A2MG

AACT

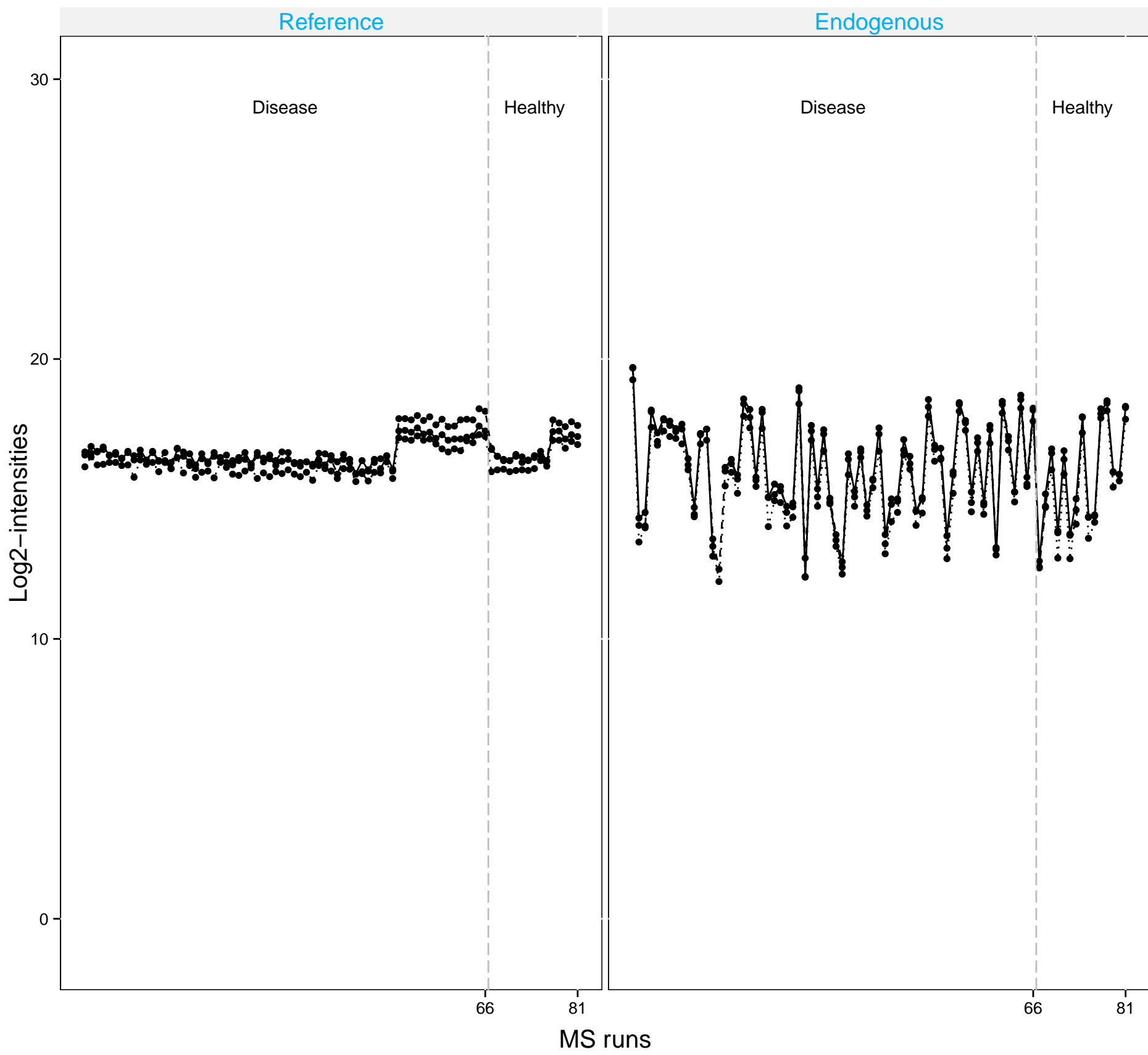**# peptide: 1** ● EIGELYLPK_2_y4_1 ● EIGELYLPK_2_y5_1 ● EIGELYLPK_2_y7_1
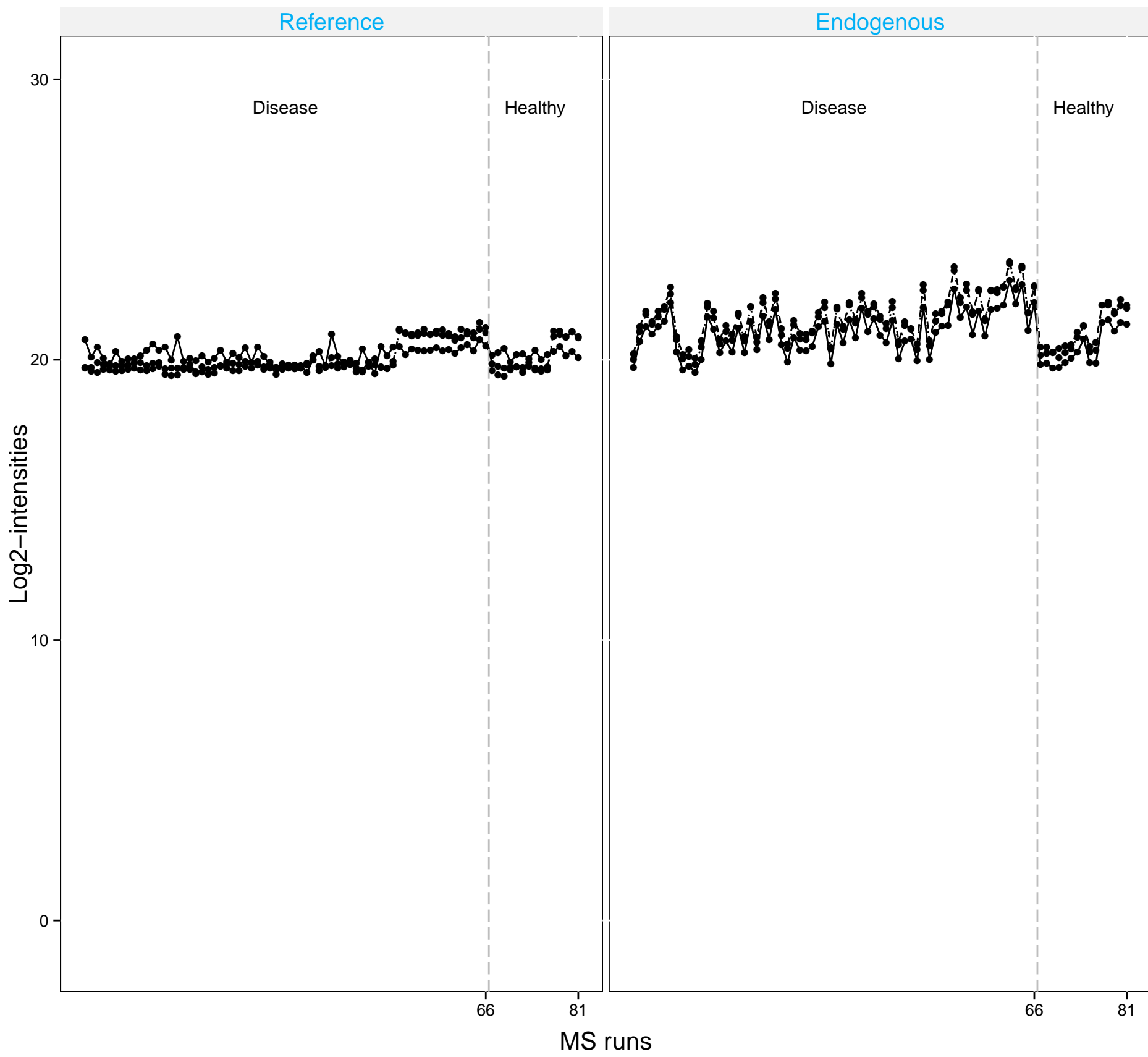
APOA

 # peptide: 1 ●— NPDAVAAPYCYTR_2_b4_1 ●— NPDAVAAPYCYTR_2_y6_1 ●— NPDAVAAPYCYTR_2_y7_1
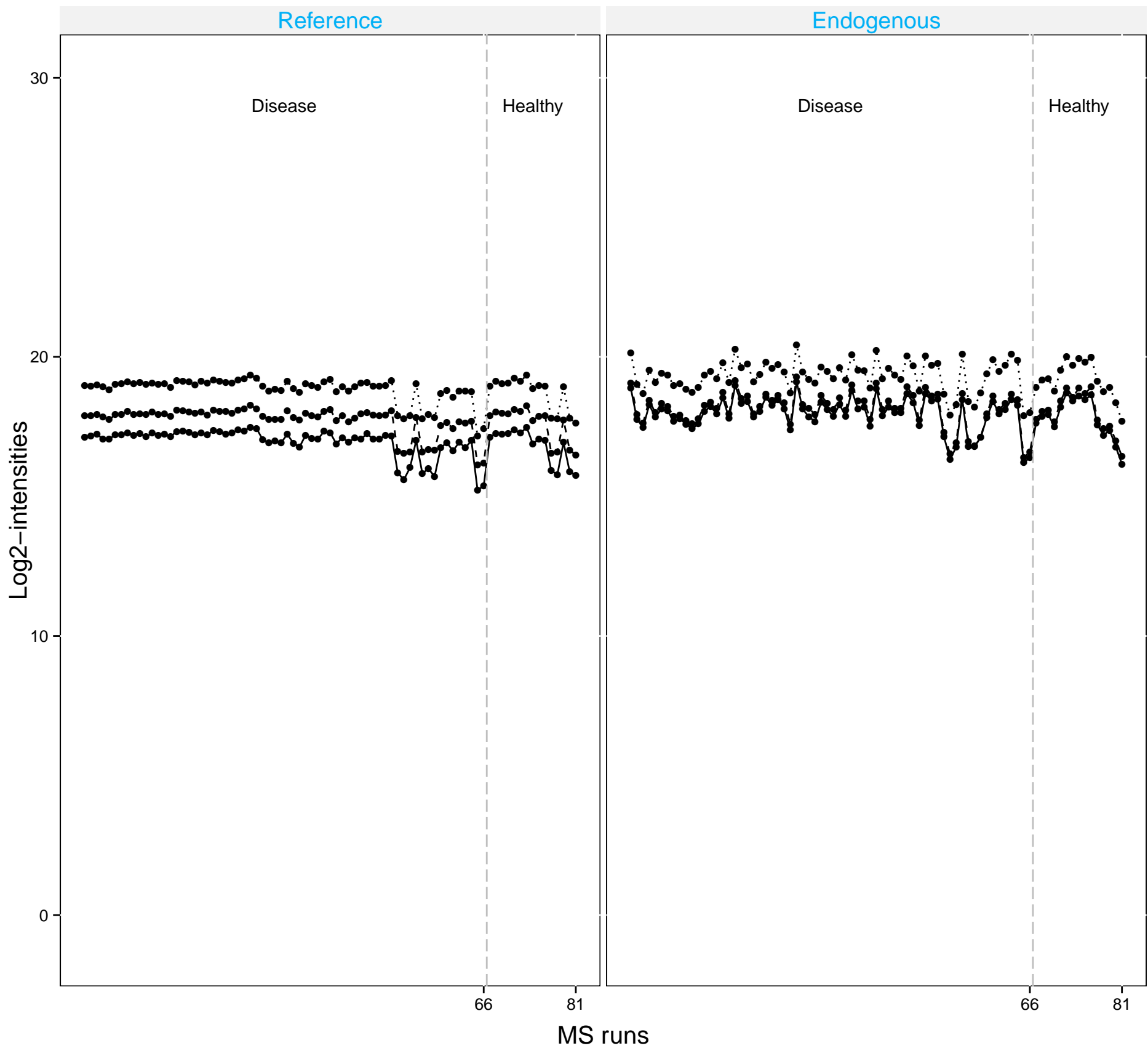
APOA1
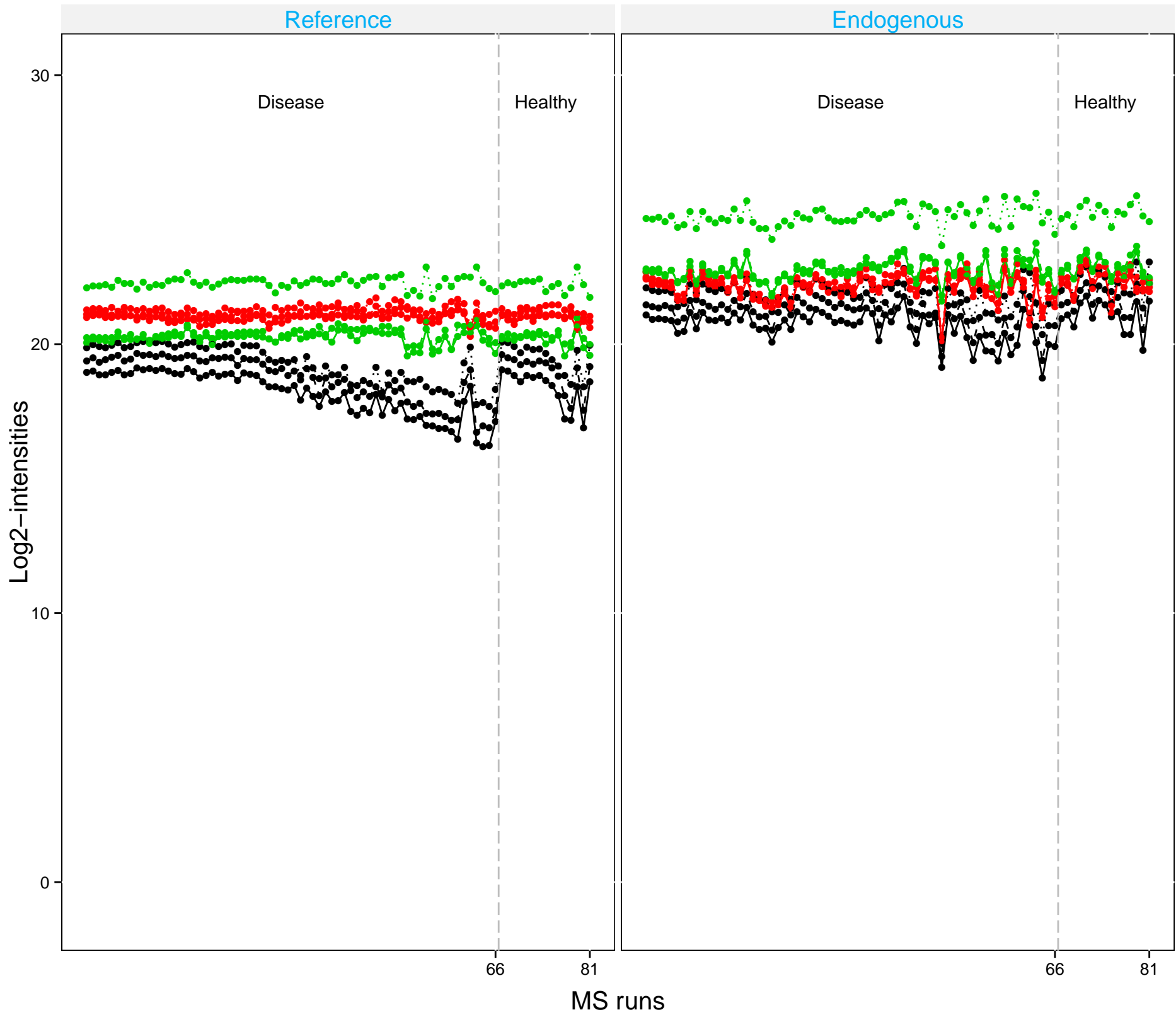
APOB

peptide: 1 —●— FSVPAGIVIPSFQALTAR_3_y4_1 —●— FSVPAGIVIPSFQALTAR_3_y5_1 —●— FSVPAGIVIPSFQALTAR_3_y9_1

APOE

peptide: 3

Legend:
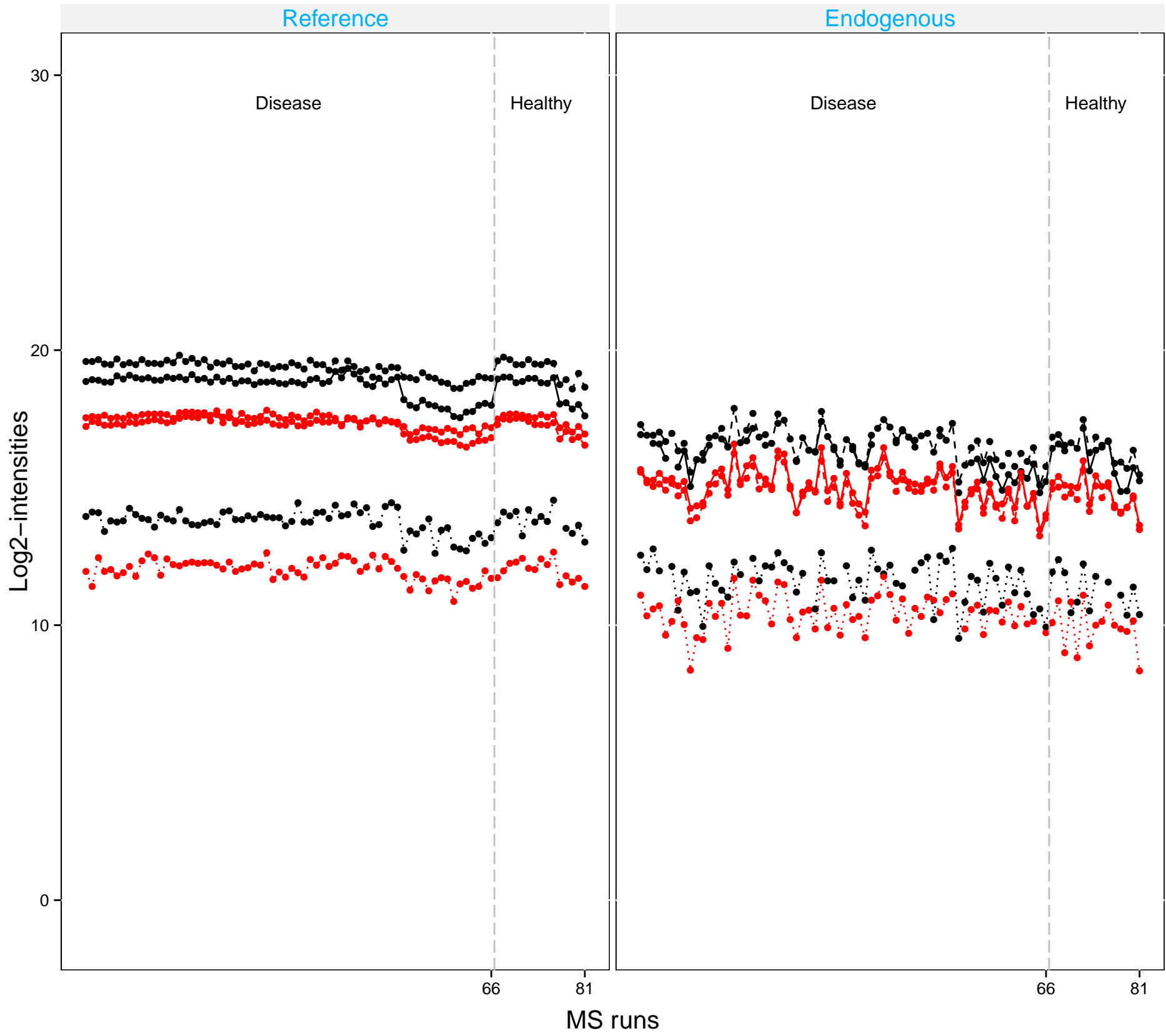- AATVGSLAGQPLQER_2_y5_1
- AATVGSLAGQPLQER_2_y7_1
- AATVGSLAGQPLQER_2_y8_1
- SELEEQLTPVAEETR_2_y3_1
- SELEEQLTPVAEETR_2_y7_1
- SELEEQLTPVAEETR_2_y8_1
- SWFEPLVEDMQR_2_y3_1
- SWFEPLVEDMQR_2_y6_1
- SWFEPLVEDMQR_2_y8_1

Reference | Endogenous

Disease | Healthy

Log2−intensities

MS runs
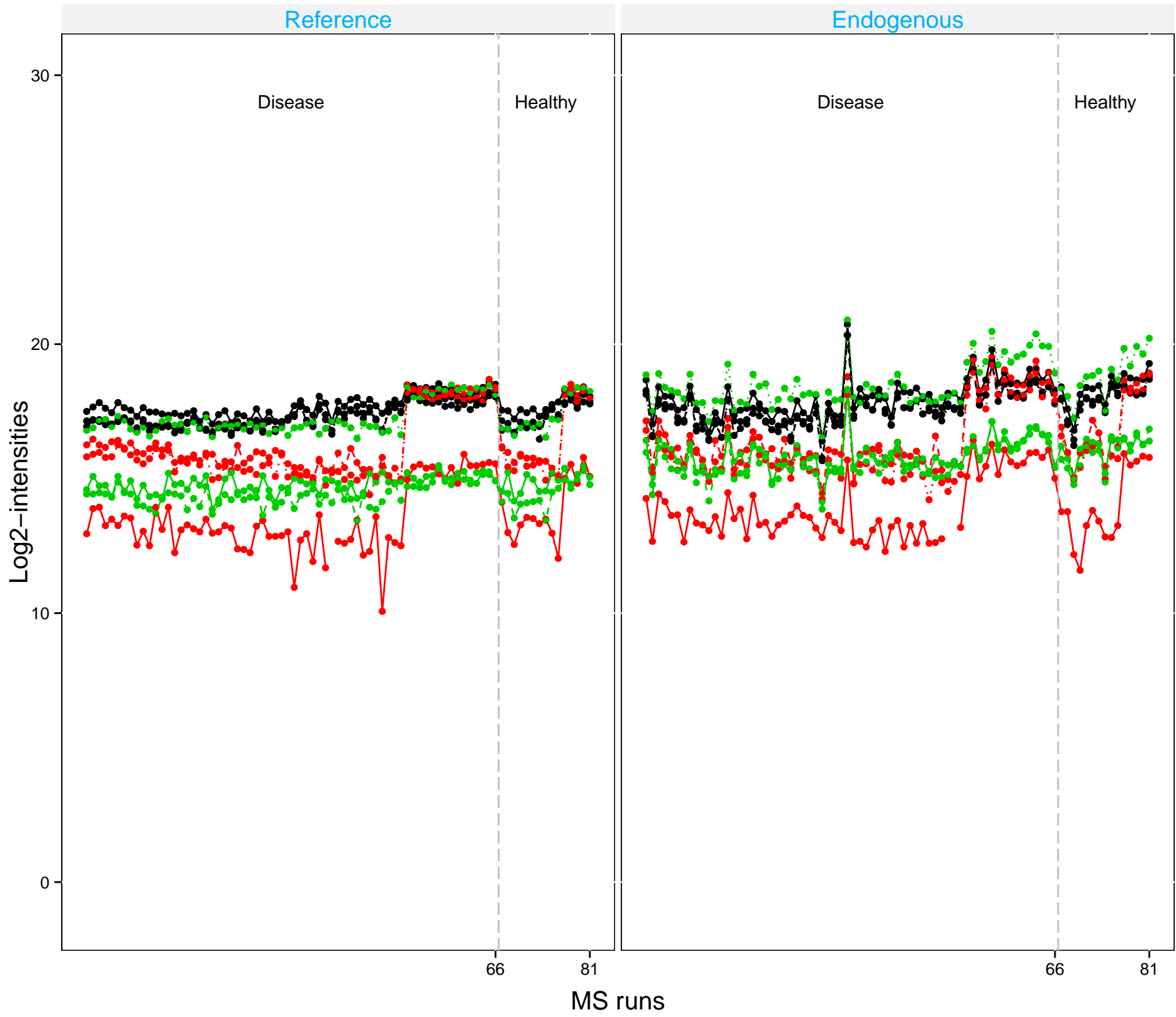
B2MG

Legend:
- VEHSDLSFSK_3_y3_1
- VEHSDLSFSK_3_y4_1
- VEHSDLSFSK_3_y7_1
- VNHVTLSQPK_3_y5_1
- VNHVTLSQPK_3_y6_1
- VNHVTLSQPK_3_y7_1
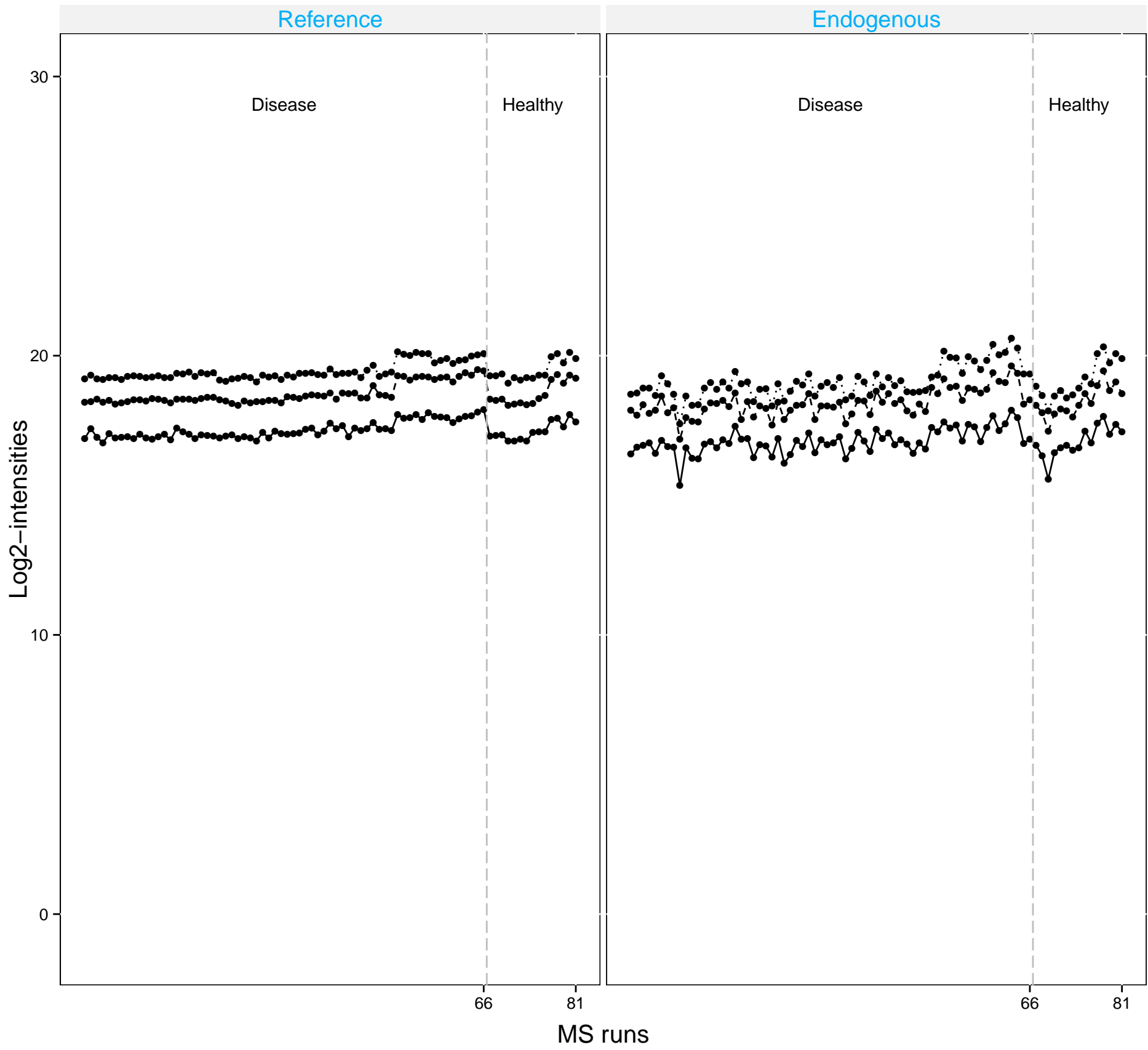
 # peptide: 2

Reference — Disease | Healthy

Endogenous — Disease | Healthy

Log2-intensities

MS runs

C1QA

C1QB

C1QC
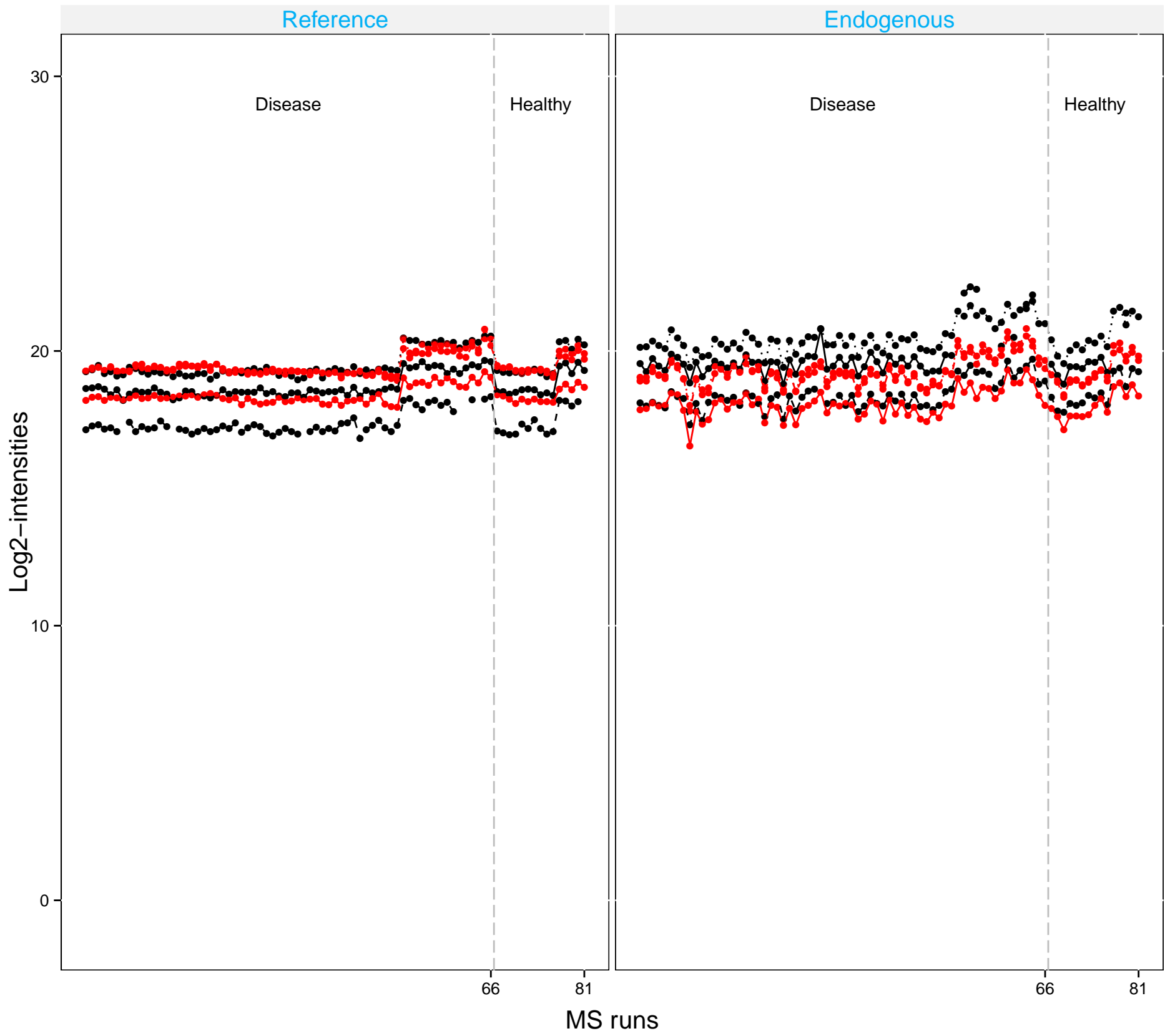
C1R

peptide: 1   ●— YTTTMGVNTYK_2_y6_1   ●— YTTTMGVNTYK_2_y7_1   ●— YTTTMGVNTYK_2_y8_1
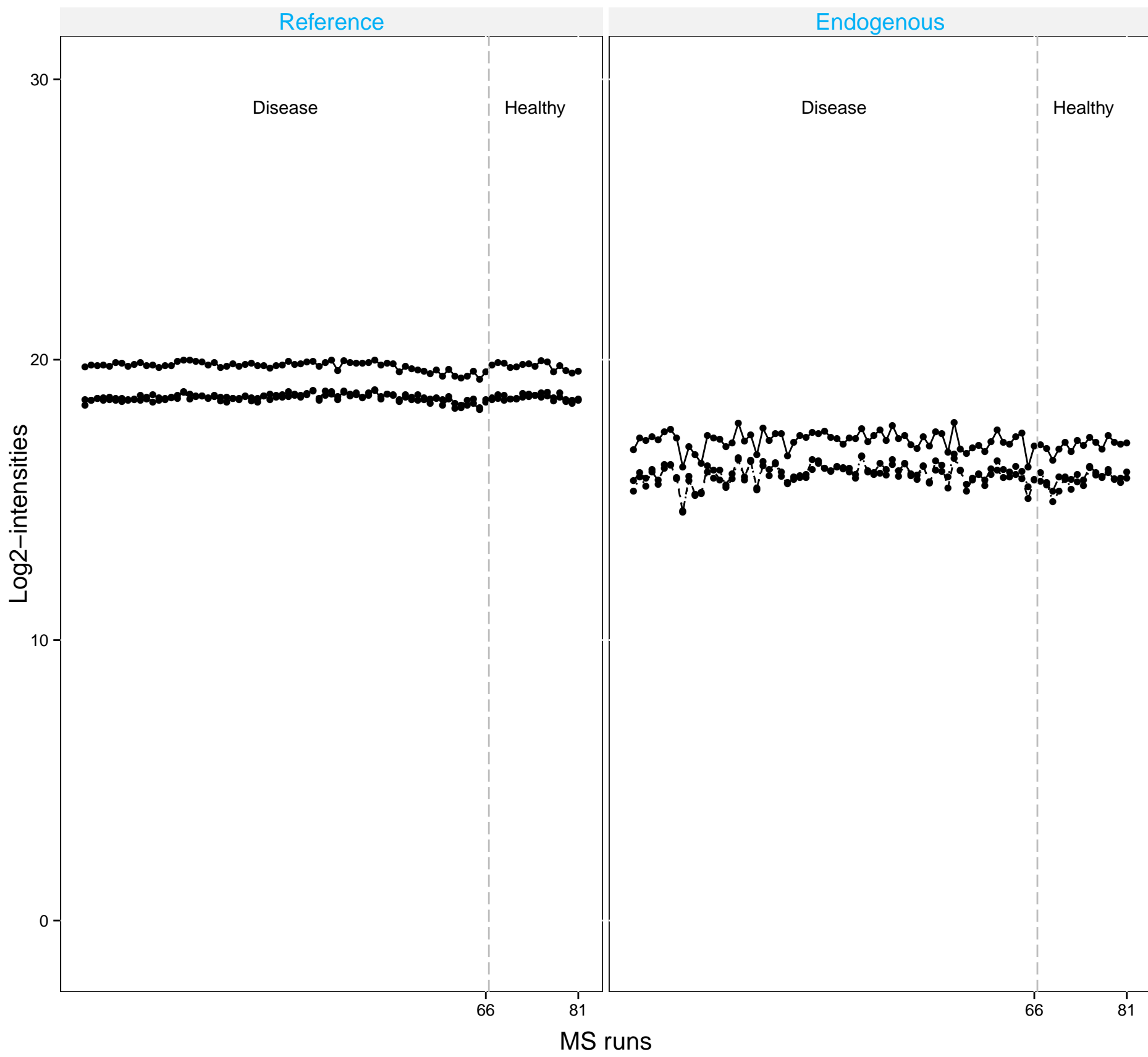
CADH5
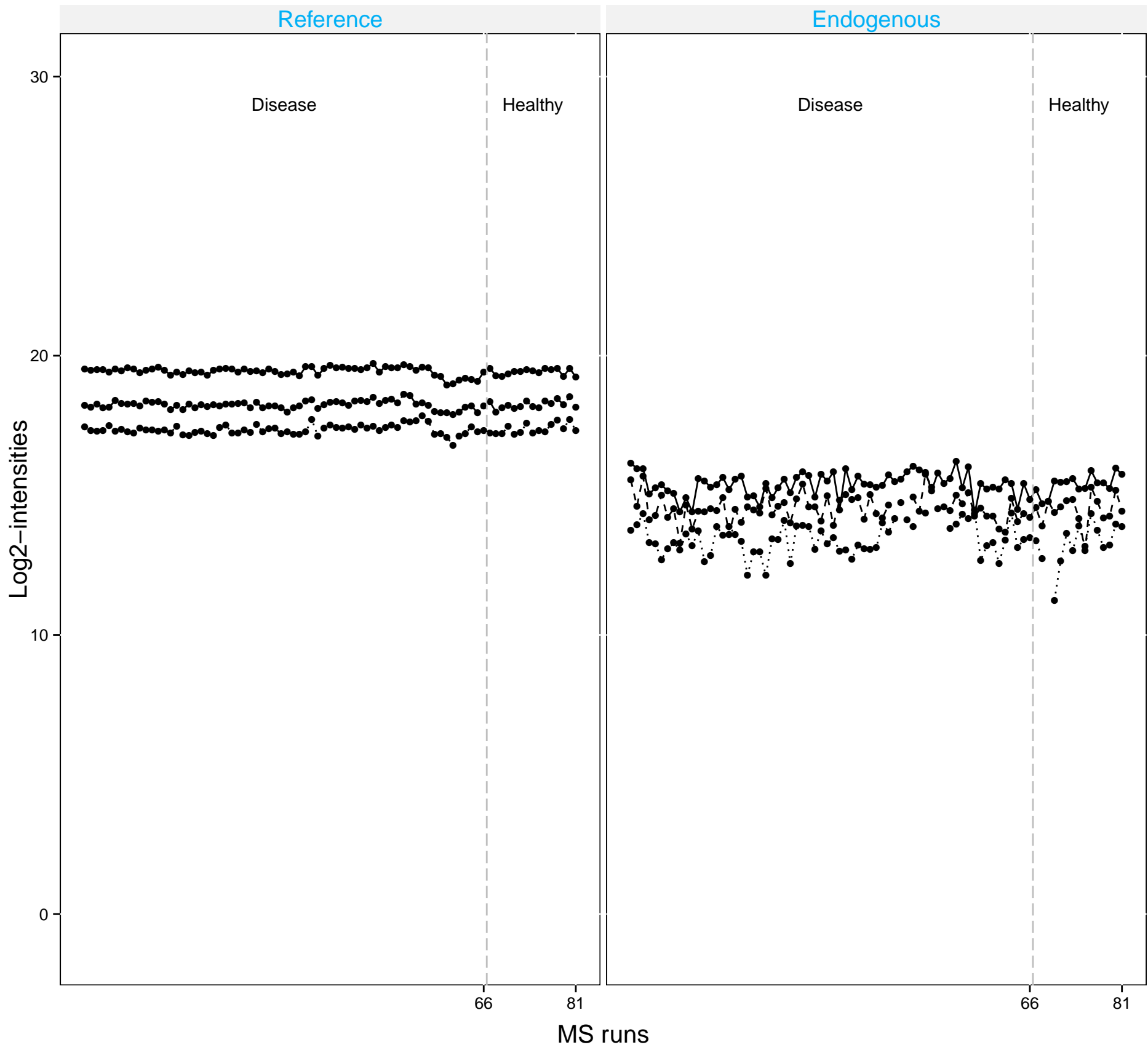
CCNH

 # peptide: 1 — HWTFSSEEQLAR_3_b4_1 — HWTFSSEEQLAR_3_y4_1 — HWTFSSEEQLAR_3_y5_1
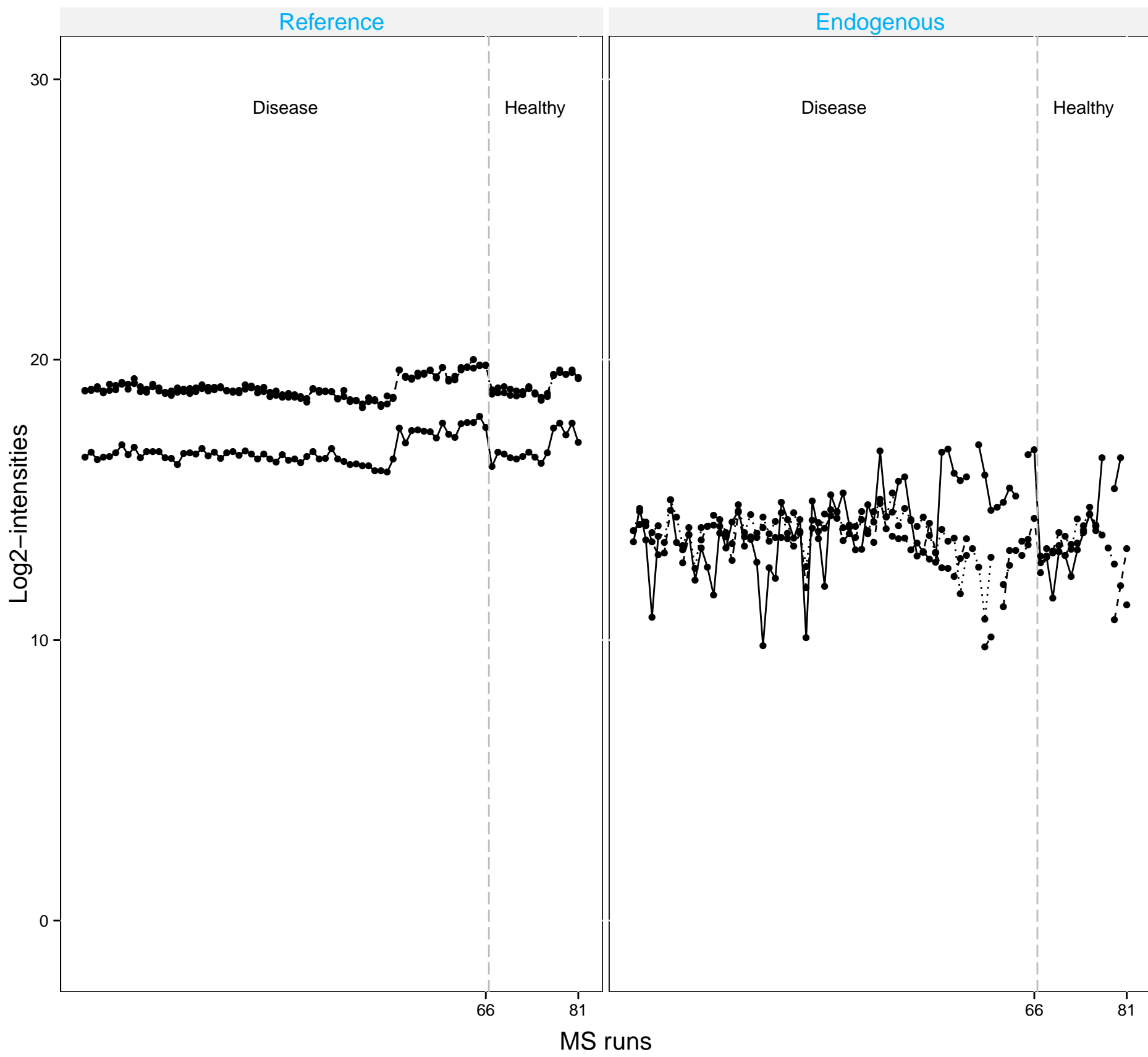
CFAB

CLUS

CO3
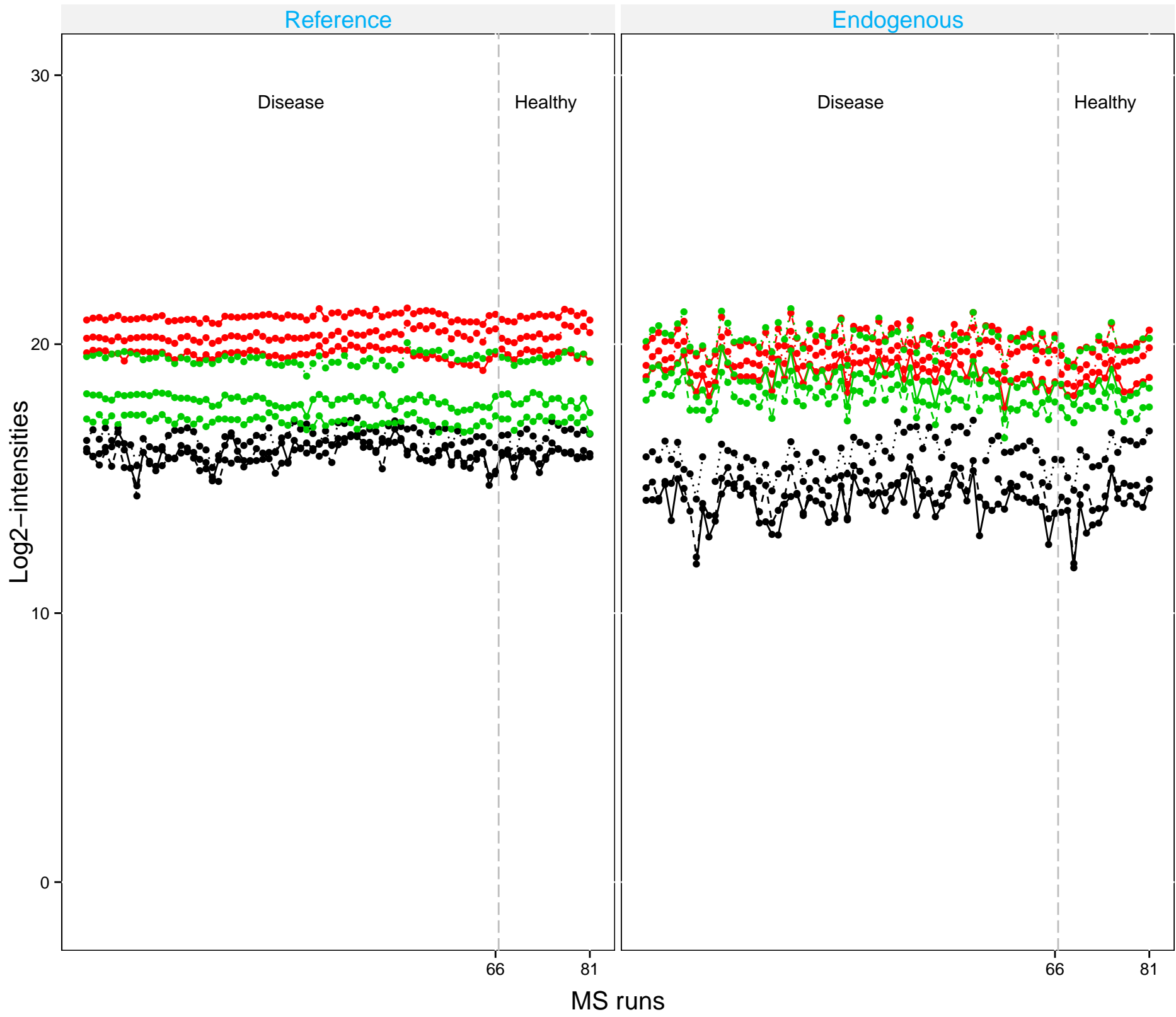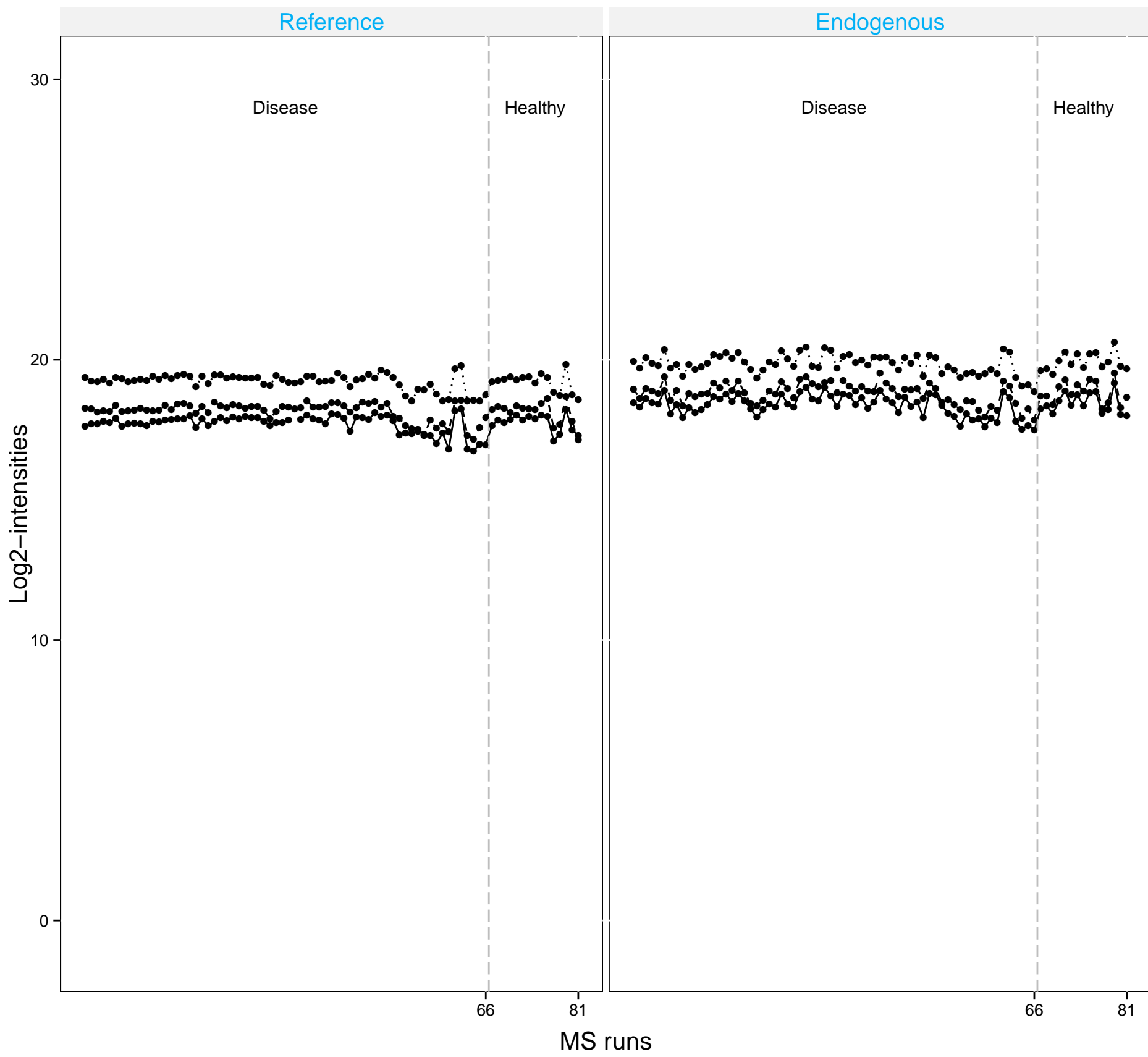
Legend:
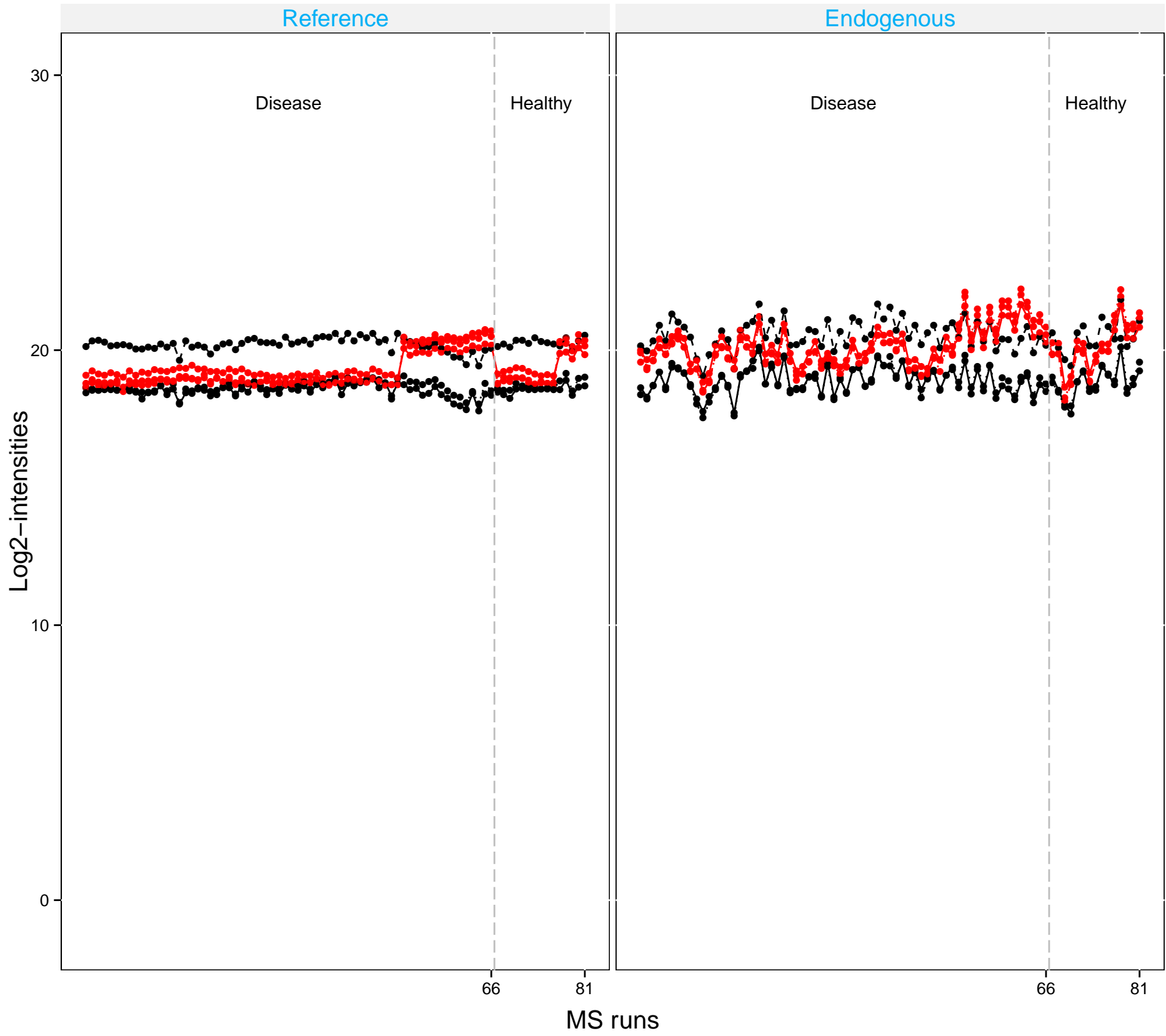- IHWESASLLR_3_b3_1
- IHWESASLLR_3_y4_1
- ILLQGTPVAQMTEDAVDAER_3_y7_1
- IHWESASLLR_3_y3_1
- ILLQGTPVAQMTEDAVDAER_3_y5_1
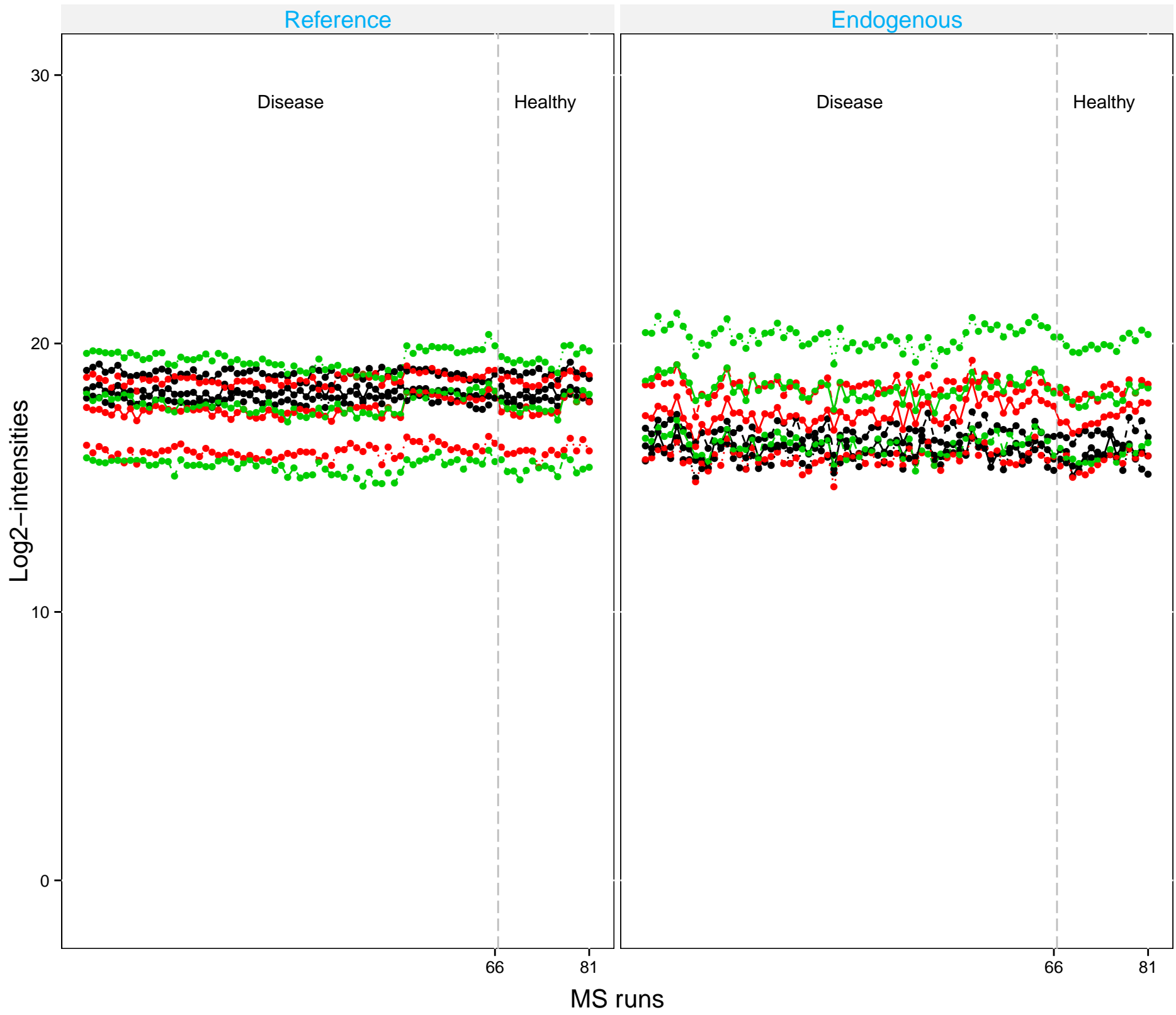- ILLQGTPVAQMTEDAVDAER_3_y8_1

 # peptide: 2

Reference — Disease / Healthy

Endogenous — Disease / Healthy

Log2-intensities

MS runs
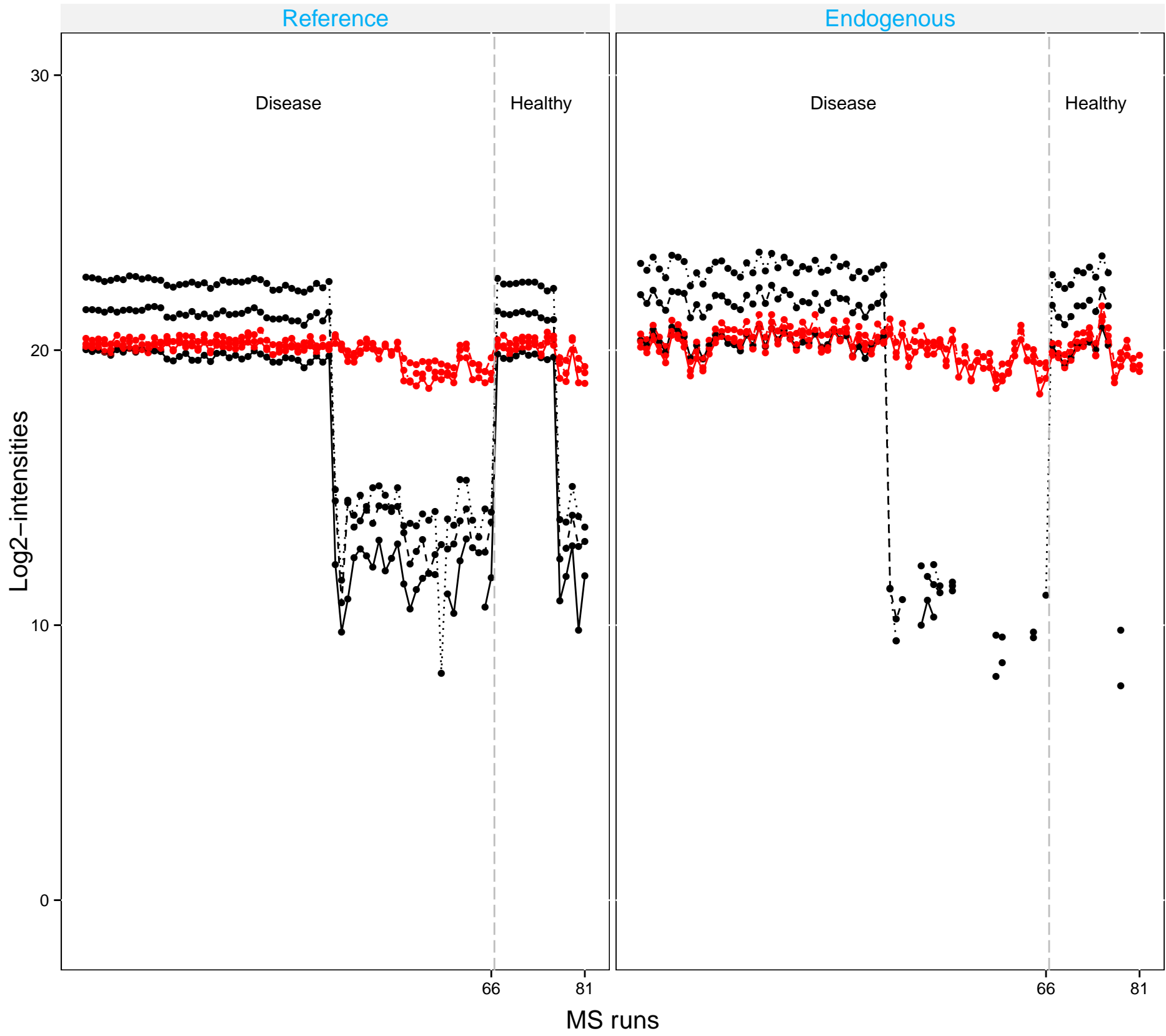
CO4A

Legend:
- GSFEFPVGDAVSK_2_b4_1
- GSFEFPVGDAVSK_2_y8_1
- GSFEFPVGDAVSK_2_y9_1
- LGQYASPTAK_2_y4_1
- LGQYASPTAK_2_y6_1
- LGQYASPTAK_2_y7_1

peptide: 2

Reference | Endogenous

Disease / Healthy

Log2-intensities

MS runs

CO5

CO6
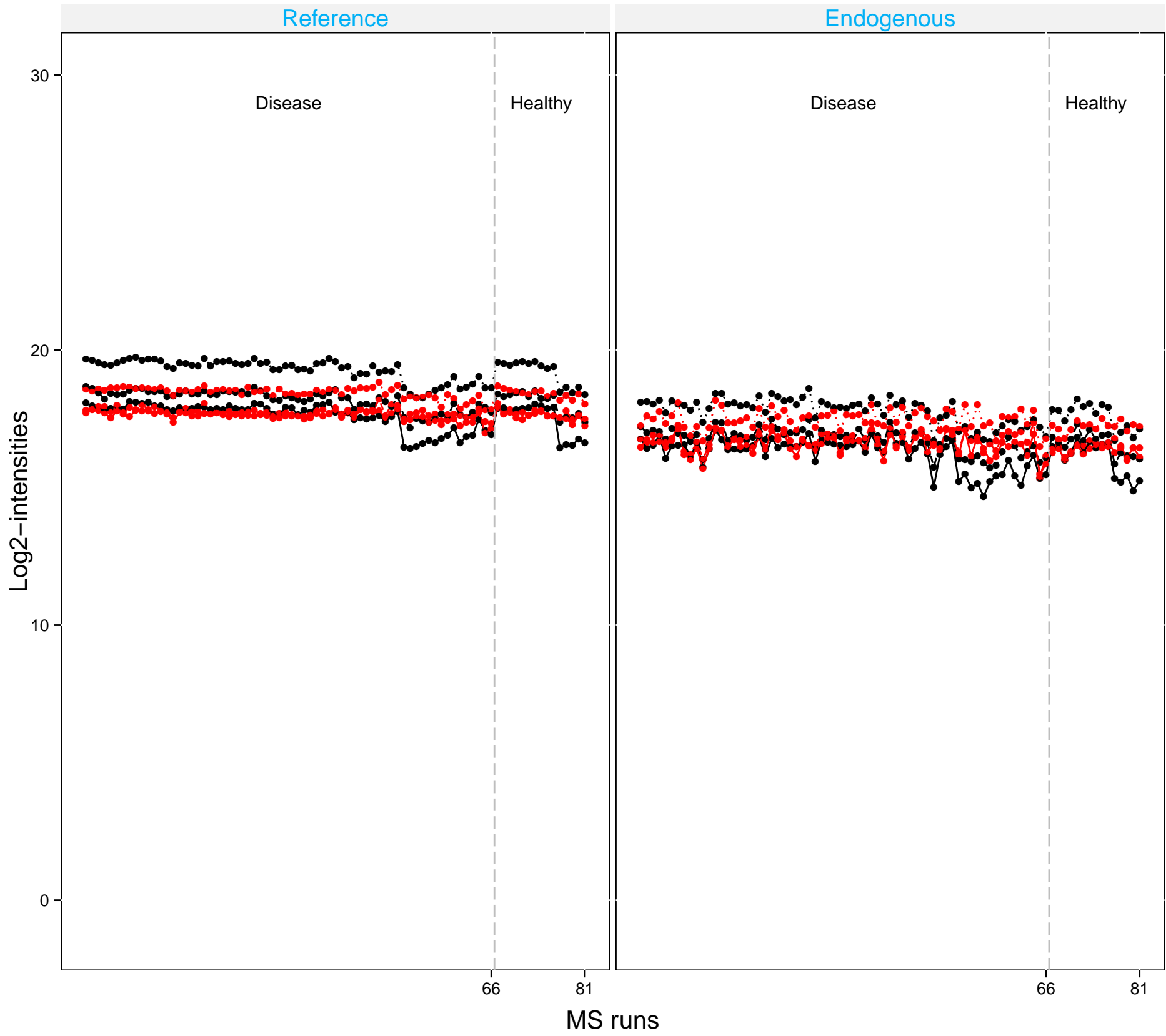
CO7

peptide: 1 ——●— YSAWAESVTNLPQVIK_2_y5_1 ——●— YSAWAESVTNLPQVIK_2_y7_1 ——●— YSAWAESVTNLPQVIK_2_y8_1
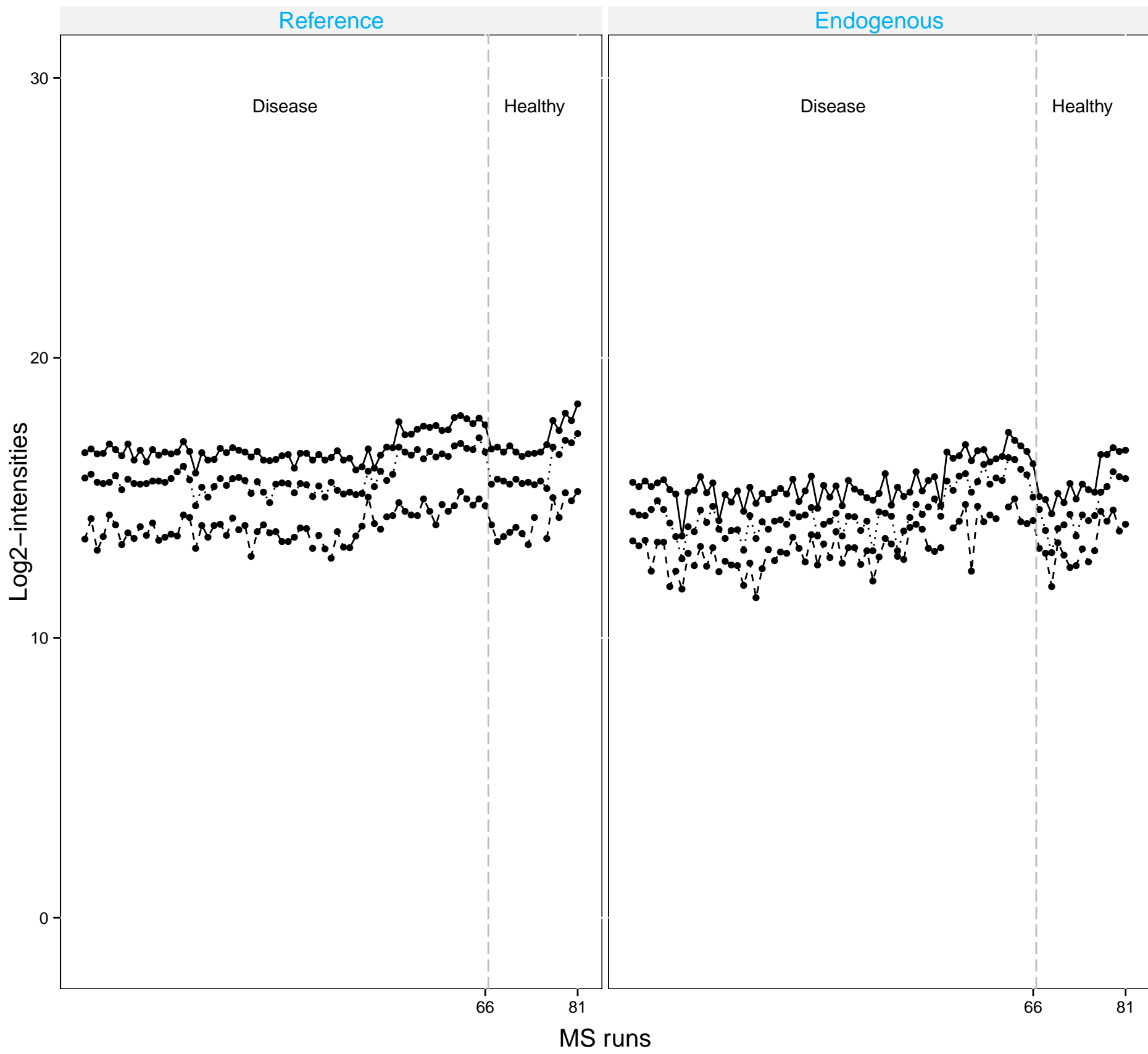
FN1

GELS

HPT
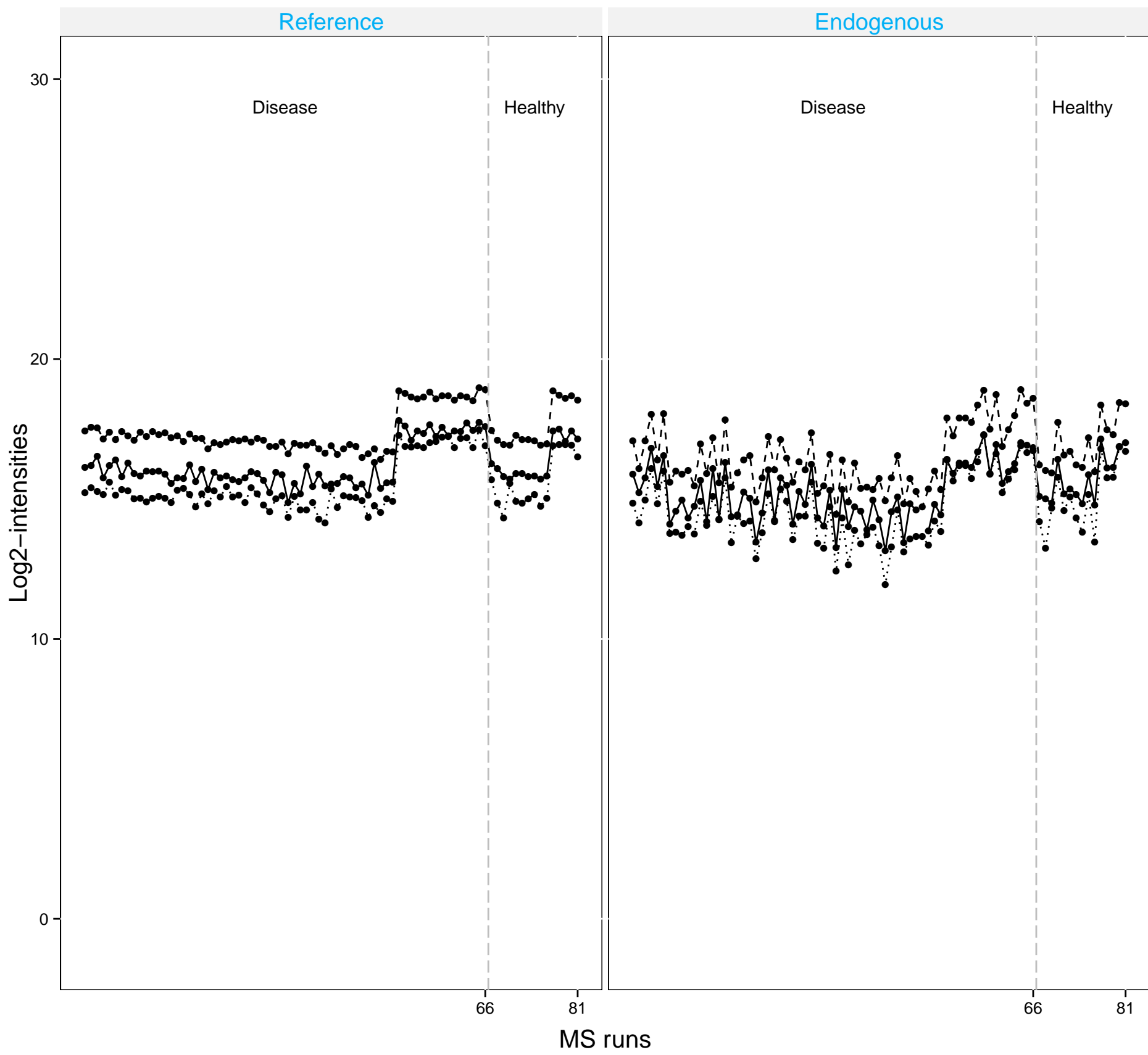
 # peptide: 4
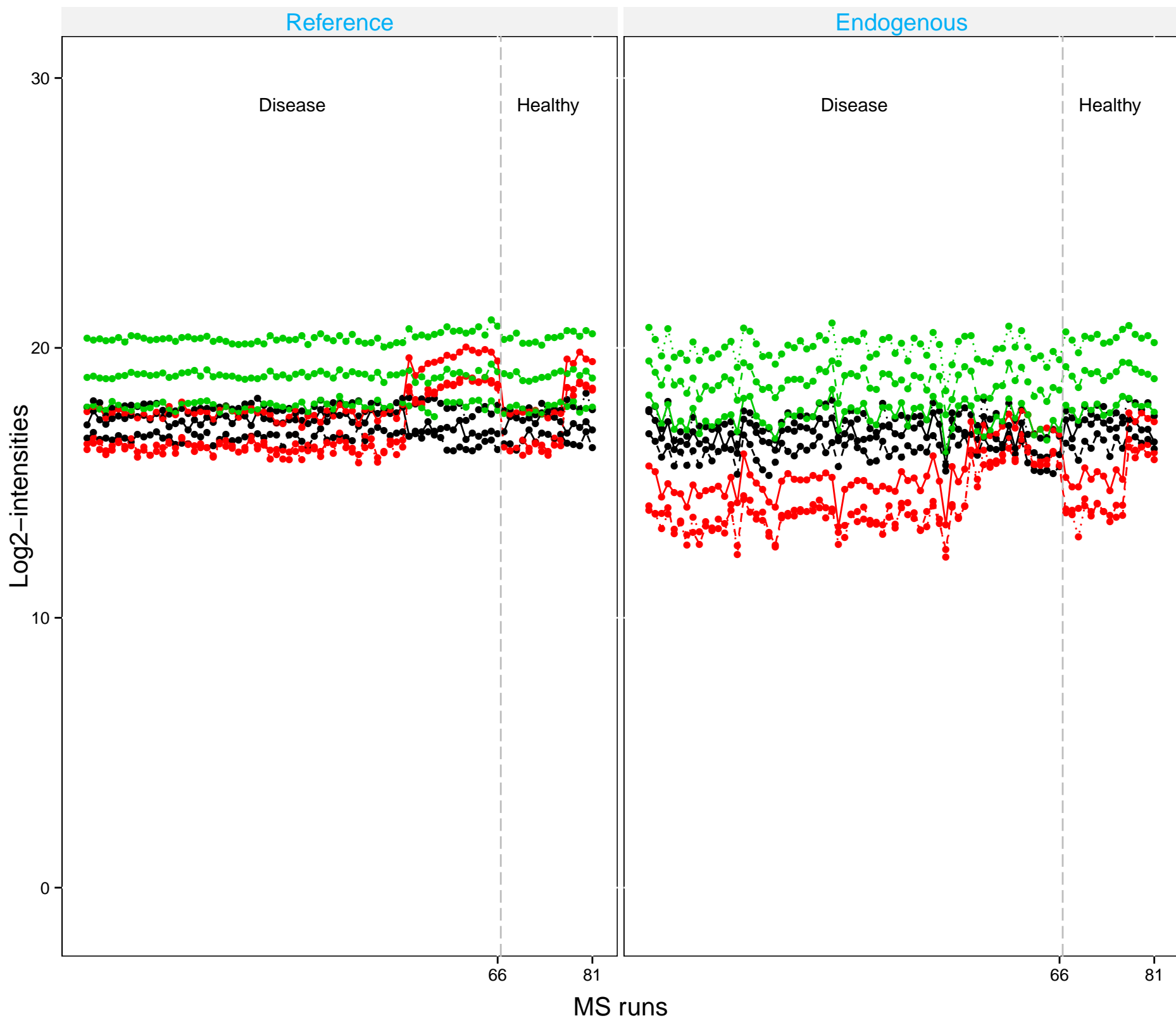
Legend:
- DIAPTLTLYVGK_2_y6_1
- DIAPTLTLYVGK_2_y9_1
- DIAPTLTLYVGK_2_y9_2
- VTSIQDWVQK_2_y4_1
- VTSIQDWVQK_2_y5_1
- VTSIQDWVQK_2_y6_1
- VVLHPNYSQVDIGLIK_3_y4_1
- VVLHPNYSQVDIGLIK_3_y5_1
- VVLHPNYSQVDIGLIK_3_y6_1
- YVMLPVADQDQCIR_2_b4_1
- YVMLPVADQDQCIR_2_y4_1
- YVMLPVADQDQCIR_2_y7_1
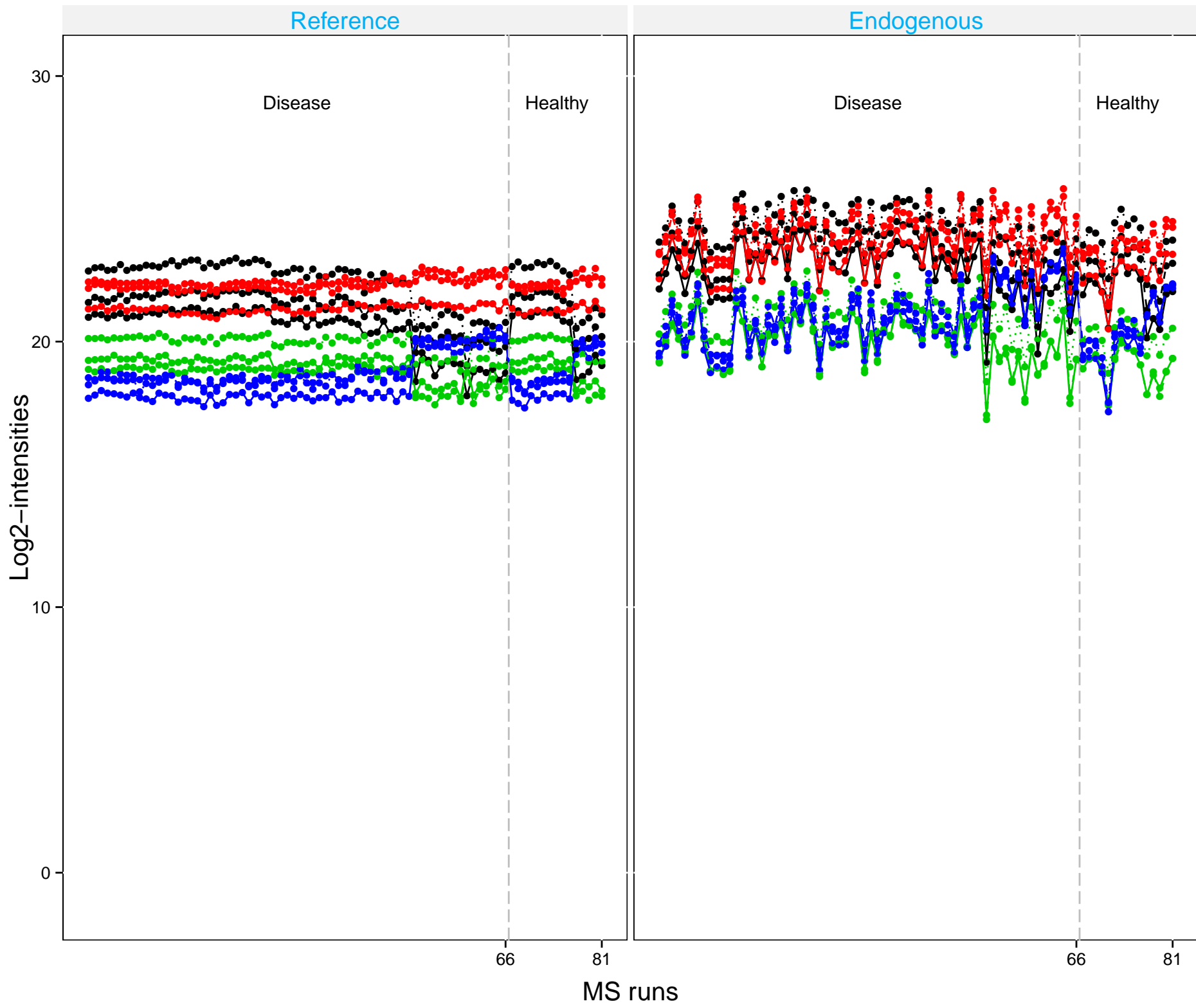
Reference | Endogenous

Disease | Healthy

Log2-intensities

MS runs

IC1

IGF2

ITIH4
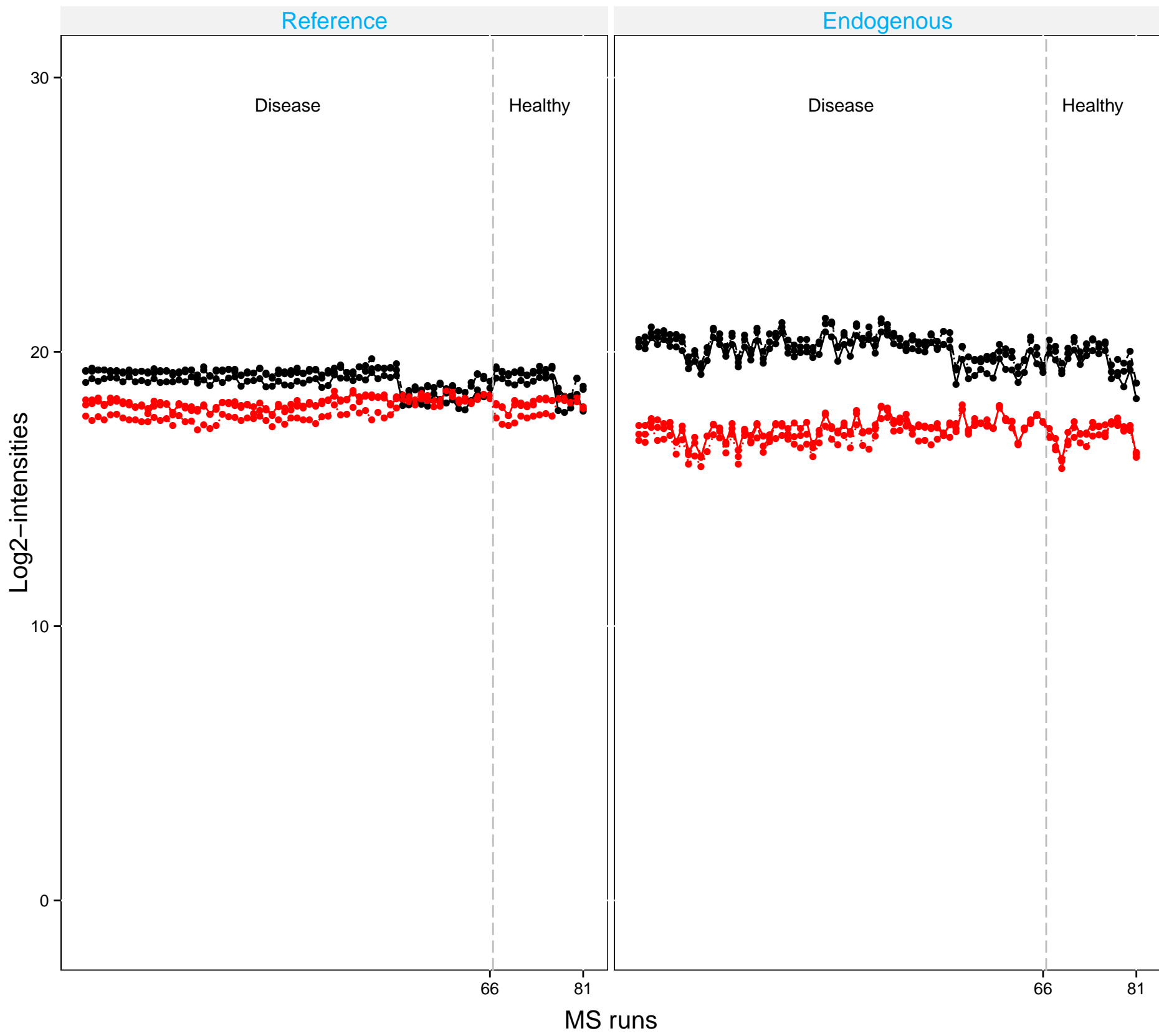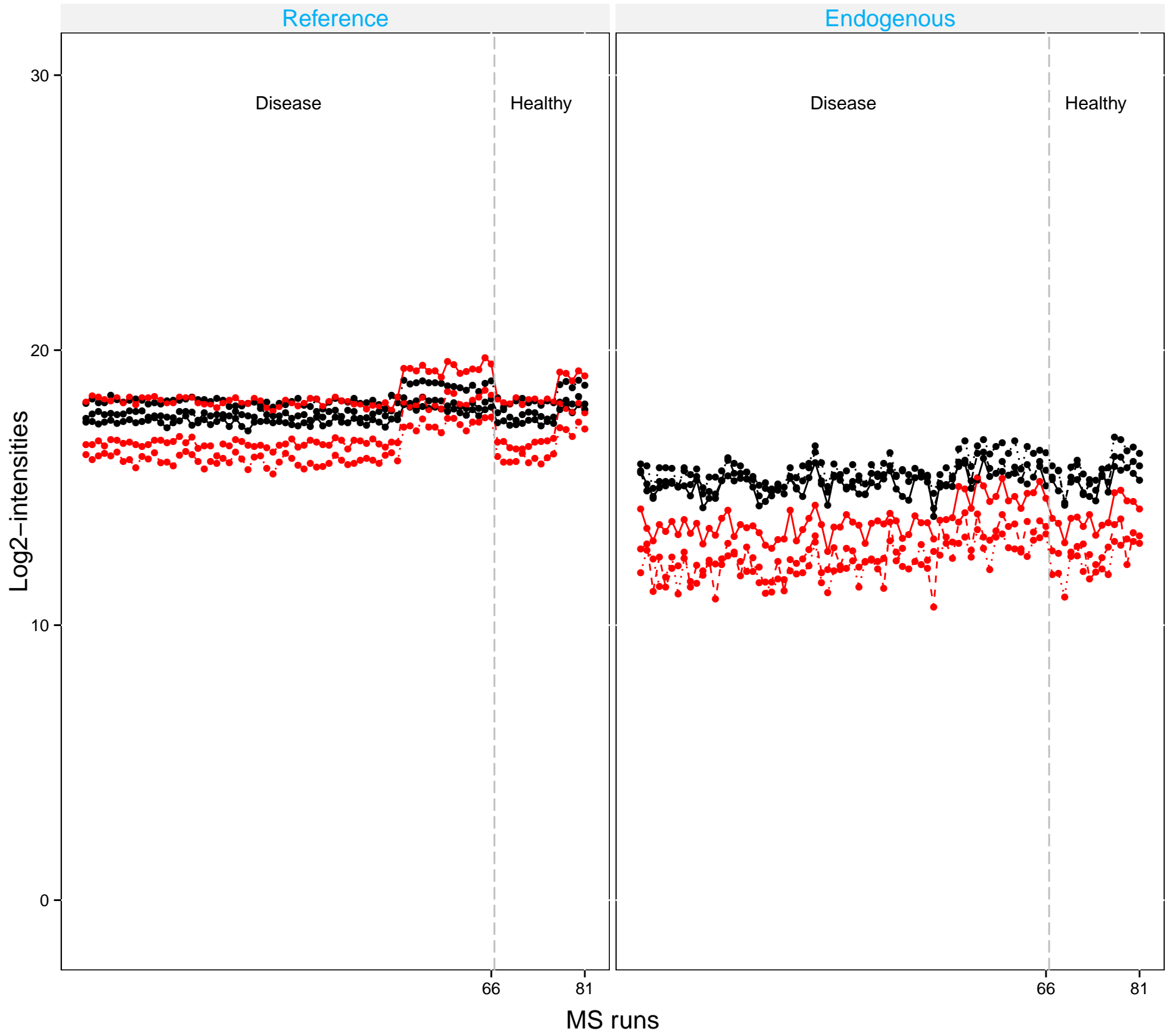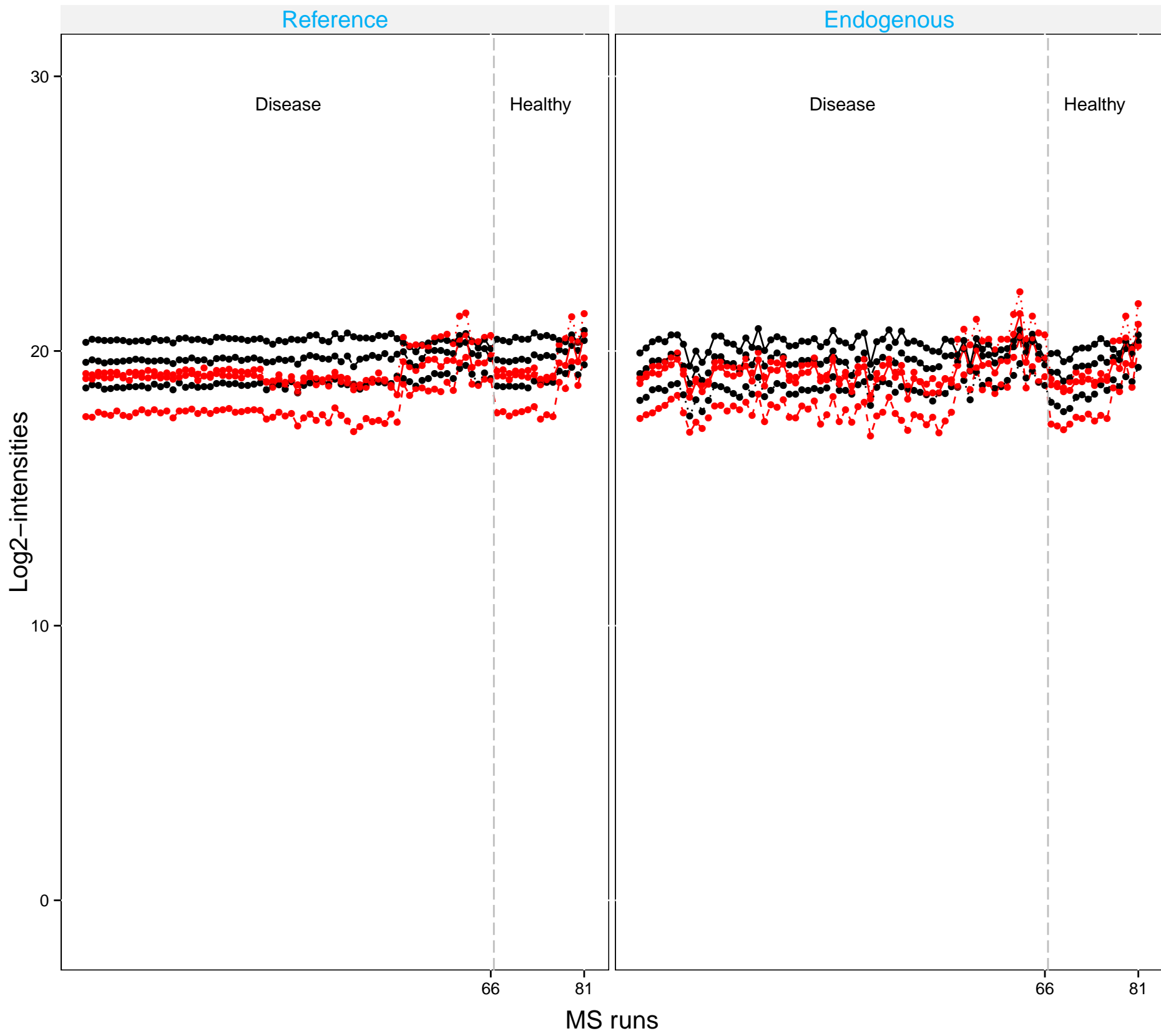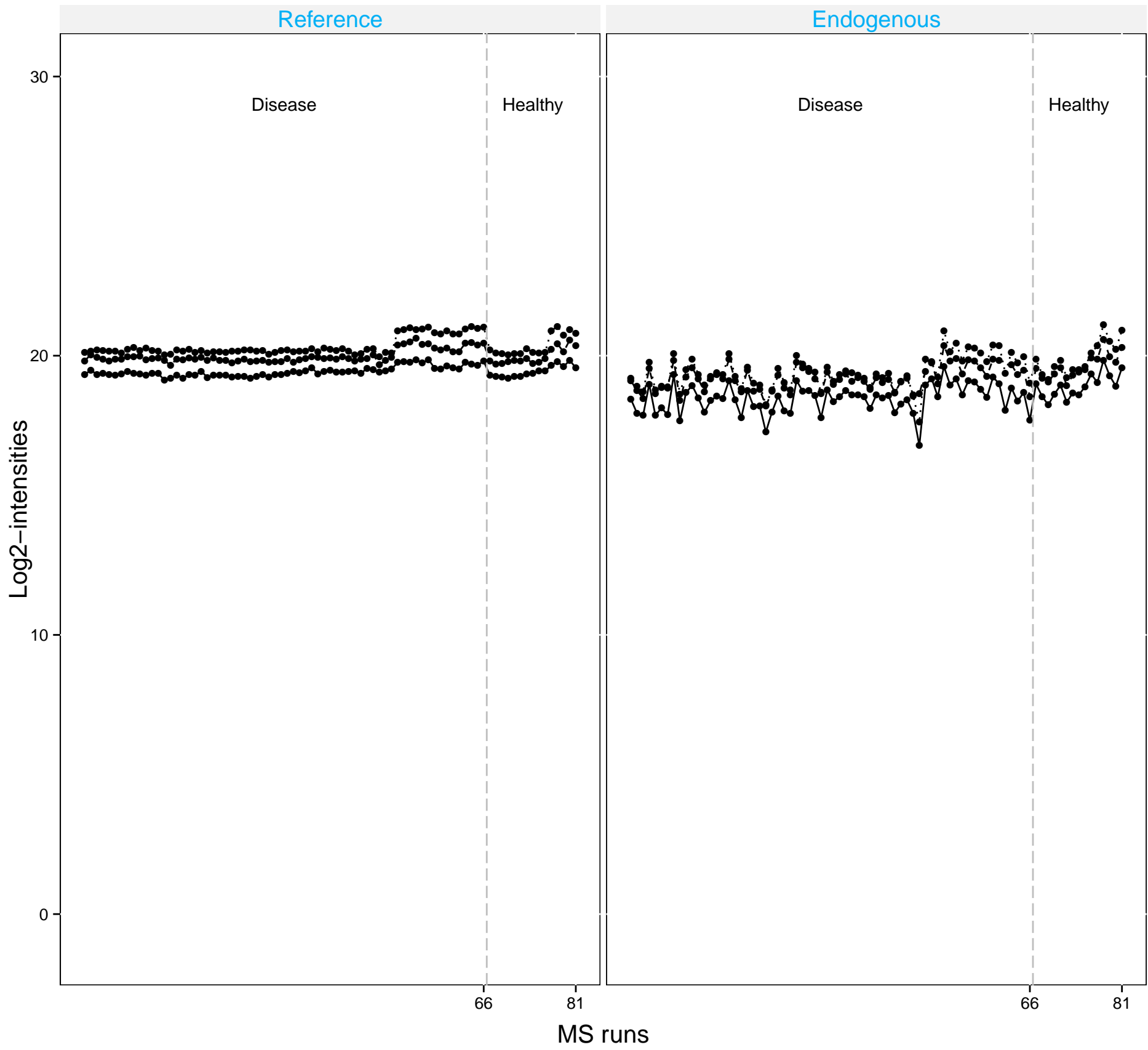
PLMN

peptide: 3

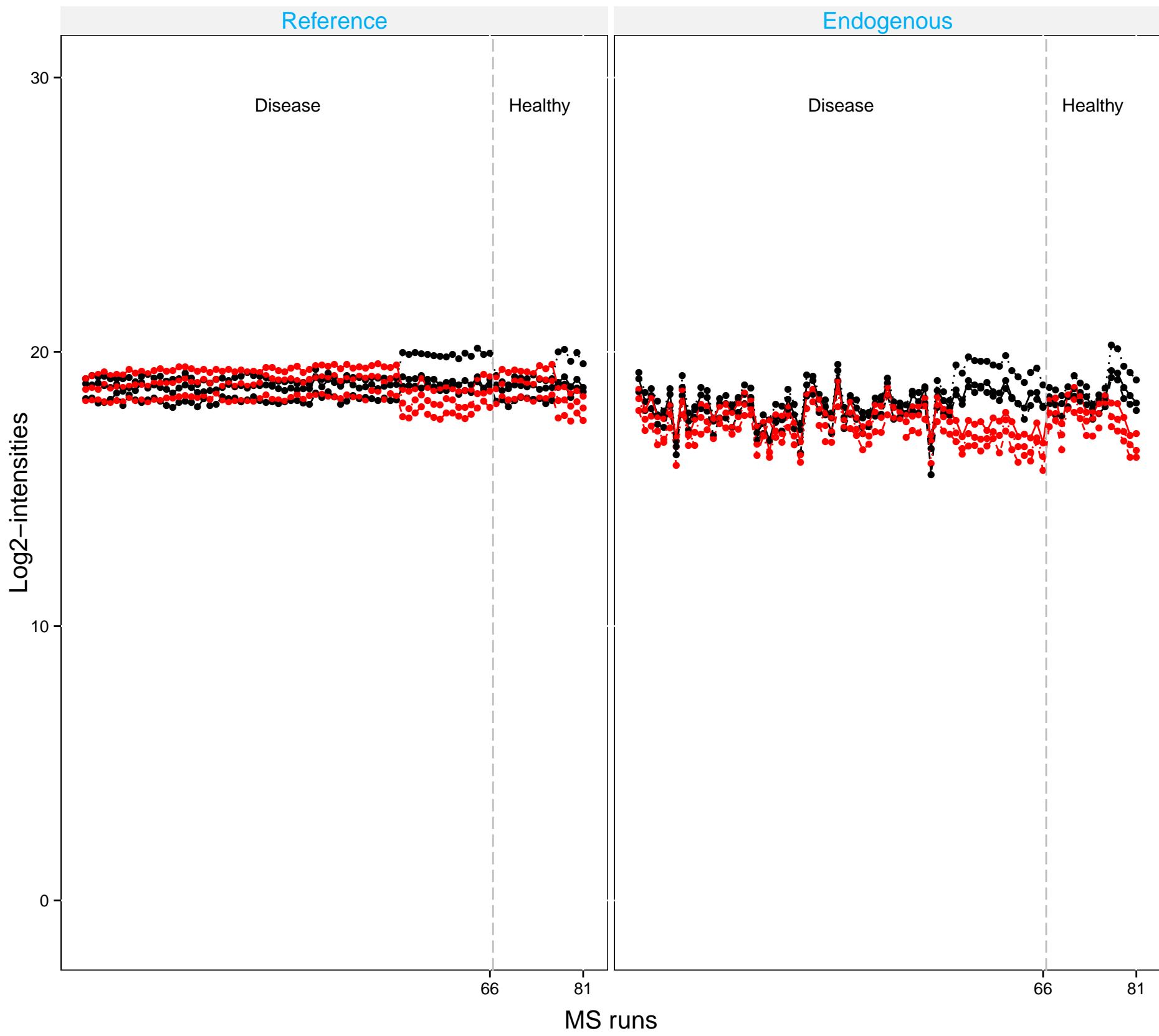|		
FSPATHPSEGLEENYCR_3_b11_2	HSIFTPETNPR_2_y6_1	QLGAGSIEECAAK_2_y6_1
FSPATHPSEGLEENYCR_3_y3_1	HSIFTPETNPR_2_y7_1	QLGAGSIEECAAK_2_y7_1
FSPATHPSEGLEENYCR_3_y6_1	HSIFTPETNPR_2_y8_1	QLGAGSIEECAAK_2_y9_1
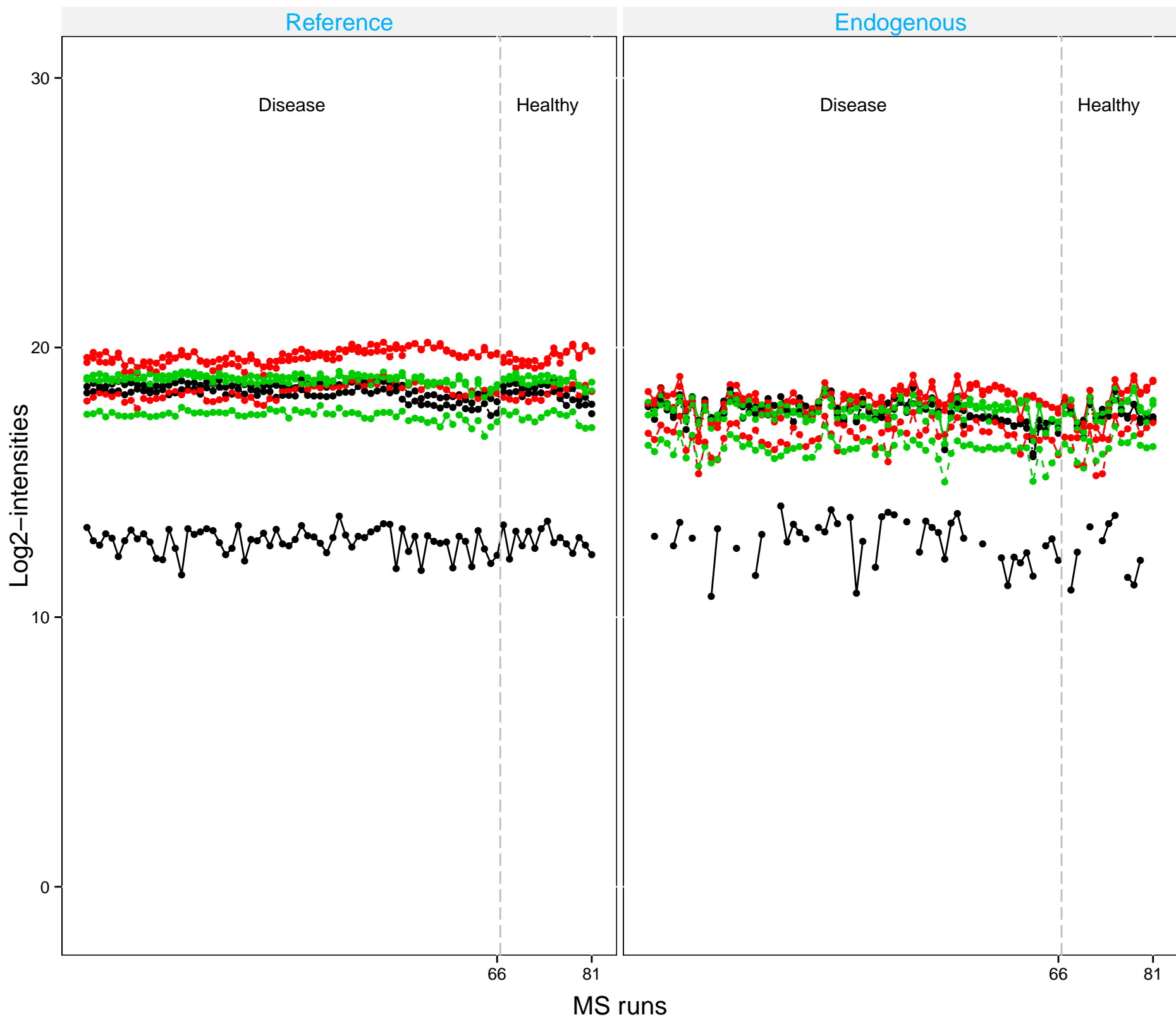
PON1

RET4

Legend:
- FSGTWYAMAK_2_y5_1
- FSGTWYAMAK_2_y6_1
- FSGTWYAMAK_2_y8_1
- LIVHNGYCDGR_3_y4_1
- LIVHNGYCDGR_3_y6_1
- LIVHNGYCDGR_3_y7_1

peptide: 2

SHBG

 # peptide: 1 •— IALGGLLFPASNLR_2_y6_1 •— IALGGLLFPASNLR_2_y7_1 •— IALGGLLFPASNLR_2_y8_1

TETN

THBG

THRB

TRFE

TTHY

VTNC

peptide: 4

DVWGIEGPIDAAFTR_2_b4_1	FEDGVLDPDYPR_2_y5_1	GQYCYELDEK_2_y7_1
DVWGIEGPIDAAFTR_2_y8_1	FEDGVLDPDYPR_2_y7_1	VDTVDPPYPR_2_y3_1
DVWGIEGPIDAAFTR_2_y9_1	GQYCYELDEK_2_y3_1	VDTVDPPYPR_2_y4_1
FEDGVLDPDYPR_2_y3_1	GQYCYELDEK_2_y6_1	VDTVDPPYPR_2_y5_1