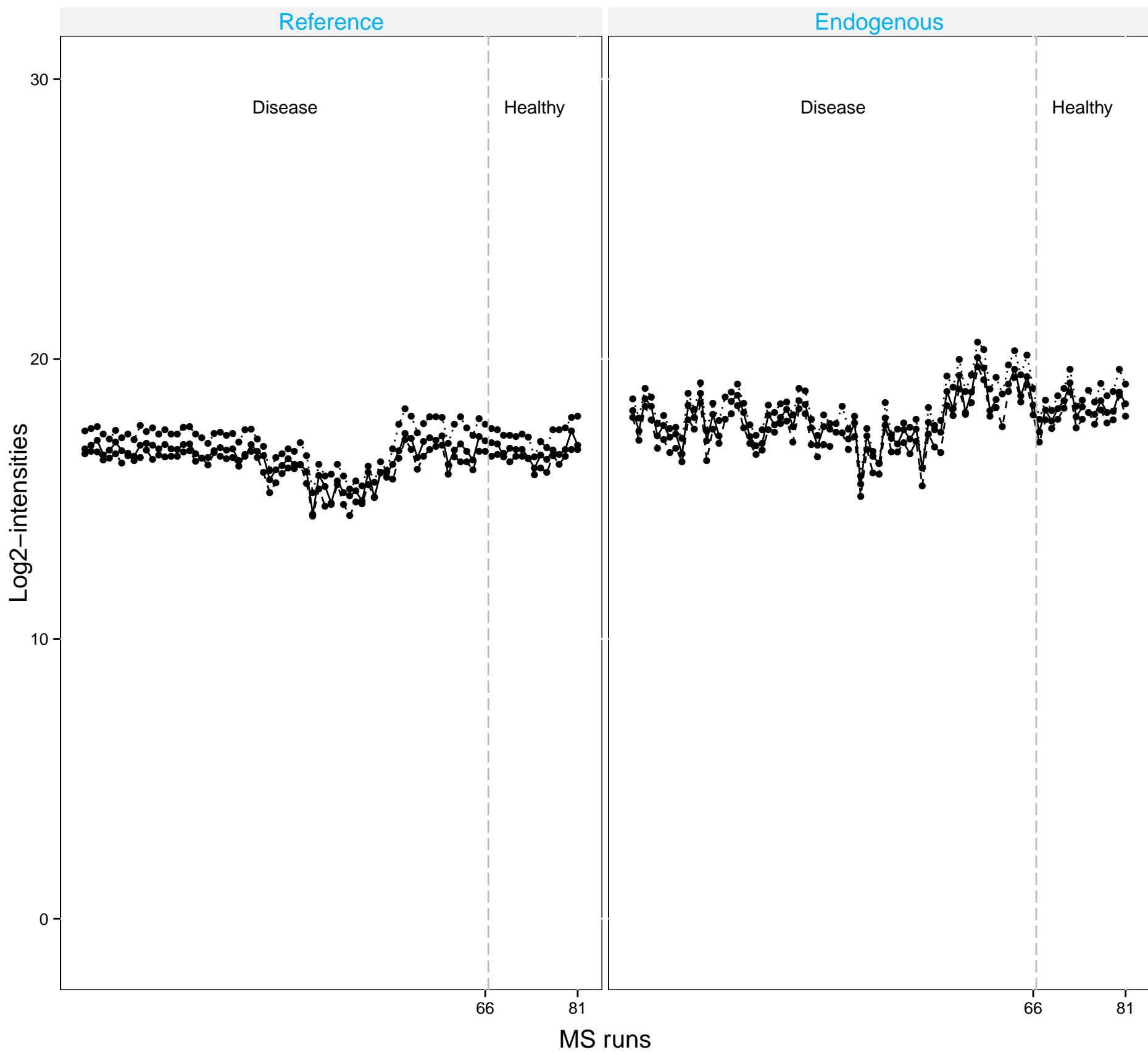


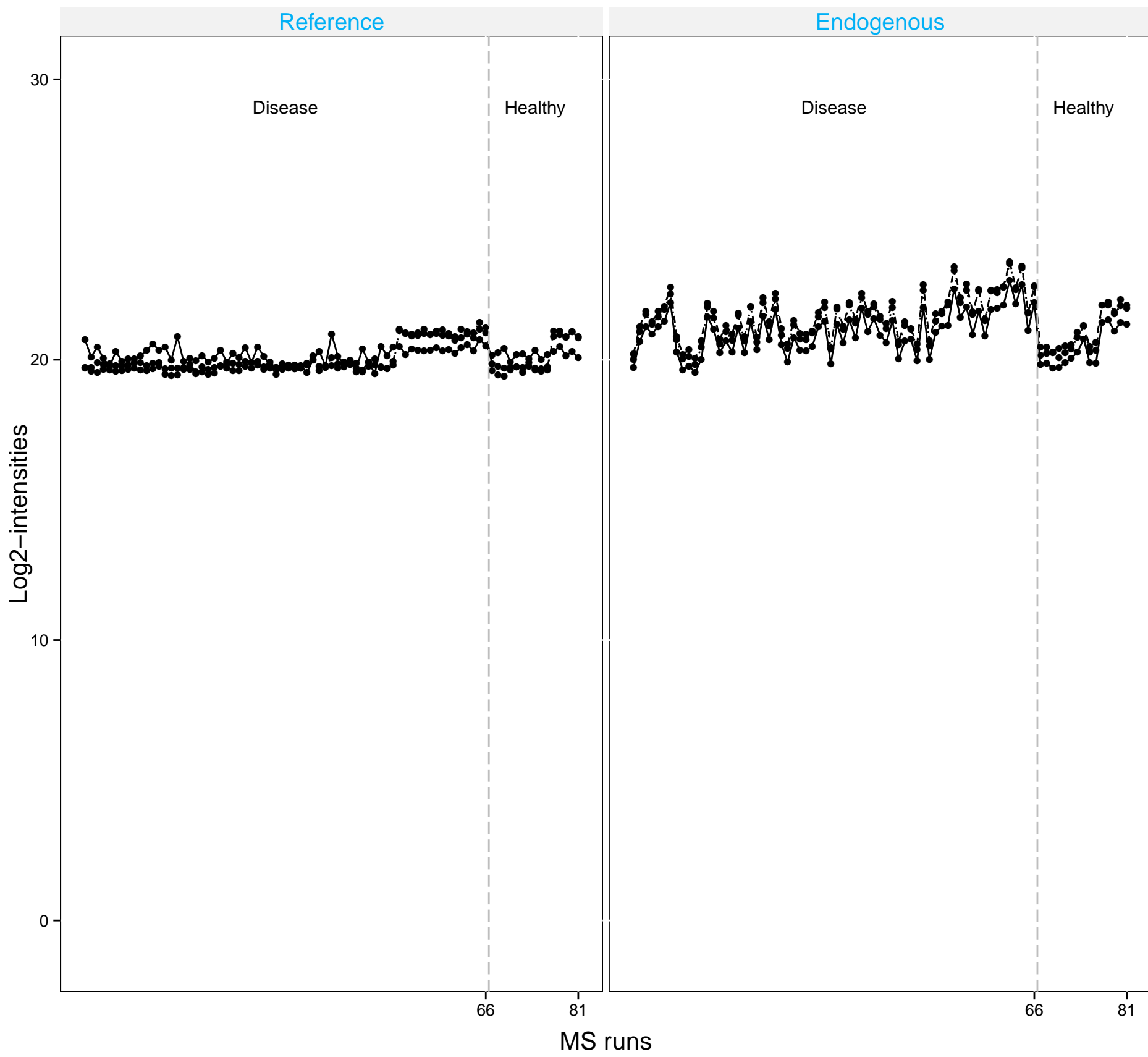
A2MG

peptide: 1 — SSSNEEVMLTVQVK_2_y5_1 — SSSNEEVMLTVQVK_2_y6_1 — SSSNEEVMLTVQVK_2_y8_1



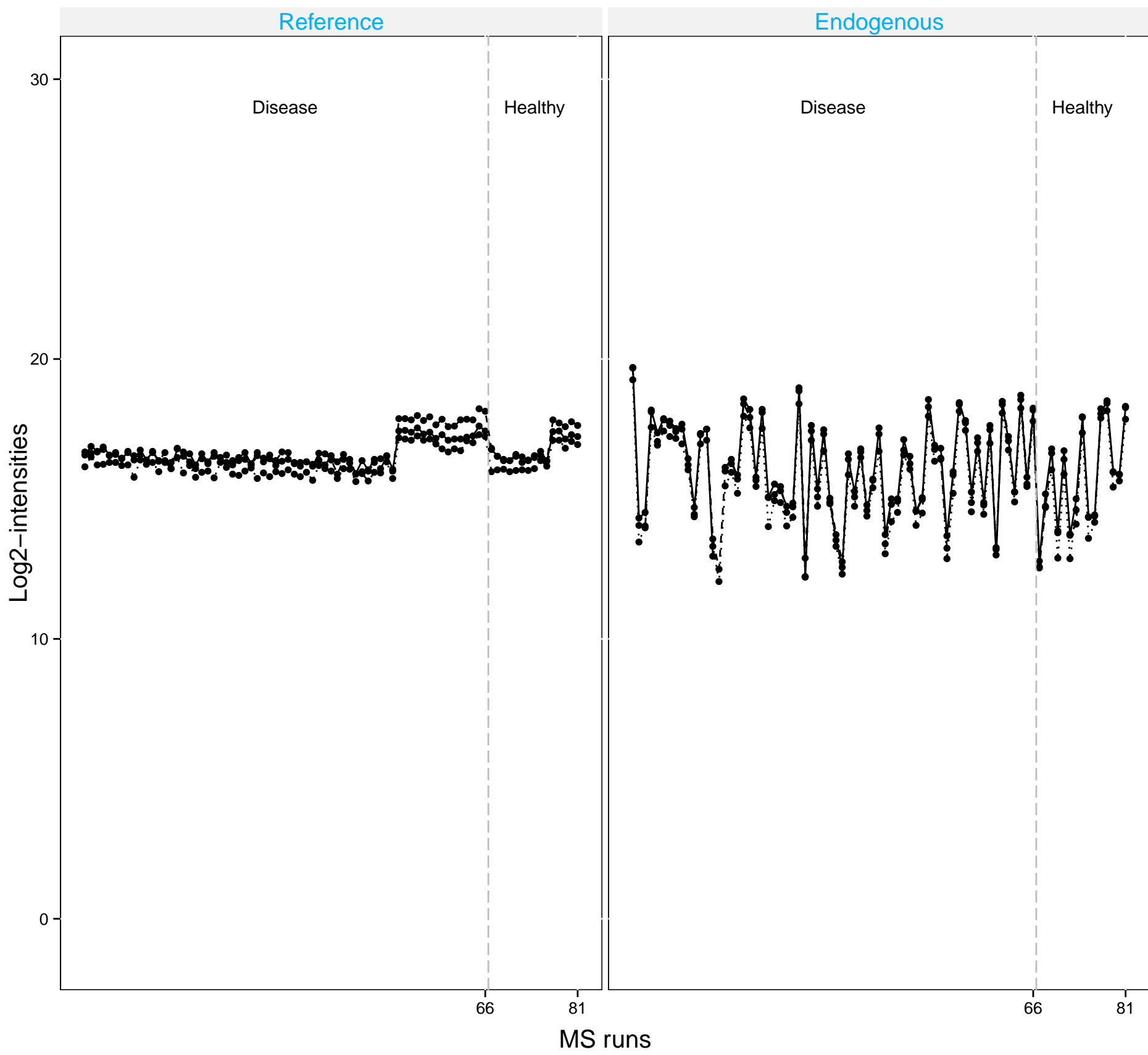
AACT

peptide: 1 — EIGELYLPK_2_y4_1 — EIGELYLPK_2_y5_1 — EIGELYLPK_2_y7_1



APOA

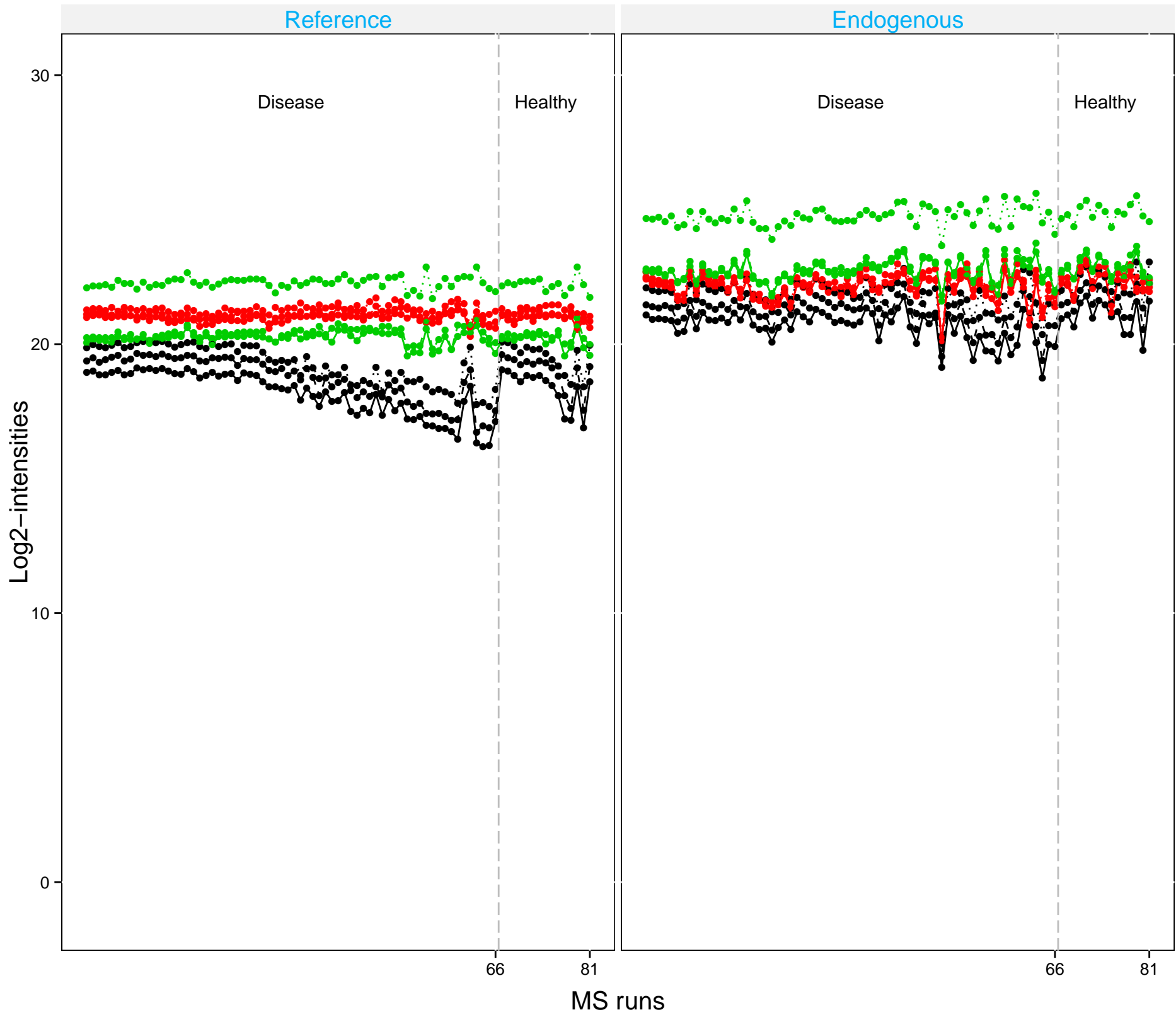
peptide: 1 —●— NPDAVAAPYCYTR_2_b4_1 —●— NPDAVAAPYCYTR_2_y6_1 —●— NPDAVAAPYCYTR_2_y7_1



APOA1

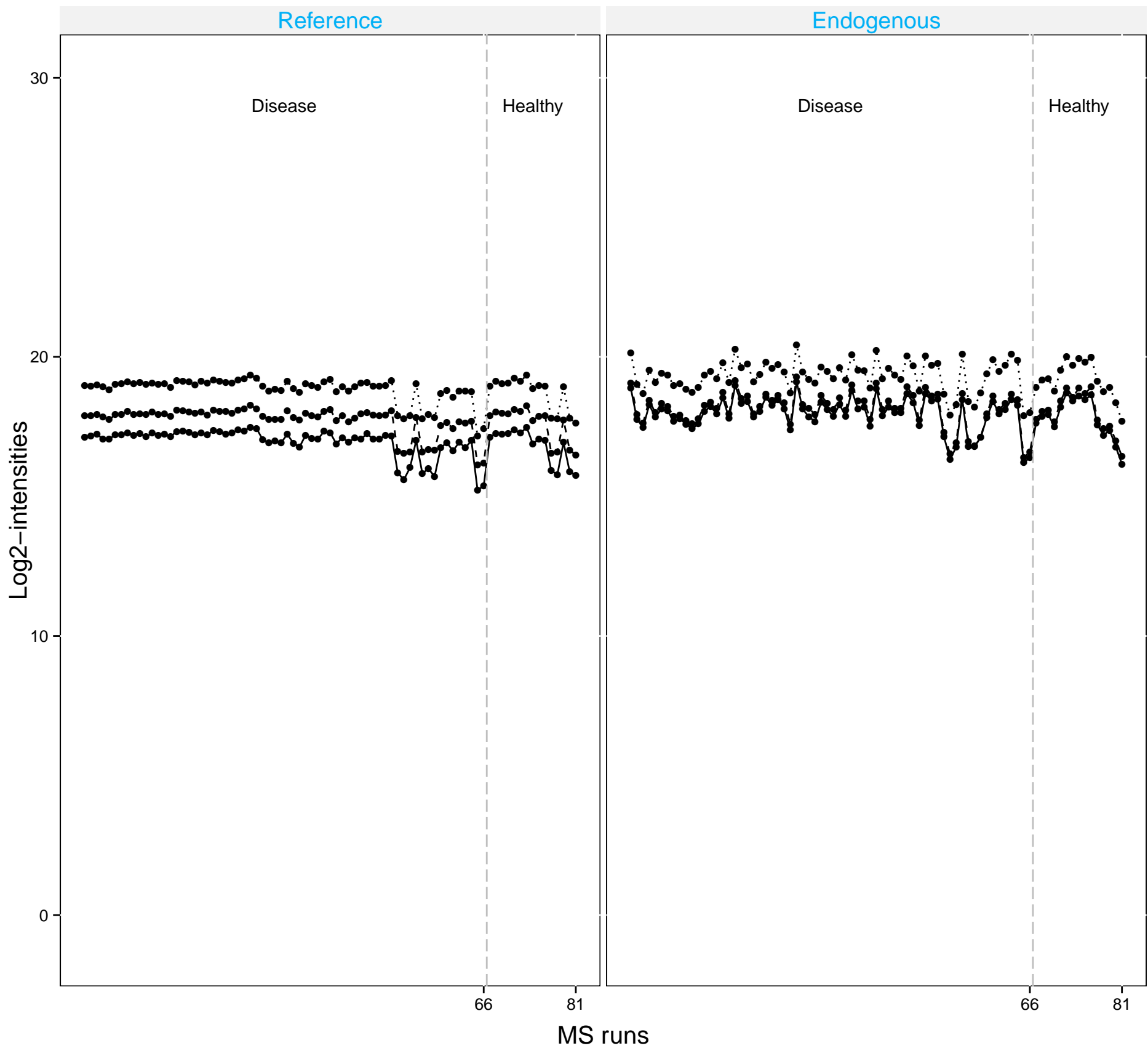
peptide: 3

DLATVYVDVLK_2_y5_1 DYVSQFEGSALGK_2_y6_1 VSFLSALEEYTK_2_y6_1
DLATVYVDVLK_2_y7_1 DYVSQFEGSALGK_2_y7_1 VSFLSALEEYTK_2_y7_1
DLATVYVDVLK_2_y8_1 DYVSQFEGSALGK_2_y8_1 VSFLSALEEYTK_2_y8_1



APOB

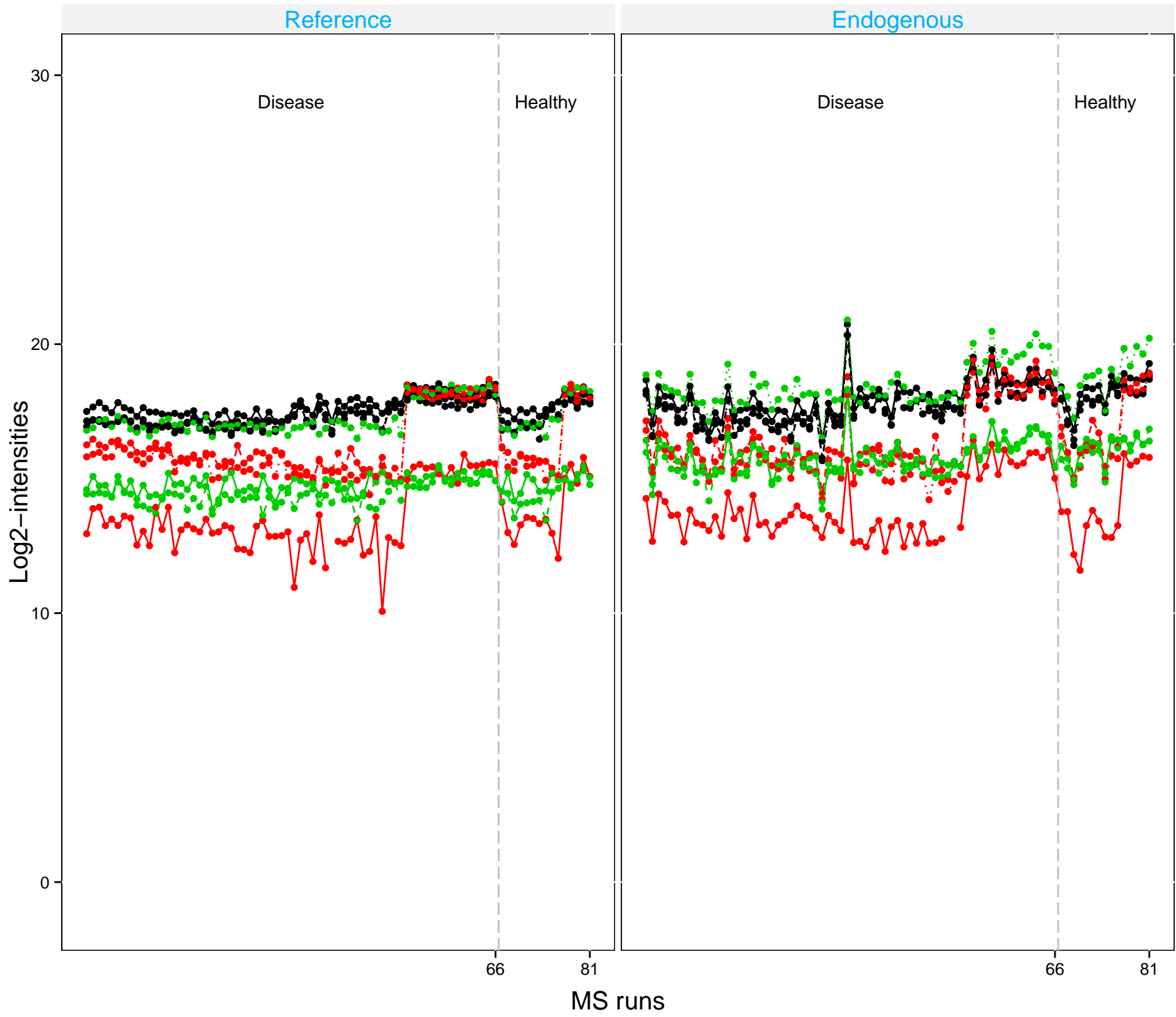
peptide: 1 —●— FSVFAGIVIPSFQALTAR_3_y4_1 —●— FSVFAGIVIPSFQALTAR_3_y5_1 —●— FSVFAGIVIPSFQALTAR_3_y9_1



APOE

peptide: 3

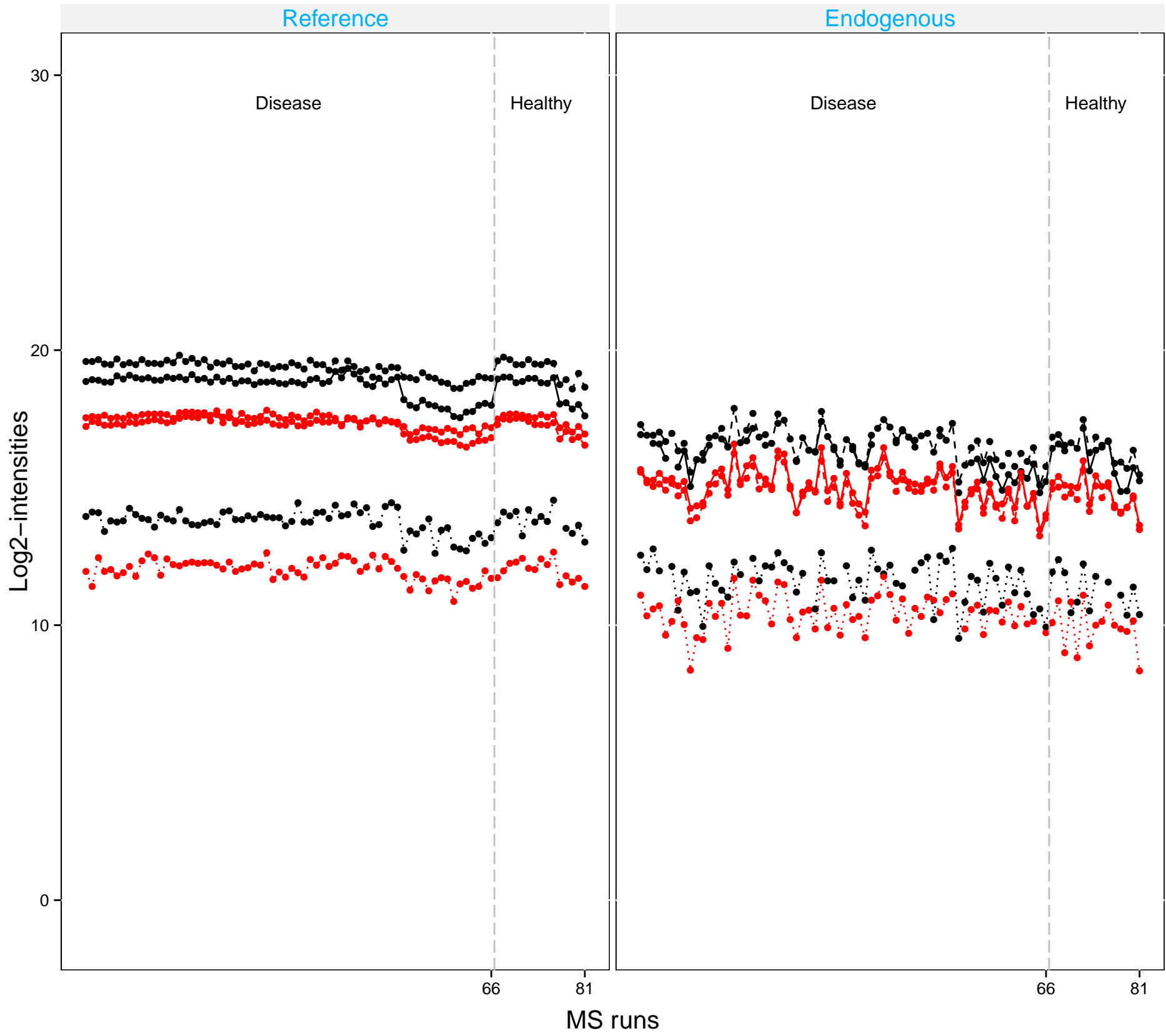
—●— AATVGSLAGQPLQER_2_y5_1 —●— SELEEQLTPVAEETR_2_y3_1 —●— SWFEPLVEDMQR_2_y3_1
—●— AATVGSLAGQPLQER_2_y7_1 —●— SELEEQLTPVAEETR_2_y7_1 —●— SWFEPLVEDMQR_2_y6_1
—●— AATVGSLAGQPLQER_2_y8_1 —●— SELEEQLTPVAEETR_2_y8_1 —●— SWFEPLVEDMQR_2_y8_1



B2MG

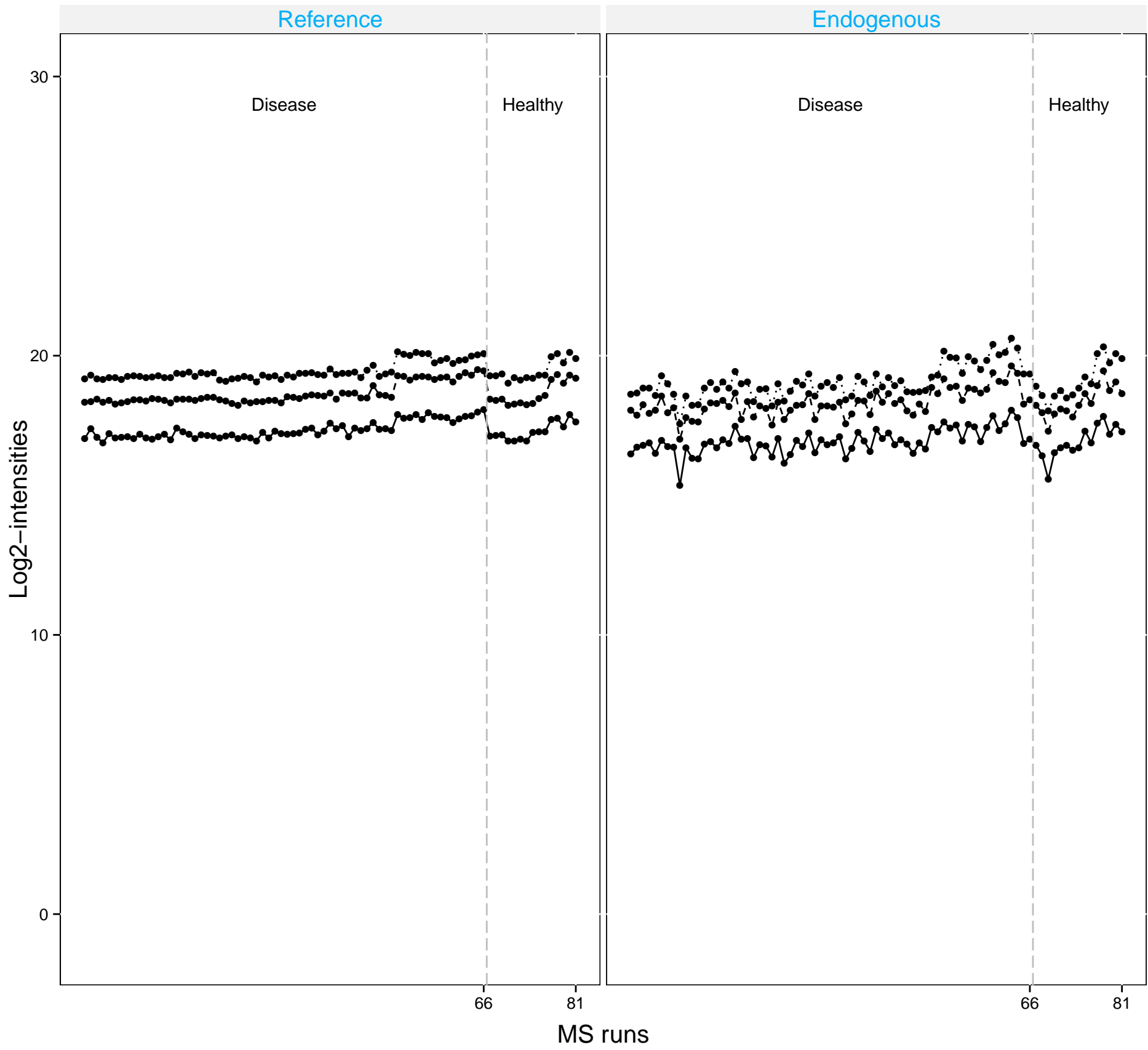
peptide: 2

VEHSDLSFSK_3_y3_1 VEHSDLSFSK_3_y7_1 VNHVTLSQPK_3_y6_1
VEHSDLSFSK_3_y4_1 VNHVTLSQPK_3_y5_1 VNHVTLSQPK_3_y7_1



C1QA

peptide: 1 ●— SLGFCDTTNK_2_y5_1 ●— SLGFCDTTNK_2_y6_1 ●— SLGFCDTTNK_2_y8_1



C1QB

peptide: 2

GNLCVNLMR_2_y4_1

GNLCVNLMR_2_y6_1

LEQGENVFLQATDK_2_y5_1

GNLCVNLMR_2_y5_1

LEQGENVFLQATDK_2_y4_1

LEQGENVFLQATDK_2_y7_1

Reference

Endogenous

Disease

Healthy

Disease

Healthy

Log2-intensities

30

20

10

0

66

81

MS runs

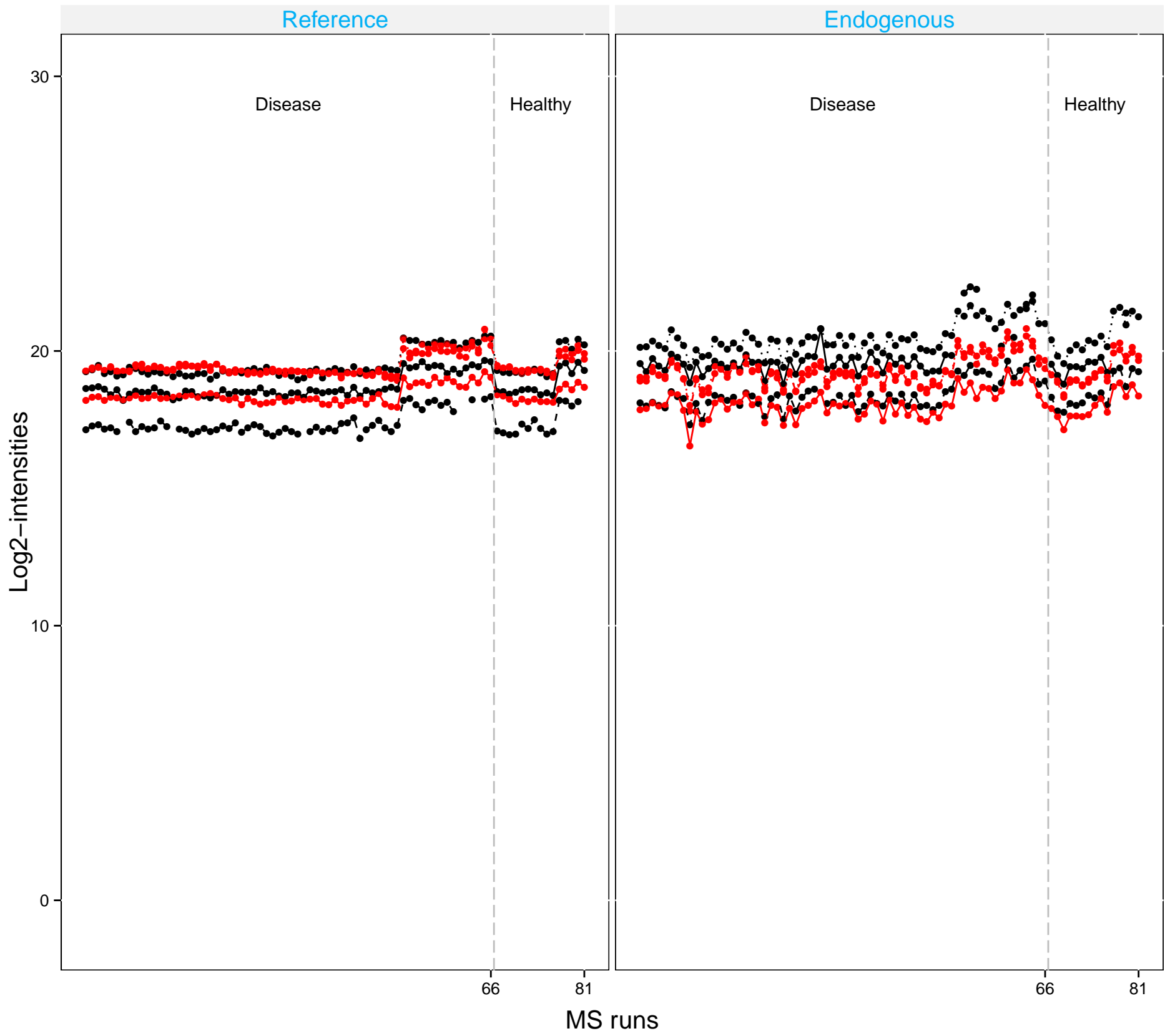
66

81

C1QC

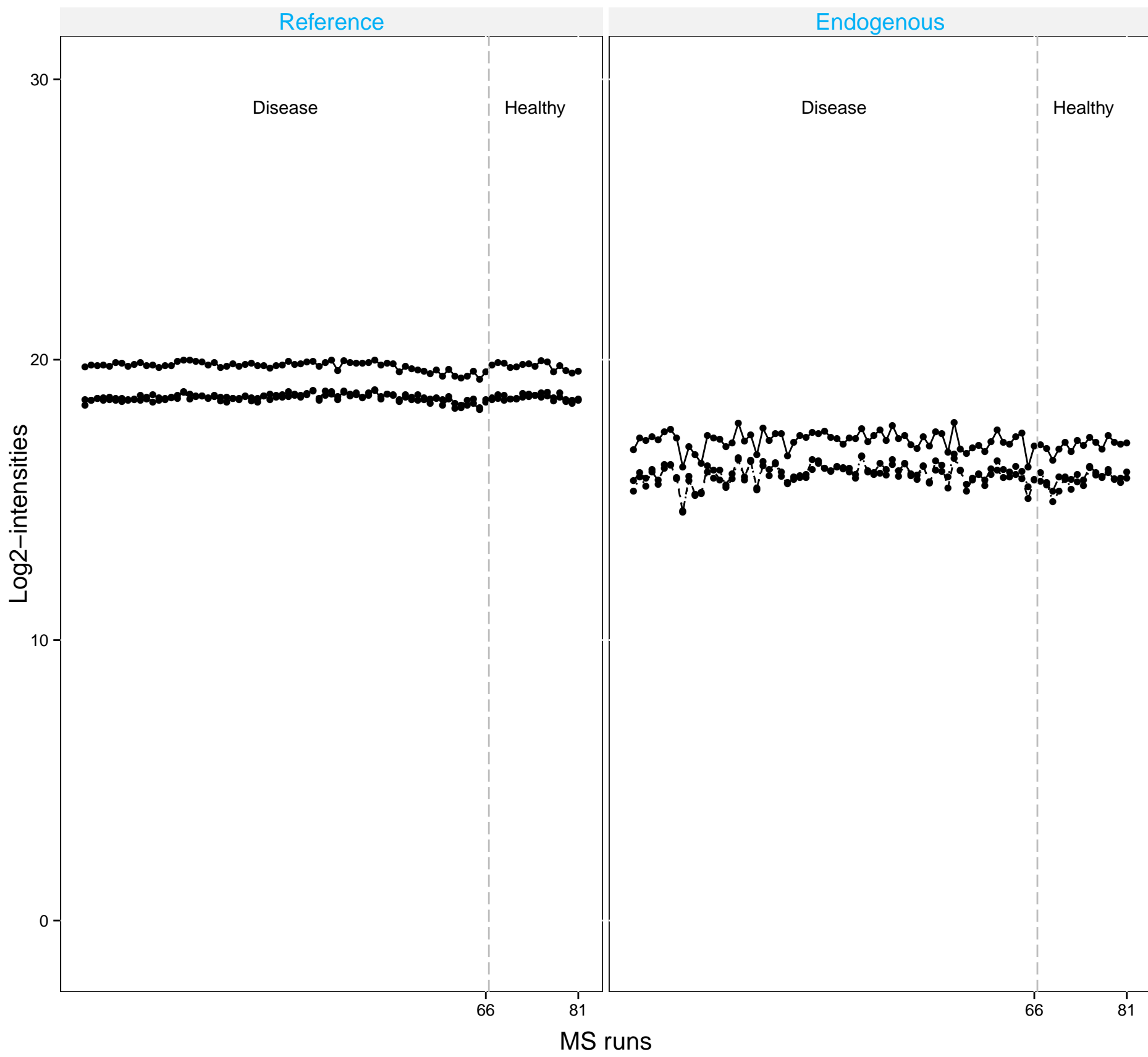
peptide: 2

—●— FQSVFTVTR_2_y5_1 —●— FQSVFTVTR_2_y7_1 —●— QTHQPPAPNSLIR_3_y6_1
—●— FQSVFTVTR_2_y6_1 —●— QTHQPPAPNSLIR_3_y5_1 —●— QTHQPPAPNSLIR_3_y6_2



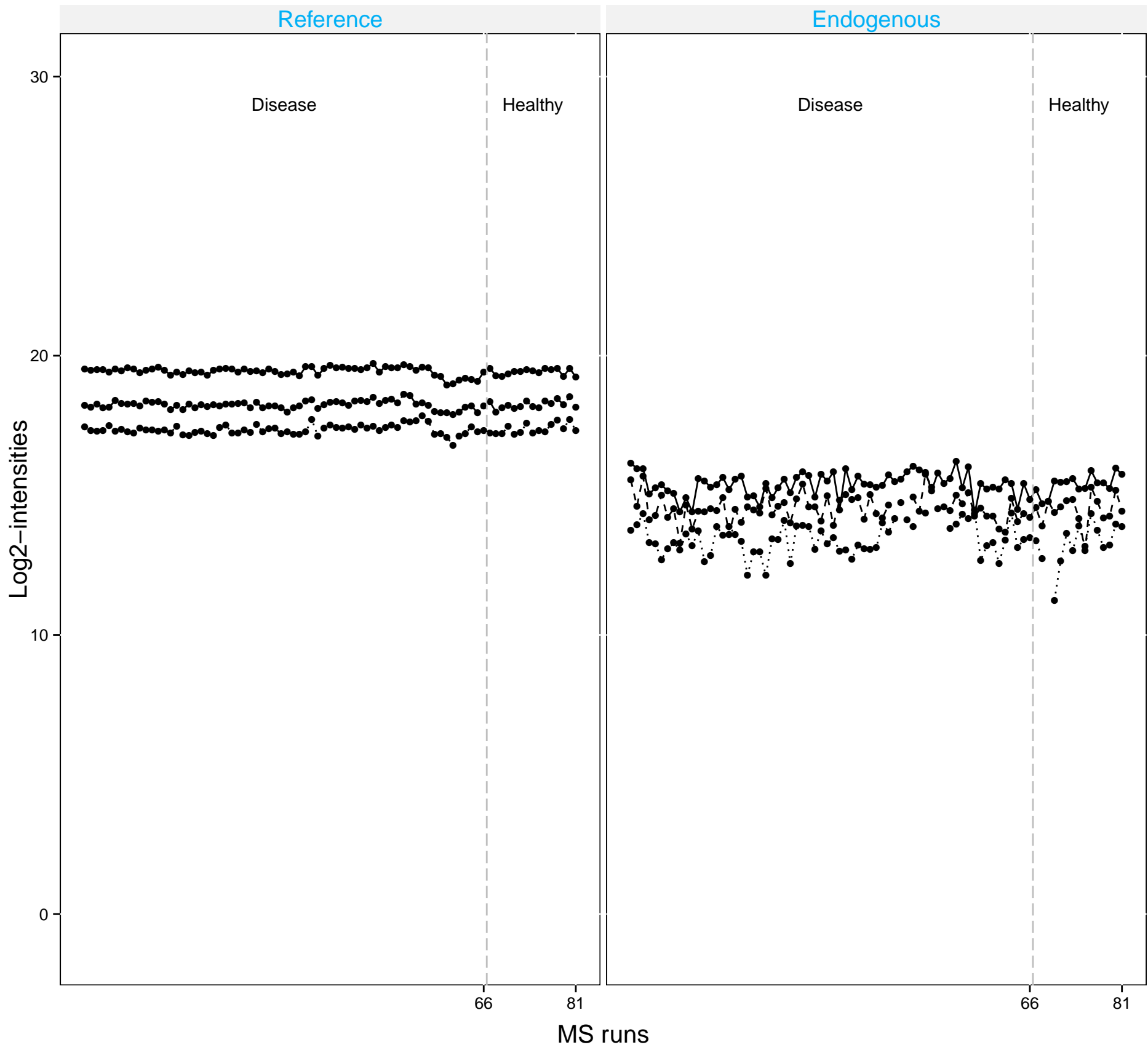
C1R

peptide: 1 —●— YTTTMGVNTYK_2_y6_1 —●— YTTTMGVNTYK_2_y7_1 —●— YTTTMGVNTYK_2_y8_1



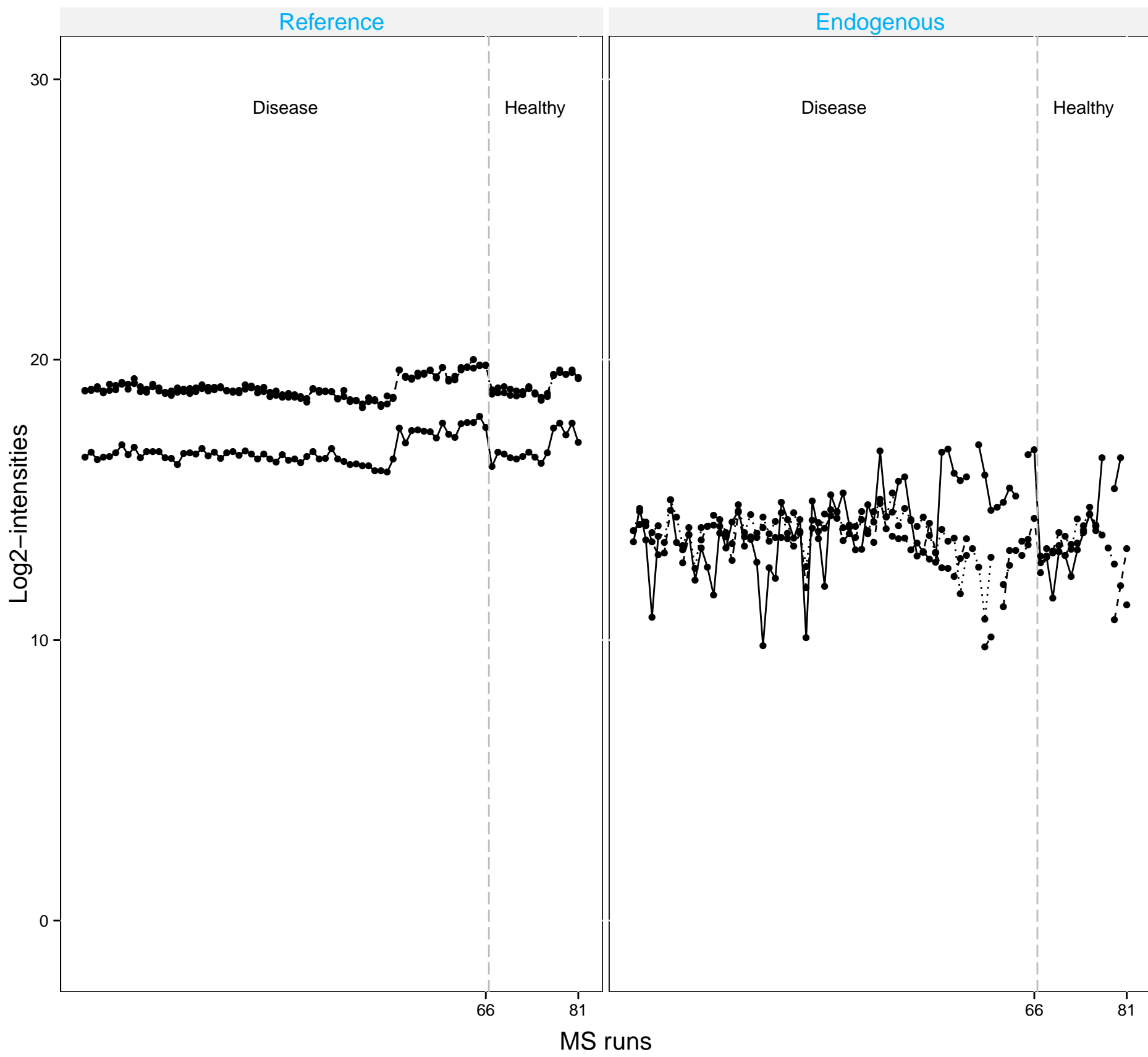
CADH5

peptide: 1 —●— YTFVVPEDTR_2_y5_1 —●— YTFVVPEDTR_2_y6_1 —●— YTFVVPEDTR_2_y8_1



CCNH

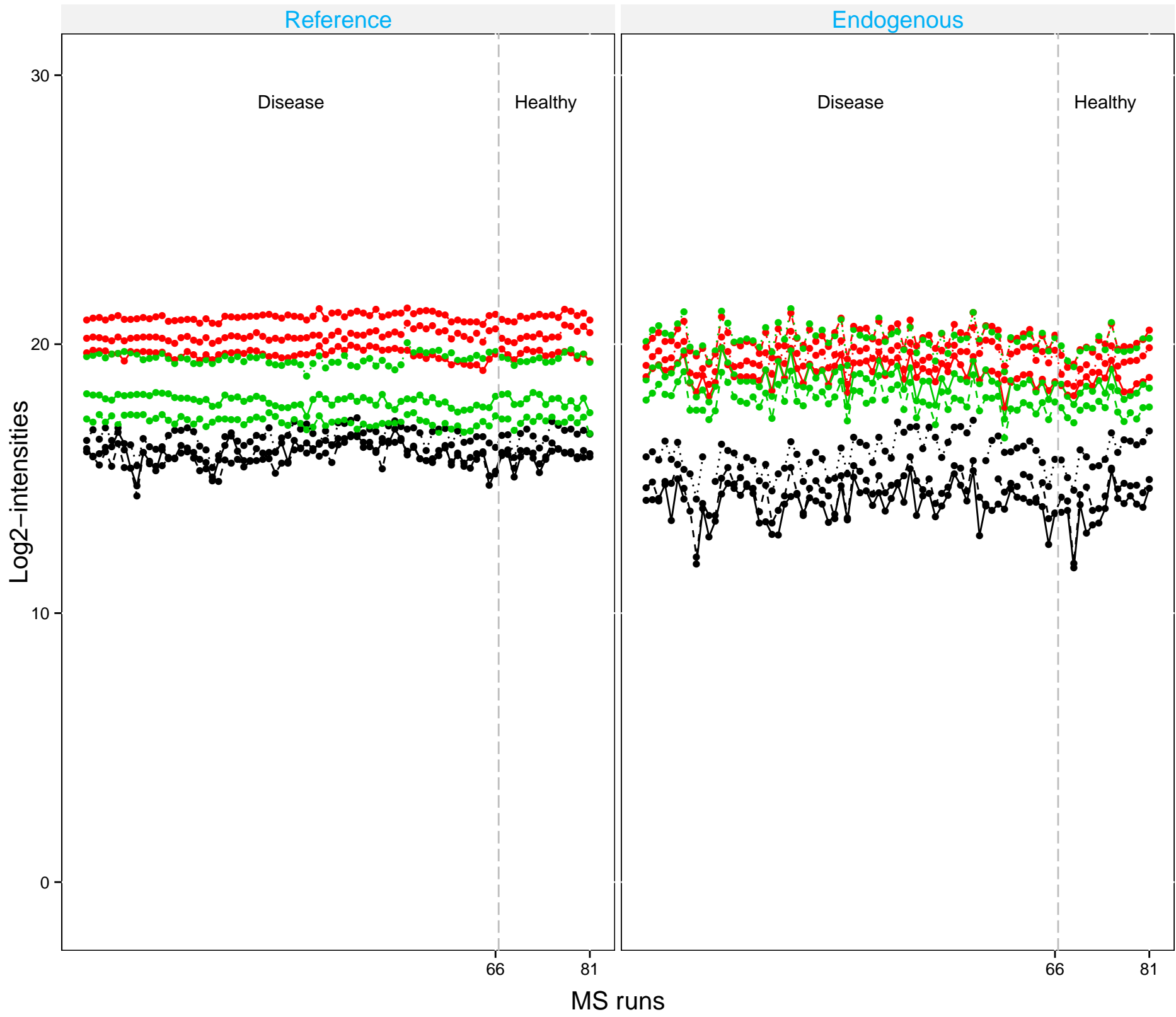
peptide: 1  HWTFSSEQLAR_3_b4_1  HWTFSSEQLAR_3_y4_1  HWTFSSEQLAR_3_y5_1



CFAB

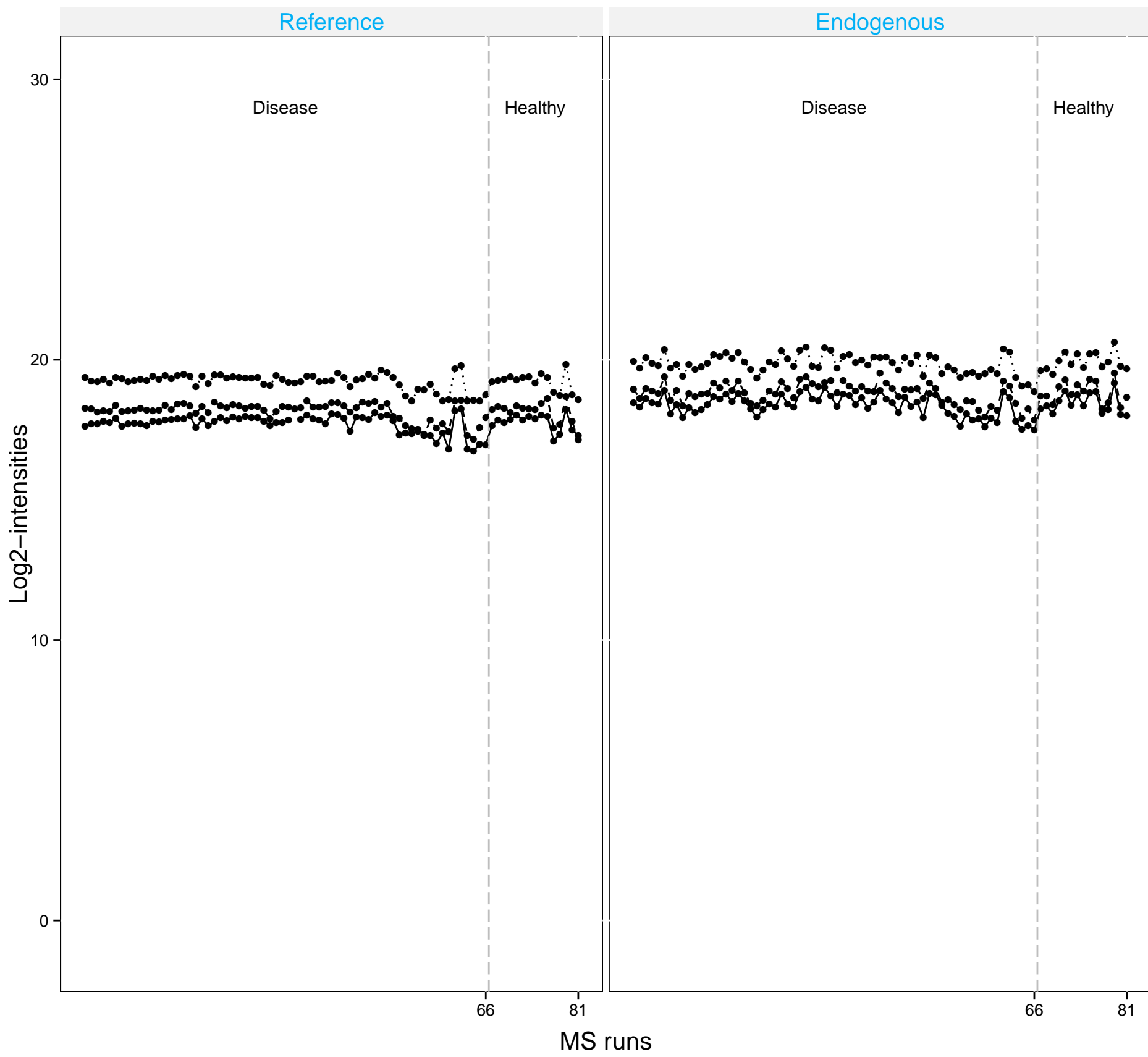
peptide: 3

—●— QLNEINYEDHK_2_y4_1 —●— YGLVTYATYPK_2_y3_1 —●— YGQTIRPICLPCTEGTTR_3_y6_1
—●— QLNEINYEDHK_2_y5_1 —●— YGLVTYATYPK_2_y7_1 —●— YGQTIRPICLPCTEGTTR_3_y7_1
—●— QLNEINYEDHK_2_y6_1 —●— YGLVTYATYPK_2_y8_1 —●— YGQTIRPICLPCTEGTTR_3_y8_1



CLUS

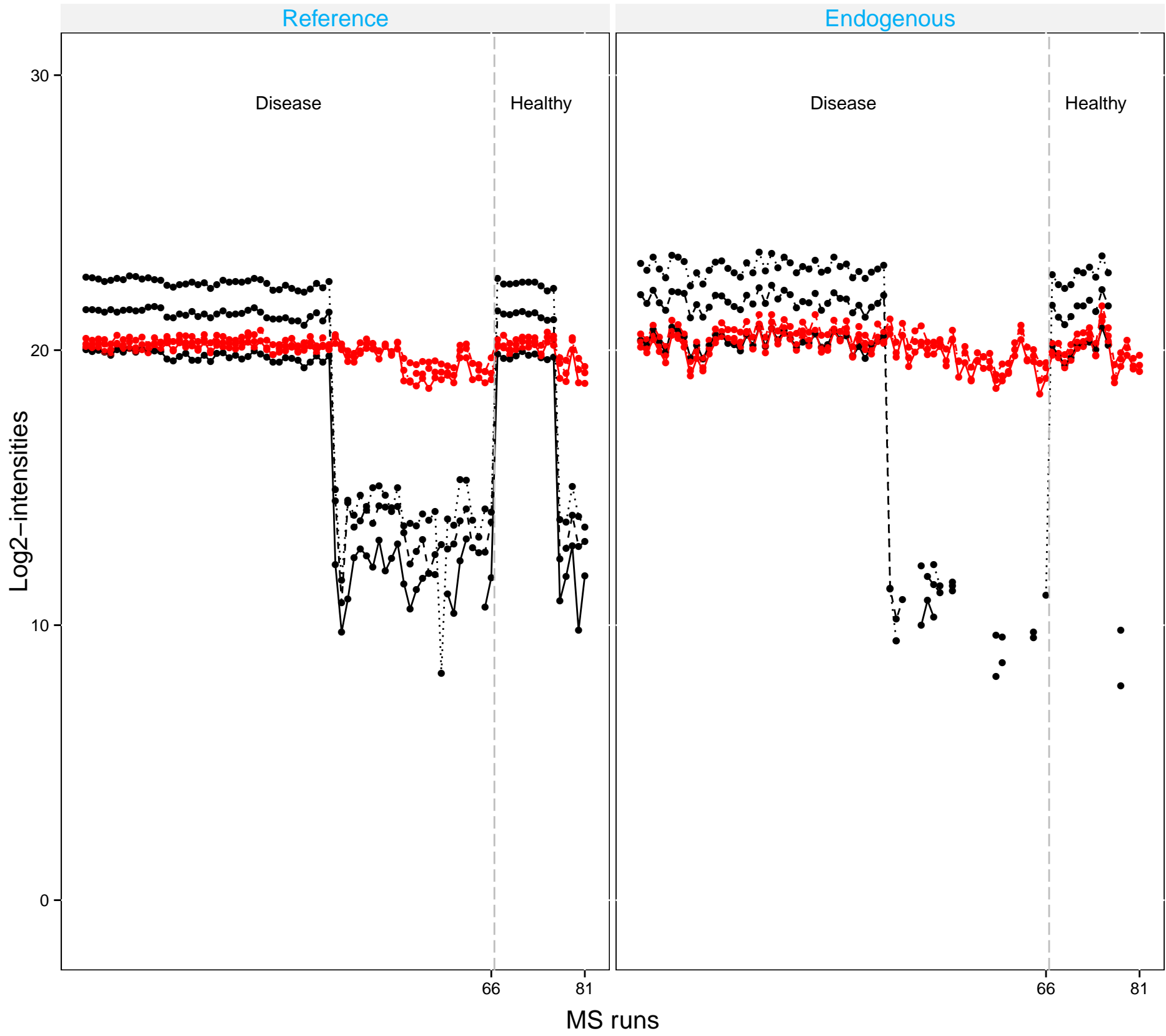
peptide: 1 ASSIIDELFQDR_2_b4_1 ASSIIDELFQDR_2_y4_1 ASSIIDELFQDR_2_y7_1



CO3

peptide: 2

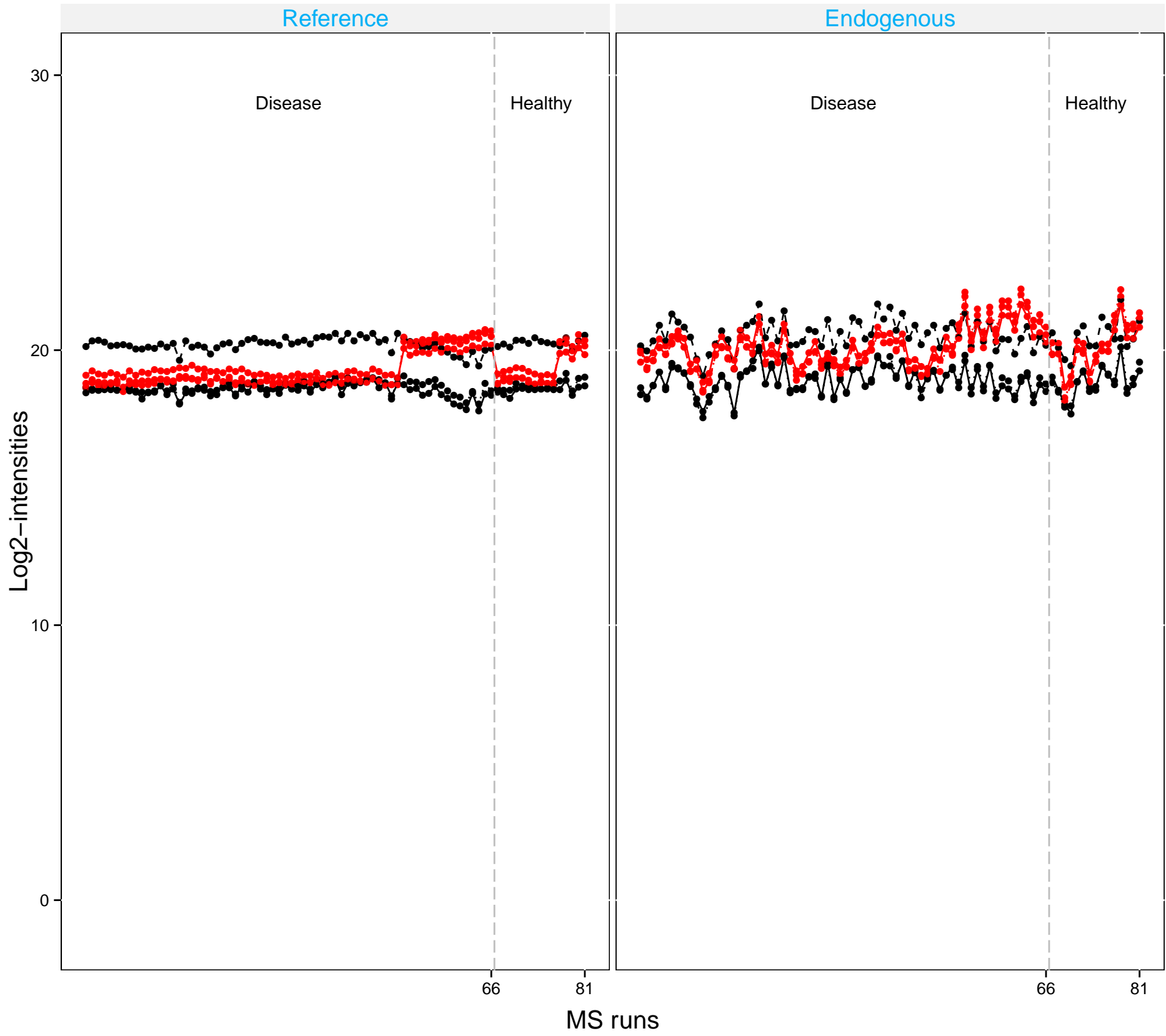
—●— IHWESASLLR_3_b3_1 —●— IHWESASLLR_3_y4_1 —●— ILLQGTPVAQMTEDAVDAER_3_y7_1
—●— IHWESASLLR_3_y3_1 —●— ILLQGTPVAQMTEDAVDAER_3_y5_1 —●— ILLQGTPVAQMTEDAVDAER_3_y8_1



CO4A

peptide: 2

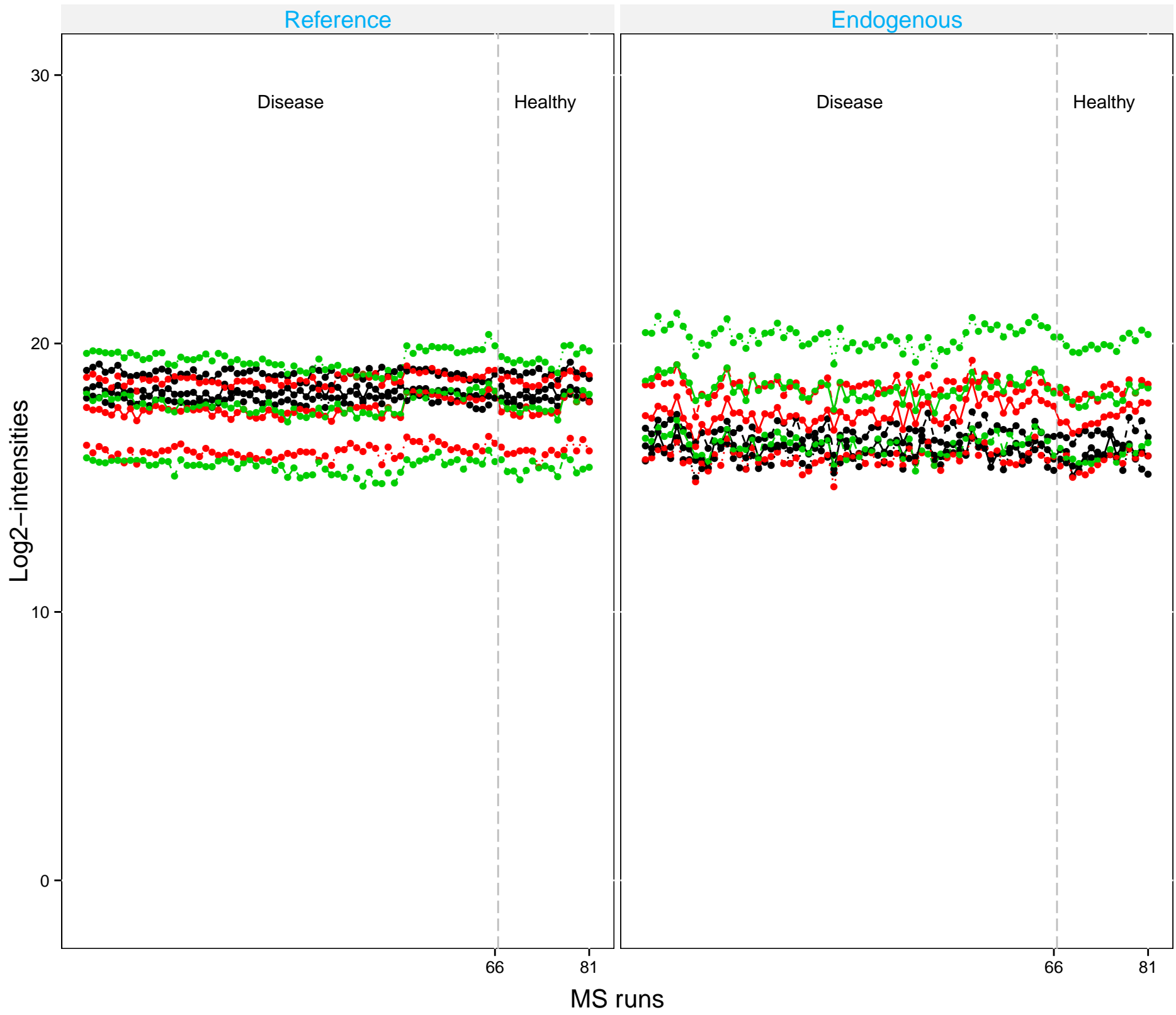
GSFEFPVGDAVSK_2_b4_1 GSFEFPVGDAVSK_2_y9_1 LGQYASPTAK_2_y6_1
GSFEFPVGDAVSK_2_y8_1 LGQYASPTAK_2_y4_1 LGQYASPTAK_2_y7_1



CO5

peptide: 3

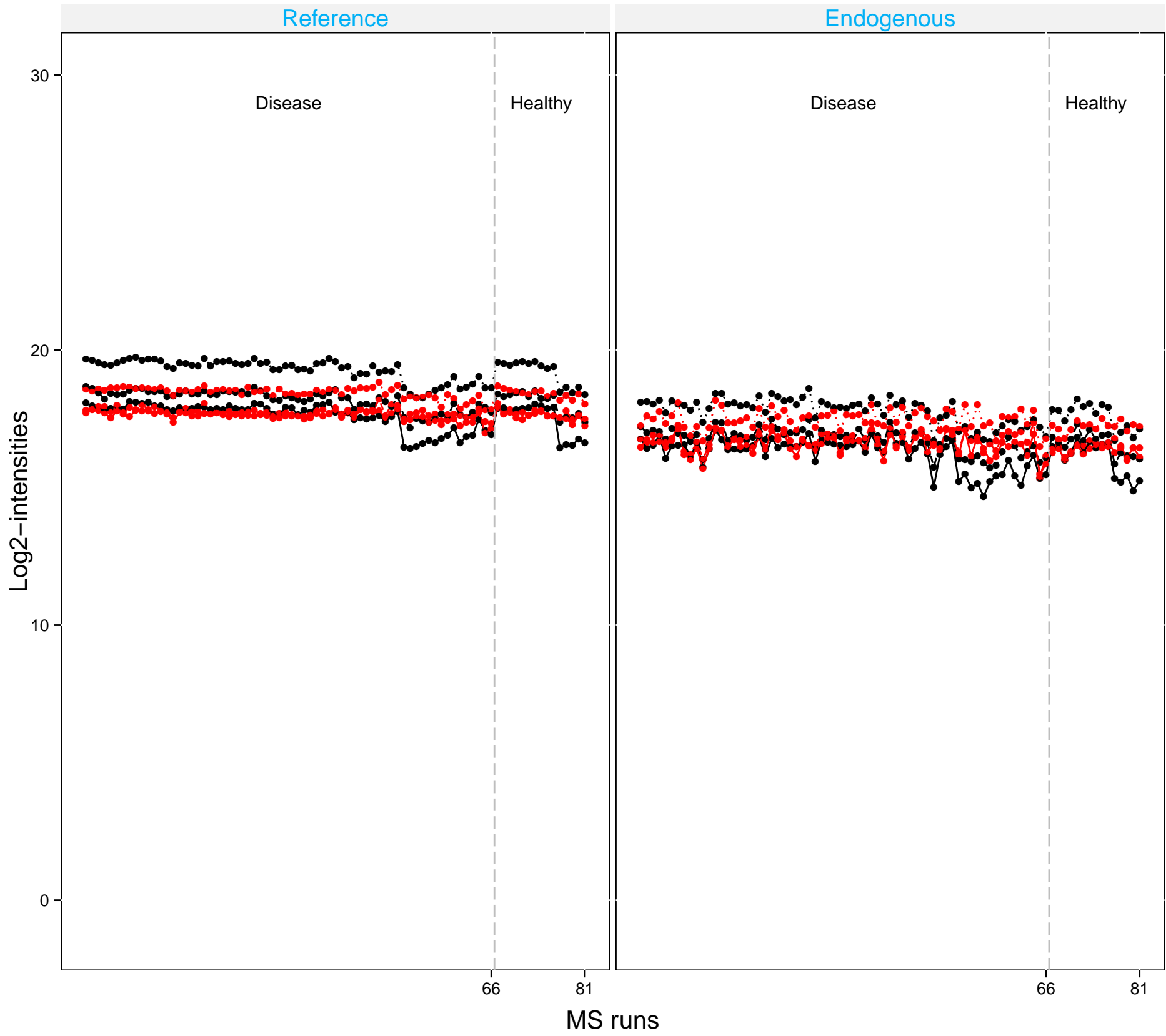
—●— FQNSAILTIQPK_2_y5_1 —●— TDAPDLPEENQAR_2_y10_2 —●— TLLPVSKPEIR_3_y4_1
—●— FQNSAILTIQPK_2_y6_1 —●— TDAPDLPEENQAR_2_y7_1 —●— TLLPVSKPEIR_3_y7_1
—●— FQNSAILTIQPK_2_y7_1 —●— TDAPDLPEENQAR_2_y8_1 —●— TLLPVSKPEIR_3_y8_2



CO6

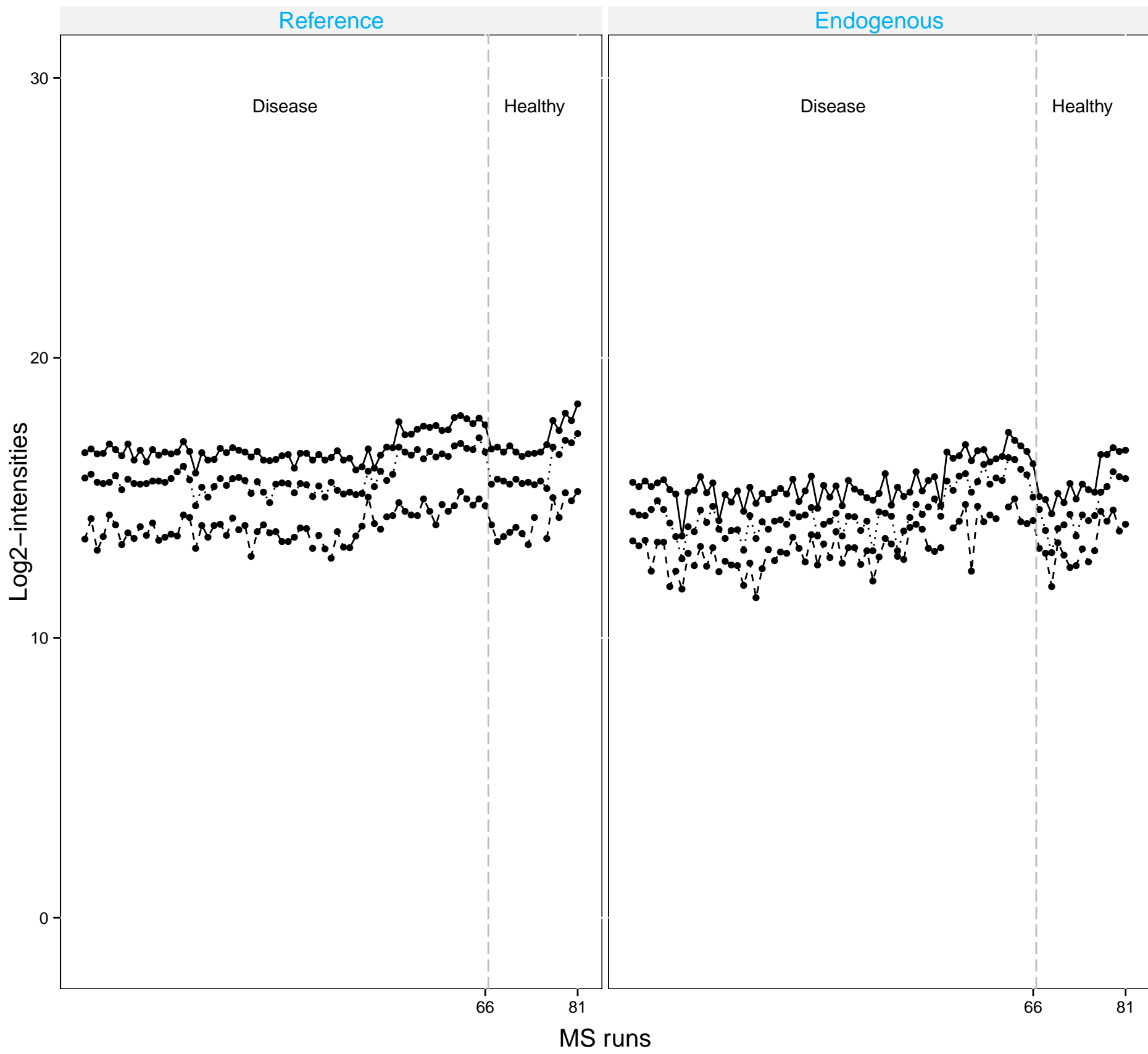
peptide: 2

DLHLSDVFLK_3_b3_1 DLHLSDVFLK_3_y3_1 IGESIELTCPK_2_y5_1
DLHLSDVFLK_3_b6_1 IGESIELTCPK_2_y4_1 IGESIELTCPK_2_y6_1






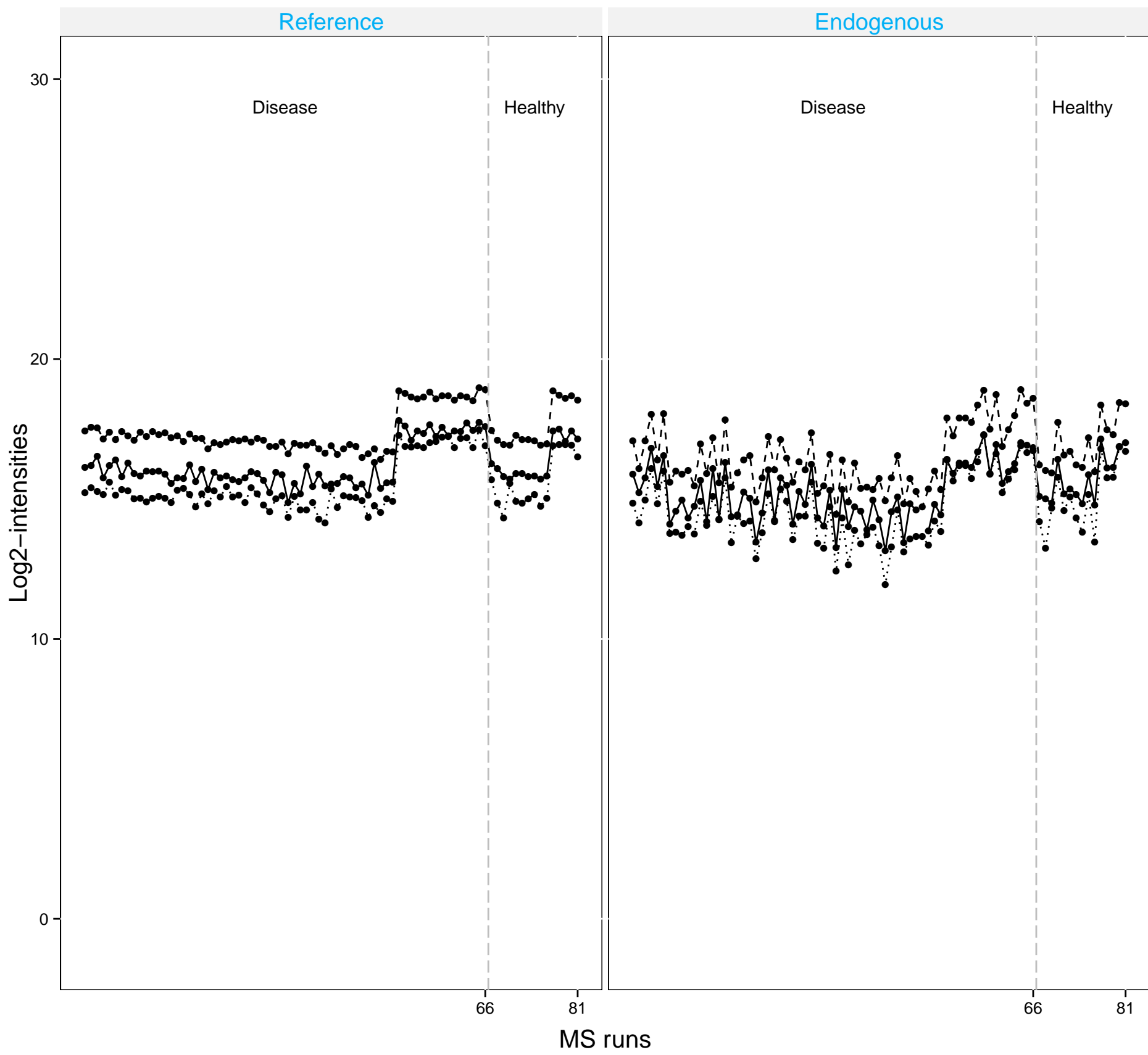
CO7

peptide: 1 —●— YSAWAESVTNLPQVIK_2_y5_1 —●— YSAWAESVTNLPQVIK_2_y7_1 —●— YSAWAESVTNLPQVIK_2_y8_1



FN1

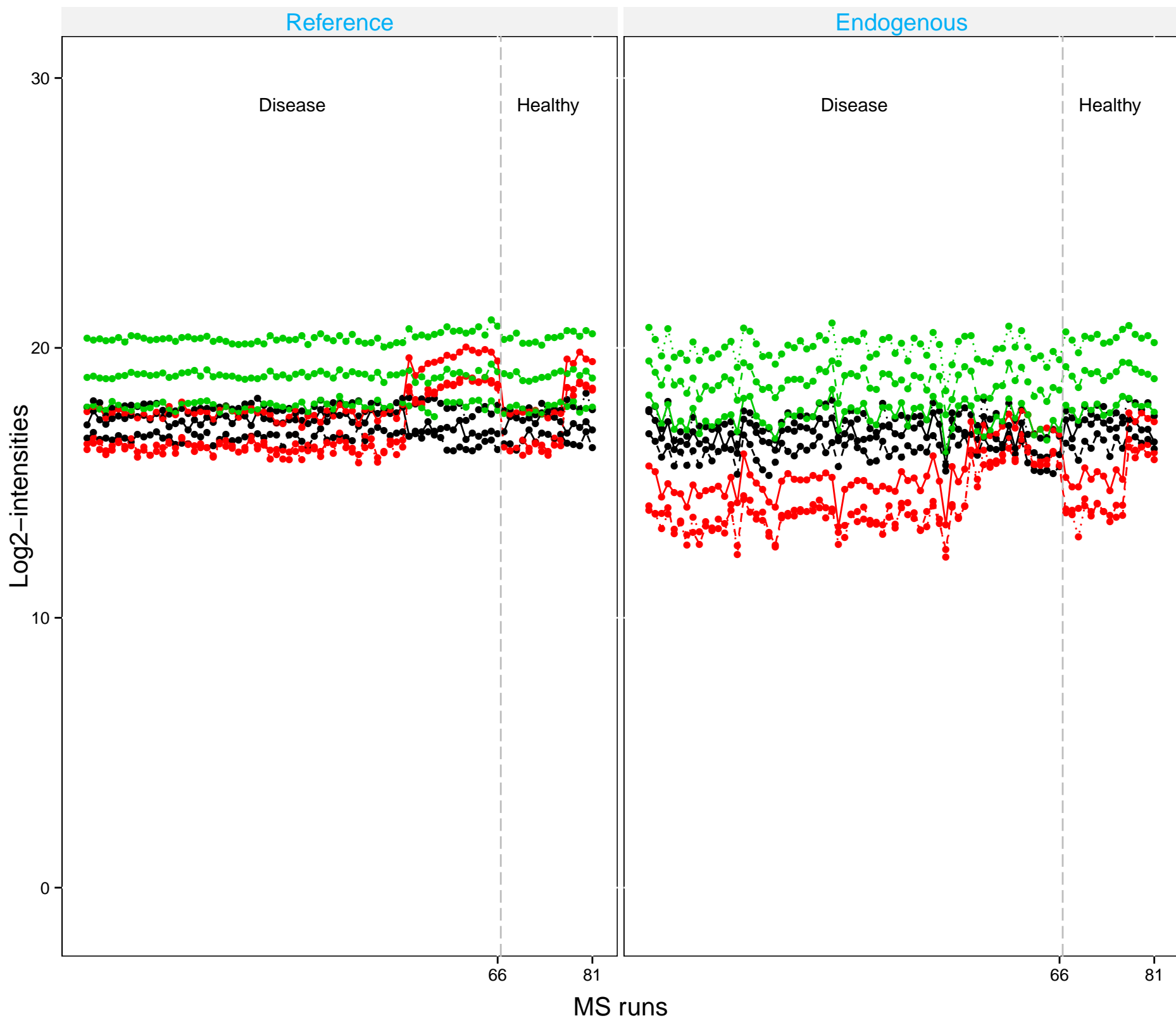
peptide: 1  SYTITGLQPGTDYK_2_b3_1  SYTITGLQPGTDYK_2_y6_1  SYTITGLQPGTDYK_2_y9_1



GELS

peptide: 3

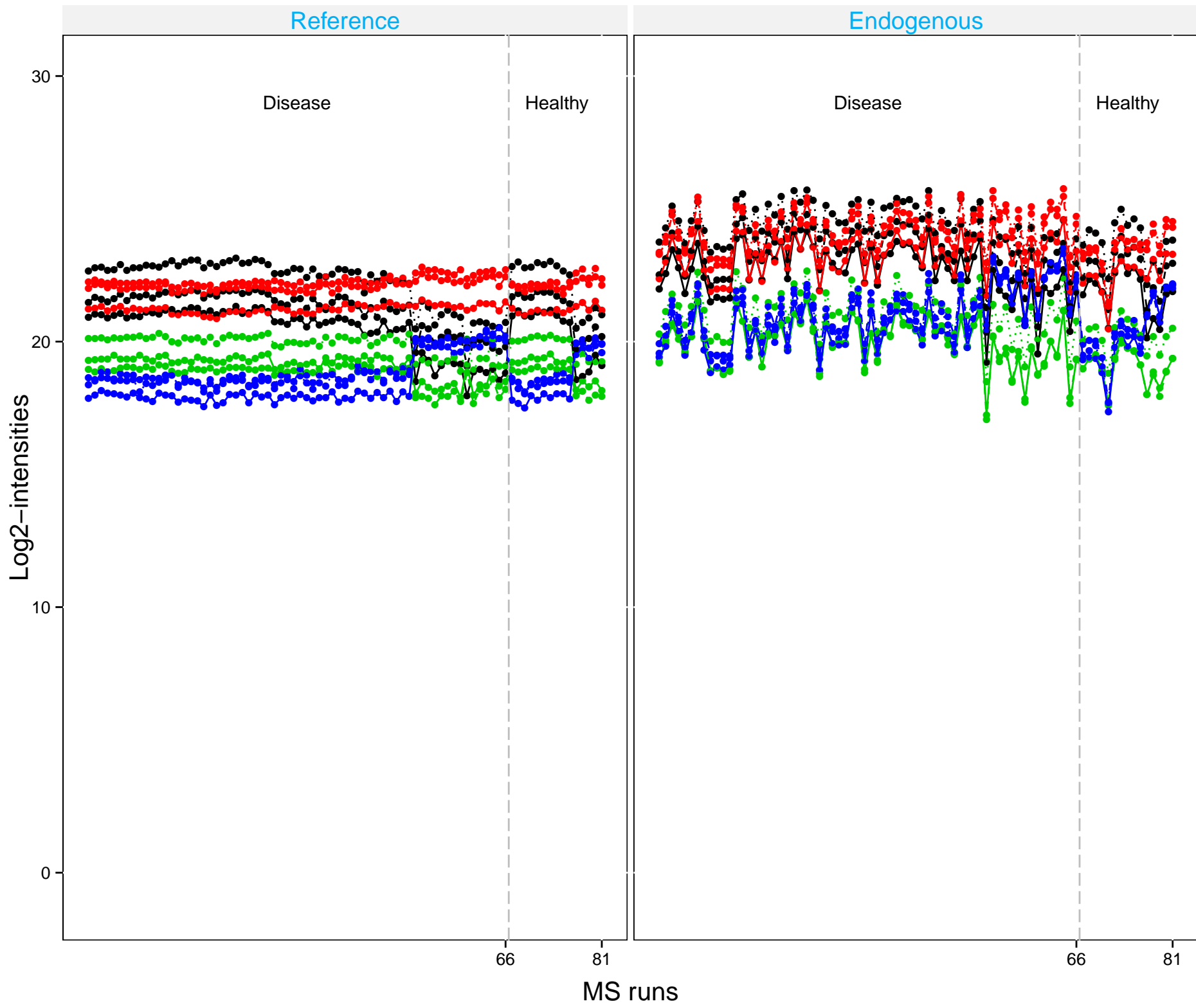
—●— AGALNSNDAFVLK_2_y3_1 —●— EVQGFESATFLGYFK_2_y4_1 —●— HVVPNEVVVQR_3_b5_1
—●— AGALNSNDAFVLK_2_y7_1 —●— EVQGFESATFLGYFK_2_y7_1 —●— HVVPNEVVVQR_3_b6_1
—●— AGALNSNDAFVLK_2_y8_1 —●— EVQGFESATFLGYFK_2_y8_1 —●— HVVPNEVVVQR_3_y4_1



HPT

peptide: 4

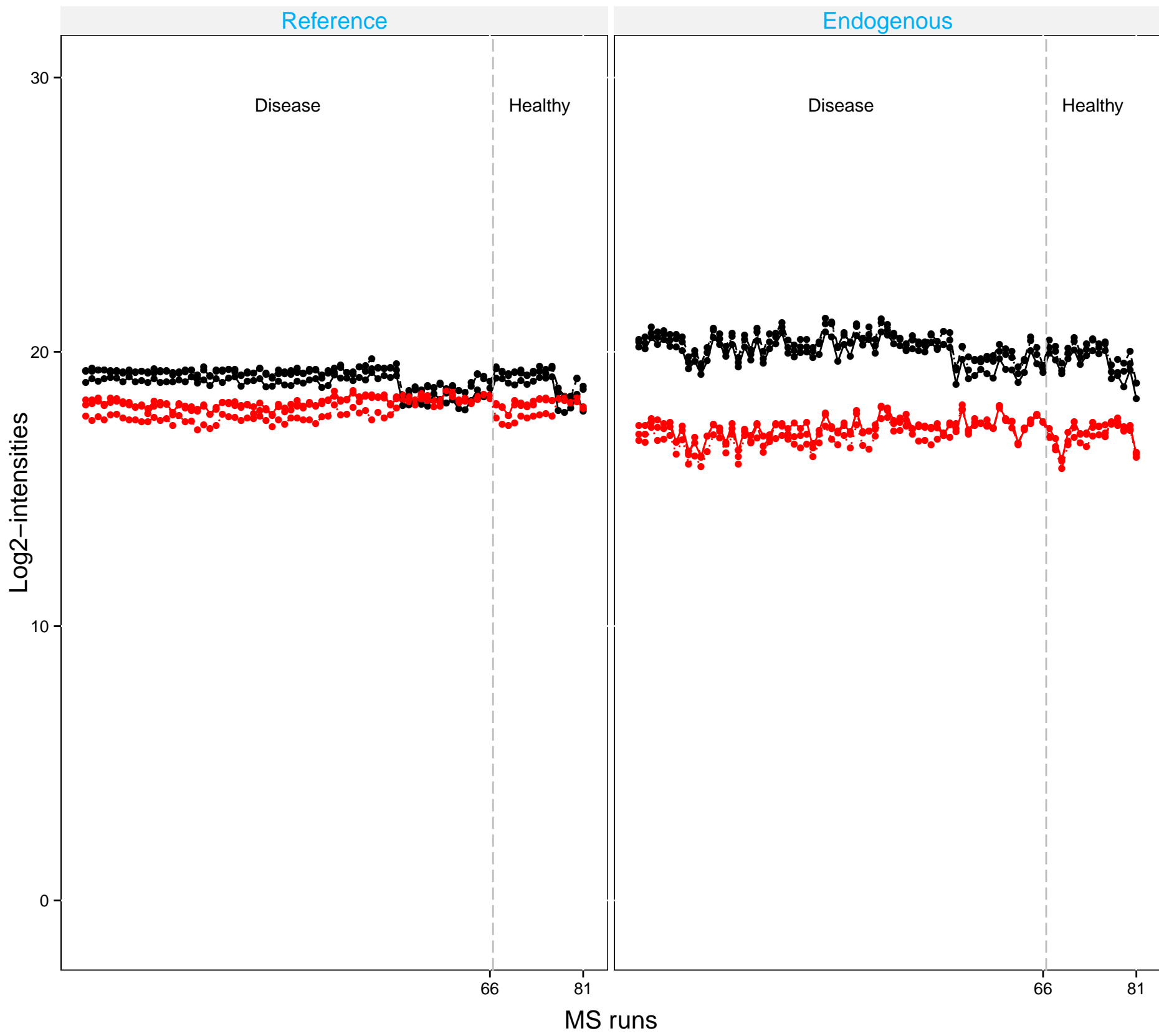
DIAPTLTLYVGK_2_y6_1 VTSIQDWVQK_2_y5_1 VVLHPNYSQVDIGLIK_3_y6_1
DIAPTLTLYVGK_2_y9_1 VTSIQDWVQK_2_y6_1 YVMLPVADQDQCIR_2_b4_1
DIAPTLTLYVGK_2_y9_2 VVLHPNYSQVDIGLIK_3_y4_1 YVMLPVADQDQCIR_2_y4_1
VTSIQDWVQK_2_y4_1 VVLHPNYSQVDIGLIK_3_y5_1 YVMLPVADQDQCIR_2_y7_1



IC1

peptide: 2

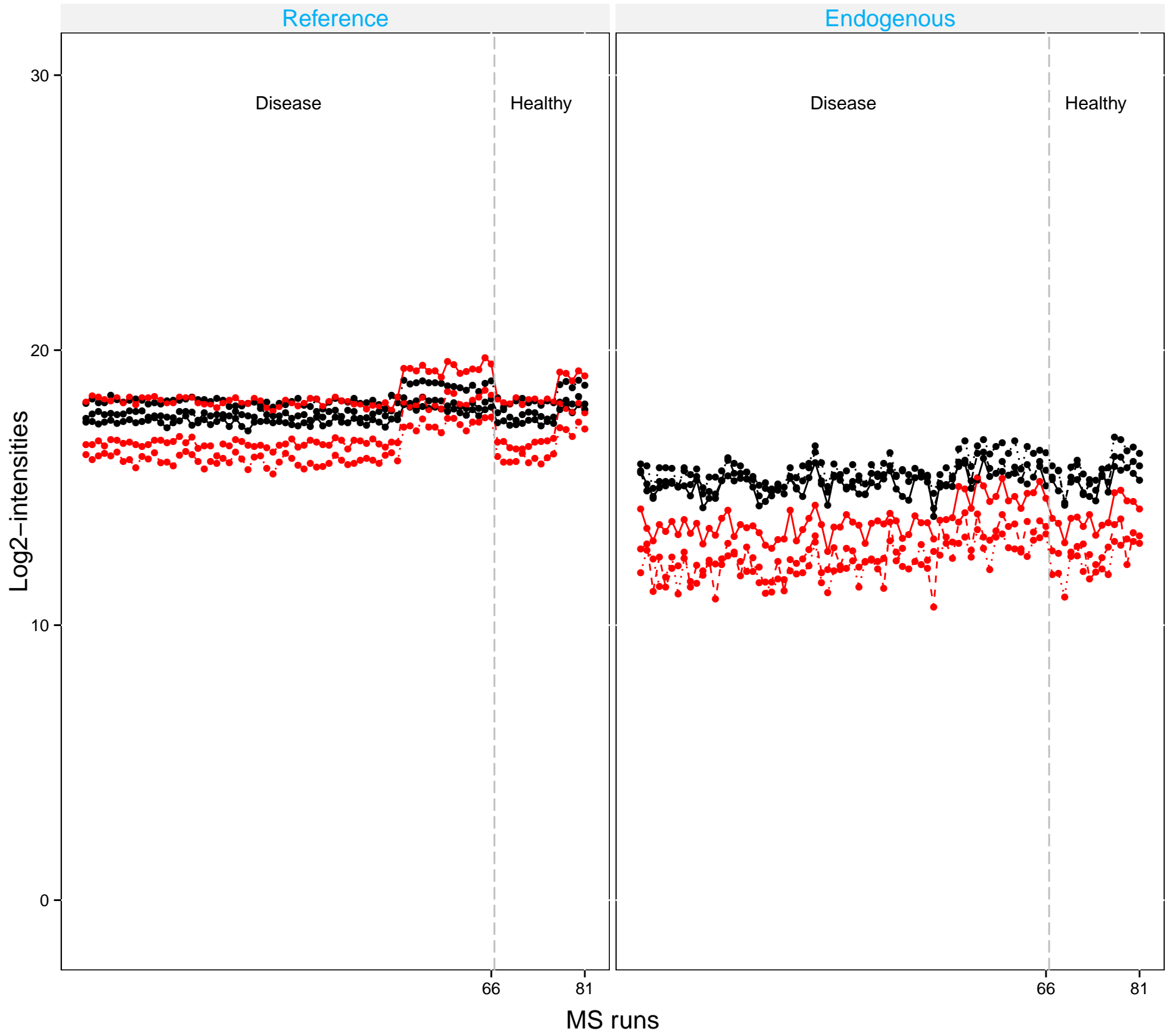
GVTSVSQIFHSPDLAIR_3_y4_1 GVTSVSQIFHSPDLAIR_3_y7_1 VTTSQDMLSIMEK_2_y5_1
GVTSVSQIFHSPDLAIR_3_y6_1 VTTSQDMLSIMEK_2_y3_1 VTTSQDMLSIMEK_2_y8_1



IGF2

peptide: 2

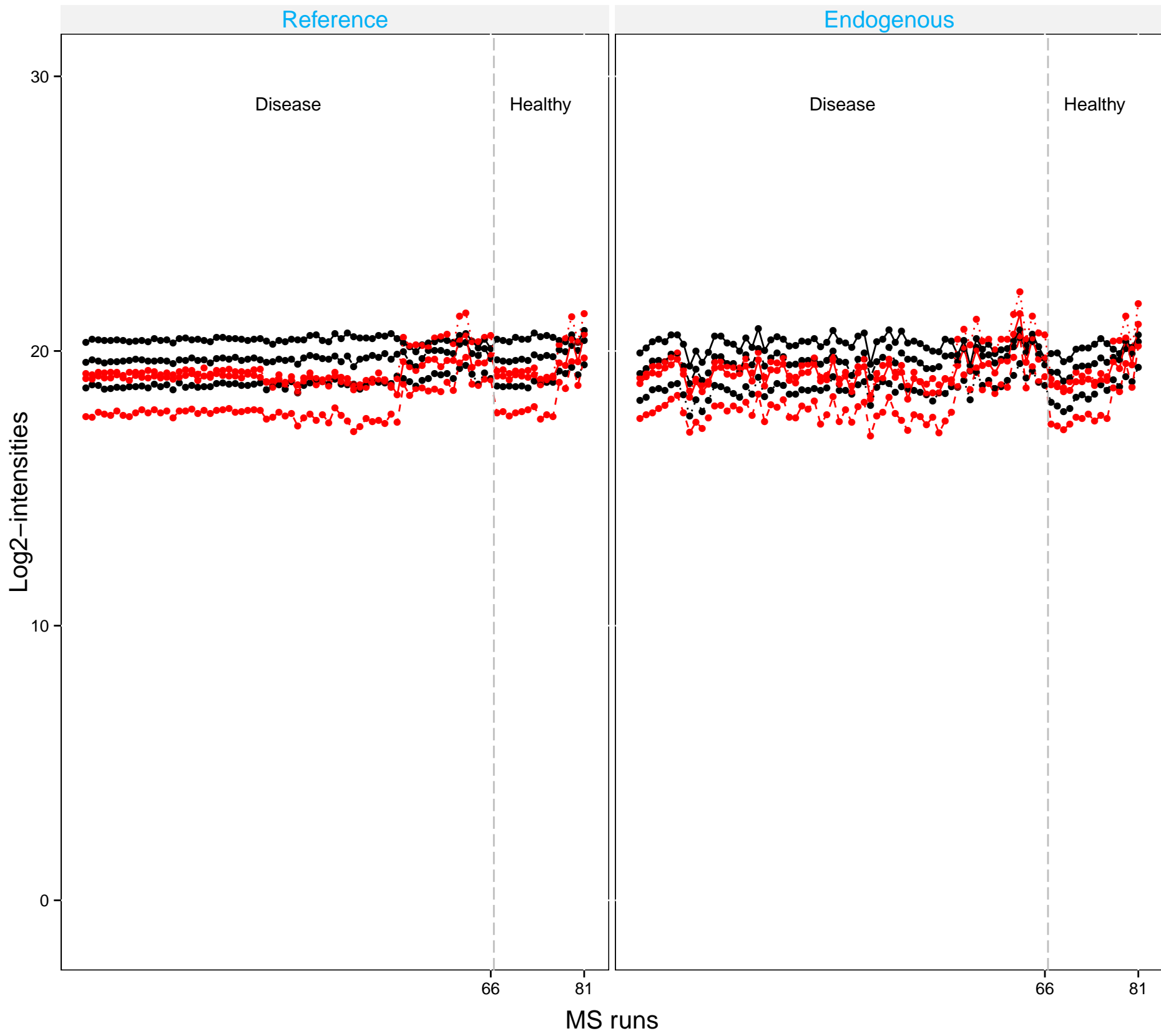
—●— GIVEECCFR_2_y4_1 —●— GIVEECCFR_2_y6_1 —●— SCDLALLETYCATPAK_2_b5_1
—●— GIVEECCFR_2_y5_1 —●— SCDLALLETYCATPAK_2_b3_1 —●— SCDLALLETYCATPAK_2_y4_1



ITI4

peptide: 2

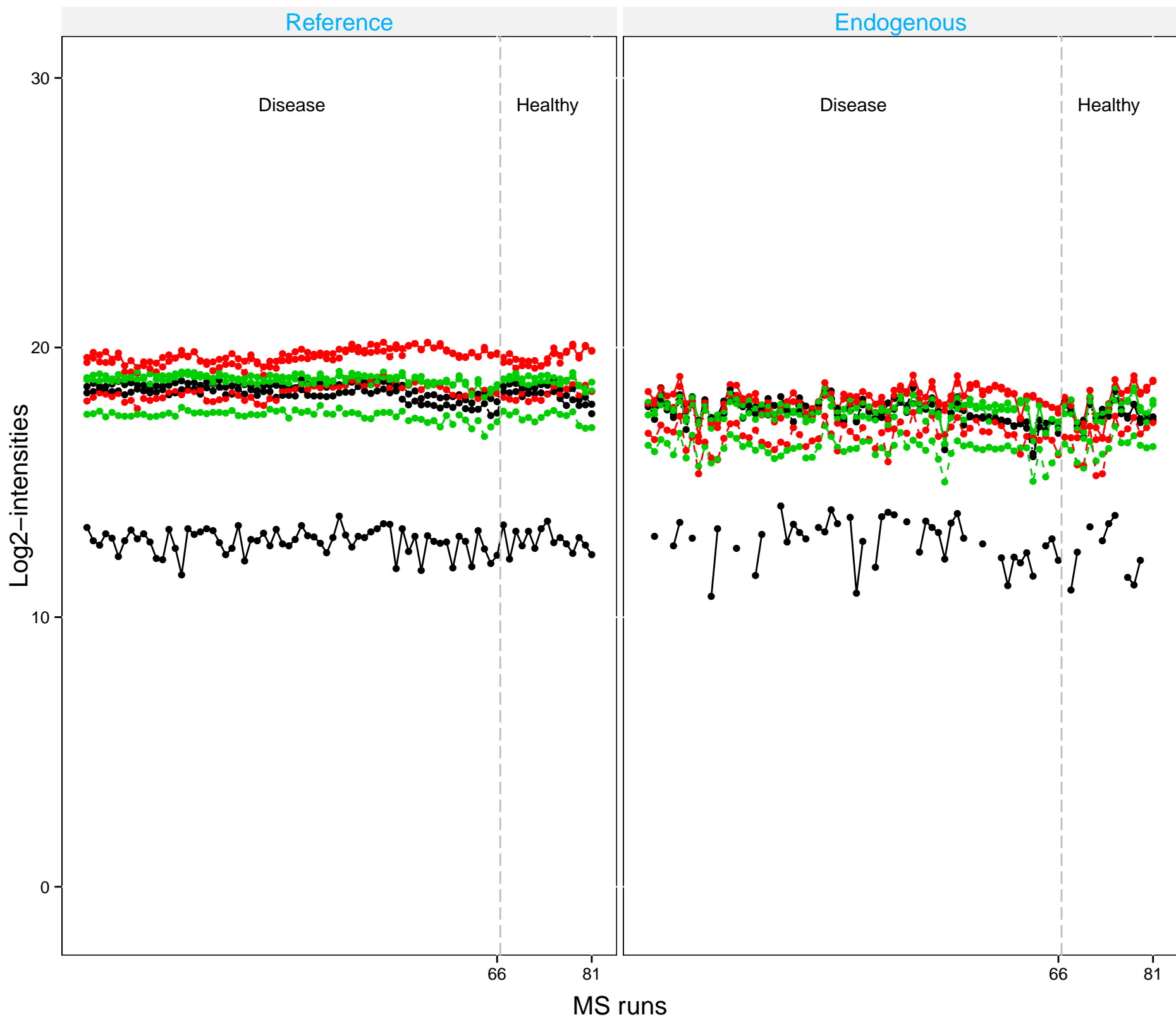
ETLFSVMPGLK_2_y4_1 ETLFSVMPGLK_2_y8_1 LGVYELLK_2_y7_1
ETLFSVMPGLK_2_y7_1 LGVYELLK_2_y5_1 LGVYELLK_2_y8_1



PLMN

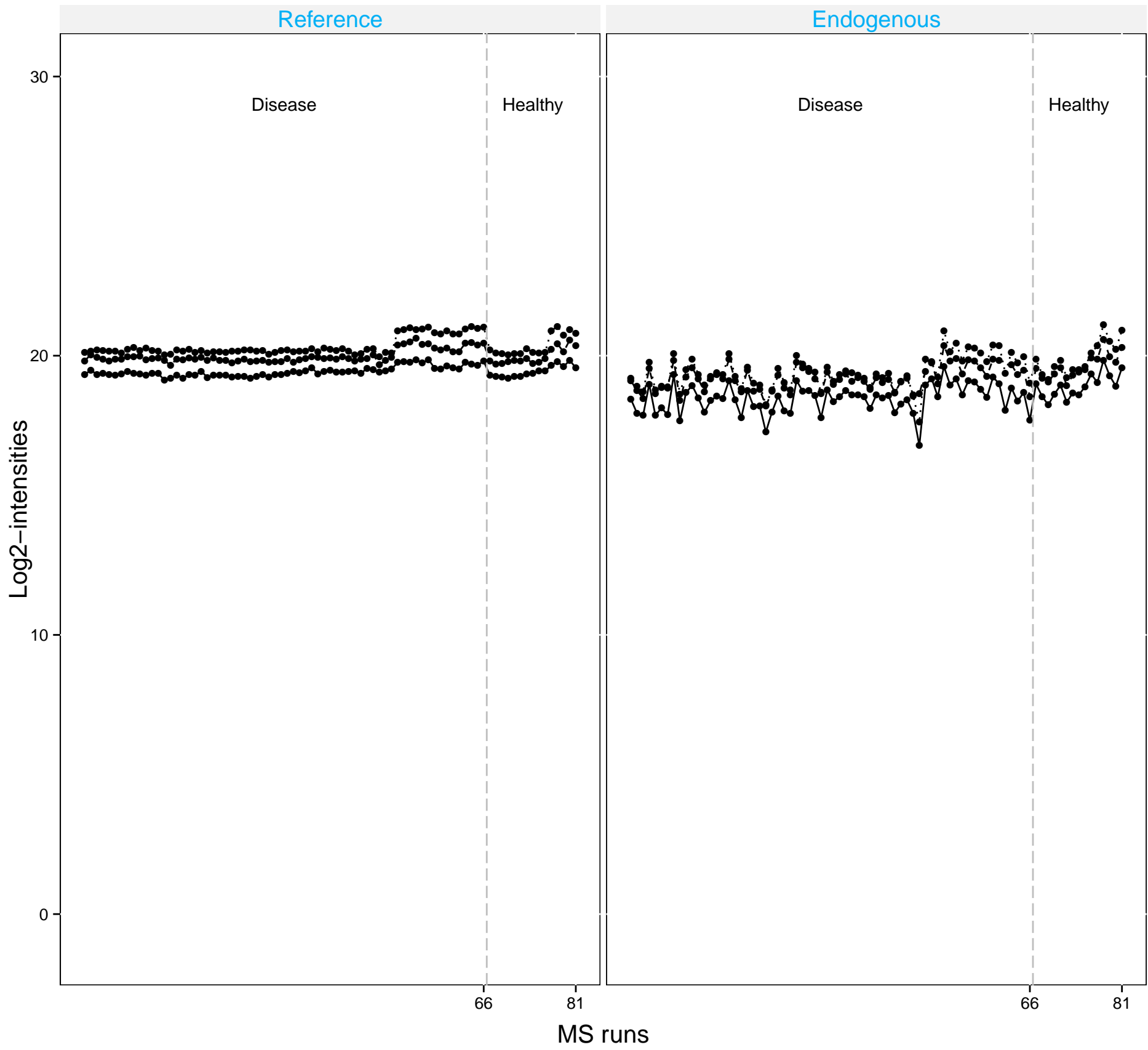
peptide: 3

● FSPATHPSEGLEENYCR_3_b11_2 ● HSIFTPETNPR_2_y6_1 ● QLGAGSIEECAAK_2_y6_1
● FSPATHPSEGLEENYCR_3_y3_1 ● HSIFTPETNPR_2_y7_1 ● QLGAGSIEECAAK_2_y7_1
● FSPATHPSEGLEENYCR_3_y6_1 ● HSIFTPETNPR_2_y8_1 ● QLGAGSIEECAAK_2_y9_1



PON1

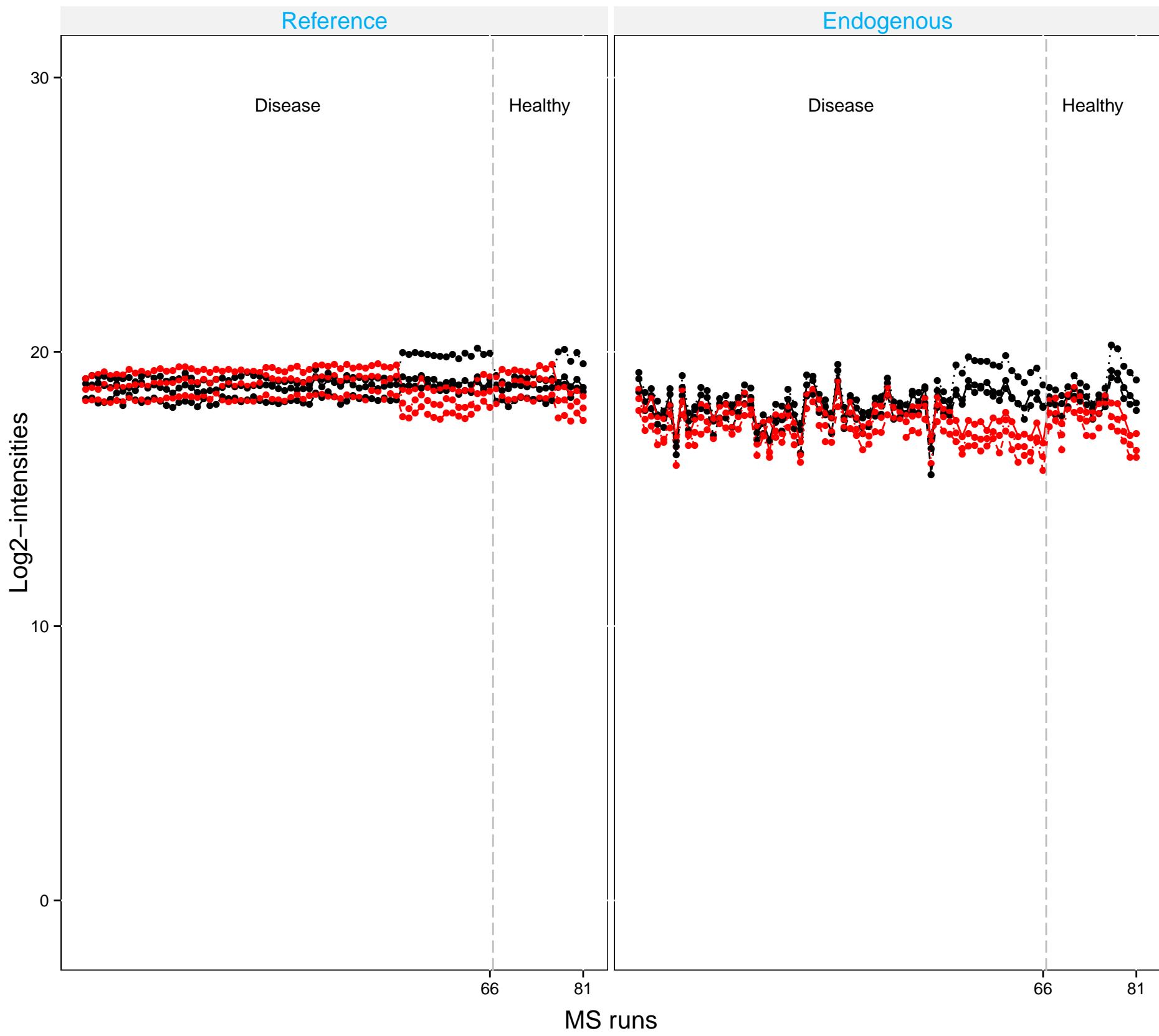
peptide: 1 IQNILTEEPK_2_y5_1 IQNILTEEPK_2_y6_1 IQNILTEEPK_2_y8_1



RET4

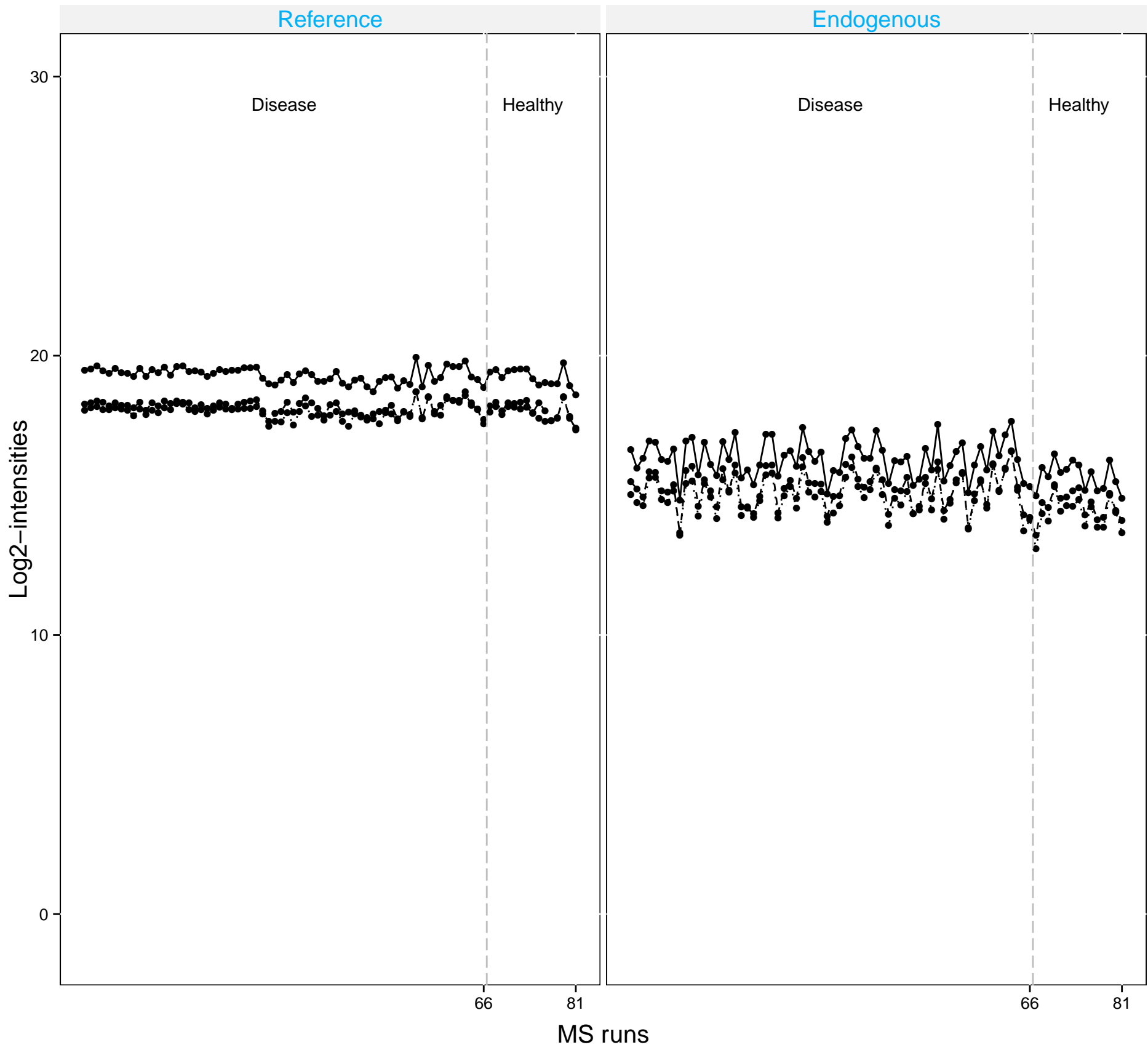
peptide: 2

—●— FSGTWYAMAK_2_y5_1 —●— FSGTWYAMAK_2_y8_1 —●— LIVHNGYCDGR_3_y6_1
—●— FSGTWYAMAK_2_y6_1 —●— LIVHNGYCDGR_3_y4_1 —●— LIVHNGYCDGR_3_y7_1



SHBG

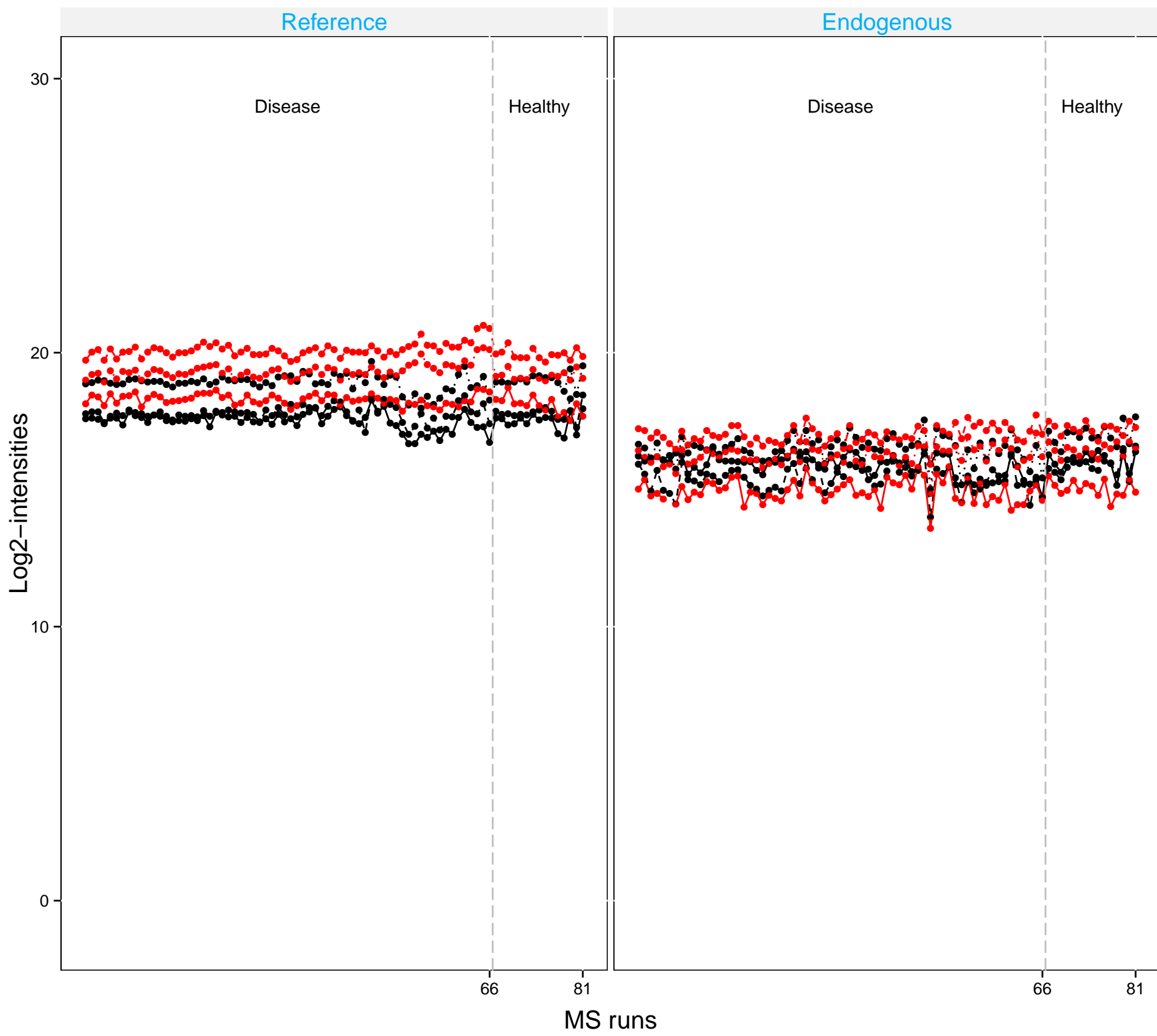
peptide: 1 —●— IALGGLLPASNLR_2_y6_1 —●— IALGGLLPASNLR_2_y7_1 —●— IALGGLLPASNLR_2_y8_1



TETN

peptide: 2

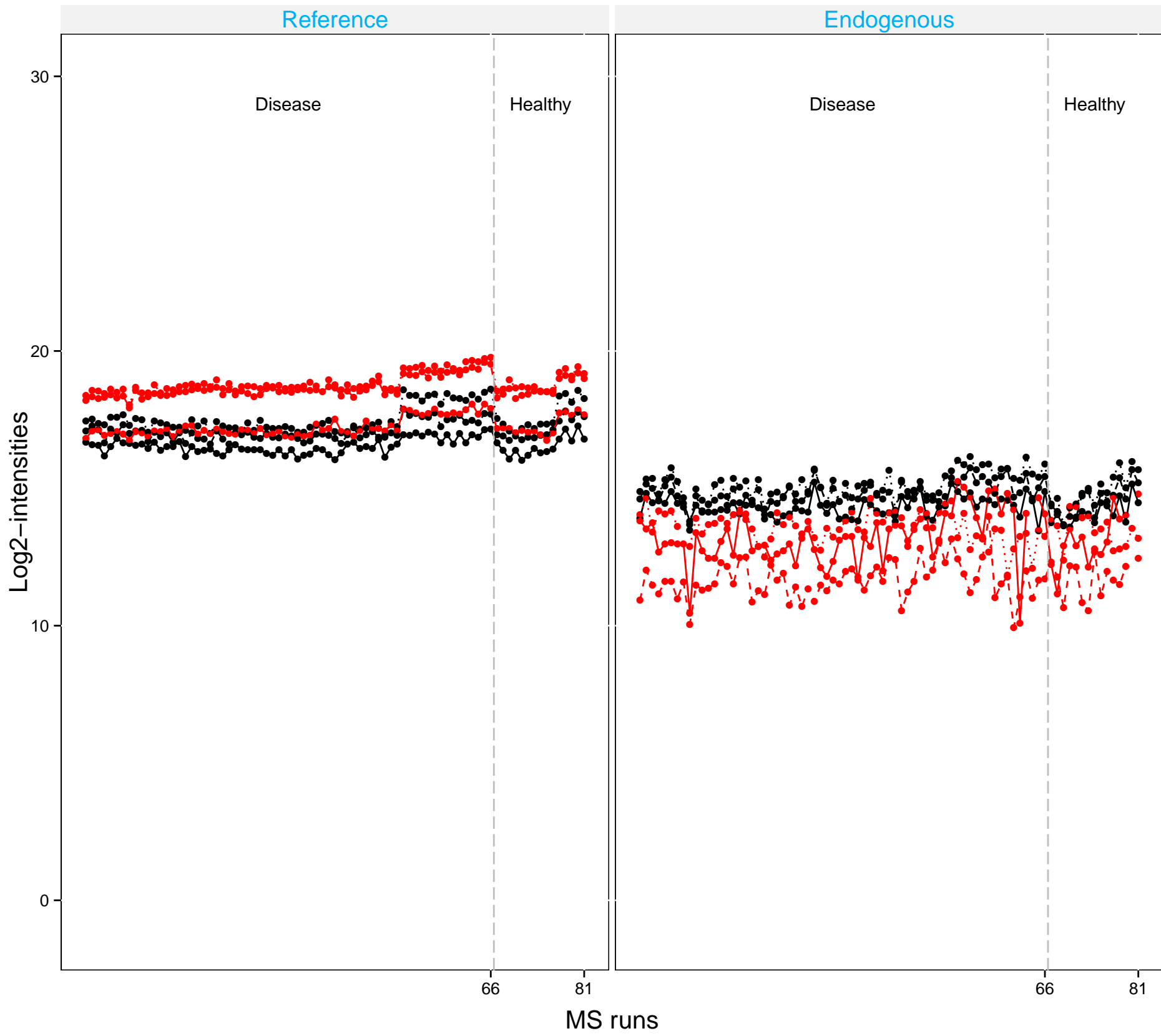
—●— LDTLAQEVALLK_2_y4_1 —●— LDTLAQEVALLK_2_y8_1 —●— TFHEASEDCISR_3_y3_1
—●— LDTLAQEVALLK_2_y7_1 —●— TFHEASEDCISR_3_b3_1 —●— TFHEASEDCISR_3_y4_1



THBG

peptide: 2

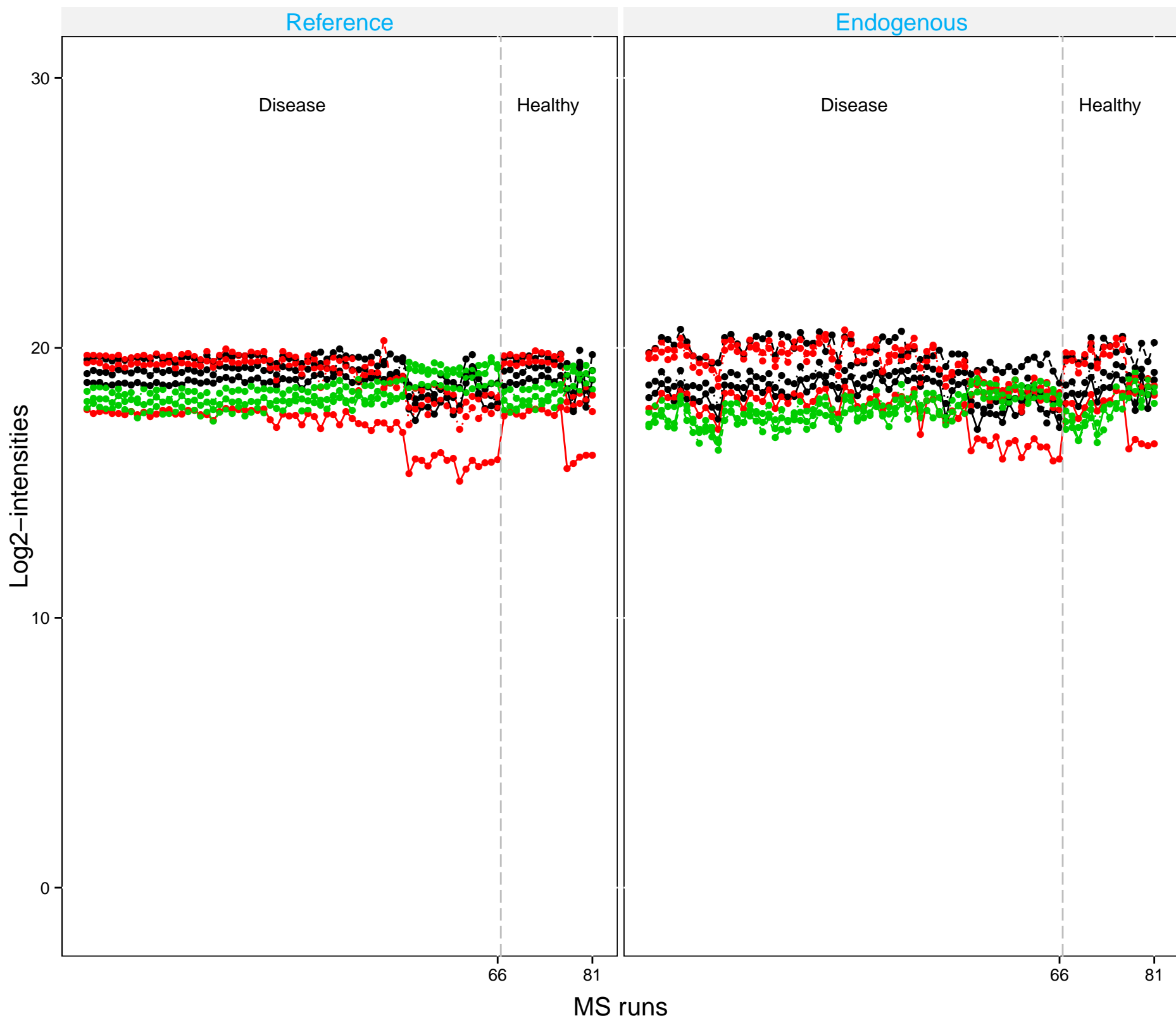
EGQMESVEAAMSSK_2_y4_1 EGQMESVEAAMSSK_2_y9_1 VTACHSSQPNATLYK_3_y7_1
EGQMESVEAAMSSK_2_y7_1 VTACHSSQPNATLYK_3_y6_1 VTACHSSQPNATLYK_3_y7_2



THRB

peptide: 3

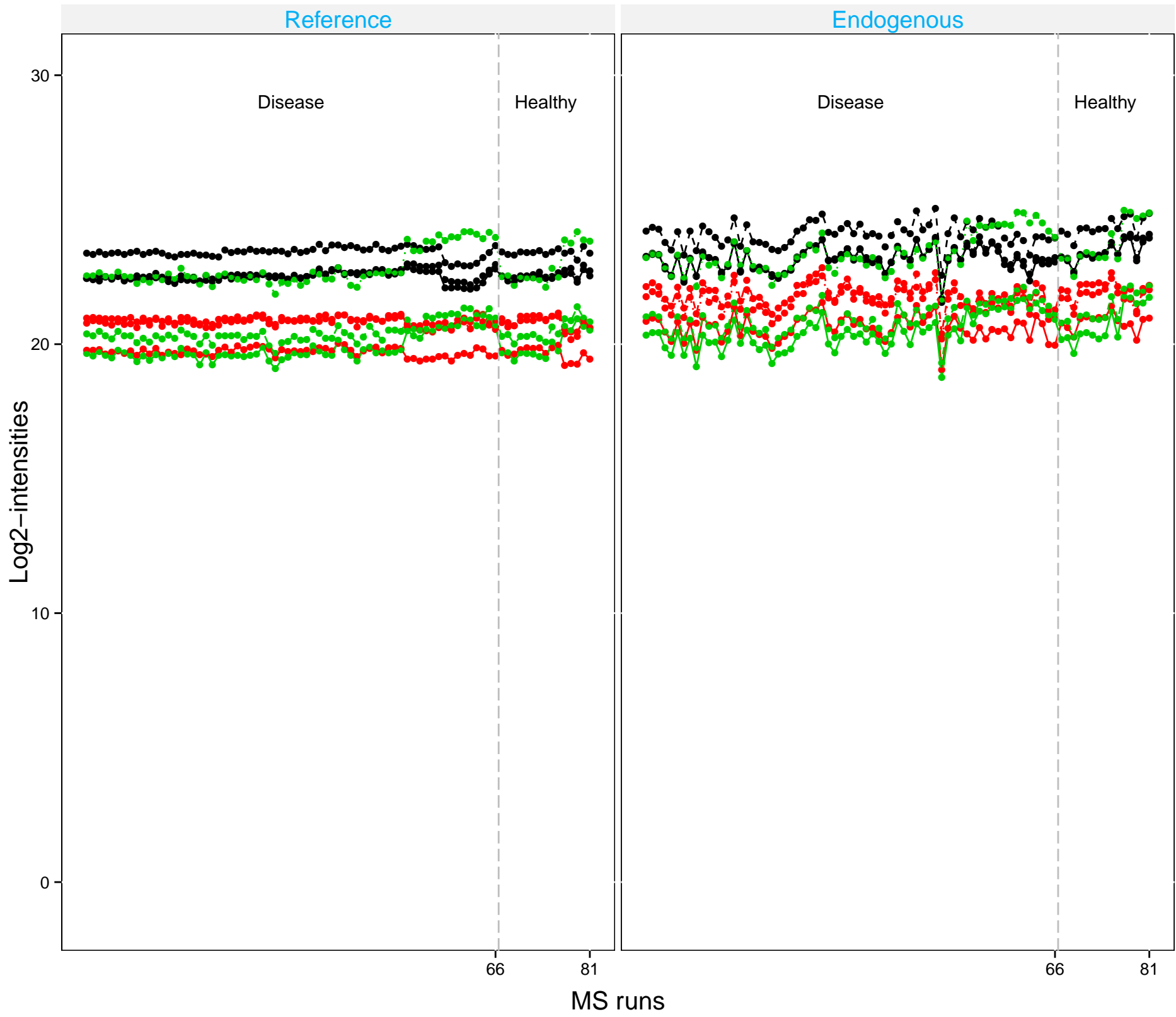
—●— HQDFNSAVQLVENFCR_3_b7_1 —●— LAVTTHGLPCLAWASAAK_3_y4_1 —●— TATSEYQTFNPR_2_y3_1
—●— HQDFNSAVQLVENFCR_3_y5_1 —●— LAVTTHGLPCLAWASAAK_3_y6_1 —●— TATSEYQTFNPR_2_y4_1
—●— HQDFNSAVQLVENFCR_3_y6_1 —●— LAVTTHGLPCLAWASAAK_3_y7_1 —●— TATSEYQTFNPR_2_y7_1



TRFE

peptide: 3

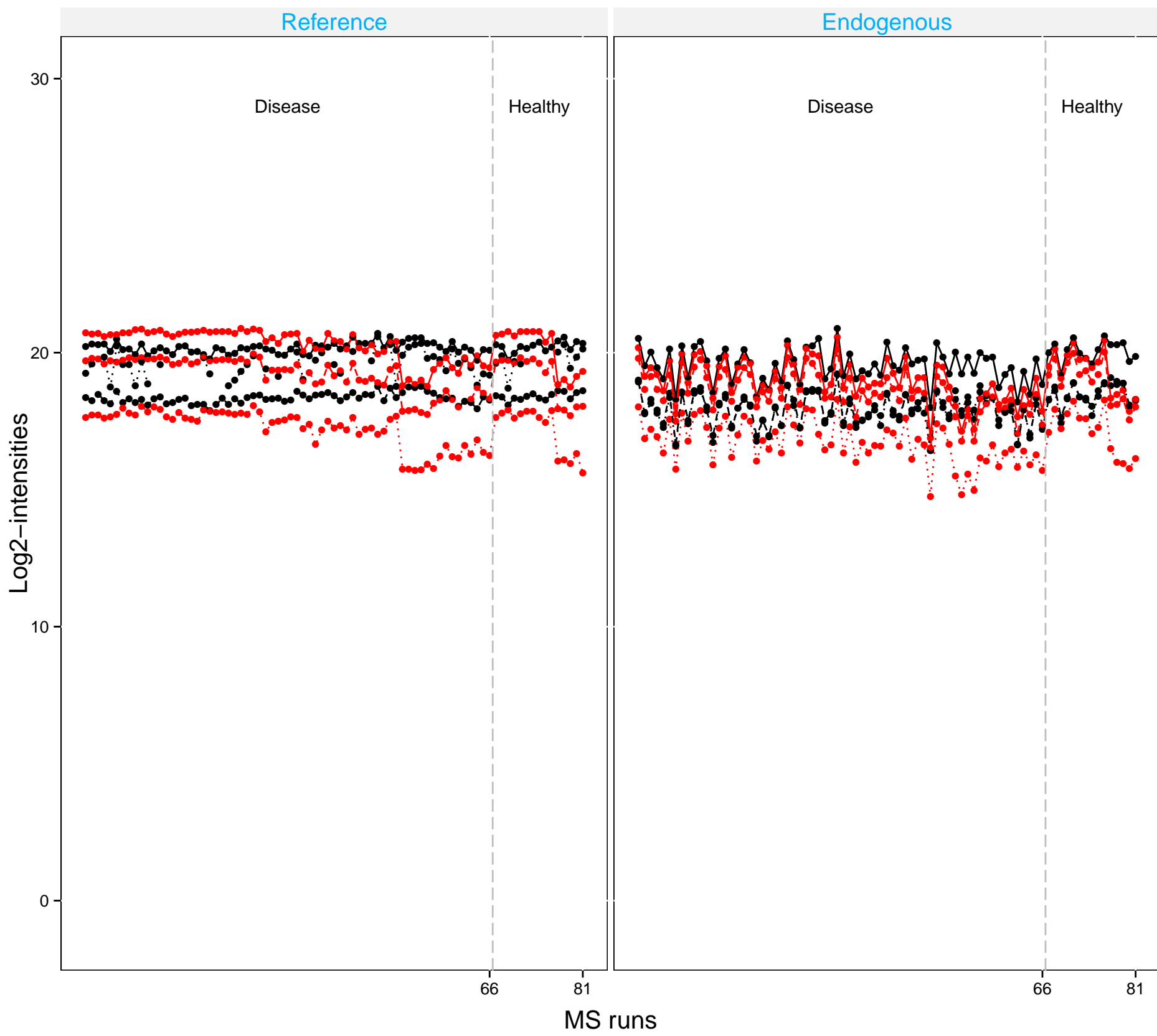
EGYYGYTGAFR_2_y5_1 MYLGYEYVTAIR_2_b3_1 TAGWNIPMGLLYNK_2_b5_1
EGYYGYTGAFR_2_y7_1 MYLGYEYVTAIR_2_y6_1 TAGWNIPMGLLYNK_2_y3_1
EGYYGYTGAFR_2_y8_1 MYLGYEYVTAIR_2_y7_1 TAGWNIPMGLLYNK_2_y8_1



TTHY

peptide: 2

—●— AADDTWEPFASGK_2_y6_1 —●— AADDTWEPFASGK_2_y8_1 —●— GSPAINVAVHVFR_3_y5_1
—●— AADDTWEPFASGK_2_y7_1 —●— GSPAINVAVHVFR_3_y4_1 —●— GSPAINVAVHVFR_3_y7_2



VTNC

peptide: 4

DVWGIEGPIDAAFTR_2_b4_1 FEDGVLPDPYPR_2_y5_1 GQYCYELDEK_2_y7_1
DVWGIEGPIDAAFTR_2_y8_1 FEDGVLPDPYPR_2_y7_1 VDTVDPYPYPR_2_y3_1
DVWGIEGPIDAAFTR_2_y9_1 GQYCYELDEK_2_y3_1 VDTVDPYPYPR_2_y4_1
FEDGVLPDPYPR_2_y3_1 GQYCYELDEK_2_y6_1 VDTVDPYPYPR_2_y5_1

