
Cdc42
Kras
Ppp3cb
Gnai3
Ppp3r1
Nck1
Ppp3cc
Cfl2
Gnai2
Rac1
Rhoa
Arhgef12
Efnb2
Plxna2
Plcg1
Sema3g
Rps6
Abhd17a
Srgap2
Pik3cb
Rock2
Rasa1
Pik3r1
Sema6d
Itgb1
Fyn
Pak2
Mapk1
Pik3r3
Nfatc3
Pik2
Limk2
Plxnb2
Efna1
Rai11
Cfl1
Nras
Pdk1
Ephb1
Cxcl4
Nxn2
Sema4c
Abhd3
Sema6a
Epha2
L1cam
Sema3f
Rnd1
Enah
Camk2g
Ptpn11
Sema7a
Abi1
Gsk3b
Ssh2
Ephb4
Ssh3
Ntng2
Plxna1
Ilk
Mapk3
Rras
Pak4
Sema6b
Smo
Cdk5
Pard6a
Lmx1
Pik3r2
Ryk
Rhod
Dpysl2
Plcg2
Fes
Boc
Ntn1
Efnb1
hclust (*, "complete")



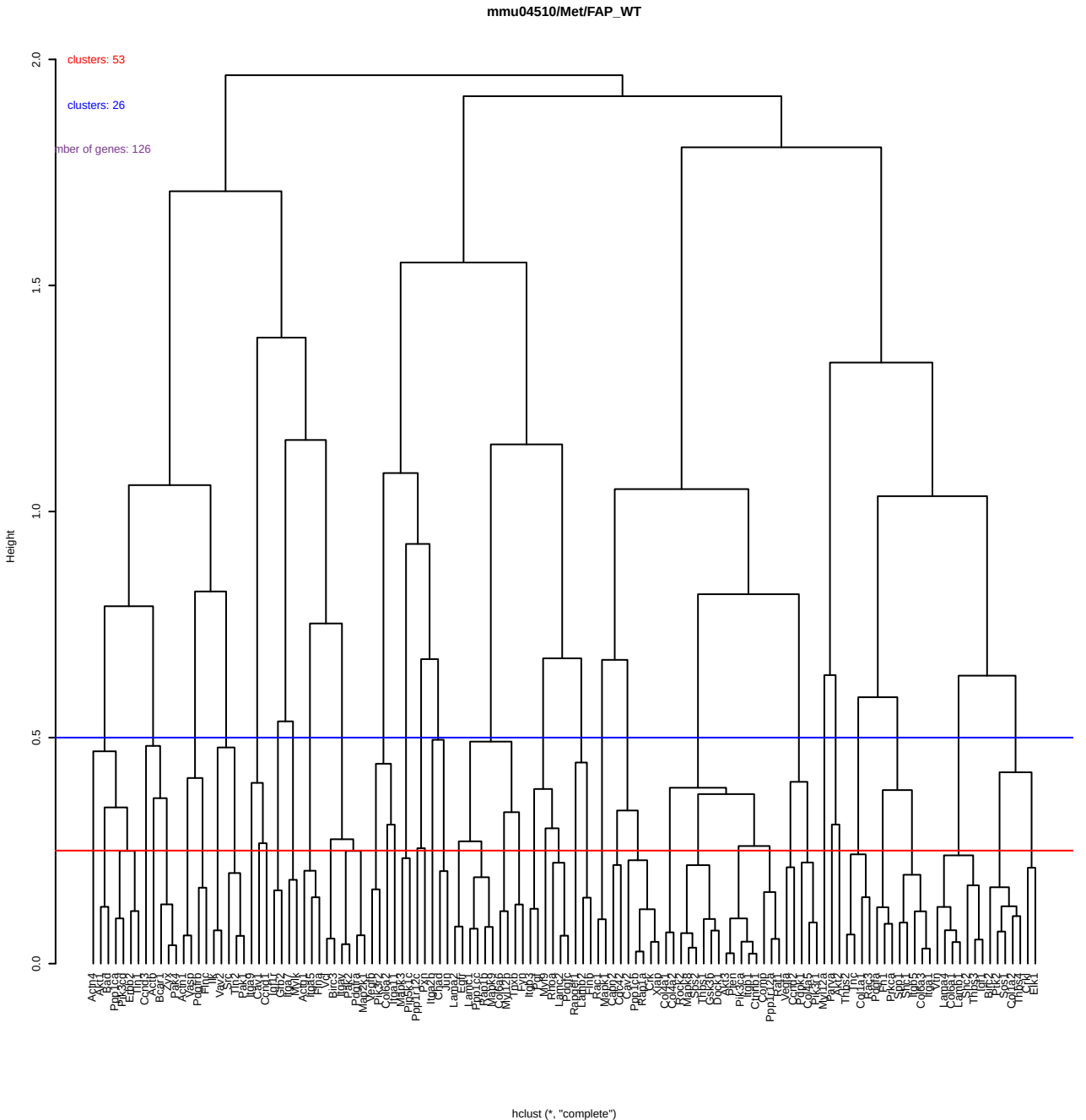
hclust (*, "complete")

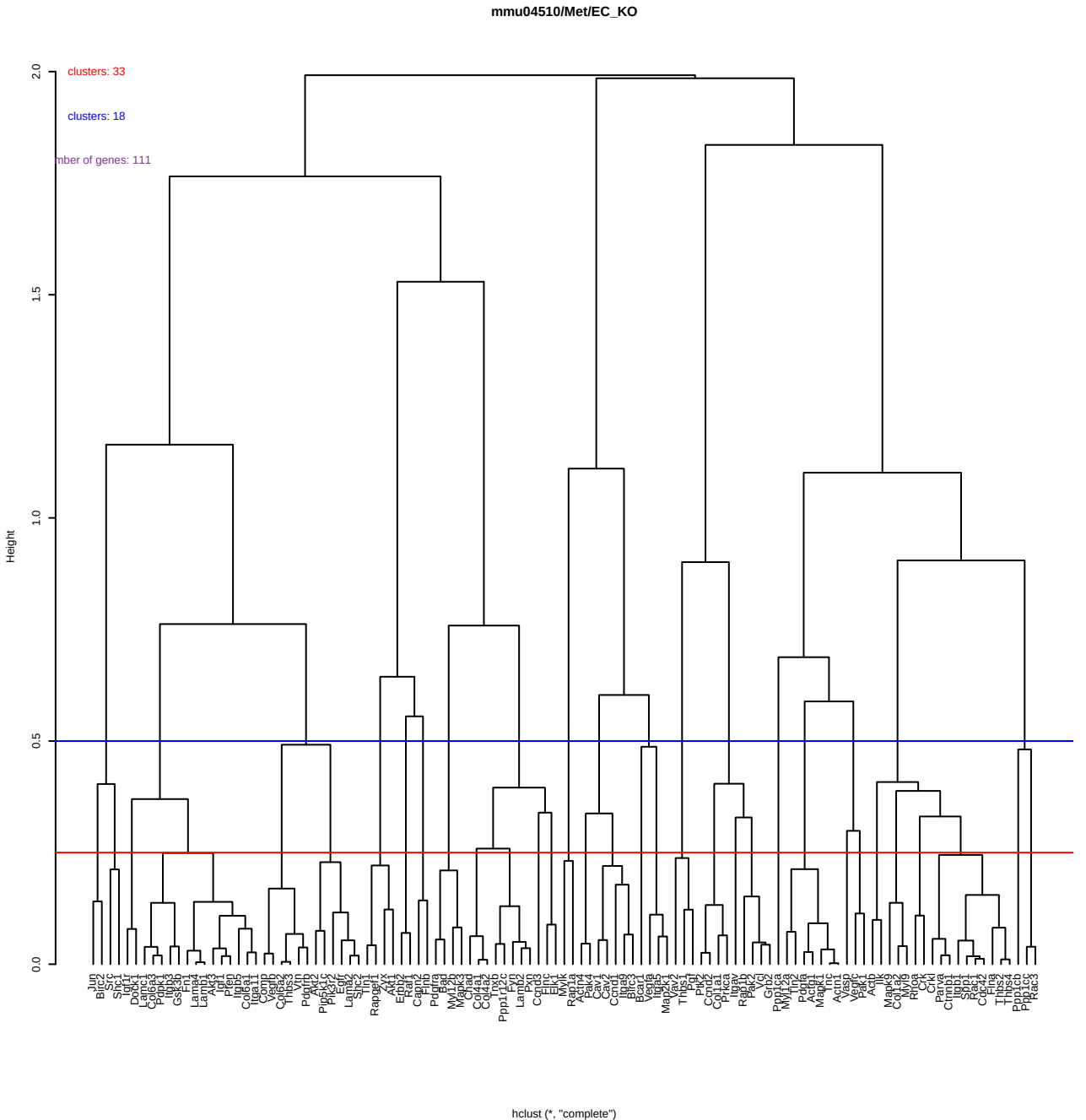

mmu04360/Met/EC_KO
clusters: 31
clusters: 17
mber of genes: 80
Height
2.0
1.5
1.0
0.5
0.0
Pak1
Cfl1
Limk2
Ilk
Ryk
Gnai3
Cfl2
Ntn4
Efnb2
Pik2
Prkca
Ppp3ca
Ppp3cb
Nfatc3
Robo1
Wnt5a
Kras
Ppp3r1
Pak2
Rac3
Ncf1
Nras
Unc5b
Rhoa
Rac1
Cdc42
Enah
Mapk1
Ptpn11
Itgb1
Efnb1
Cxcl12
Ppp3cc
Camk2g
Ntn1
Limk1
Pak4
Slit3
Rnd1
Pard6a
Fyn
Sema6d
Mapk3
Plxna1
Sema4c
Ablim1
Boc
Fes
Rras
Srgap1
Nrp1
Cdk5
Camk2d
Sema3b
Dpysl2
Arhgef12
Ssh2
Abl1
Pik3r2
Plxnb2
Ssh1
Gnai2
Rat1
Efna1
Ephb3
Nck2
Plcg1
Ephb2
Gsk3b
Sema6c
Plxna3
Srgap2
Ssh3
Nfatc4
Src
Rgs3
Rhod
Pard6g
Ephb4
Smo
hclust (*, "complete")

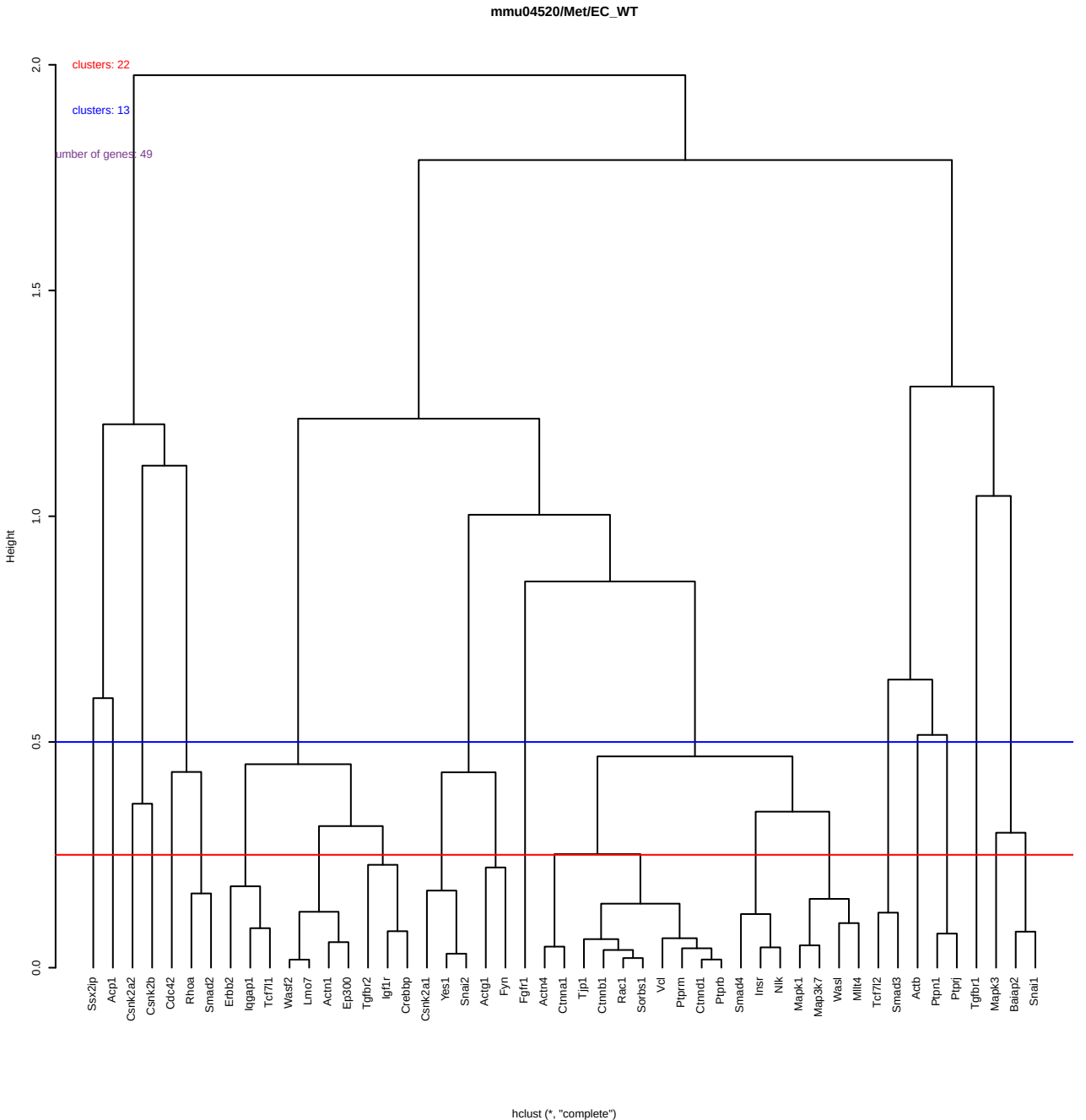

mmu04510/Met/EC_WT
clusters: 38
clusters: 22
mber of genes: 127
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

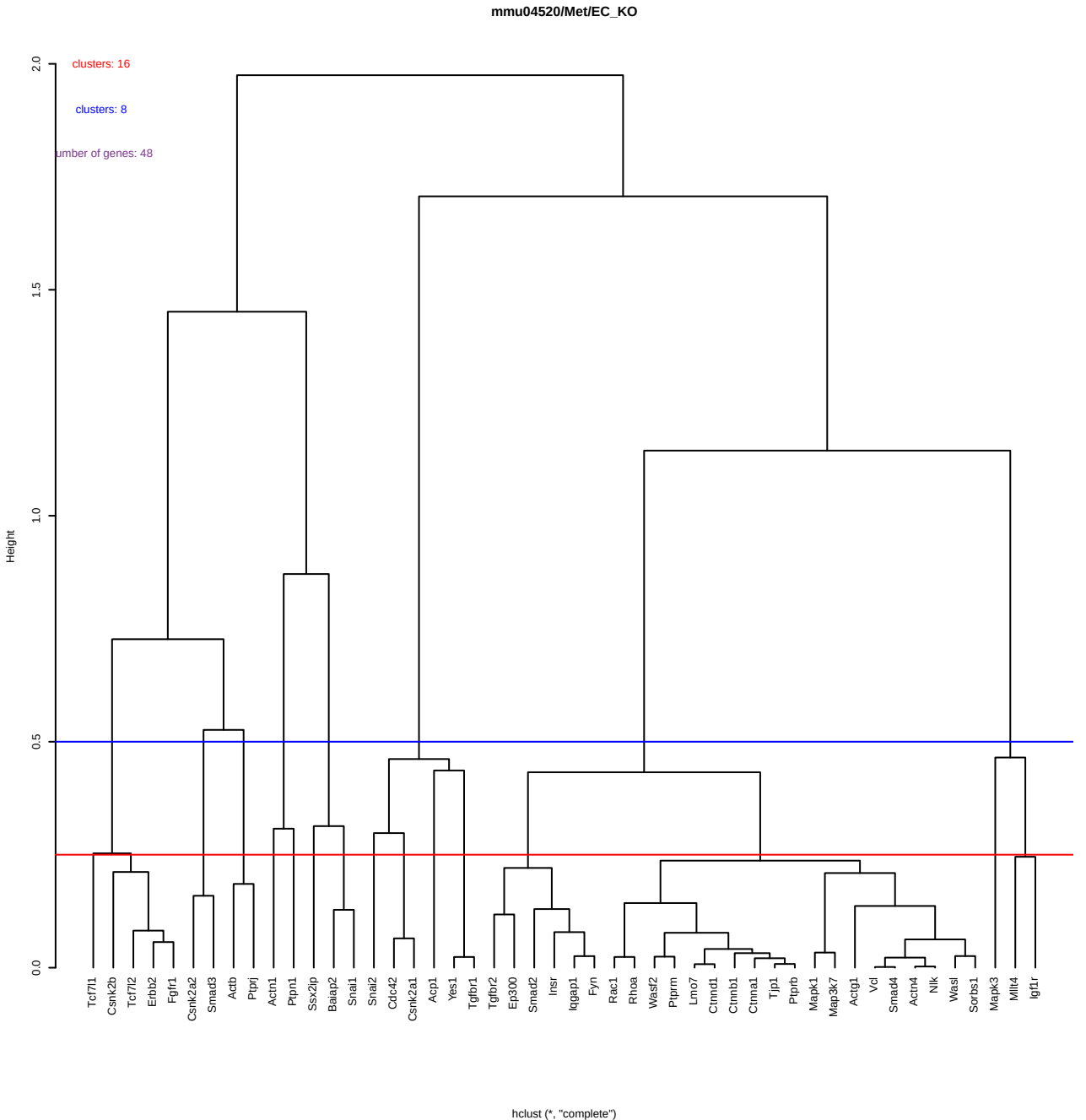


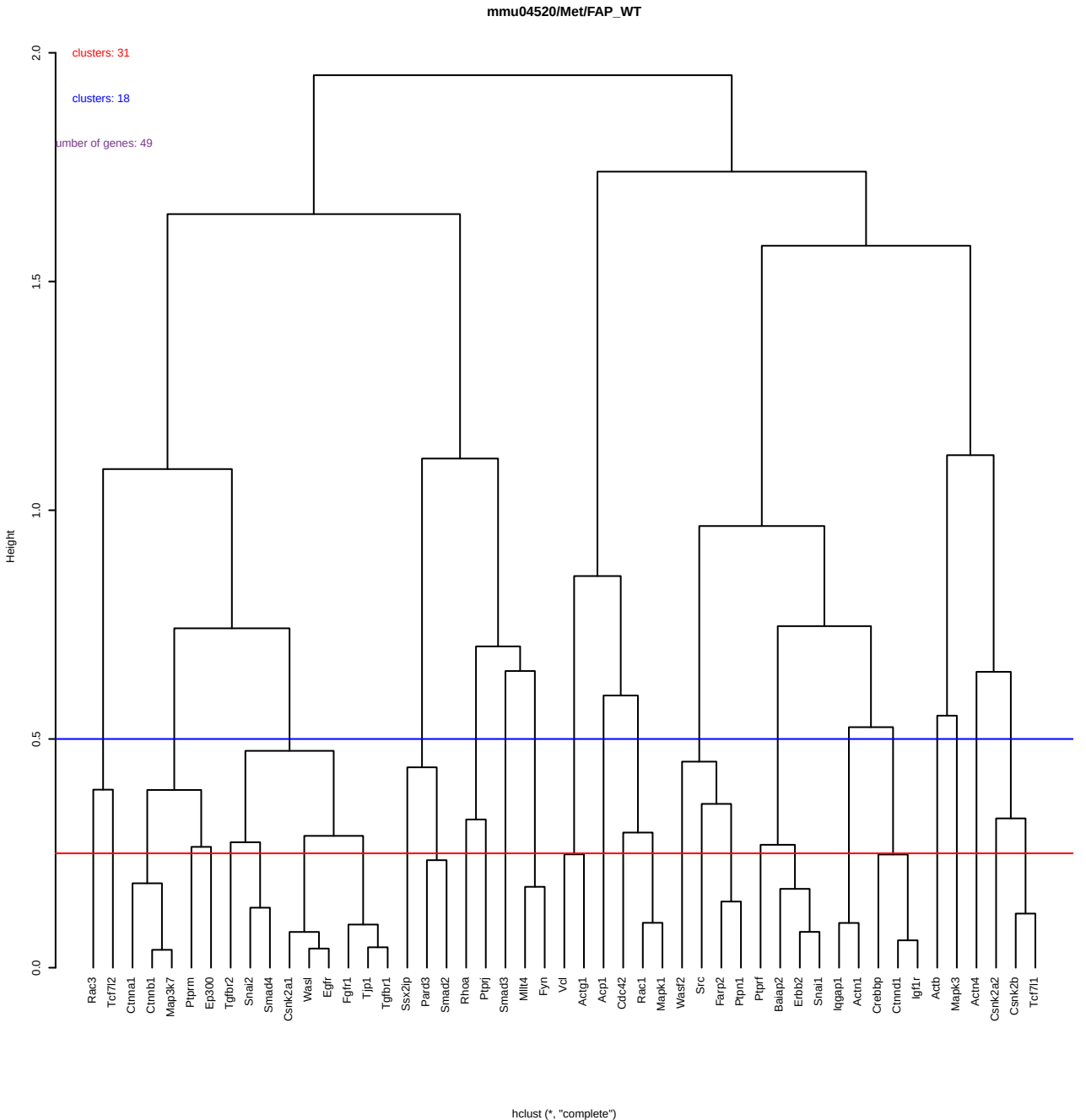
hclust (*, "complete")

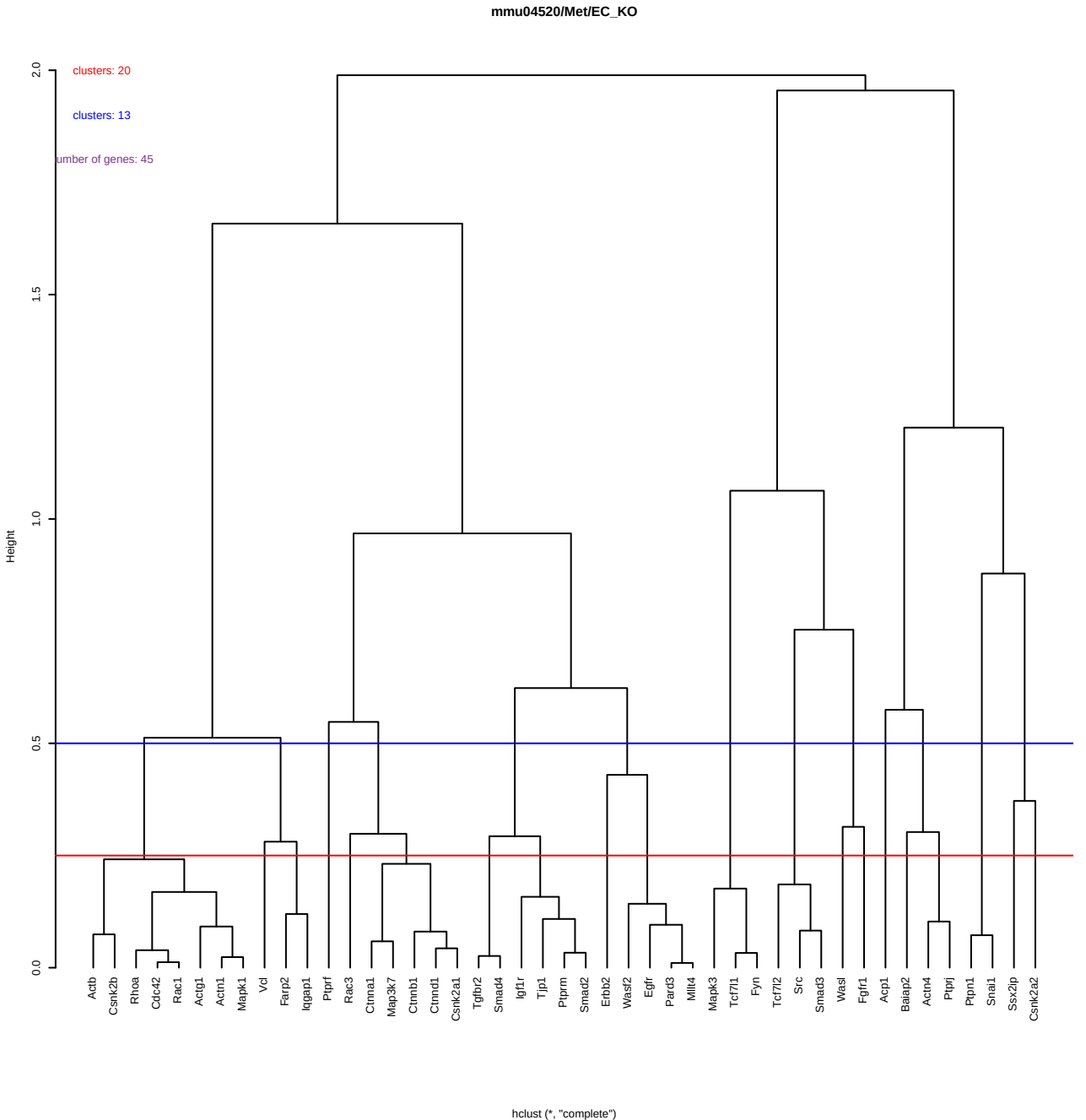

mmu04510/Met/FAP_WT
clusters: 53
clusters: 26
mber of genes: 126
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

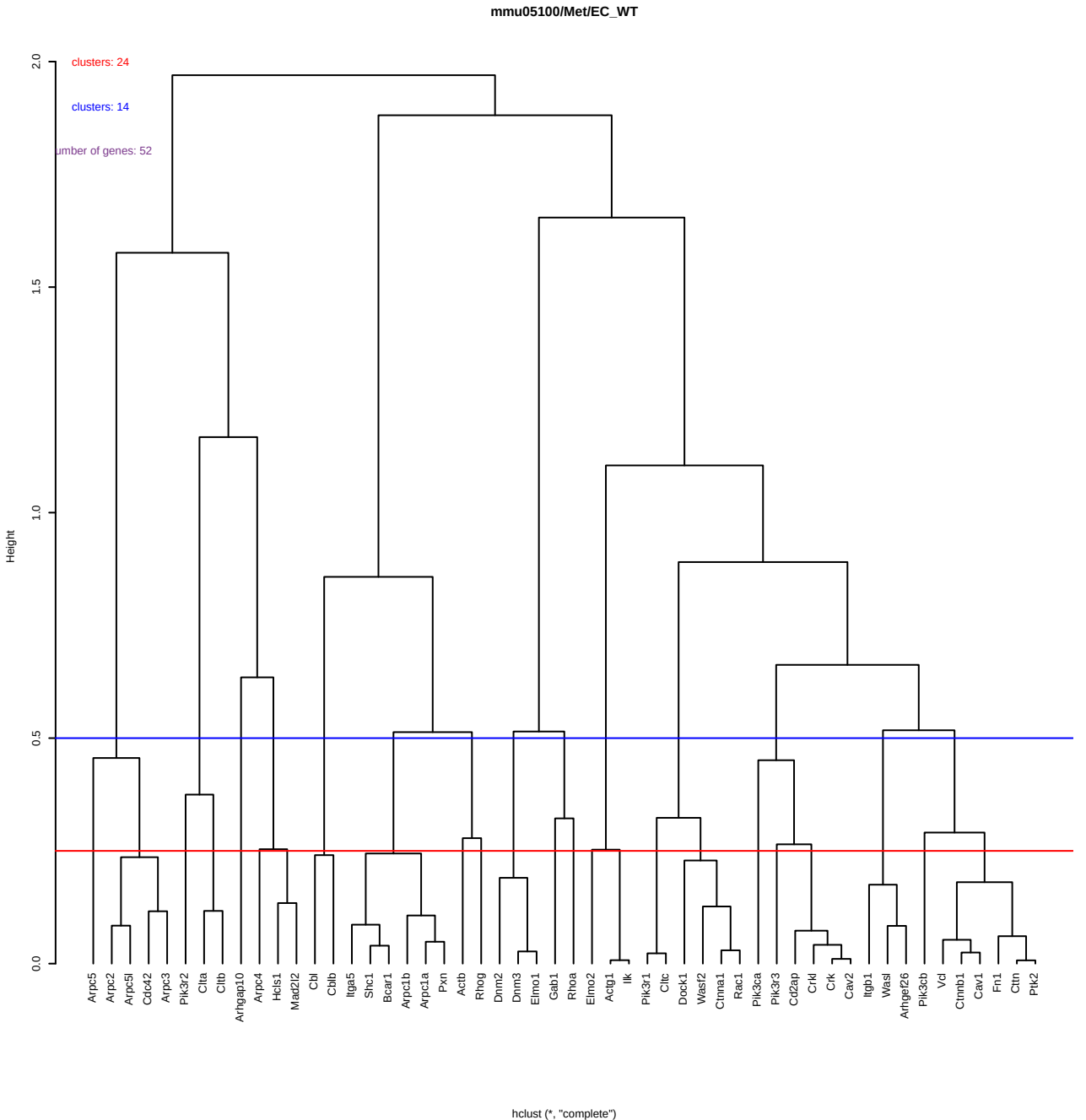

mmu04510/Met/EC_KO
clusters: 33
clusters: 18
mber of genes: 111
Height
2.0
1.5
1.0
0.5
0.0
Jun
Birc2
Src
Sgk1
Idh1
Dok1
Lamc1
Cdc42
Pdpk1
Itgb3
Gsk3b
Fn1
Lama4
Lamb1
Akt3
Jak1
Pfkp
Itgb5
Cdh1
Itga11
Comp
Vegfa
Cxcl12
Thbs3
Vtn
Ptpn11
Akt2
Pik3r1
Pik3r2
Lama5
Shc2
Tjp1
Rapgef1
Fyx
Akt1
Ezr
Rac1
Capn2
Prkcb
Itgb
Bad
Myl12b
Mmp2
Cfl1
Col4a1
Col4a2
Ppp1r12c
Fyn
Lamb2
Lamc2
Cnn3
Fn1
Elnc
Mylk
Rap1a
Actn4
Pdk1
Cav1
Cav2
Cnn1
Cnn2
Birc3
Bcar1
Vegfa
Itga9
Map2k1
Tjp1
Tnfrsf1a
Pgf
Pik3r2
Col1a2
Prkca
Itgav
Rac2
Pak2
Vcl
Gnb2
Gnb1
Ppp1r12a
Myl12a
Tln2
Actn1
Actg1
Mapk1
Inc
Actn2
Vasp
Vegfc
Pak1
Actn3
Ilk
Mapk9
Cxcl9
Myl9
Rhoa
Crk
Crk1
Parva
Ctnnb1
Sgk1
Spp1
Rac1
Cdc42
Fn1
Thbs2
Thbs4
Ppp1cb
Ppp1cc
Rac3
hclust (*, "complete")

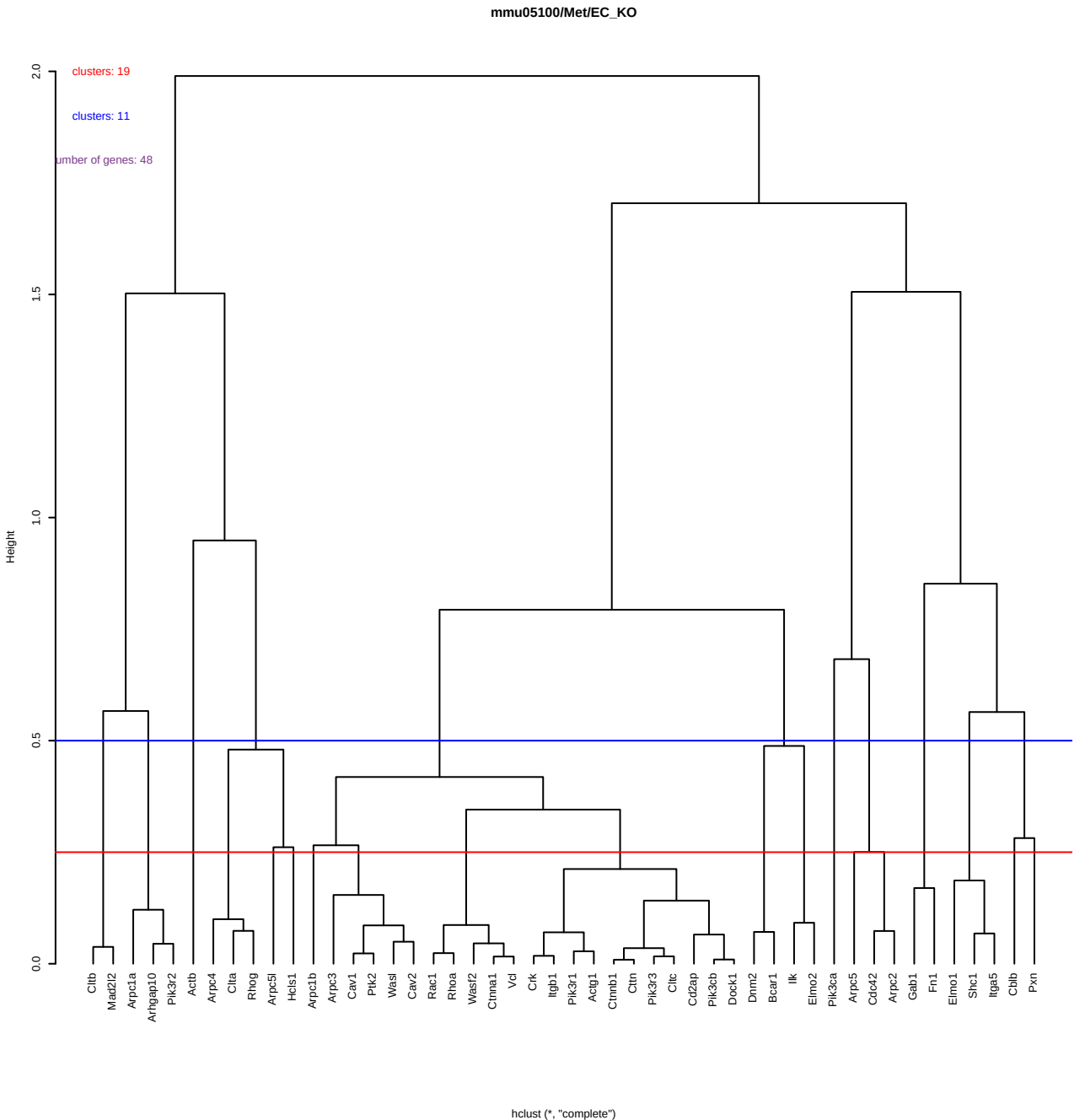

mmu04520/Met/EC_WT
clusters: 22
clusters: 13
umber of genes: 49
Height
2.0
1.5
1.0
0.5
0.0
Ssx2ip
Acp1
Csnk2a2
Csnk2b
Cdc42
Rhoa
Smad2
Erbb2
Iqgap1
Tcf7l1
Wasf2
Lmo7
Actn1
Ep300
Tgfbr2
Igf1r
Crebbp
Csnk2a1
Yes1
Snai2
Actg1
Fyn
Fgfr1
Actn4
Ctnna1
Tjp1
Ctnnb1
Rac1
Sorbs1
Vcl
Ptprm
Ctnnd1
Ptprb
Smad4
Insr
Nlk
Mapk1
Map3k7
Wasl
Mll4
Tcf7l2
Smad3
Actb
Ptpn1
Ptprj
Tgfbr1
Mapk3
Baiap2
Snai1
hclust (*, "complete")

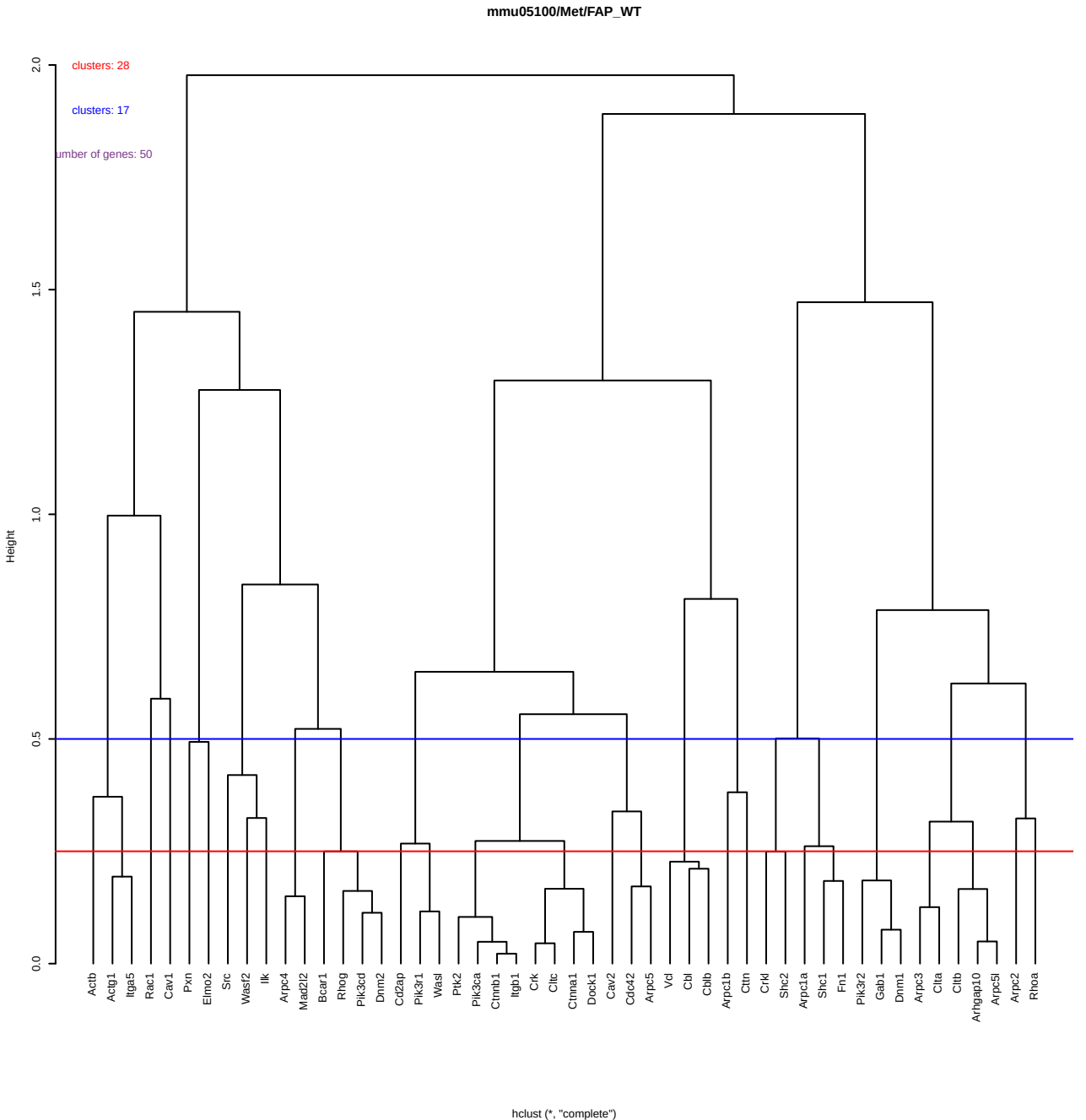

mmu04520/Met/EC_KO
clusters: 16
clusters: 8
umber of genes: 48
Height
2.0
1.5
1.0
0.5
0.0
Tcf7l1
Csnk2b
Tcf7l2
Erbb2
Fgfr1
Csnk2a2
Smad3
Actb
Ptprj
Actn1
Ptpn1
Ssx2ip
Baiap2
Snai1
Snai2
Cdc42
Csnk2a1
Acp1
Yes1
Tgfbr1
Tgfbr2
Ep300
Smad2
Insr
Iqgap1
Fyn
Rac1
Rhoa
Wasf2
Ptprm
Lmo7
Ctnnd1
Ctnnb1
Ctnna1
Tjp1
Ptprb
Mapk1
Map3k7
Actg1
Vcl
Smad4
Actn4
Nlk
Wasl
Sorbs1
Mapk3
Mllt4
Igf1r
hclust (*, "complete")

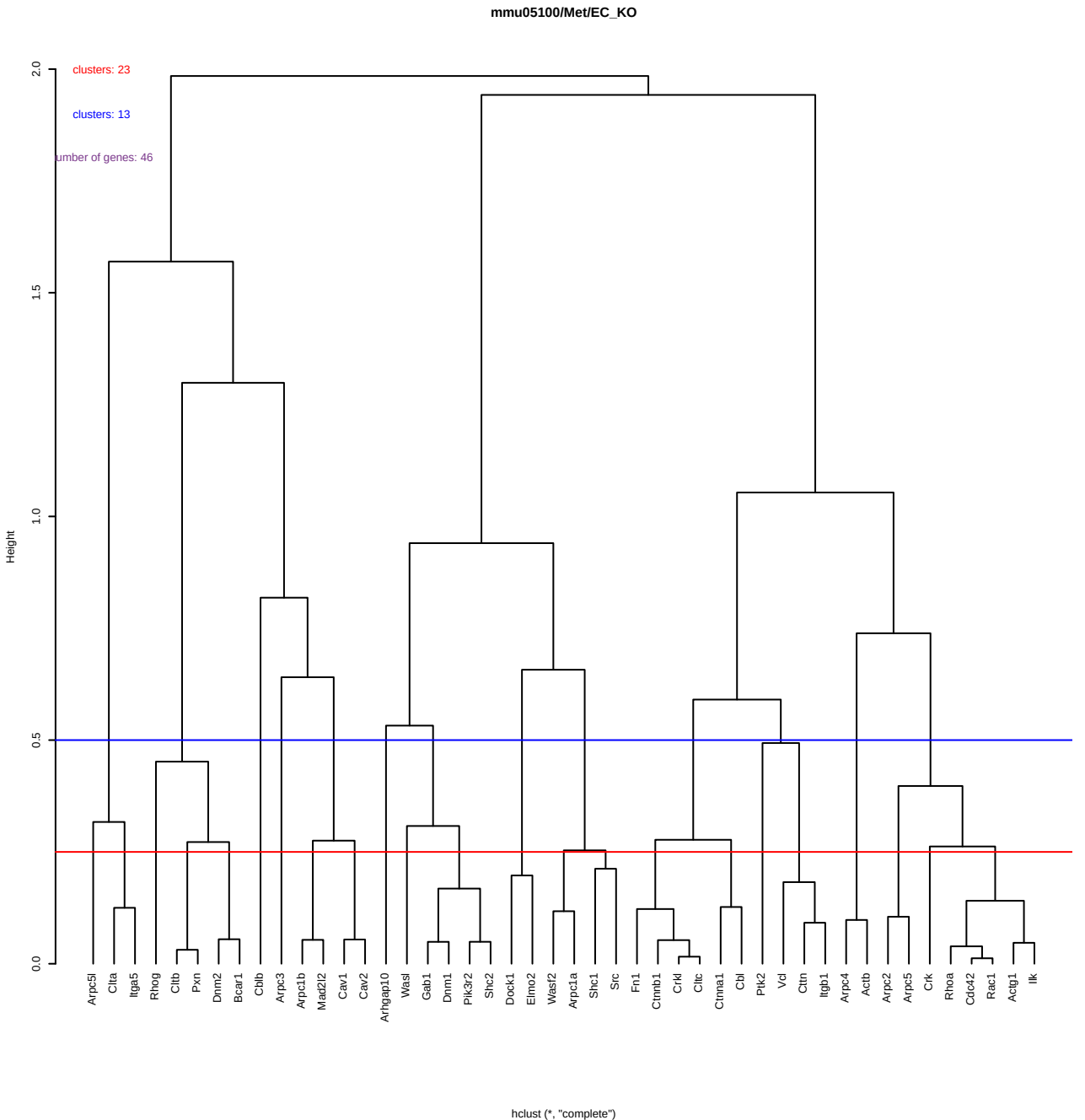

mmu04520/Met/FAP_WT
clusters: 31
clusters: 18
umber of genes: 49
Height
2.0
1.5
1.0
0.5
0.0
Rac3
Tcf7l2
Ctnna1
Ctnnb1
Map3k7
Ptprm
Ep300
Tgfbr2
Snai2
Smad4
Csnk2a1
Wasl
Egfr
Fgfr1
Tjp1
Tgfbr1
Ssx2ip
Pard3
Smad2
Rhoa
Ptprj
Smad3
Mllt4
Fyn
Vcl
Actg1
Acp1
Cdc42
Rac1
Mapk1
Wasf2
Src
Farp2
Ptpn1
Ptprf
Baiap2
Erbb2
Snai1
Iqgap1
Actn1
Crebbp
Ctnnd1
Igf1r
Actb
Mapk3
Actn4
Csnk2a2
Csnk2b
Tcf7l1
hclust (*, "complete")

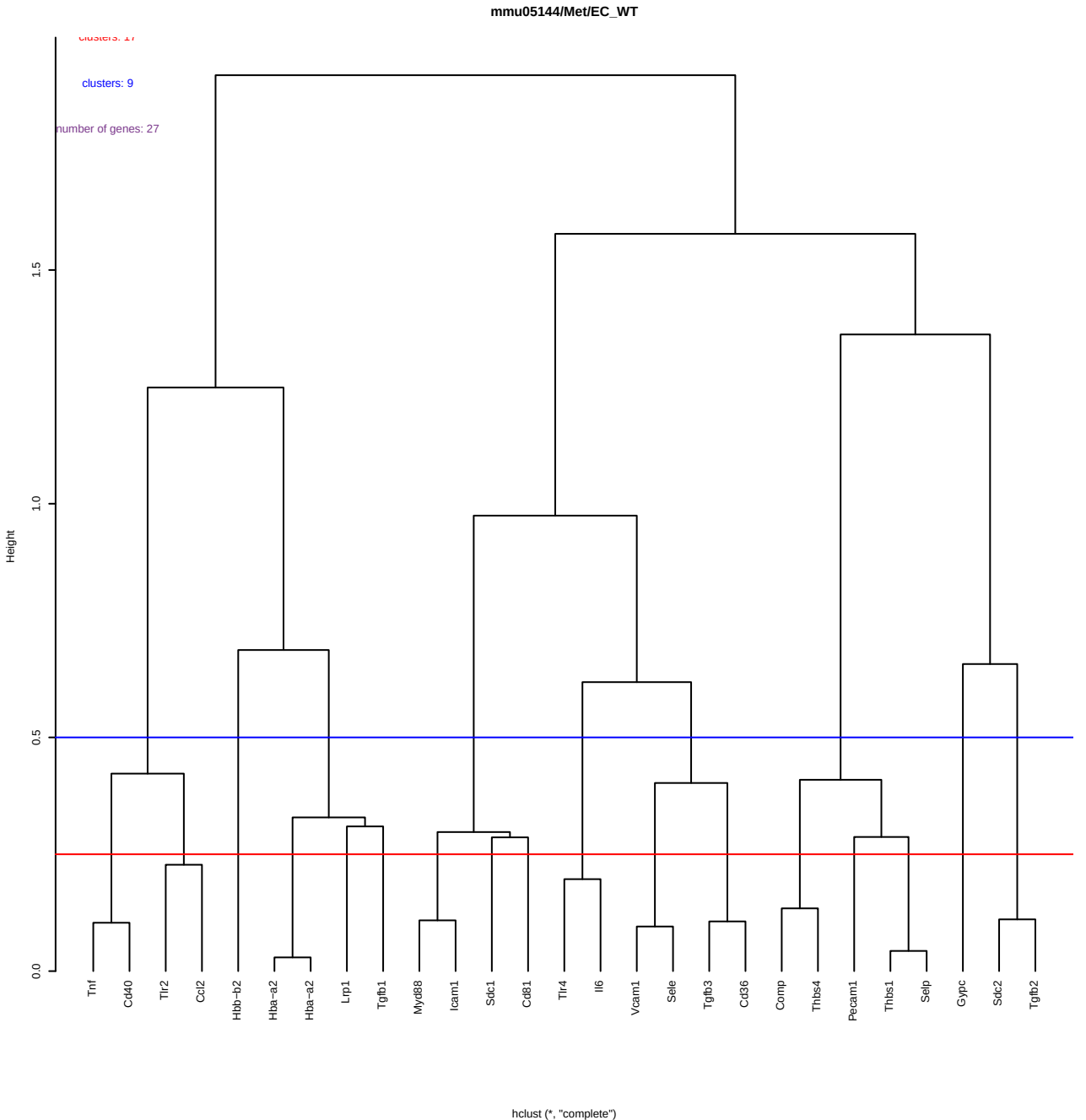

mmu04520/Met/EC_KO
clusters: 20
clusters: 13
umber of genes: 45
Height
2.0
1.5
1.0
0.5
0.0
Actb
Csnk2b
Rhoa
Cdc42
Rac1
Actg1
Actn1
Mapk1
Vcl
Farp2
Iqgap1
Ptprf
Rac3
Ctnna1
Map3k7
Ctnnb1
Ctnnd1
Csnk2a1
Tgfbr2
Smad4
Igf1r
Tjp1
Ptprm
Smad2
Erbb2
Wasf2
Egfr
Pard3
Mllt4
Mapk3
Tcf7l1
Fyn
Tcf7l2
Src
Smad3
Wasl
Fgfr1
Acp1
Baiap2
Actn4
Ptprj
Ptpn1
Snai1
Ssx2ip
Csnk2a2
hclust (*, "complete")

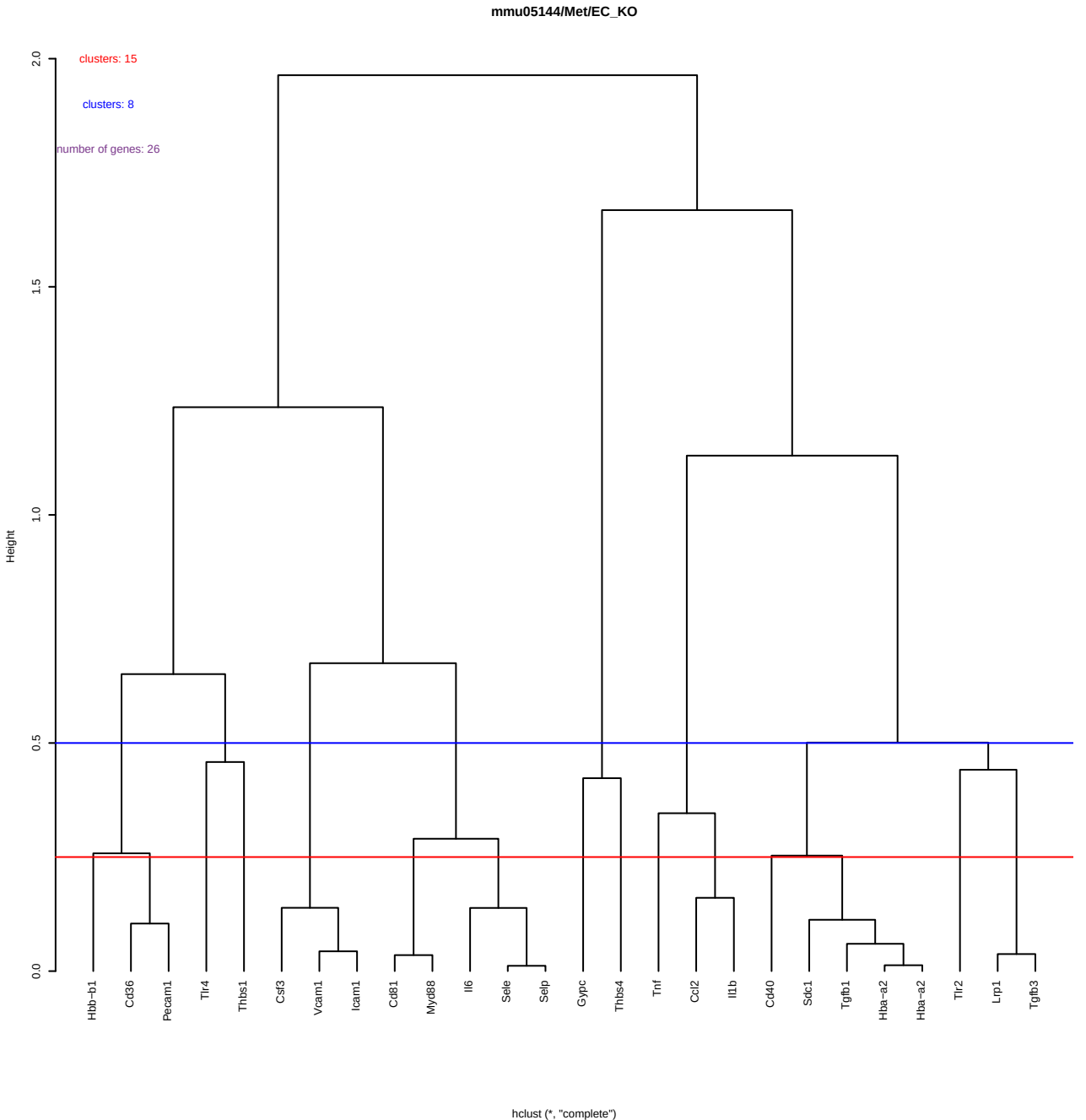

mmu05100/Met/EC_WT
clusters: 24
clusters: 14
umber of genes: 52
Height
2.0
1.5
1.0
0.5
0.0
Arpc5
Arpc2
Arpc5l
Cdc42
Arpc3
Pik3r2
Clta
Cltb
Arhgap10
Arpc4
Hcls1
Mad2l2
Ctl
Cblb
Itga5
Shc1
Bcar1
Arpc1b
Arpc1a
Pxn
Actb
Rhog
Dnm2
Dnm3
Elmo1
Gab1
Rhoa
Elmo2
Actg1
Ilk
Pik3r1
Cltc
Dock1
Wasf2
Ctnna1
Rac1
Pik3ca
Pik3r3
Cd2ap
Crkl
Crk
Cav2
Itgb1
Wasl
Arhgef26
Pik3cb
Vcl
Ctnnb1
Cav1
Fn1
Ctn
Ptk2
hclust (*, "complete")

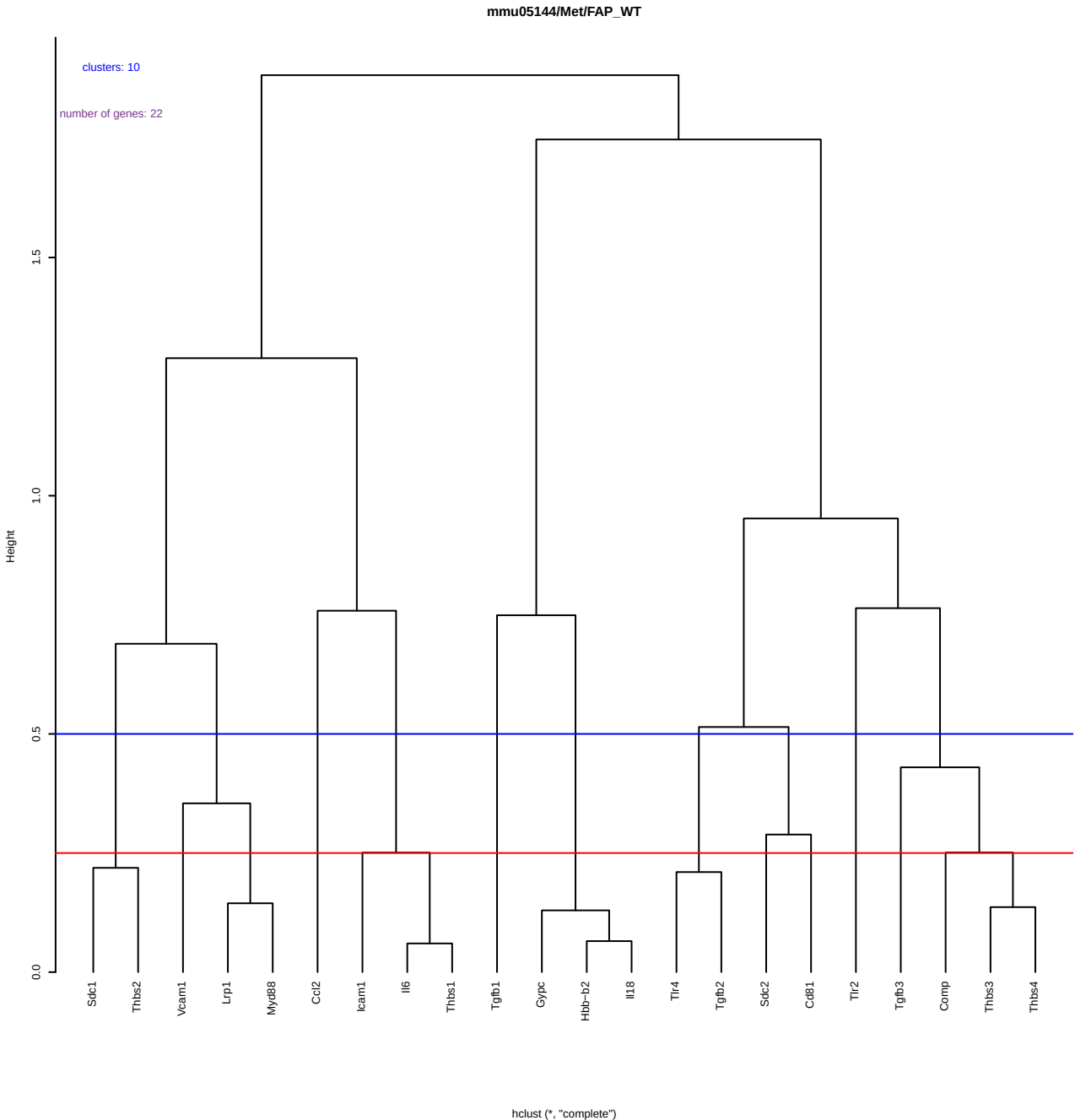

mmu05100/Met/EC_KO
clusters: 19
clusters: 11
umber of genes: 48
Height
2.0
1.5
1.0
0.5
0.0
Cltb
Mad2l2
Arpc1a
Arhgap10
Pik3r2
Actb
Arpc4
Clta
Rhog
Arpc5l
Hcls1
Arpc1b
Arpc3
Cav1
Ptk2
Wasl
Cav2
Rac1
Rhoa
Wasf2
Ctnna1
Vcl
Crk
Itgb1
Pik3r1
Actg1
Ctnnb1
Cttn
Pik3r3
Cltc
Cd2ap
Pik3cb
Dock1
Dnm2
Bcar1
Ilk
Elmo2
Pik3ca
Arpc5
Cdc42
Arpc2
Gab1
Fn1
Elmo1
Shc1
Itga5
Cblb
Pxn
hclust (*, "complete")

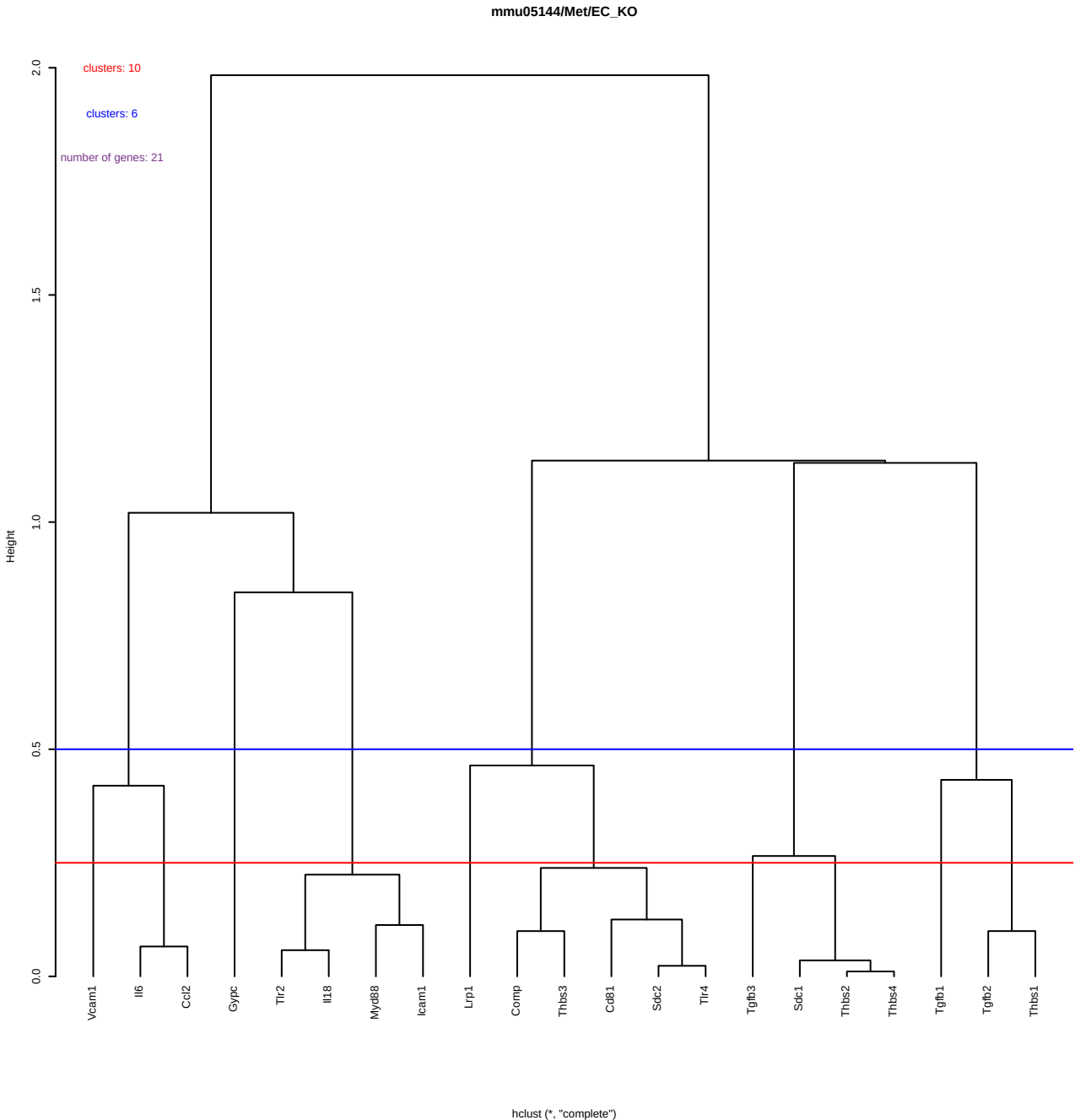

mmu05100/Met/FAP_WT
clusters: 28
clusters: 17
umber of genes: 50
Height
2.0
1.5
1.0
0.5
0.0
Actb
Actg1
Itga5
Rac1
Cav1
Pxn
Elmo2
Src
Wasf2
Ilk
Arpc4
Mad2l2
Bcar1
Rhog
Pik3cd
Dnm2
Cd2ap
Pik3r1
Wasl
Ptk2
Pik3ca
Ctnnb1
Itgb1
Crk
Cltc
Ctnna1
Dock1
Cav2
Cdc42
Arpc5
Vcl
Cbl
Cblb
Arpc1b
Cttn
Crkl
Shc2
Arpc1a
Shc1
Fn1
Pik3r2
Gab1
Dnm1
Arpc3
Clta
Cltb
Arhgap10
Arpc5l
Arpc2
Rhoa
hclust (*, "complete")

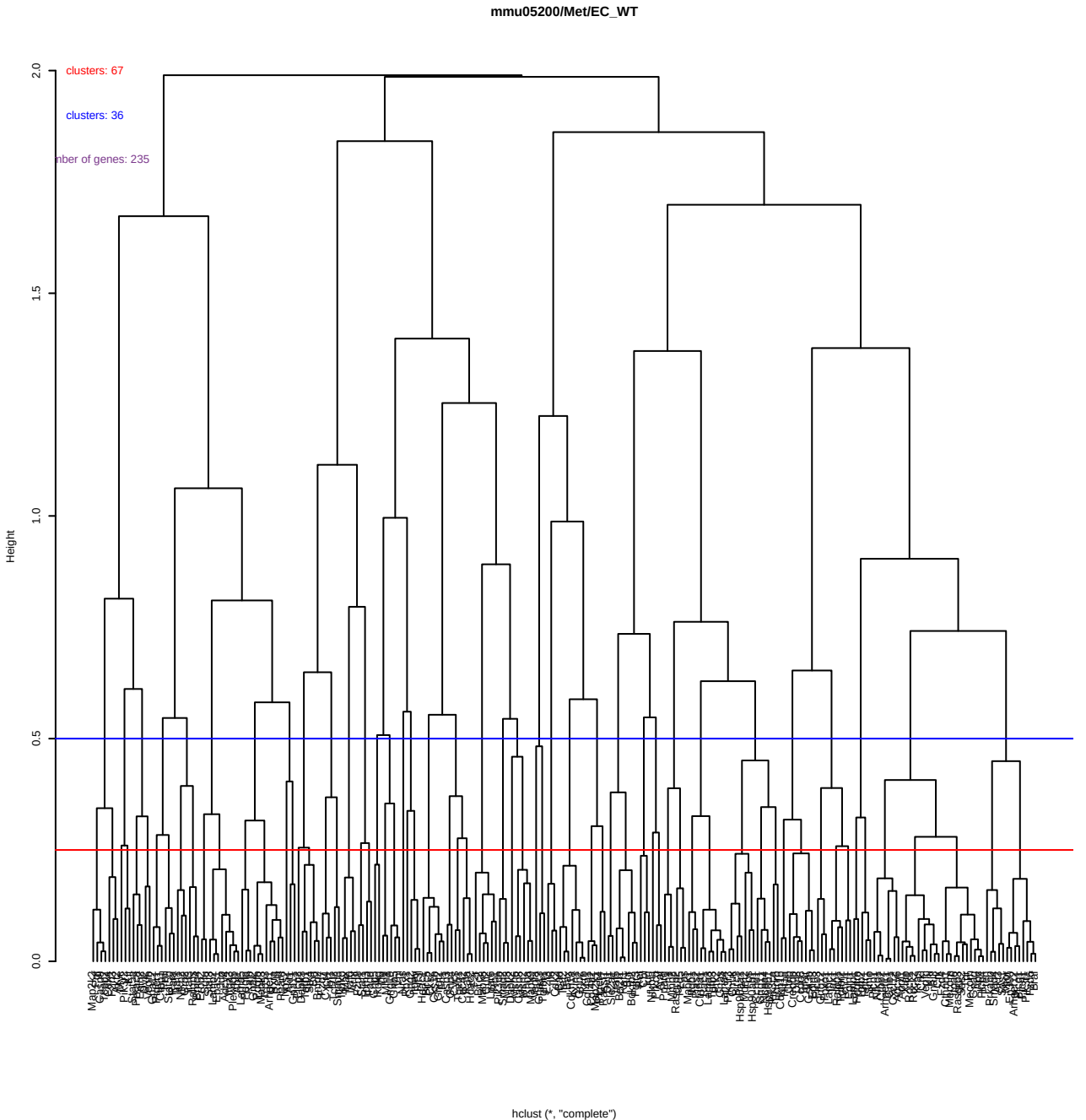

mmu05100/Met/EC_KO
clusters: 23
clusters: 13
umber of genes: 46
Height
2.0
1.5
1.0
0.5
0.0
Arpc5l
Clta
Itga5
Rhog
Cltb
Pxn
Dnm2
Bcar1
Cblb
Arpc3
Arpc1b
Madd2l2
Cav1
Cav2
Arhgap10
Wasl
Gab1
Dnm1
Pik3r2
Shc2
Dock1
Elmo2
Wasf2
Arpc1a
Shc1
Src
Fn1
Ctnnb1
Crkl
Cltc
Ctnna1
Cbl
Ptk2
Vcl
Cttn
Itgb1
Arpc4
Actb
Arpc2
Arpc5
Crk
Rhoa
Cdc42
Rac1
Actg1
Ilk
hclust (*, "complete")

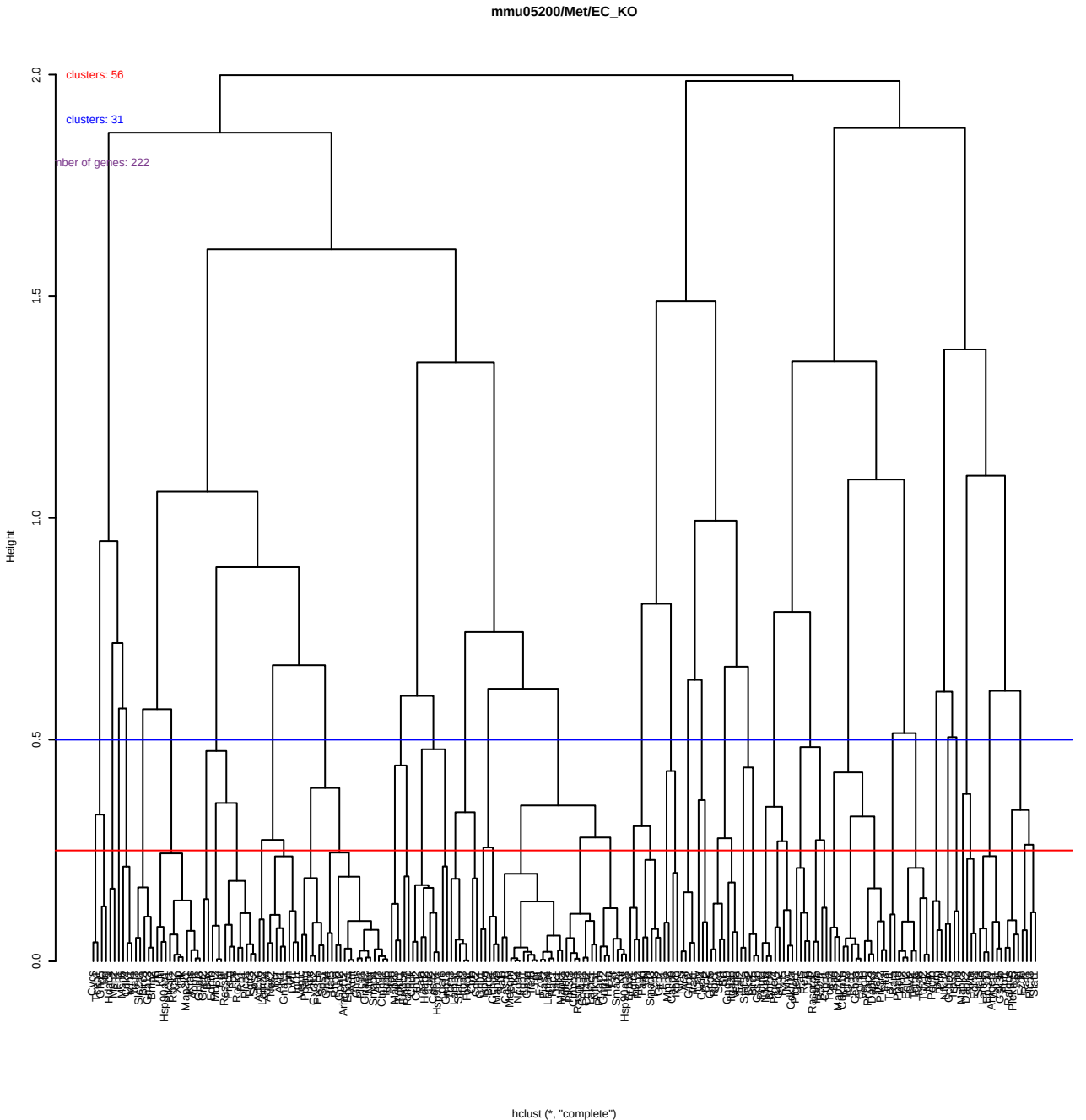

mmu05144/Met/EC_WT
clusters: 17
clusters: 9
number of genes: 27
Height
0.0
0.5
1.0
1.5
Tnf
Cd40
Tlr2
Ccl2
Hbb−b2
Hba−a2
Hba−a2
Lrp1
Tgfb1
Myd88
Icam1
Sdc1
Cd81
Tlr4
Il6
Vcam1
Sele
Tgfb3
Cd36
Comp
Thbs4
Pecam1
Thbs1
Selp
Gypc
Sdc2
Tgfb2
hclust (*, "complete")

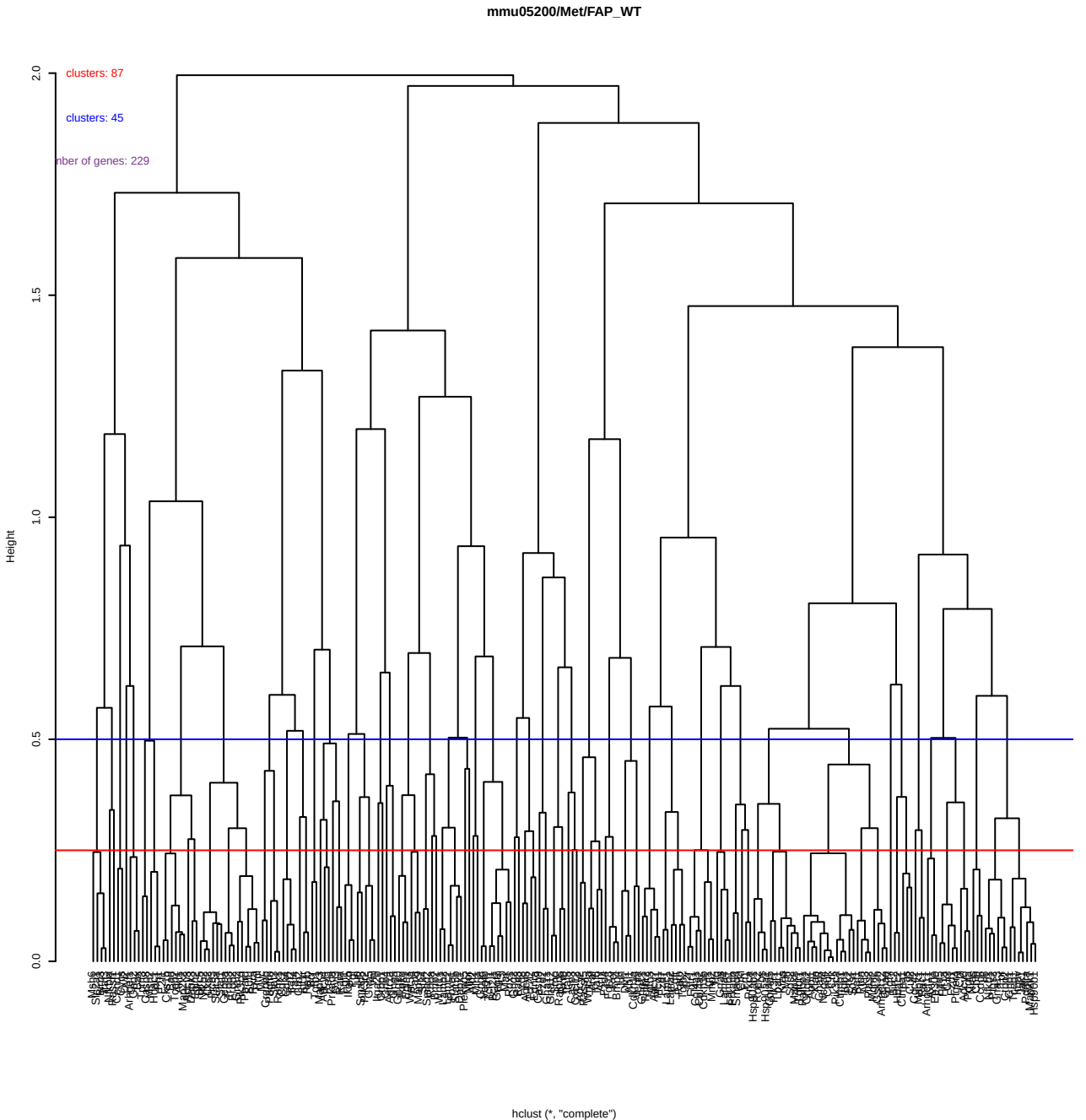

mmu05144/Met/EC_KO
clusters: 15
clusters: 8
number of genes: 26
Height
0.0
0.5
1.0
1.5
2.0
Hbb−b1
Cd36
Pecam1
Tlr4
Thbs1
Csf3
Vcam1
Icam1
Cd81
Myd88
Il6
Sele
Selp
Gypc
Thbs4
Tnf
Ccl2
Il1b
Cd40
Sdc1
Tgfb1
Hba−a2
Hba−a2
Tlr2
Lrp1
Tgfb3
hclust (*, "complete")

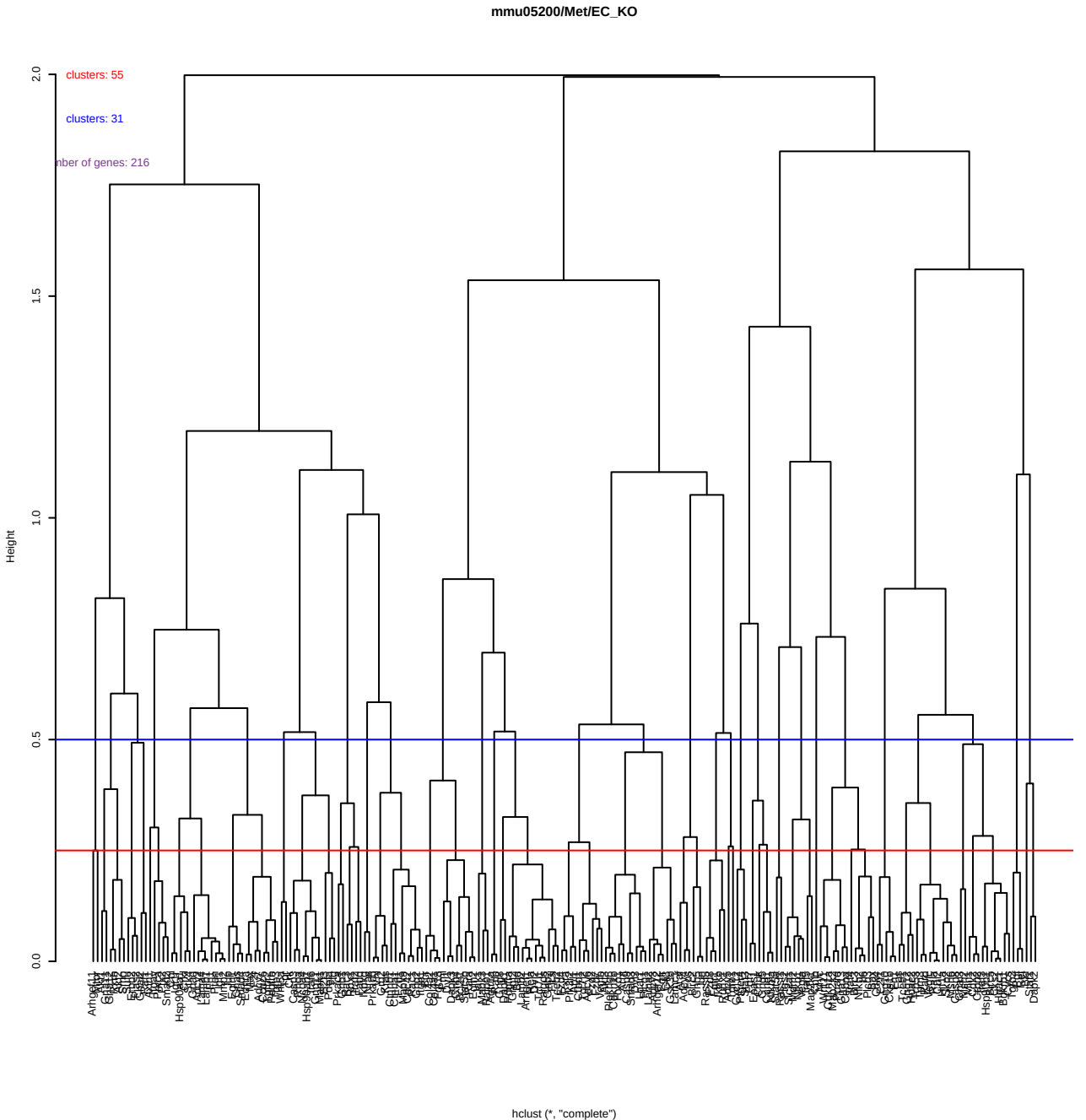

mmu05144/Met/FAP_WT
clusters: 10
number of genes: 22
Height
0.0
0.5
1.0
1.5
Sdc1
Thbs2
Vcam1
Lrp1
Myd88
Ccl2
Icam1
Il6
Thbs1
Tgfb1
Gypc
Hbb−b2
Il18
Tlr4
Tgfb2
Sdc2
Cd81
Tlr2
Tgfb3
Comp
Thbs3
Thbs4
hclust (*, "complete")

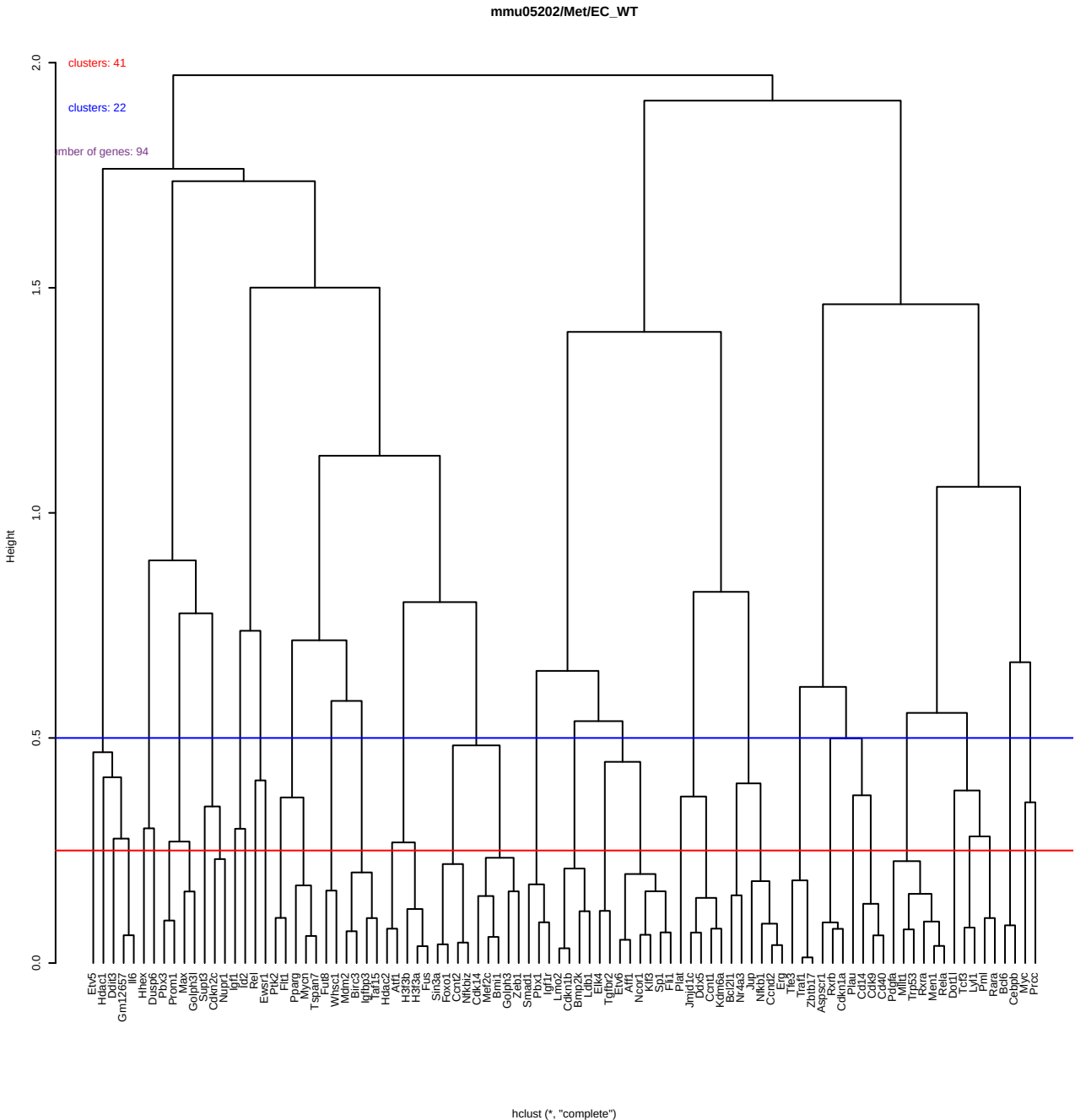

mmu05144/Met/EC_KO
clusters: 10
clusters: 6
number of genes: 21
Height
2.0
1.5
1.0
0.5
0.0
Vcam1
Il6
Ccl2
Gypc
Tlr2
Il18
Myd88
Icam1
Lrp1
Comp
Thbs3
Cd81
Sdc2
Tlr4
Tgfb3
Sdc1
Thbs2
Thbs4
Tgfb1
Tgfb2
Thbs1
hclust (*, "complete")

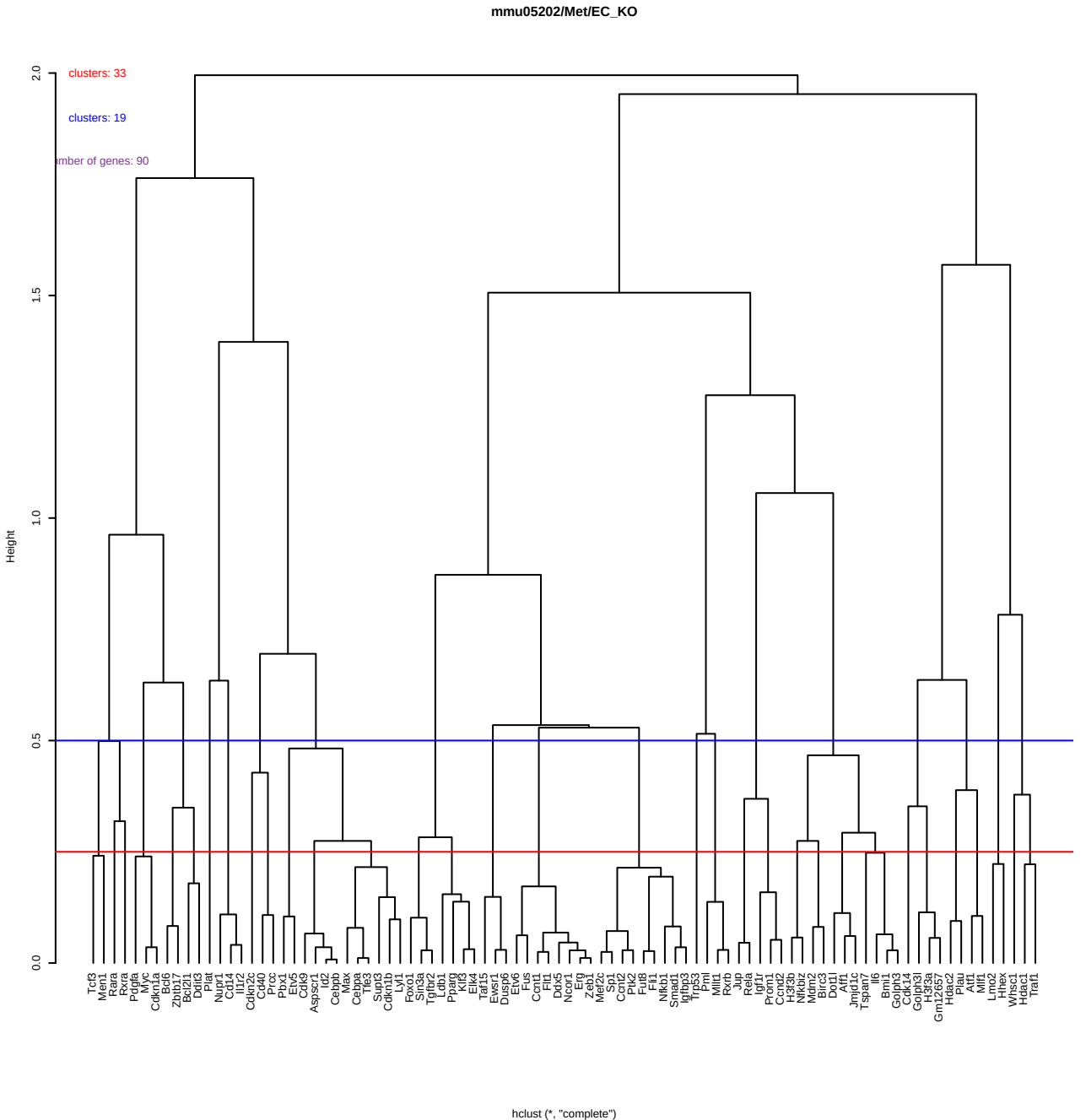

mmu05200/Met/EC_WT
clusters: 67
clusters: 36
nber of genes: 235
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

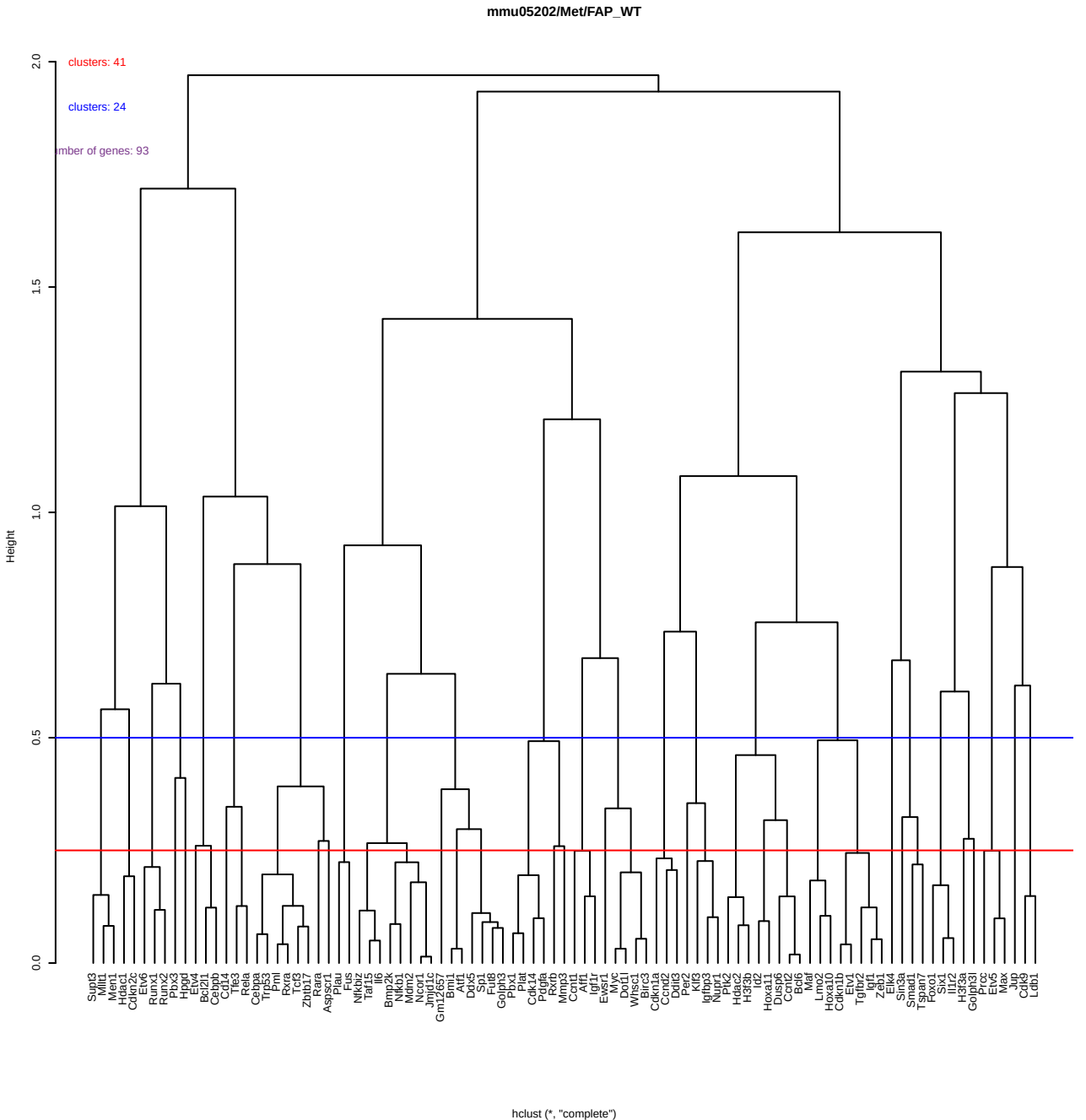

mmu05200/Met/EC_KO
clusters: 56
clusters: 31
nber of genes: 222
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

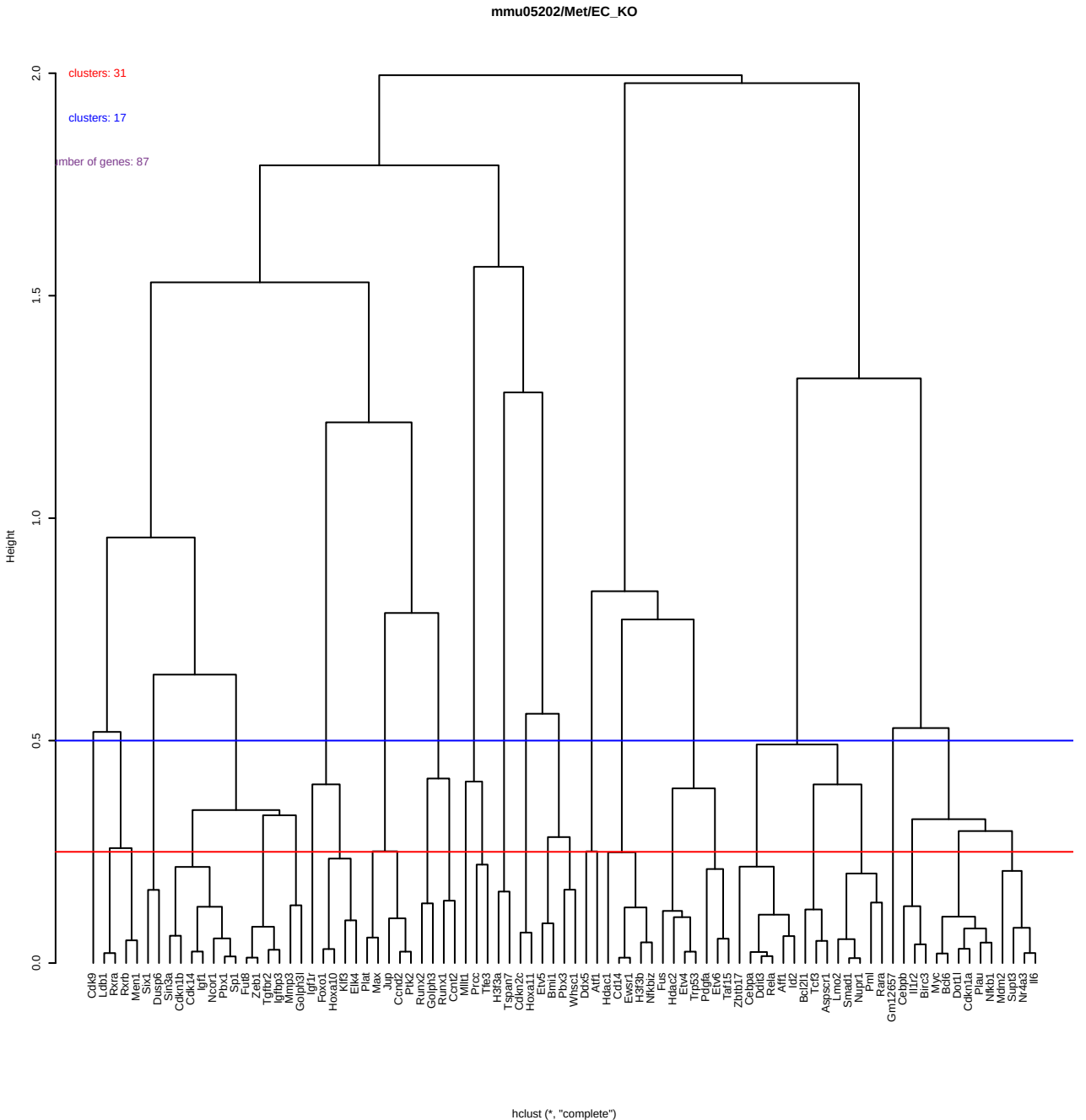

mmu05200/Met/FAP_WT
clusters: 87
clusters: 45
nber of genes: 229
Height
0.0
0.5
1.0
1.5
2.0
hclust (*, "complete")

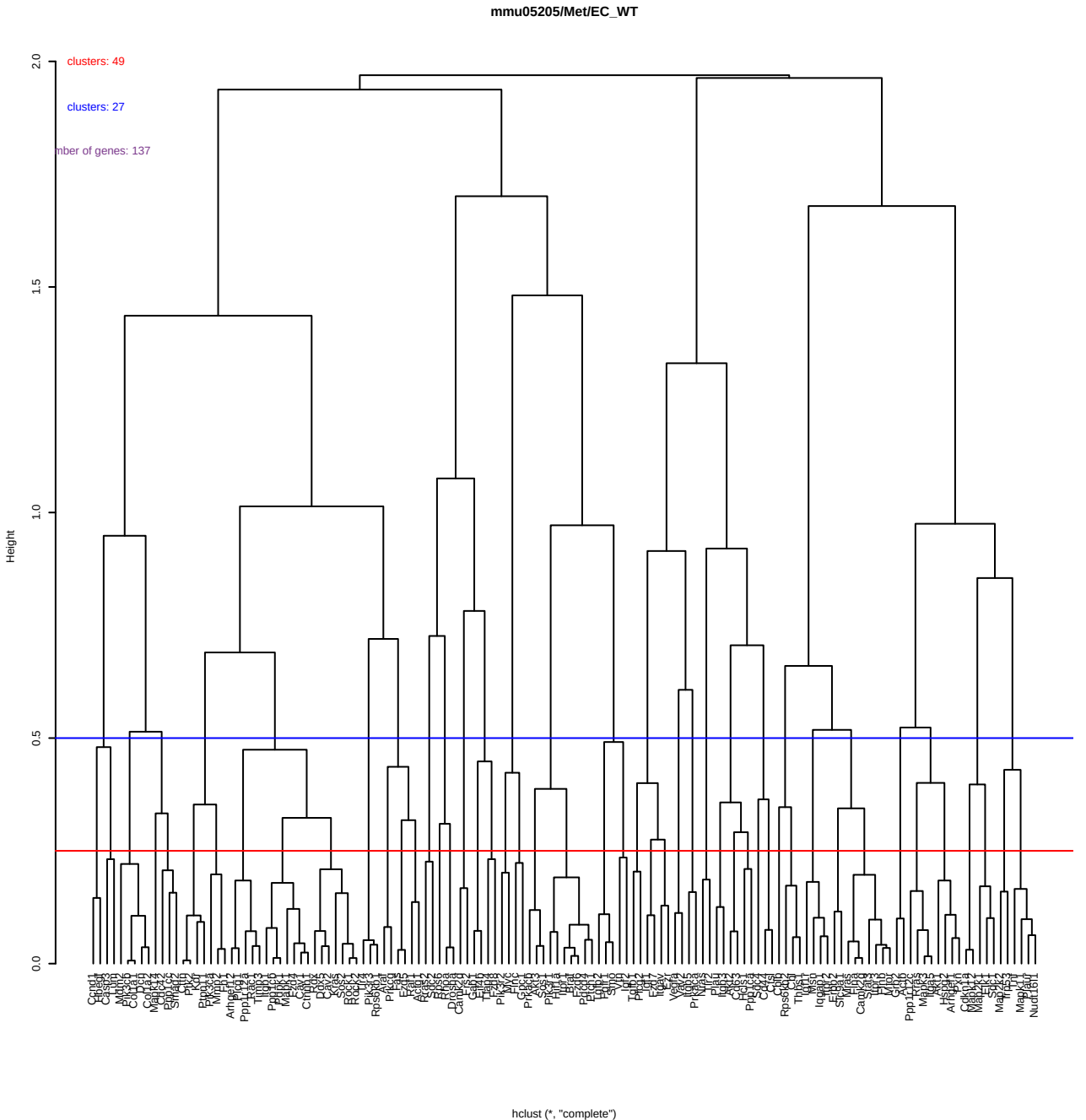

mmu05200/Met/EC_KO
clusters: 55
clusters: 31
nber of genes: 216
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

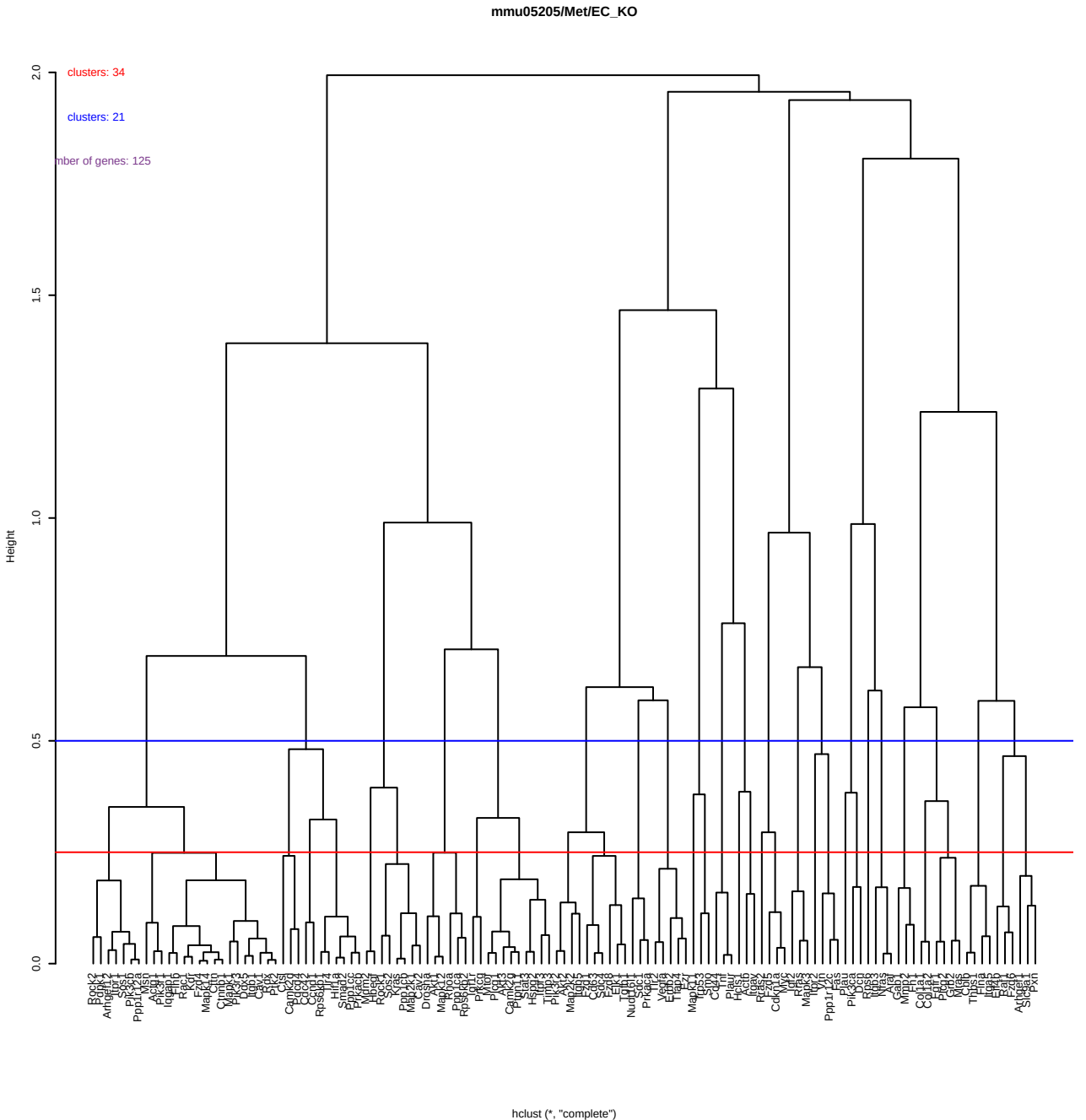

mmu05202/Met/EC_WT
clusters: 41
clusters: 22
mber of genes: 94
Height
2.0
1.5
1.0
0.5
0.0
Etv5
Hdac1
Ddit3
Gm12657
Il6
Hhex
Dbp3
Pbx3
Prom1
Max
Golph3l
Sup3
Cdkn2c
Nupr1
Igf1
Id2
Rei
Ewsr1
Pik2
Fit1
Pxdng
Myor
Tspan7
Fut8
Whsc1
Mdm2
Birc3
Ikbkb
Tgfb1
Hdac2
Atf1
H3f3b
H3f3a
Fus
Slx3a
Foxo1
Ccn1
Nfkbiz
Cdk14
Mef2c
Bmi1
Gzmm
Zfp13
Smad1
Pbx1
Igf1r
Lmo2
Cdkn1b
Bmp2k
Lhx1
Elk4
Tgfbr2
Etv6
Atf1
Nccr1
Klf3
Sp1
Fit1
Plat
Jmjd1c
Ddx5
Ccnt1
Kmt2a
Ikbkn
Bcl7l
Nr4a3
Jup
Nfkb1
Ccnd2
Eg
Tle3
Tnfrsf1
Zbtb17
Aspscr1
Rxrb
Cdkn1a
Plau
Cdk4
Cdk14
Cd40
Pdgfa
Mll1
Trp53
Rxra
Mef2a
Rel1
Dot1l
Tcf3
Lyl1
Pml
Rara
Bcl6
Cebpb
Myc
Prc
hclust (*, "complete")

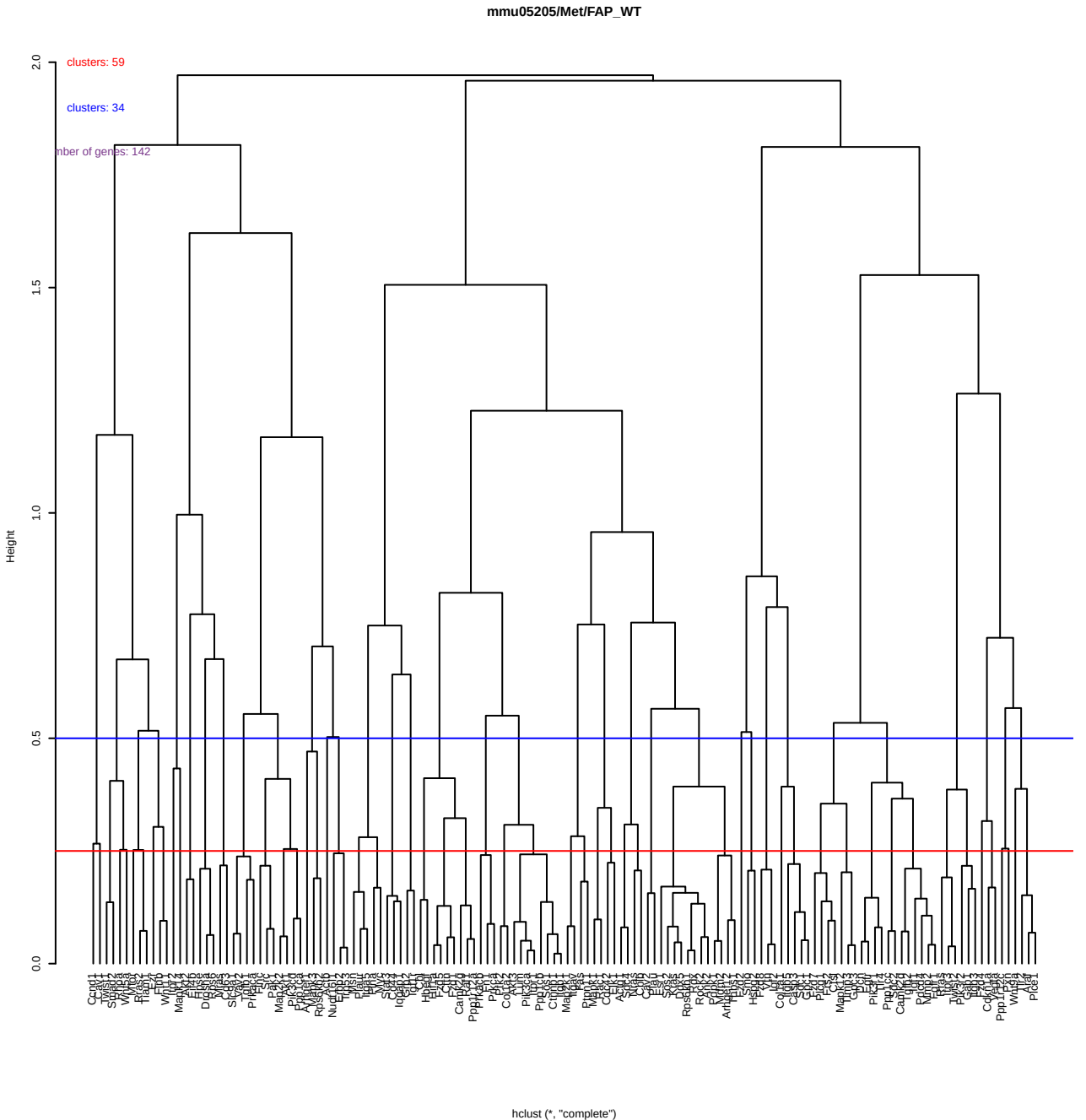

mmu05202/Met/EC_KO
clusters: 33
clusters: 19
mber of genes: 90
Height
0.0
0.5
1.0
1.5
2.0
Tcf3
Men1
Rara
Rxra
Pdgfa
Myc
Cdkn1a
Bcl6
Zbtb17
Bcl2l1
Ddit3
Plat
Nupr1
Cd14
Il1r2
Cdkn2c
Cd40
Prcc
Phx1
Etv1
Cik9
Aspscr1
Id2
Cebpb
Max
Cebpa
Tle3
Sun13
Cdkn1b
Lyl1
Foxo1
Sin3a
Tgfbr2
Ldb1
Pparg
Klf3
Elk4
Tef15
Ewsr1
Dusp6
Etv6
Fus
Ccnt1
Flt1
Ddx5
Ncor1
Erg
Zfat1
Mef2c
Spi1
Ccnt2
Ptk2
Fut8
Fli1
Nfkb1
Smad1
Igfbp3
Tfrc3
Pm
Mll1
Rxrb
Jup
Rela
Igf1r
Prom1
Ccnd2
Cxcr2
Nfkb2
Mdm2
Birc3
Dot1l
Aff1
Jmjd1c
Tspan7
Il6
Bmi1
Gbp3
Cdk14
Golph3l
H3f3a
Gm12657
Hdac2
Plau
Atf1
Mll1
Lmo2
Hhex
Whsc1
Hdac1
Traf1
hclust (*, "complete")

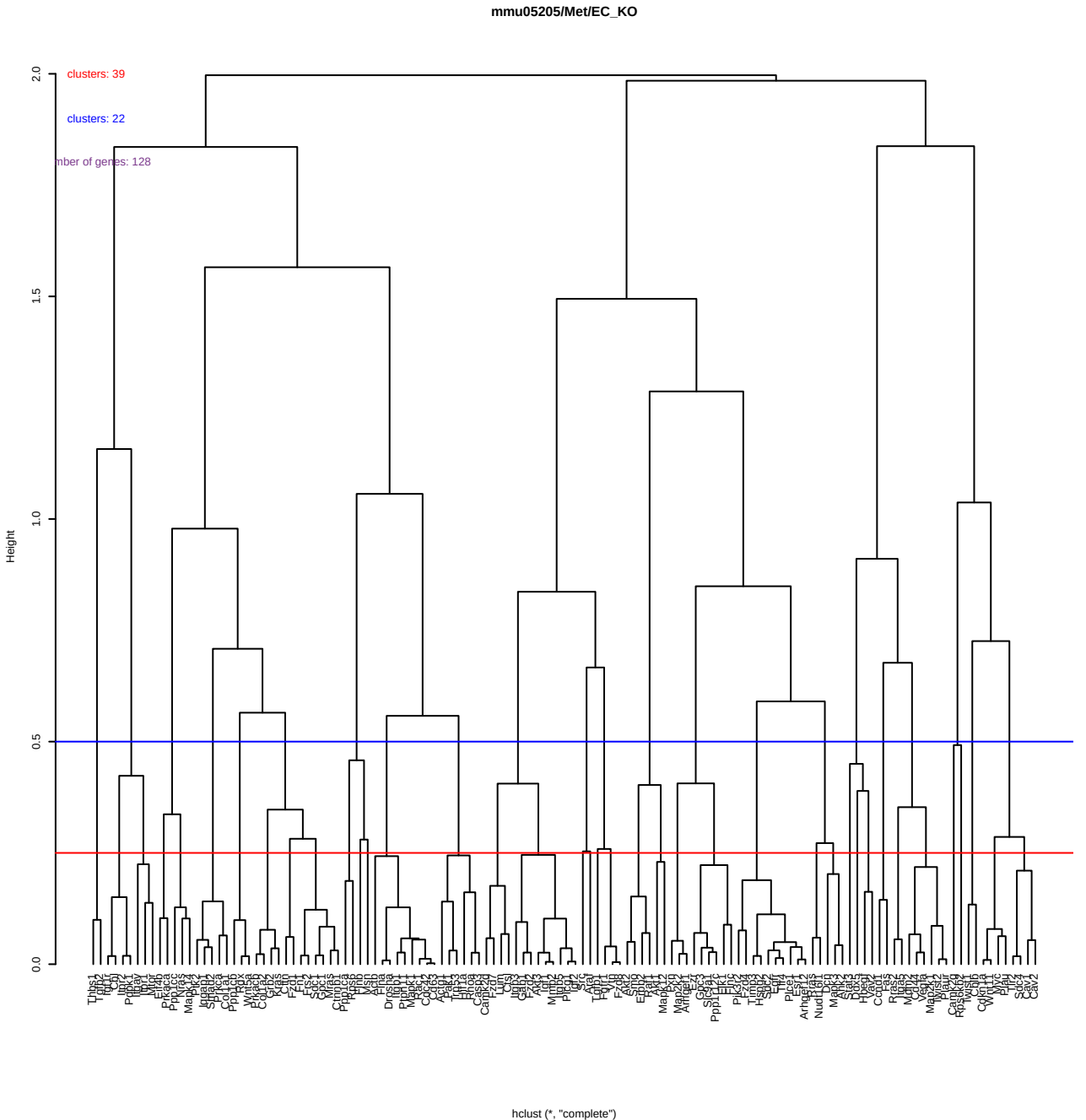

mmu05202/Met/FAP_WT
clusters: 41
clusters: 24
mber of genes: 93
Height
2.0
1.5
1.0
0.5
0.0
Supt3
Mllt1
Men1
Hdac1
Cdkn2c
Etv6
Runx1
Runx2
Pbx2
Hpgd
Etv4
Bcl2l1
Cebpb
Cd14
Tfe3
Rela
Cebpa
Trp53
Pml
Rxra
Tcf3
Zfp217
Rara
Asgscr1
Plau
Fus
Nfkbiz
Taf15
Il6
Bmp2k
Nfkb1
Mdm2
Ncor1
Jmjd1c
Gm12657
Bmi1
Atf1
Atf3
Dbx1
Sp1
Fut8
Golph3
Pbx1
Plat
Cdk14
Fgfr1a
Rxrb
Mmp3
Ccnt1
Aff1
Igf1r
Ewsr1
Myc
Dnmt1
Whsc1
Birc3
Cdkn1a
Ccnd2
Ddit3
Per2
Kdm3
Igfbp3
Nupr1
Pik2
Hdac2
H3f3b
Id2
Hoxa11
Dusp2
Ccnf2
Bcl6
Maf
Lmo2
Hoxa10
Cdkn1b
Ezh2
Tgfbr1
Igf1
Zeb1
Elk4
Sin3a
Smad1
Tspan7
Fmod
Six1
Il1r2
H3f3a
Golph3l
Prcc
Etv5
Max
Jup
Cdk9
Ldb1
hclust (*, "complete")

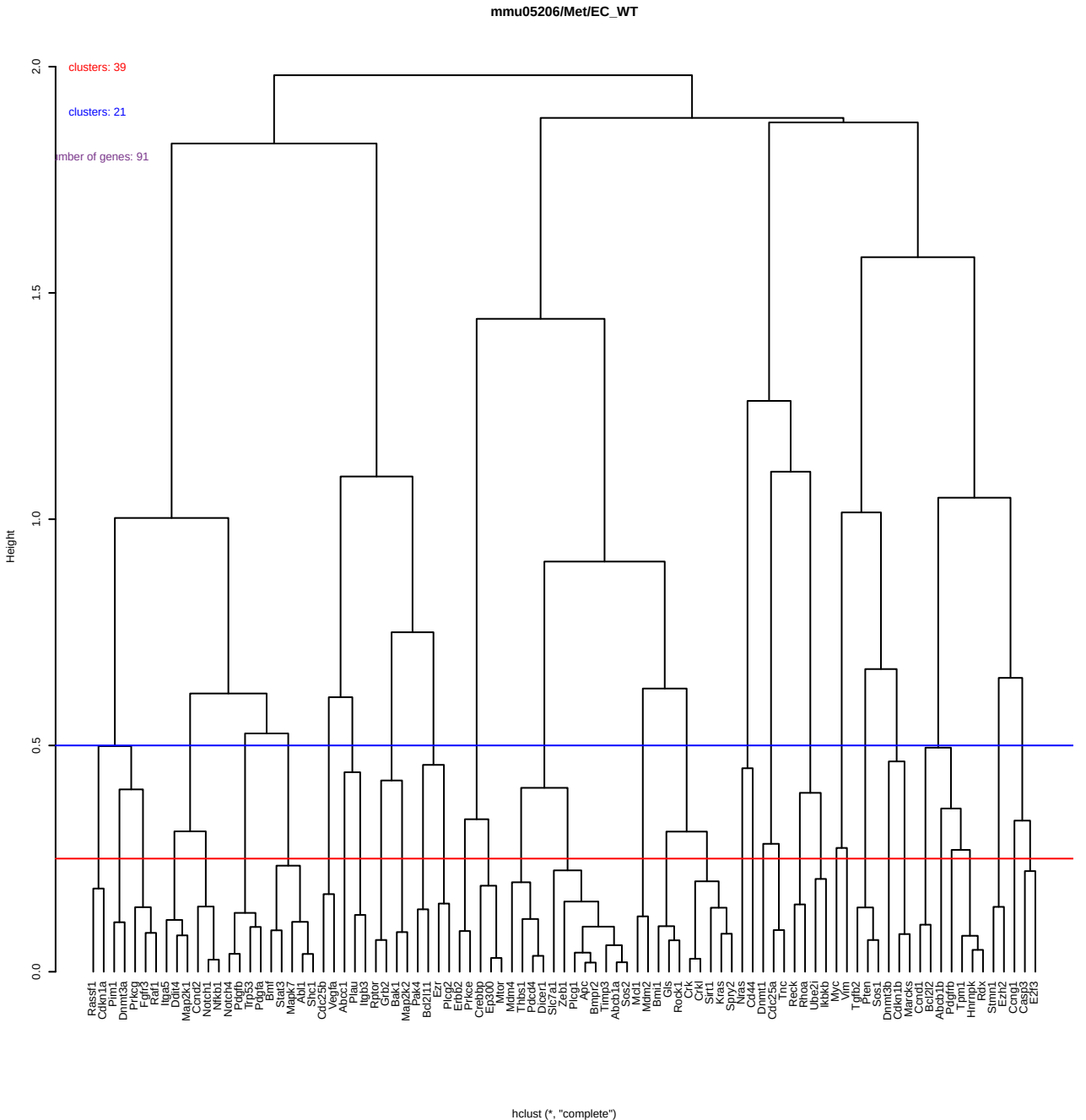

mmu05202/Met/EC_KO
clusters: 31
clusters: 17
mber of genes: 87
Height
2.0
1.5
1.0
0.5
0.0
Cdk9
Ldb1
Rxra
Rxrb
Men1
Six1
Dusp6
Sin3a
Cdkn1b
Cdk14
Igf1
Ncor1
Prox1
Sp1
Fut8
Zeb1
Tgfbr2
Igfbp3
Mmp3
Golph3l
Igf1r
Foxo1
Hoxa10
Klf3
Elk4
Plat
Max
Jup
Ccnd2
Pik2
Runx2
Golph3
Runx1
Ccnt2
Mllt1
Proc
Tle3
H3f3a
Tspan7
Cdkn2c
Hoxa1
Etv1
Bmi1
Pbx3
Whsc1
Ddx5
Atf1
Hdac1
Cd14
Ewsr1
H3f3b
Nfkbiz
Fus
Hdac2
Etv4
Trp53
Pdgfa
Etv6
Taf15
Zbtb17
Cebpa
Ddit3
Rela
Atf1
Id2
Bcl2l1
Tcf3
Aspscr1
Lmo1
Smad1
Nupr1
Pml
Rara
Gm12657
Cebpb
Il1r2
Birc3
Myc
Bdf6
Dot1l
Cdkn1a
Plau
Nfkb1
Mdm2
Supt3
Nr4a3
Il6
hclust (*, "complete")

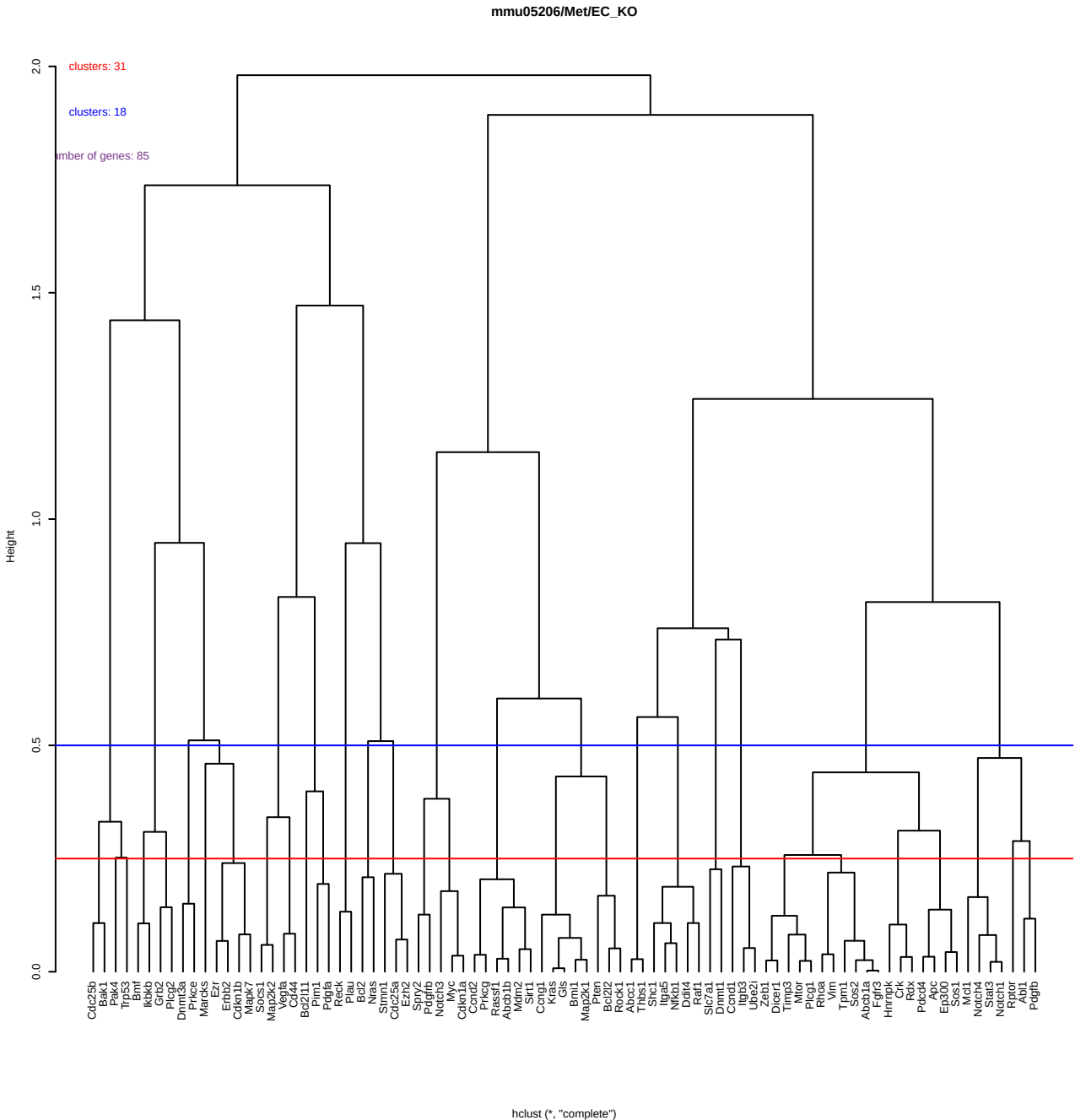

mmu05205/Met/EC_WT
clusters: 49
clusters: 27
nber of genes: 137
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

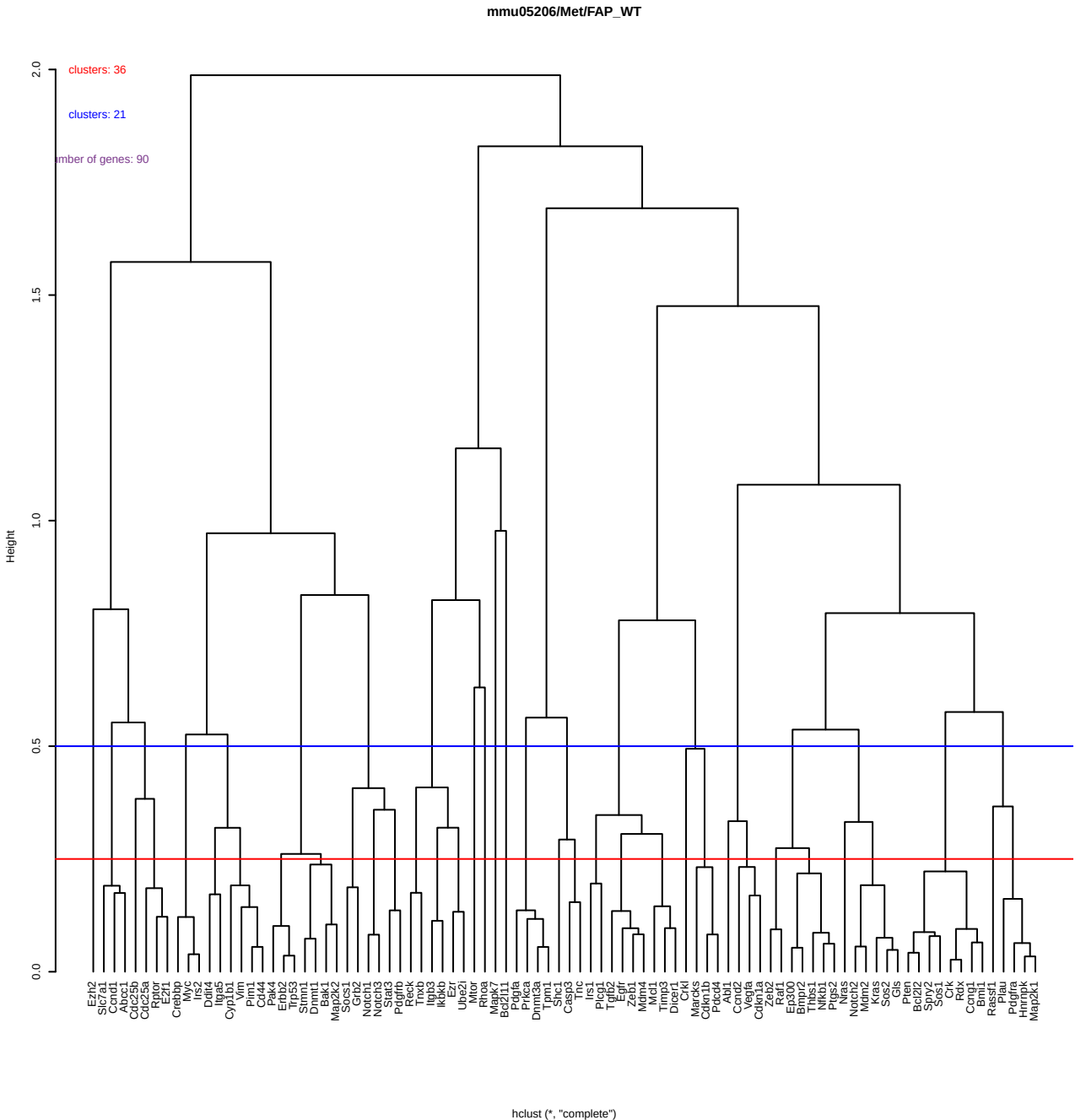

mmu05205/Met/EC_KO
clusters: 34
clusters: 21
mber of genes: 125
Height
0.0
0.5
1.0
1.5
2.0
hclust (*, "complete")

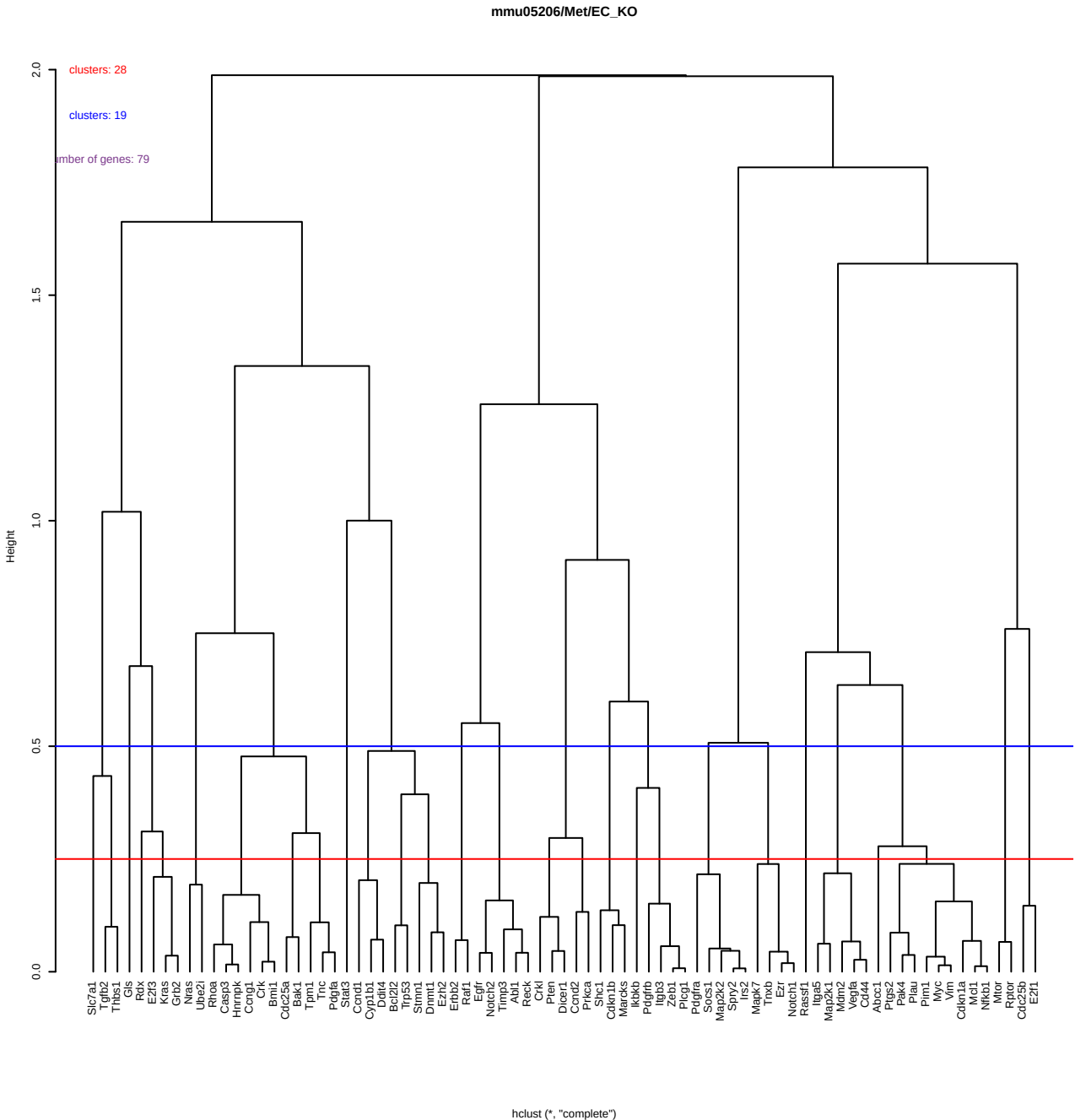

mmu05205/Met/FAP_WT
clusters: 59
clusters: 34
mber of genes: 142
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

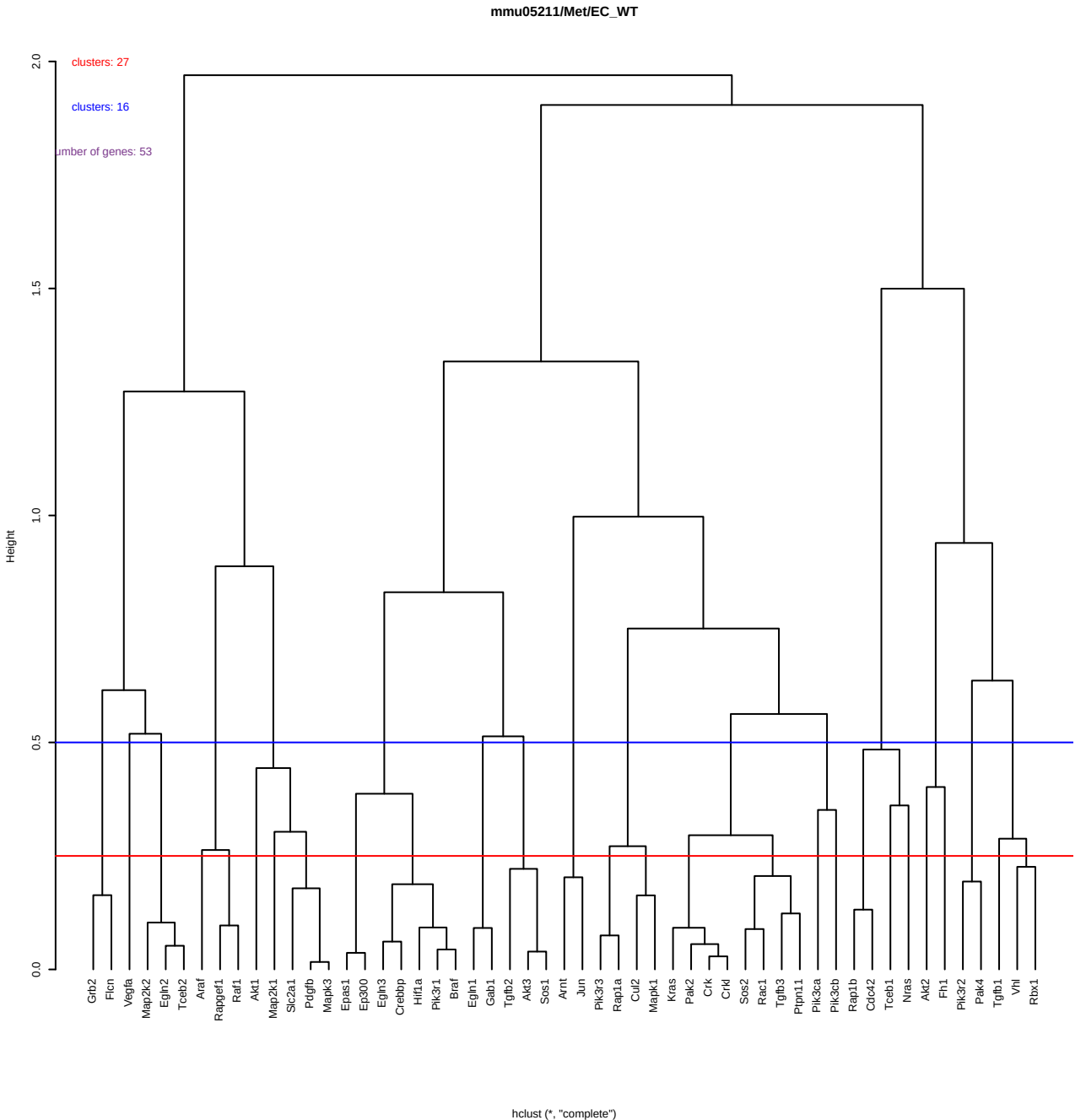

mmu05205/Met/EC_KO
clusters: 39
clusters: 22
nber of genes: 128
Height
2.0
1.5
1.0
0.5
0.0
hclust (*, "complete")

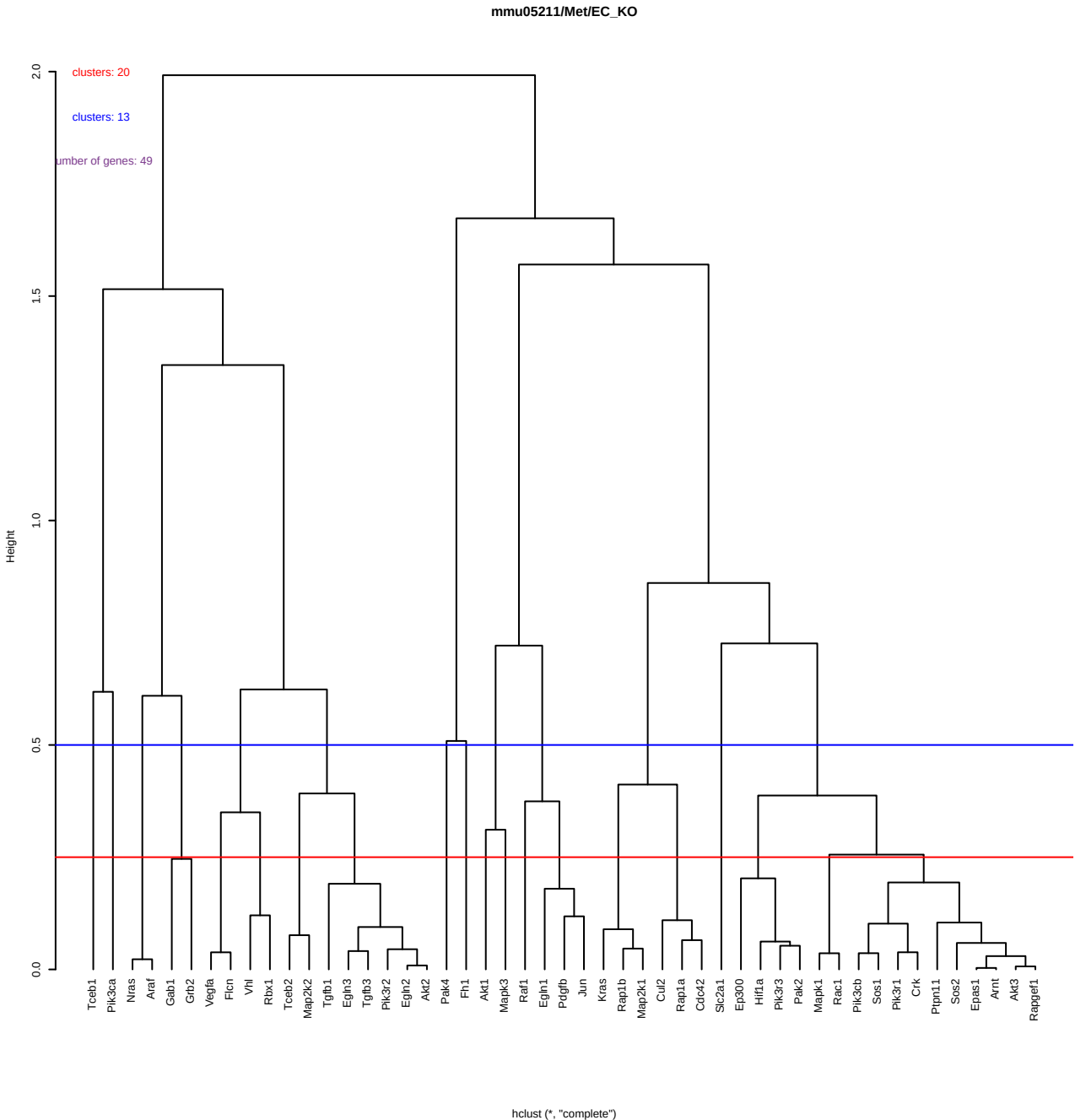

mmu05206/Met/EC_WT
clusters: 39
clusters: 21
mber of genes: 91
Height
2.0
1.5
1.0
0.5
0.0
Rassf1
Cdkn1a
Pim1
Dnmt3a
Prkcg
Fgfr3
Raf1
Itga5
Dpp4
Map2k1
Cond2
Notch1
Nfkb1
Notch4
Pdgfb
Trp53
Pdgfa
Bmp7
Stat3
Mapk7
Abl1
Shc1
Cdc25b
Vegfa
Abcc1
Rhau
Irs3
Rptor
Grb2
Bak1
Map2k2
Pak4
Bcl2l11
Ezr
Plcg2
Erbb2
Prkce
Crebbp
Ep300
Mtor
Mdm4
Tlbs1
Pdcd4
Dgcr2
Slc7a1
Zeb1
Plcg1
Apc
Bmpr2
Timp3
Abcb1a
Sos2
Mcl1
Mdm2
Bmi1
Gls
Rock1
Crk
Crkl
Sirt1
Kras
Smad2
Nras
Cd44
Dnmt1
Cdc25a
Tnc
Reck
Rhoa
Ube2i
Ikbkb
Myc
Vim
Tgfb2
Pten
Sos1
Dnmt3b
Cdkn1b
Mapk3
Mapk1
Bcl2l2
Abcb1b
Pdgfrb
Tpm1
Hnrnpk
Rbx
Stmn1
Ezh2
Ccnd1
Casp3
E2f3
hclust (*, "complete")

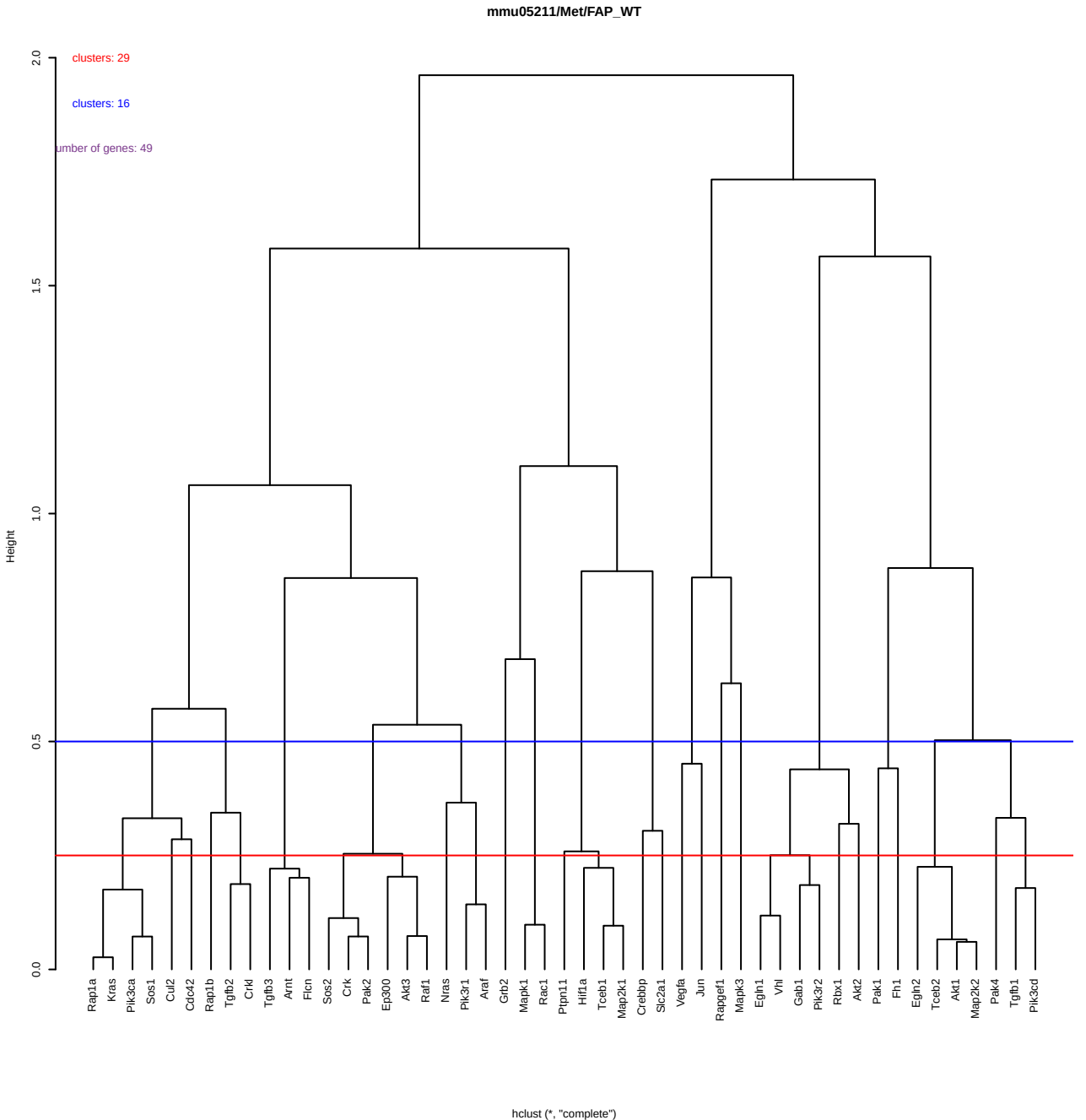

mmu05206/Met/EC_KO
clusters: 31
clusters: 18
mber of genes: 85
Height
2.0
1.5
1.0
0.5
0.0
Cdc25b
Bak1
Pak4
Trp53
Bmf
Ikbkb
Grb2
Plcg2
Dnmt3a
Prkce
Marcks
Ezr
Erbb2
Cdkn1b
Mapk7
Socs1
Map2k2
Vegfa
Cdk4
Bcl2l11
Pim1
Pdgfa
Reck
Plau
Bcl2
Nras
Stmn1
Cdc25a
Ezh2
Spry2
Pdgfrb
Notch3
Myc
Cdkn1a
Cond2
Prkcg
Rass1
Atgn1b
Mdm2
Sirt1
Ccng1
Kras
Gls
Bmi1
Map2k1
Pten
Bcl2l2
Rock1
Abcc1
Thbs1
Shc1
Itga5
Nfkb1
Ddit4
Ral1
Slc7a1
Dnmt1
Ccnd1
Itgb3
Ube2i
Dicer1
Zeb1
Timp3
Mtor
Plcg1
Rhoa
Vim
Tpm1
Sos2
Abcb1a
Fgfr3
Hnrnpk
Crk
Rdx
Pdcd4
Apc
Ep300
Sos1
Mcl1
Notch4
Stat3
Notch1
Rptor
Abl1
Pdgfb
hclust (*, "complete")

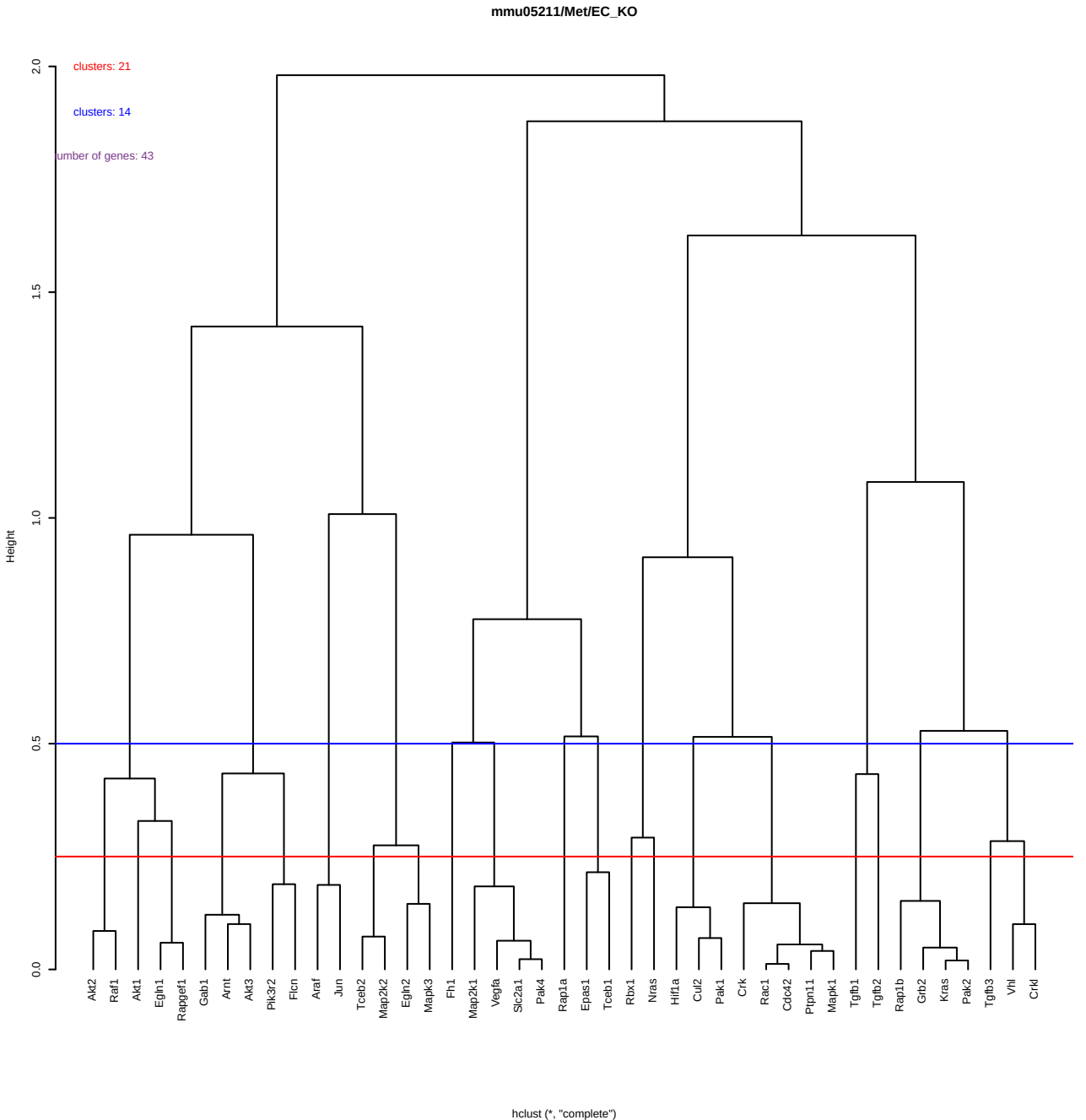

mmu05206/Met/FAP_WT
clusters: 36
clusters: 21
mber of genes: 90
Height
2.0
1.5
1.0
0.5
0.0
Ezh2
Slc7a1
Cnd1
Abcc1
Cdc25b
Cdc25a
Rpior
E2f1
Creb3l2
Myc
Irs2
Ddit4
Itga5
Cyp1b1
Vim
Pim1
Cd44
Pak4
Erbb2
Tlnrb3
Stmn1
Dnm1
Bak1
Map2k2
Socs1
Grb2
Notch1
Notch3
Stat3
Pdgfrb
Reck
Tnxb
Itgb3
Ikbkb
Ezr
Ube2i
Mtor
Rhoa
Map4k7
Bcl2l11
Pdgfa
Prkca
Dnmt3a
Tpm1
Shc1
Casp3
Tnc
Irs1
Plcg1
Tgfb2
Egfr
Zeb1
Mdm4
Mcl1
Timp3
Dicer1
Crkl
Mapk1
Cdkn1b
Pdcd4
Abl1
Cnd2
Vegfa
Cdkn1a
Zeb2
Rad1
Epsin2
Ehmt2
Tnfrsf1
Nfkb1
Ptgs2
Nras
Notch2
Mdm2
Kras
Sos2
Cfl1
Plek
Bcl2l2
Spry2
Sos1
Crk
Rdx
Cong1
Bmi1
Rassf1
Thbd
Pdgfra
Hnrnpk
Map2k1
hclust (*, "complete")

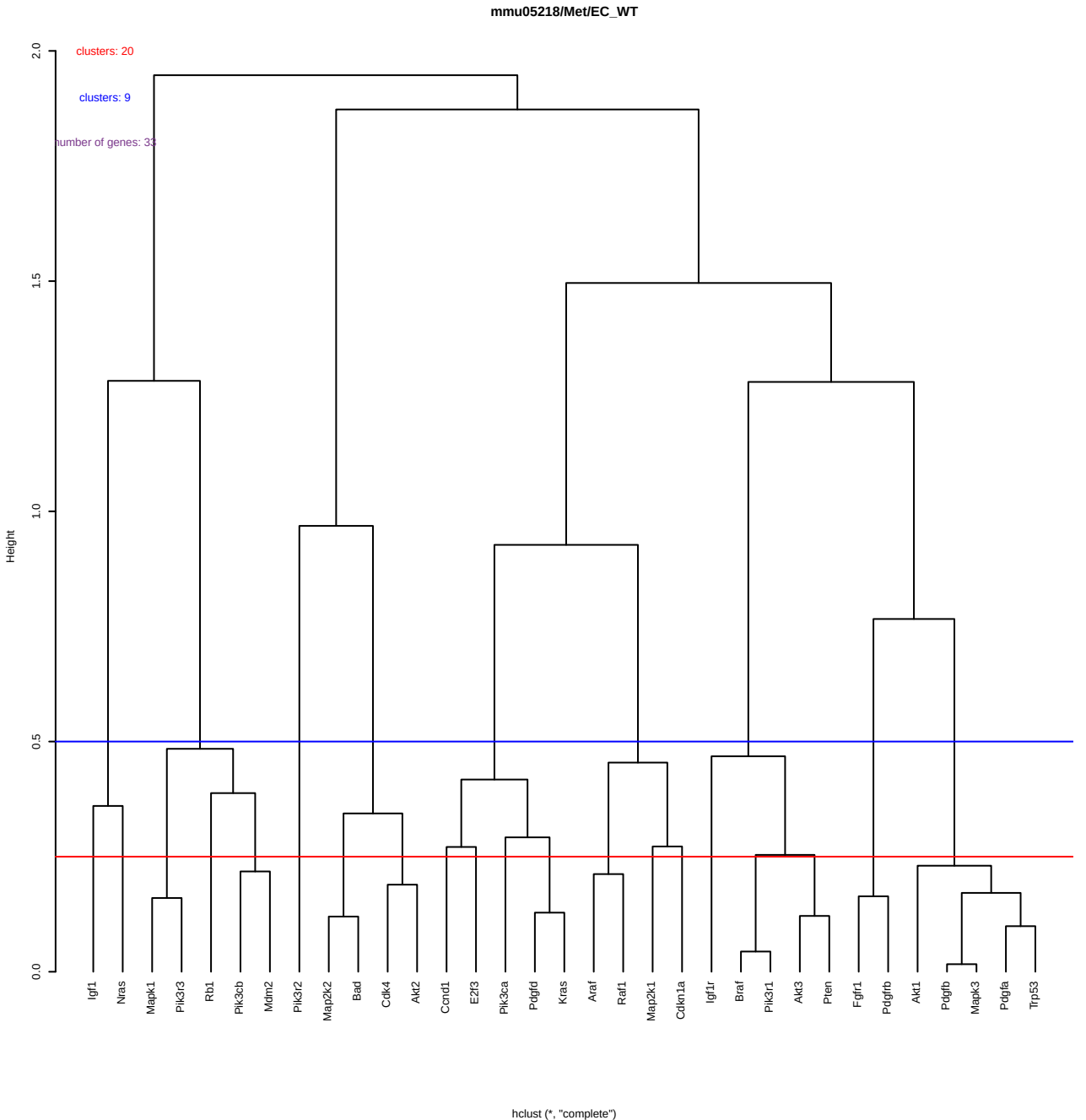

mmu05206/Met/EC_KO
clusters: 28
clusters: 19
mber of genes: 79
Height
2.0
1.5
1.0
0.5
0.0
Slc7a1
Tgfb2
Thbs1
Gls
Rdx
E2f3
Kras
Grb2
Nras
Ube2i
Rhoa
Casp3
Hnrnpk
Ccng1
Crk
Bmi1
Cdc25a
Bak1
Tpm1
Tnc
Pdgfa
Stat3
Ccnd1
Cyp1b1
Ddit4
Bcl2l2
Trp53
Stmn1
Dnmt1
Ezh2
Erbb2
Raf1
Egfr
Notch2
Timp3
Abl1
Reck
Crkl
Pten
Dicer1
Ccnd2
Prkca
Shc1
Cdkn1b
Marcks
Ikbkb
Pdgfrb
Itgb3
Zgpb1
Plcg1
Pdgfra
Socs1
Map2k2
Spry2
Irs2
Mapk7
Tnxb
Ezr
Notch1
Rassf1
Itga5
Map2k1
Mdm2
Vegfa
Cd44
Abcc1
Ptgs2
Pak4
Plau
Pim1
Myc
Vim
Cdkn1a
Mdl1
Nfkb1
Mtor
Rptor
Cdc25b
E2f1
hclust (*, "complete")

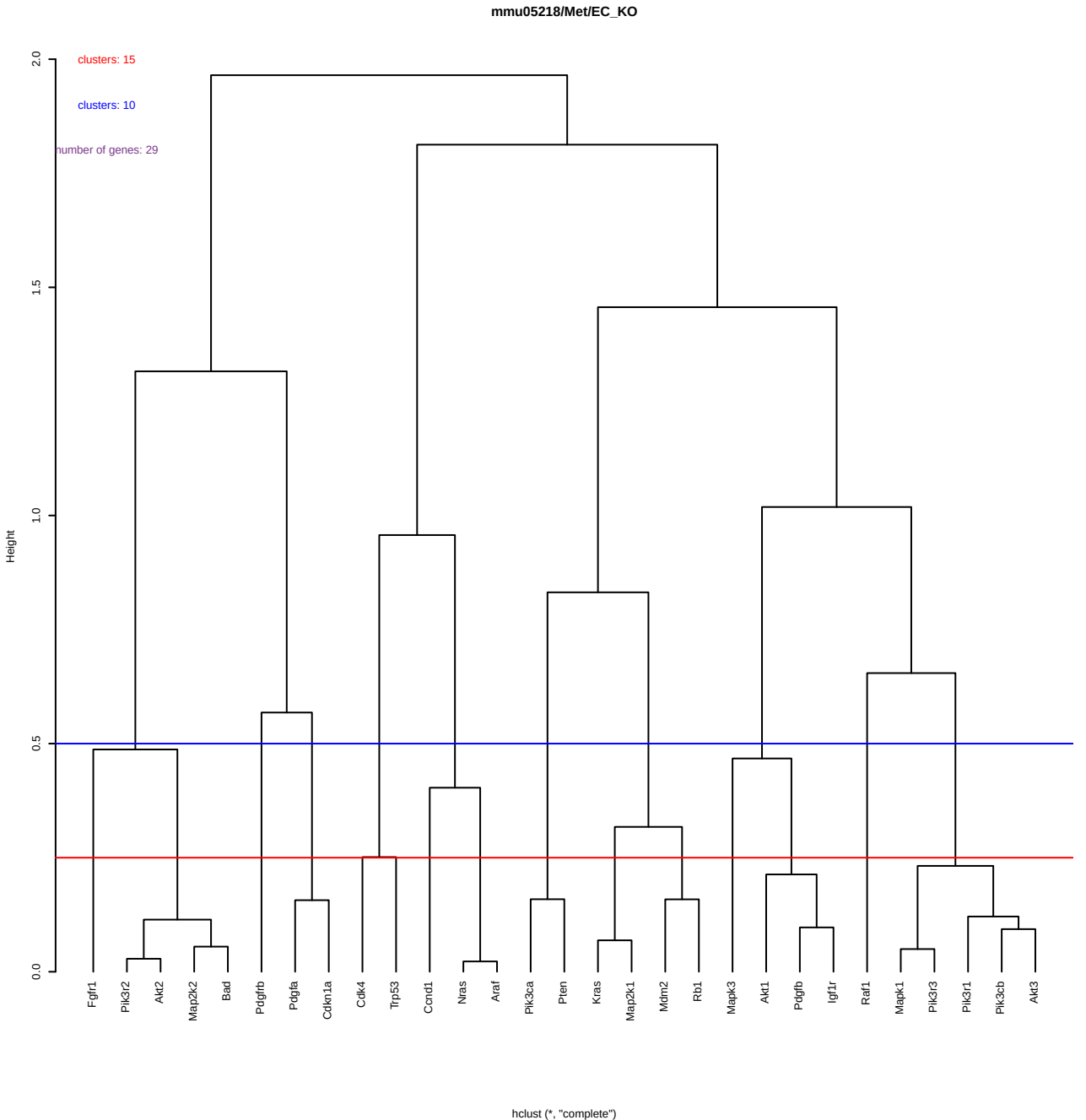

mmu05211/Met/EC_WT
clusters: 27
clusters: 16
umber of genes: 53
Height
2.0
1.5
1.0
0.5
0.0
Grb2
Flcn
Vegfa
Map2k2
Egln2
Tceb2
Araf
Rapgef1
Raf1
Akt1
Map2k1
Slc2a1
Pdgfb
Mapk3
Epas1
Ep300
Egln3
Crebbp
Hif1a
Pik3r1
Braf
Egln1
Gab1
Tgfb2
Akt3
Sos1
Arnt
Jun
Pik3r3
Rap1a
Cul2
Mapk1
Kras
Pak2
Crk
Crkl
Sos2
Rac1
Tgfb3
Ptpn11
Pik3ca
Pik3cb
Rap1b
Cdc42
Tceb1
Nras
Akt2
Fh1
Pik3r2
Pak4
Tgfb1
Vhl
Rbx1
hclust (*, "complete")

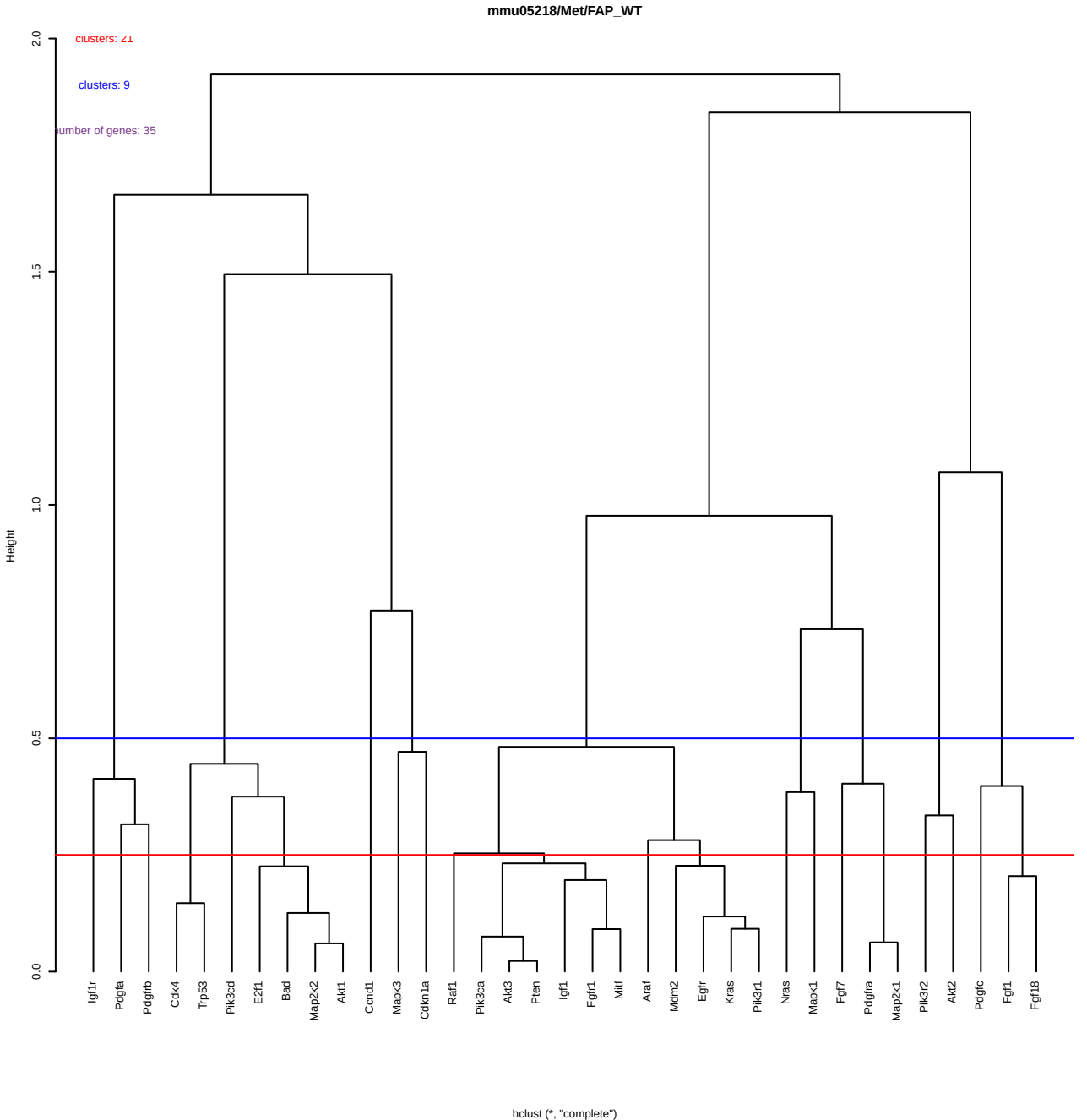

mmu05211/Met/EC_KO
clusters: 20
clusters: 13
umber of genes: 49
Height
2.0
1.5
1.0
0.5
0.0
Tceb1
Pik3ca
Nras
Araf
Gab1
Grb2
Vegfa
Flcn
Vhl
Rbx1
Tceb2
Map2k2
Tgfb1
Egln3
Tgfb3
Pik3r2
Egln2
Akt2
Pak4
Fh1
Akt1
Mapk3
Raf1
Egln1
Pdgfb
Jun
Kras
Rap1b
Map2k1
Cul2
Rap1a
Cdc42
Slc2a1
Ep300
Hif1a
Pik3r3
Pak2
Mapk1
Rac1
Pik3cb
Sos1
Pik3r1
Crk
Ptpn11
Sos2
Epas1
Arnt
Akt3
Rapgef1
hclust (*, "complete")

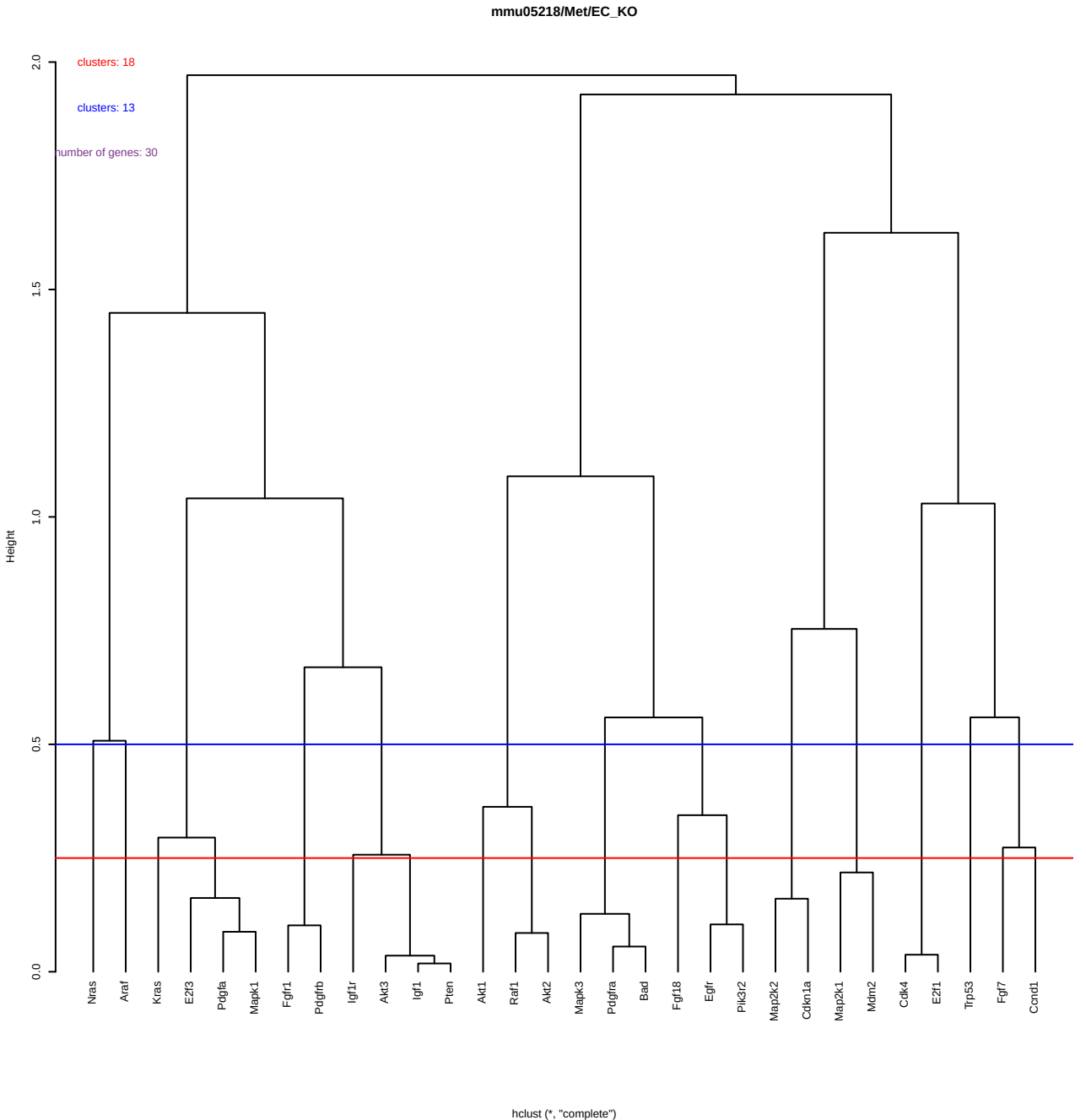

mmu05211/Met/FAP_WT
clusters: 29
clusters: 16
umber of genes: 49
Height
2.0
1.5
1.0
0.5
0.0
Rap1a
Kras
Pik3ca
Sos1
Cul2
Cdc42
Rap1b
Tgfb2
Crkl
Tgfb3
Arnt
Flcn
Sos2
Crk
Pak2
Ep300
Akt3
Raf1
Nras
Pik3r1
Araf
Grb2
Mapk1
Rac1
Ptpn11
Hif1a
Tceb1
Map2k1
Crebbp
Slc2a1
Vegfa
Jun
Rapgef1
Mapk3
Egln1
Vhl
Gab1
Pik3r2
Rbx1
Akt2
Pak1
Fh1
Egln2
Tceb2
Akt1
Map2k2
Pak4
Tgfb1
Pik3cd
hclust (*, "complete")

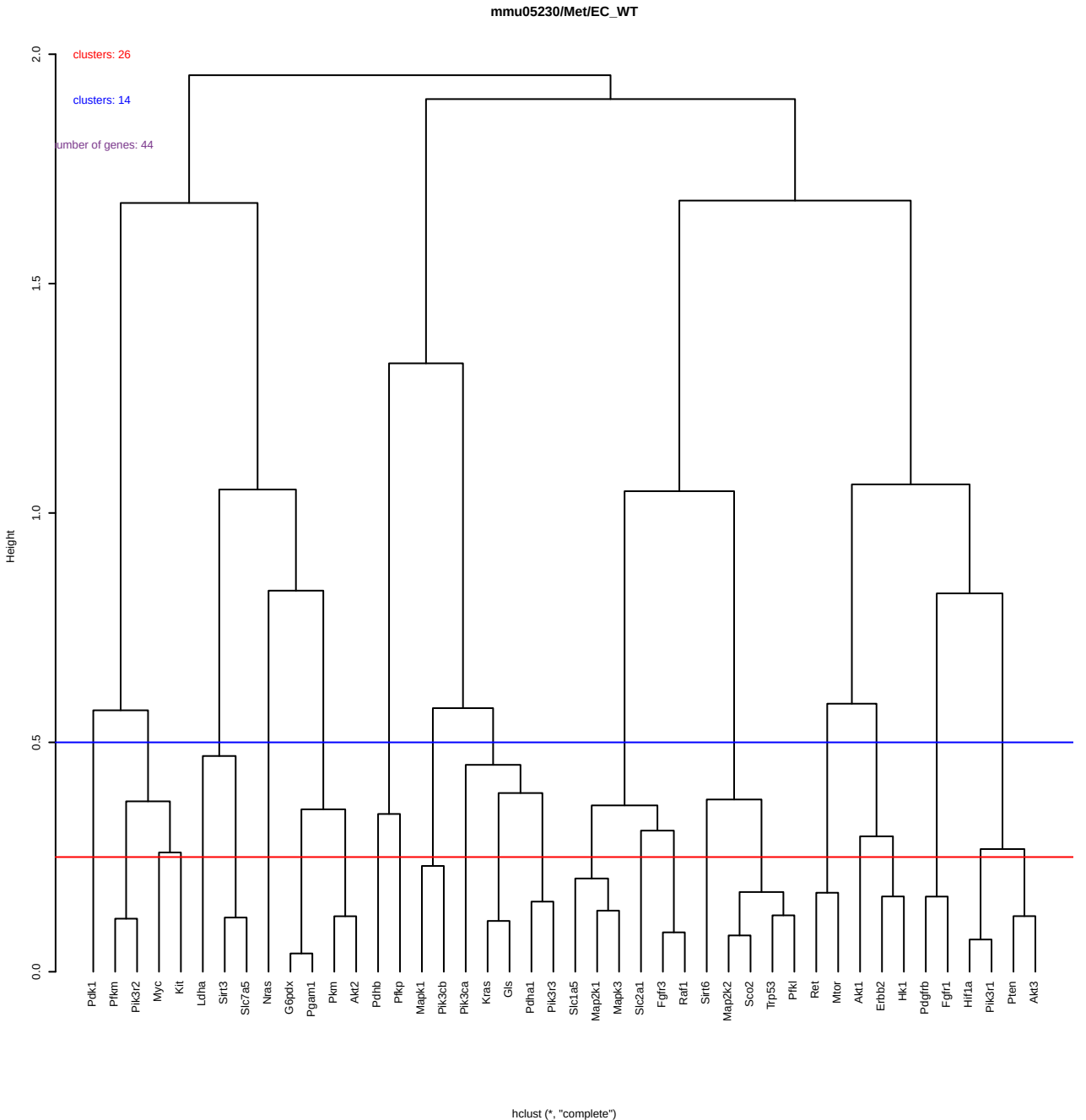

mmu05211/Met/EC_KO
clusters: 21
clusters: 14
umber of genes: 43
Height
0.0
0.5
1.0
1.5
2.0
Akt2
Raf1
Akt1
Egln1
Rapgef1
Gab1
Arnt
Akt3
Pik3r2
Flcn
Araf
Jun
Tceb2
Map2k2
Egln2
Mapk3
Fh1
Map2k1
Vegfa
Slc2a1
Pak4
Rap1a
Epas1
Tceb1
Rbx1
Nras
Hif1a
Cul2
Pak1
Crk
Rac1
Cdc42
Ptpn11
Mapk1
Tgfb1
Tgfb2
Rap1b
Grb2
Kras
Pak2
Tgfb3
Vhl
Crkl
hclust (*, "complete")

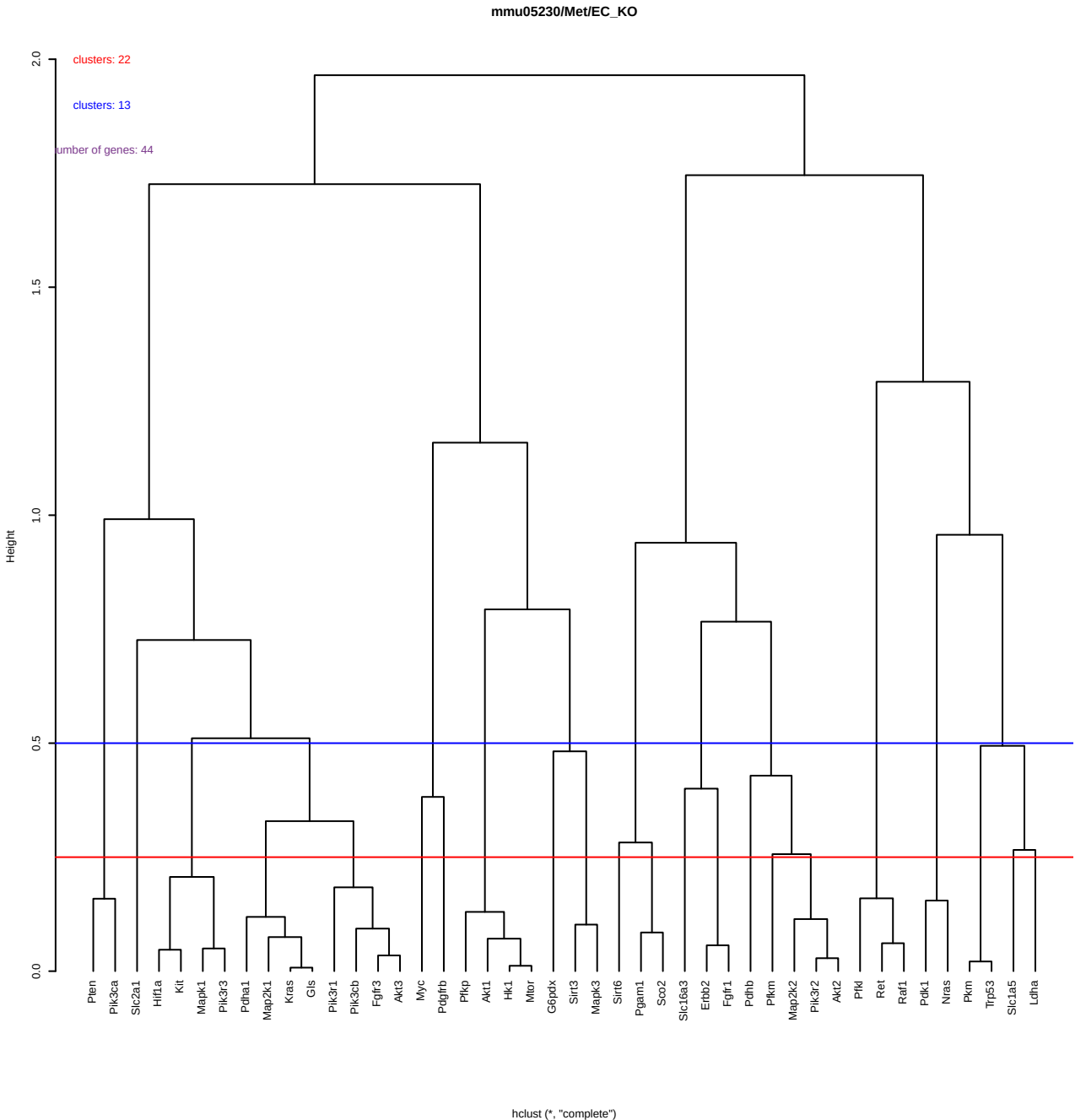

mmu05218/Met/EC_WT
clusters: 20
clusters: 9
number of genes: 33
Height
2.0
1.5
1.0
0.5
0.0
Igf1
Nras
Mapk1
Pik3r3
Rb1
Pik3cb
Mdm2
Pik3r2
Map2k2
Bad
Cdk4
Akt2
Ccnd1
E2f3
Pik3ca
Pdgfd
Kras
Araf
Raf1
Map2k1
Cdkn1a
Igf1r
Braf
Pik3r1
Akt3
Pten
Fgfr1
Pdgfrb
Akt1
Pdgfb
Mapk3
Pdgfa
Trp53
hclust (*, "complete")

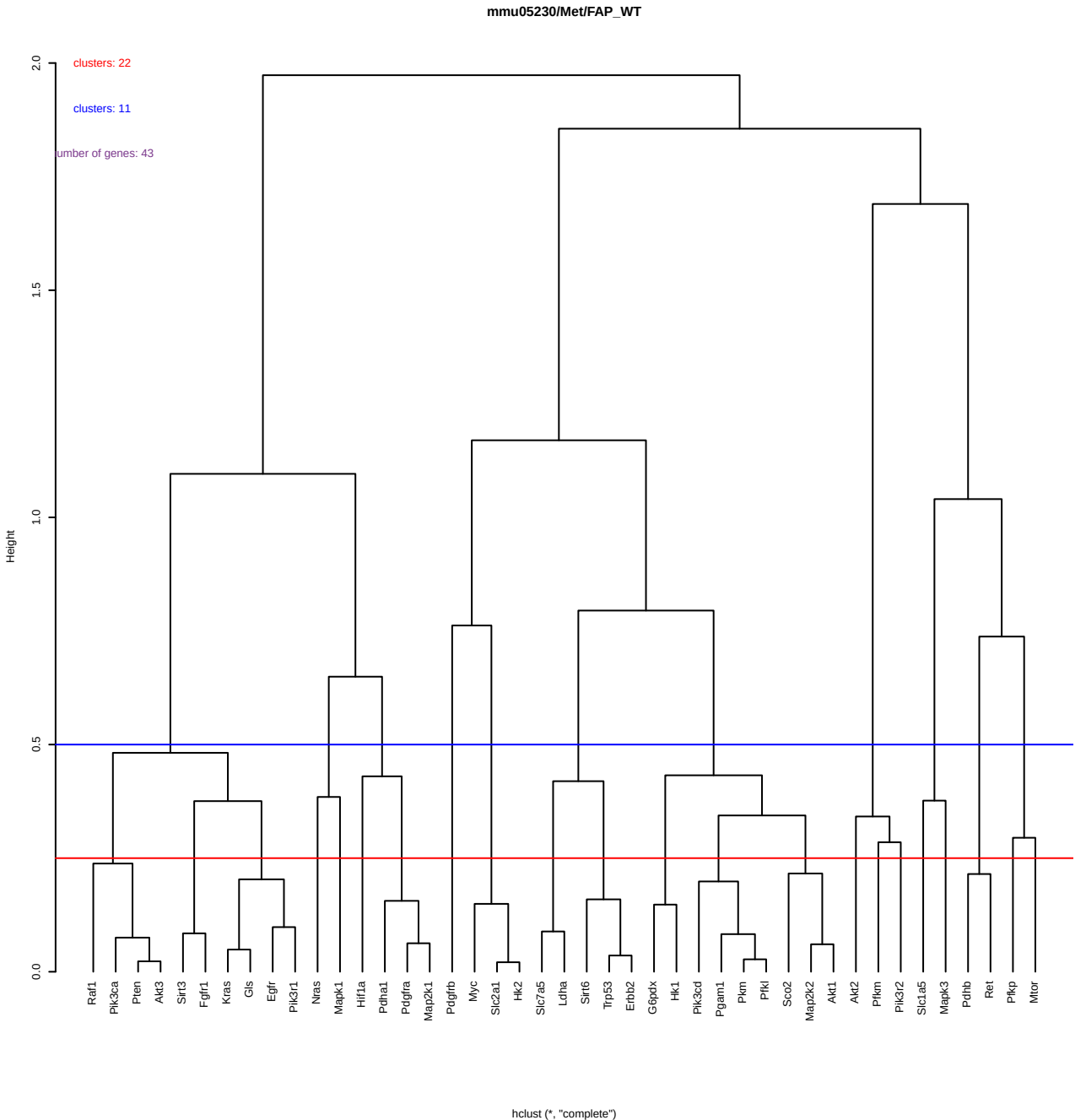

mmu05218/Met/EC_KO
clusters: 15
clusters: 10
number of genes: 29
Height
2.0
1.5
1.0
0.5
0.0
Fgfr1
Pik3r2
Akt2
Map2k2
Bad
Pdgfrb
Pdgfa
Cdkn1a
Cdk4
Trp53
Ccnd1
Nras
Araf
Pik3ca
Pten
Kras
Map2k1
Mdm2
Rb1
Mapk3
Akt1
Pdgfb
Igf1r
Raf1
Mapk1
Pik3r3
Pik3r1
Pik3cb
Akt3
hclust (*, "complete")

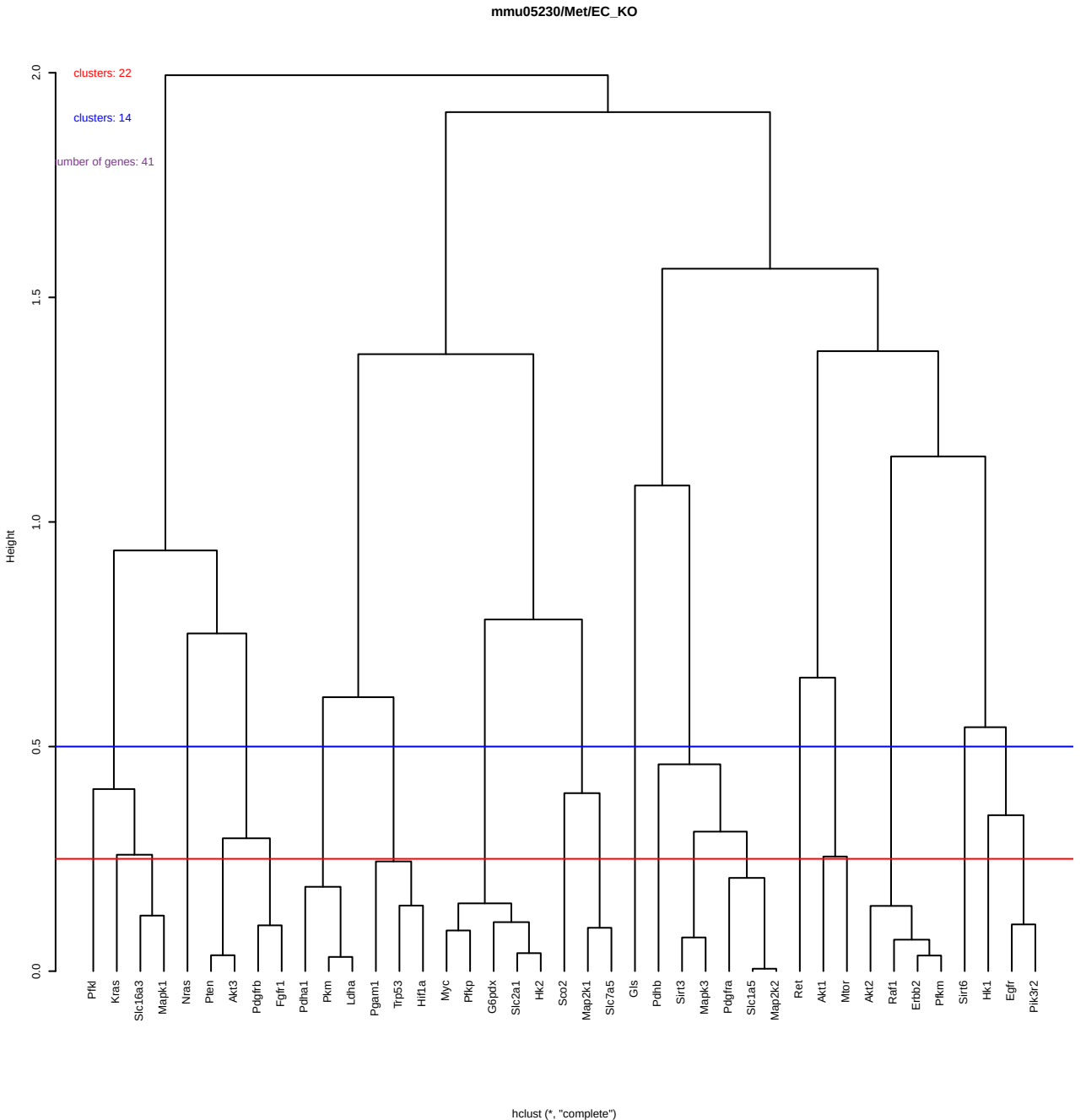

mmu05218/Met/FAP_WT
clusters: 21
clusters: 9
number of genes: 35
Height
0.0
0.5
1.0
1.5
2.0
Igf1r
Pdgfa
Pdgfrb
Cdk4
Trp53
Pik3cd
E2f1
Bad
Map2k2
Akt1
Ccnd1
Mapk3
Cdkn1a
Raf1
Pik3ca
Akt3
Pten
Igf1
Fgfr1
Mitf
Araf
Mdm2
Egfr
Kras
Pik3r1
Nras
Mapk1
Fgf7
Pdgfra
Map2k1
Pik3r2
Akt2
Pdgfc
Fgf1
Fgf18
hclust (*, "complete")


mmu05218/Met/EC_KO
clusters: 18
clusters: 13
number of genes: 30
Height
2.0
1.5
1.0
0.5
0.0
Nras
Araf
Kras
E2f3
Pdgfa
Mapk1
Fgfr1
Pdgfrb
Igf1r
Akt3
Igf1
Pten
Akt1
Raf1
Akt2
Mapk3
Pdgfra
Bad
Fgf18
Egfr
Pik3r2
Map2k2
Cdkn1a
Map2k1
Mdm2
Cdk4
E2f1
Trp53
Fgf7
Ccnd1
hclust (*, "complete")


mmu05230/Met/EC_WT
clusters: 26
clusters: 14
umber of genes: 44
Height
2.0
1.5
1.0
0.5
0.0
Pdk1
Pfkm
Pik3r2
Myc
Kit
Ldha
Sirt3
Slc7a5
Nras
G6pdx
Pgam1
Pkm
Akt2
Pdhb
Pfkp
Mapk1
Pik3cb
Pik3ca
Kras
Gls
Pdha1
Pik3r3
Slc1a5
Map2k1
Mapk3
Slc2a1
Fgfr3
Raf1
Sirt6
Map2k2
Sco2
Trp53
Pfkl
Ret
Mtor
Akt1
Erbb2
Hk1
Pdgfrb
Fgfr1
Hif1a
Pik3r1
Pten
Akt3
hclust (*, "complete")


mmu05230/Met/EC_KO
clusters: 22
clusters: 13
umber of genes: 44
Height
2.0
1.5
1.0
0.5
0.0
Pten
Pik3ca
Slc2a1
Hif1a
Kit
Mapk1
Pik3r3
Pdha1
Map2k1
Kras
Gls
Pik3r1
Pik3cb
Fgfr3
Akt3
Myc
Pdgfrb
Pfkp
Akt1
Hk1
Mtor
G6pdx
Sirt3
Mapk3
Sirt6
Pgam1
Sco2
Slc16a3
Erbb2
Fgfr1
Pdhb
Pfkm
Map2k2
Pik3r2
Akt2
Pfkl
Ret
Raf1
Pdk1
Nras
Pkm
Trp53
Slc1a5
Ldha
hclust (*, "complete")


mmu05230/Met/FAP_WT
clusters: 22
clusters: 11
umber of genes: 43
Height
2.0
1.5
1.0
0.5
0.0
Raf1
Pik3ca
Pten
Akt3
Sirt3
Fgfr1
Kras
Gls
Egfr
Pik3r1
Nras
Mapk1
Hif1a
Pdha1
Pdgfra
Map2k1
Pdgfrb
Myc
Slc2a1
Hk2
Slc7a5
Ldha
Sirt6
Trp53
Erbb2
G6pdx
Hk1
Pik3cd
Pgam1
Pkm
Pfkl
Sco2
Map2k2
Akt1
Akt2
Pfkm
Pik3r2
Slc1a5
Mapk3
Pdhb
Ret
Pfkp
Mtor
hclust (*, "complete")


mmu05230/Met/EC_KO
clusters: 22
clusters: 14
umber of genes: 41
Height
0.0
0.5
1.0
1.5
2.0
Pfkl
Kras
Slc16a3
Mapk1
Nras
Pten
Akt3
Pdgfrb
Fgfr1
Pdha1
Pkm
Ldha
Pgam1
Trp53
Hif1a
Myc
Pfkp
G6pdx
Slc2a1
Hk2
Sco2
Map2k1
Slc7a5
Gls
Pdhb
Sirt3
Mapk3
Pdgfra
Slc1a5
Map2k2
Ret
Akt1
Mtor
Akt2
Raf1
Erbb2
Pfkm
Sirt6
Hk1
Egfr
Pik3r2
hclust (*, "complete")