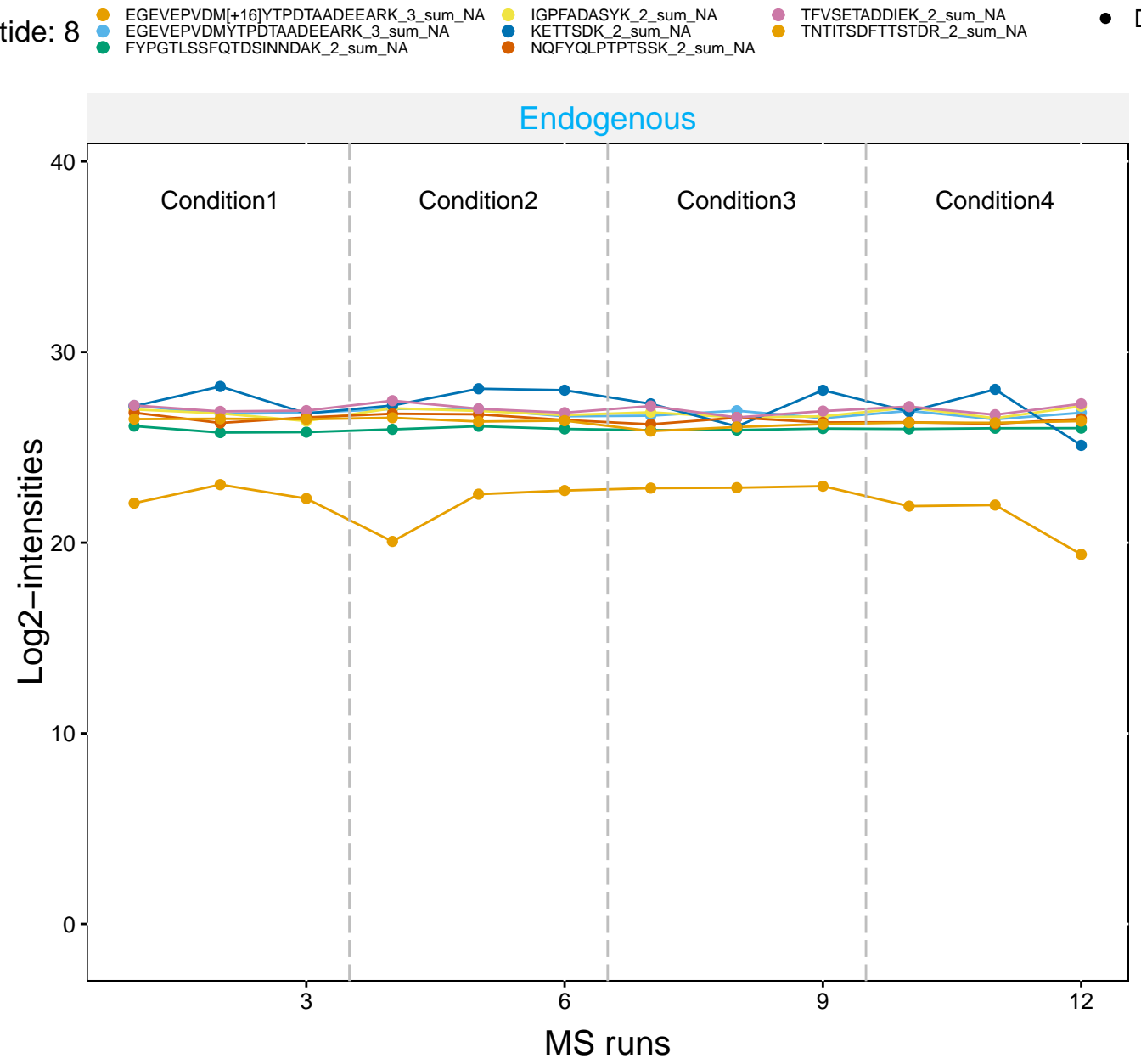


sp|D6VTK4|STE2_YEAST

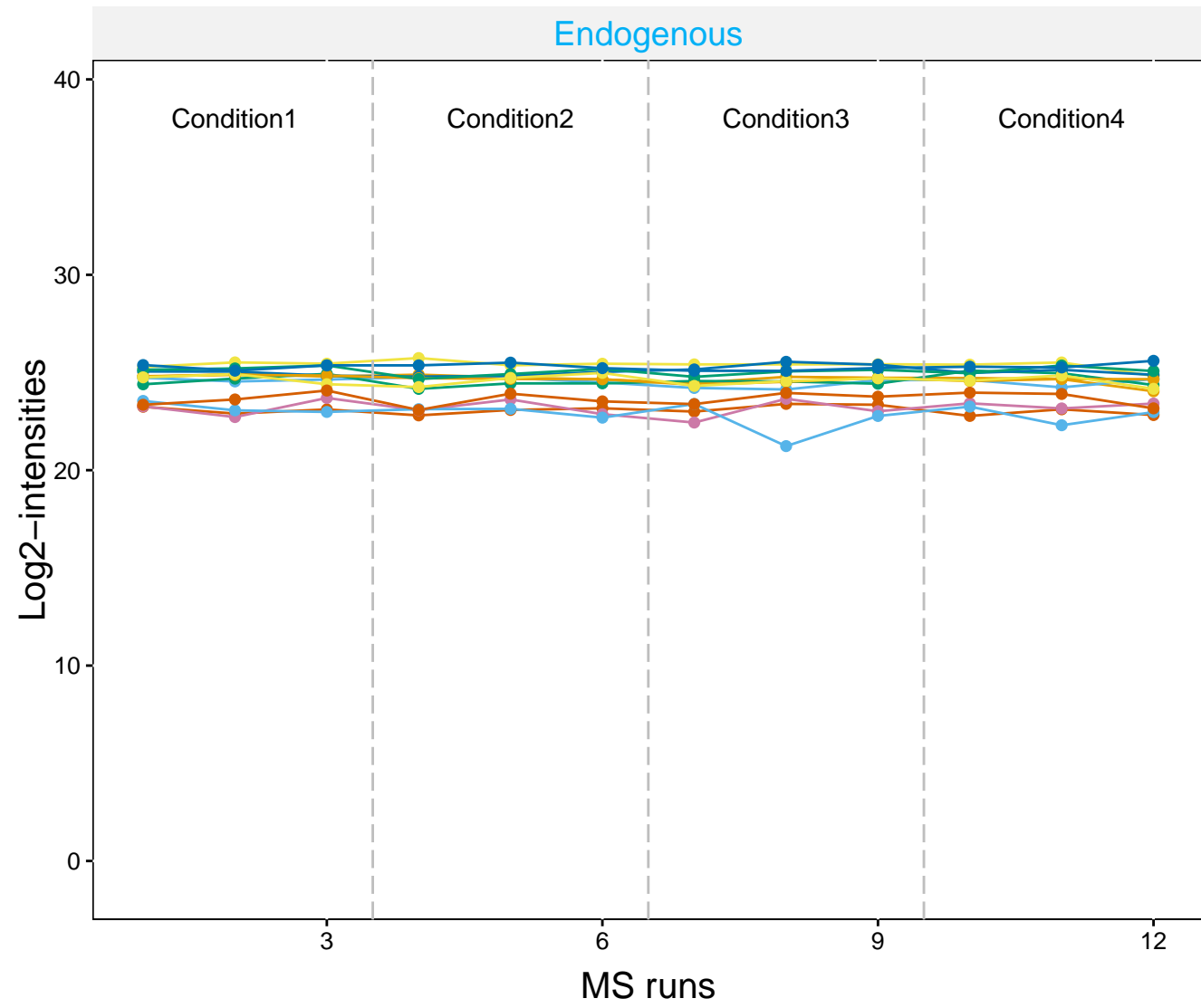
tide: 8



sp|O13297|CET1_YEAST

e: 13

- ALQSINVK_2_sum_NA
- FSIIESQTR_2_sum_NA
- GISEVTENTGK_2_sum_NA
- IAGNAVGSVVK_2_sum_NA
- IDPSFLNIIPDDDLTK_2_sum_NA
- ISLNLELPVPDNDPPEK_2_sum_NA
- LSEAANGSRPFAENLESDINQTETGQAAPIDNYK_3_sum_NA
- LSSLSYEIFEGSK_2_sum_NA
- NLLSNGSPPMNDGSDANAK_2_sum_NA
- QGNENGIASNYITQVPLQK_2_sum_NA
- SQSPISER_2_sum_NA
- TFLNNGTIIR_2_sum_NA
- VGLSTQRPR_3_sum_NA



sp|O13329|FOB1_YEAST

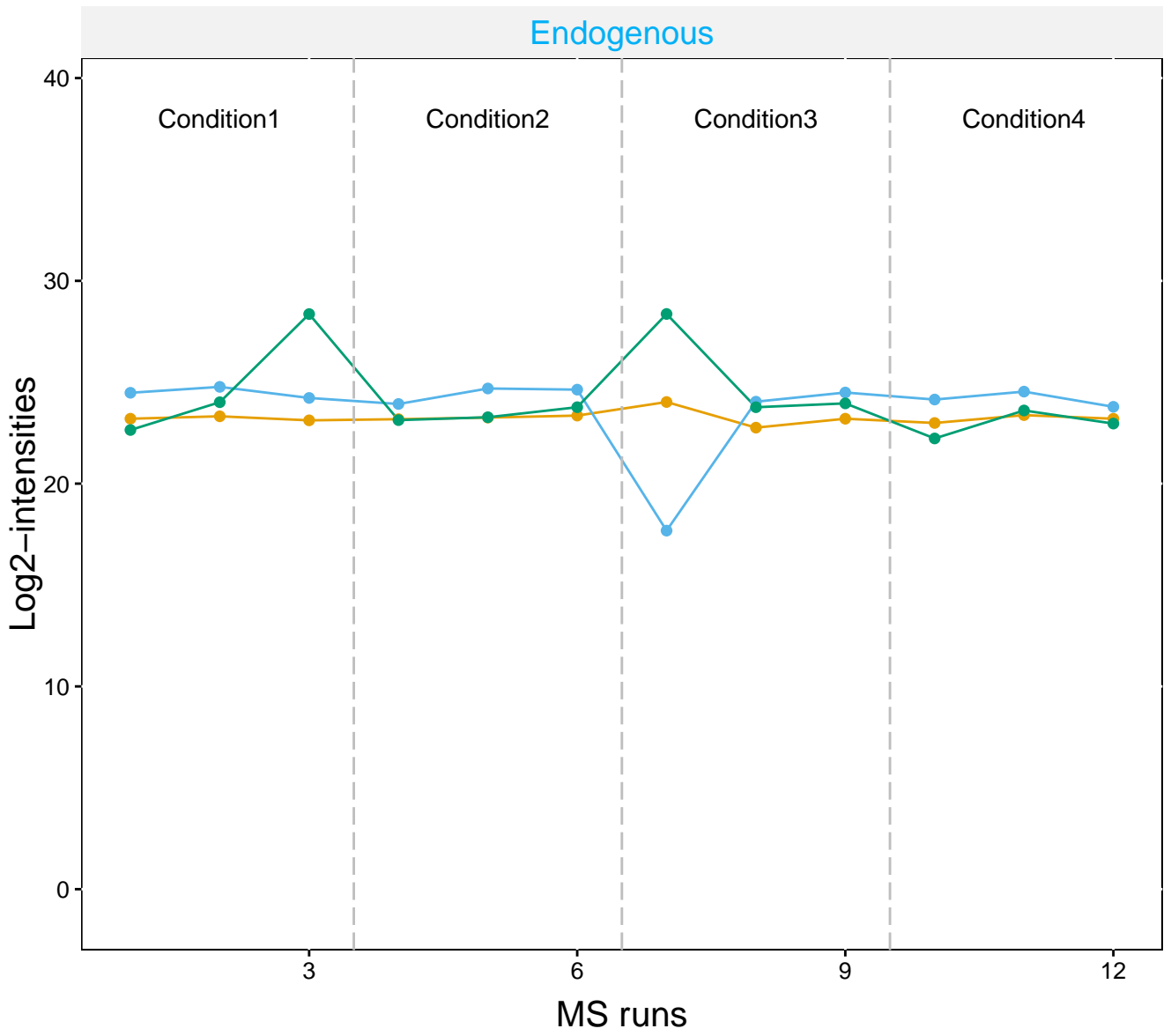
peptide: 3

● ETPIFLESLTR_2_sum_NA

● KEC[+57]LADWGK_2_sum_NA

● SLTADTLQQISDK_2_sum_NA

● Detected



sp|O13539|THP2_YEAST

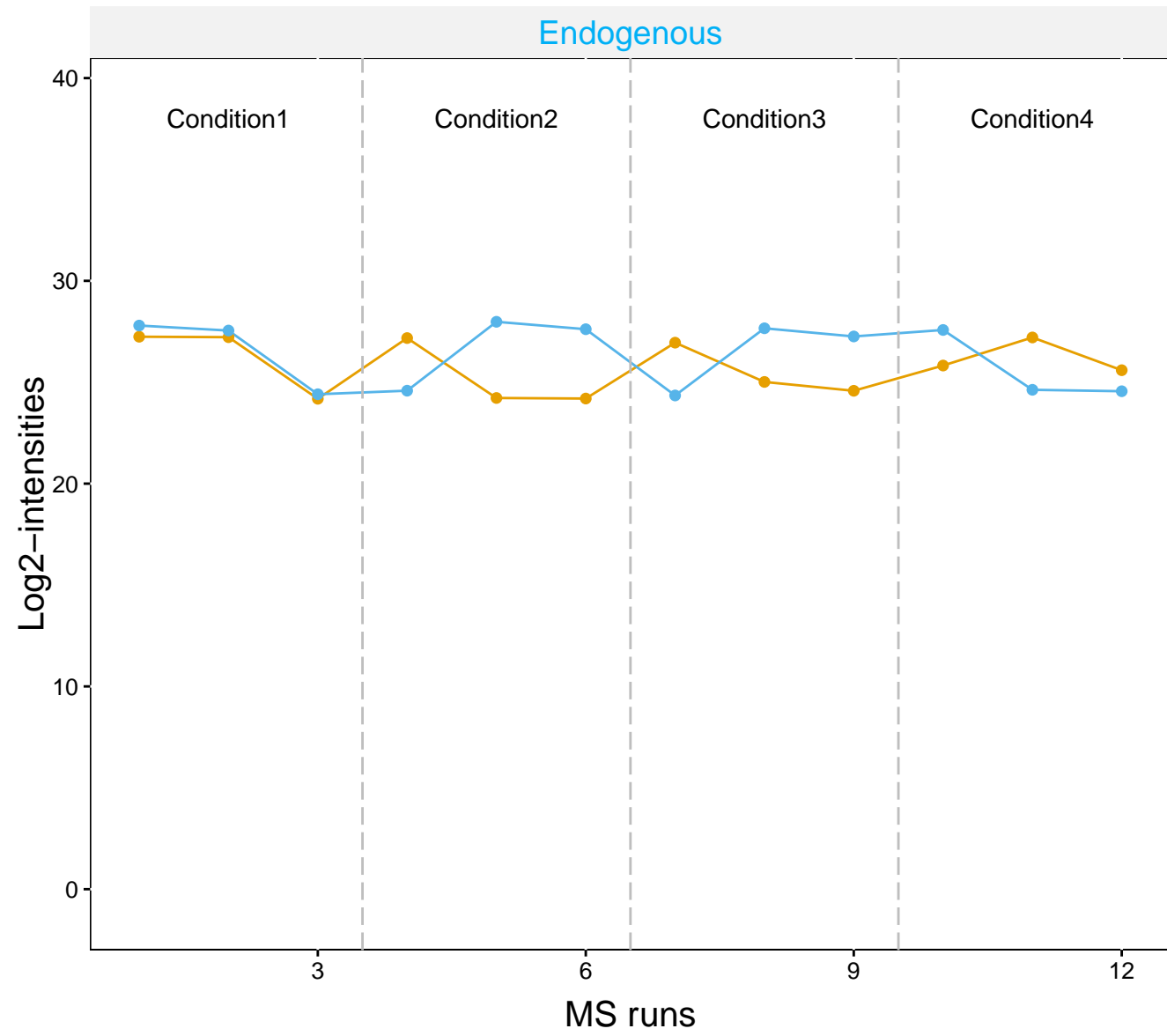
• Detected data

peptide: 2

● EAQLLEDTK_2_sum_NA

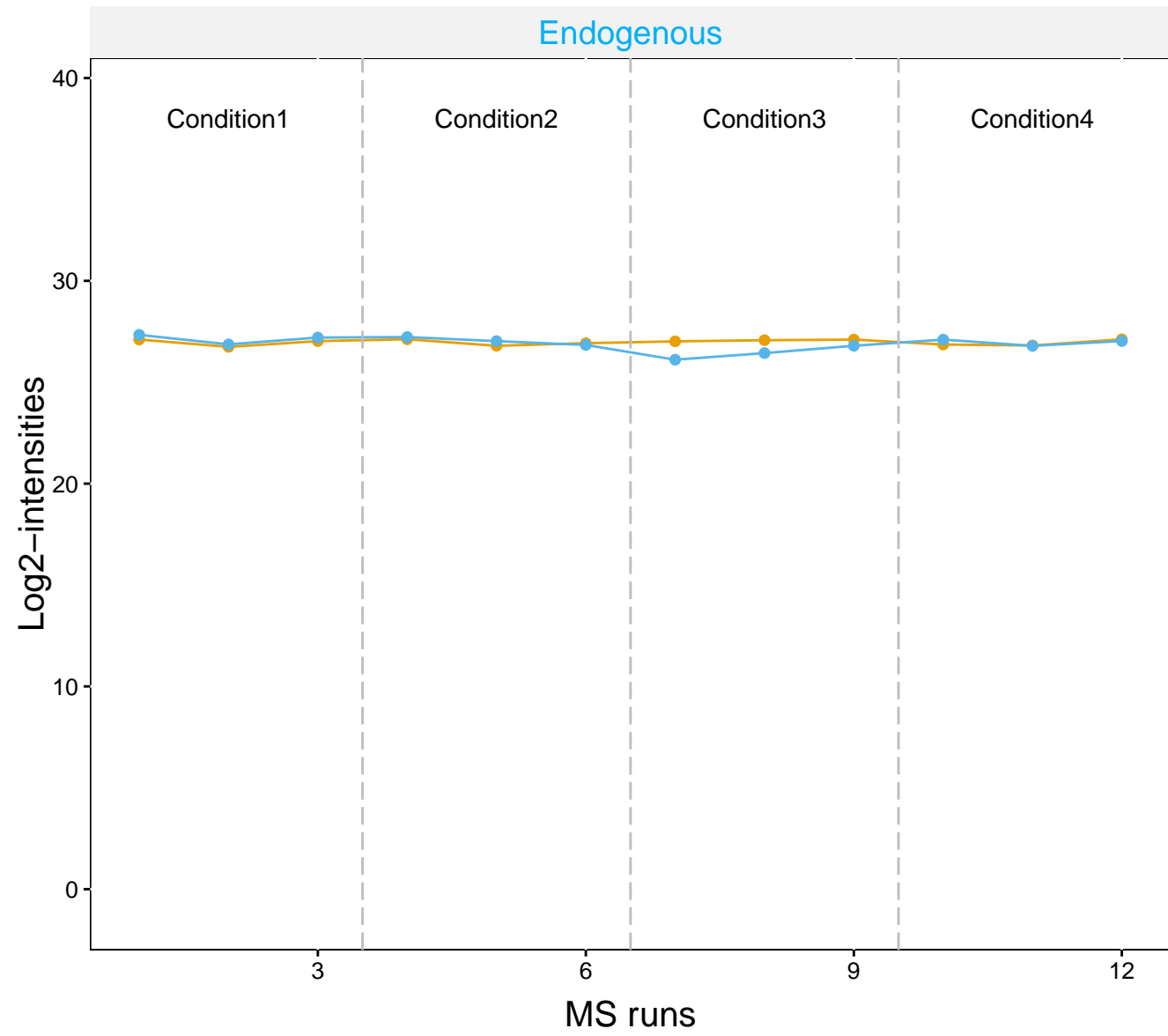
● SSDDLLETSLK_2_sum_NA

Endogenous



sp|O13547|CCW14_YEAST

ptide: 2 ● C[+57]LDSIC[+57]PNNDADAAYSAFK_2_sum_NA ● SSSTC[+57]DSLNQVTC[+57]YC[+57]EHENSAVK_3_sum_NA ● D



sp|O13563|RPN13_YEAST

peptide: 5

● AGVC[+57]EYNEDSR_2_sum_NA

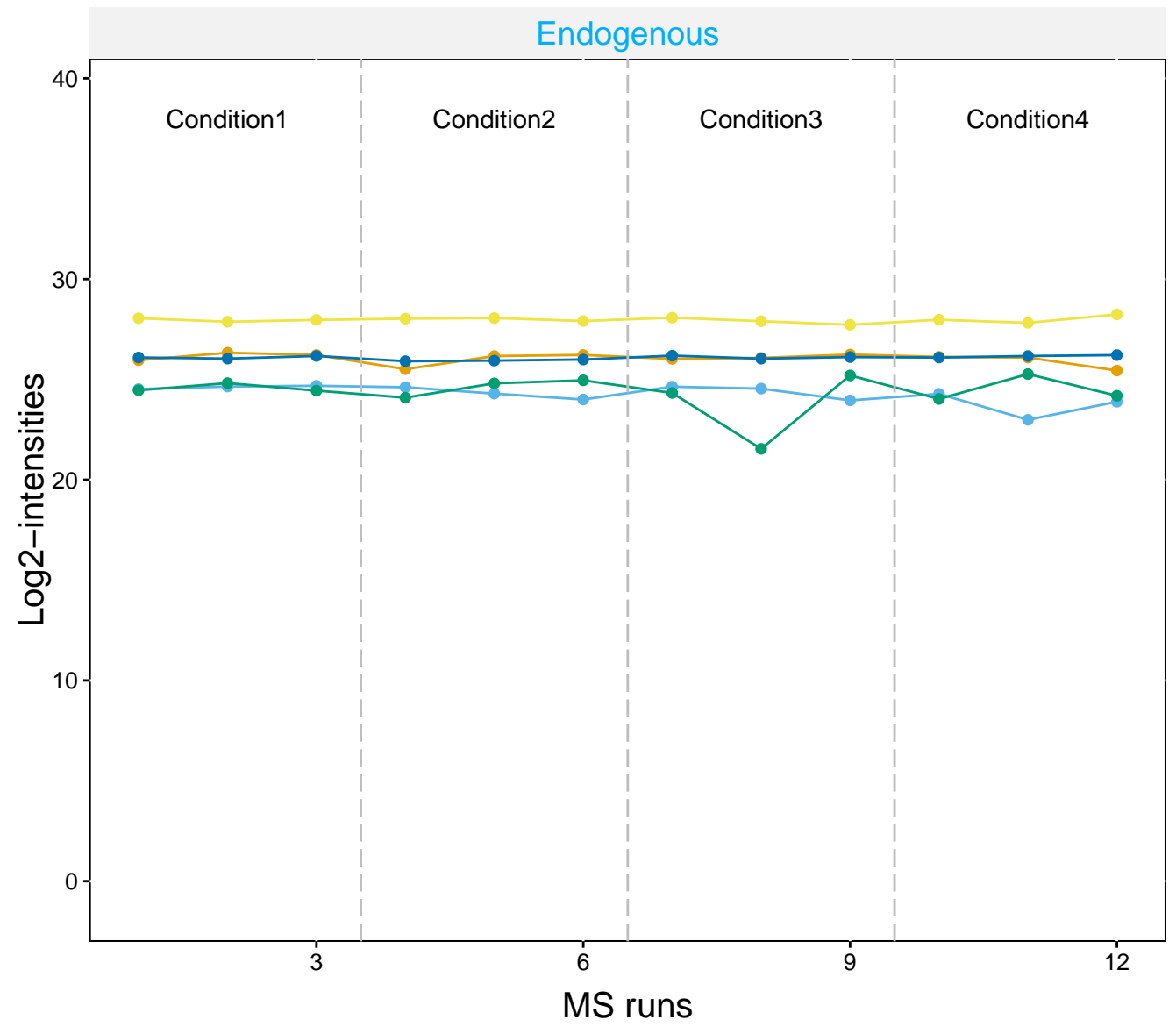
● AQDVDVSMQD_2_sum_NA

● IFALVFSSNER_2_sum_NA

● NSGNLPLNELSAK_2_sum_NA

● YFFWLQEK_2_sum_NA

● Detected



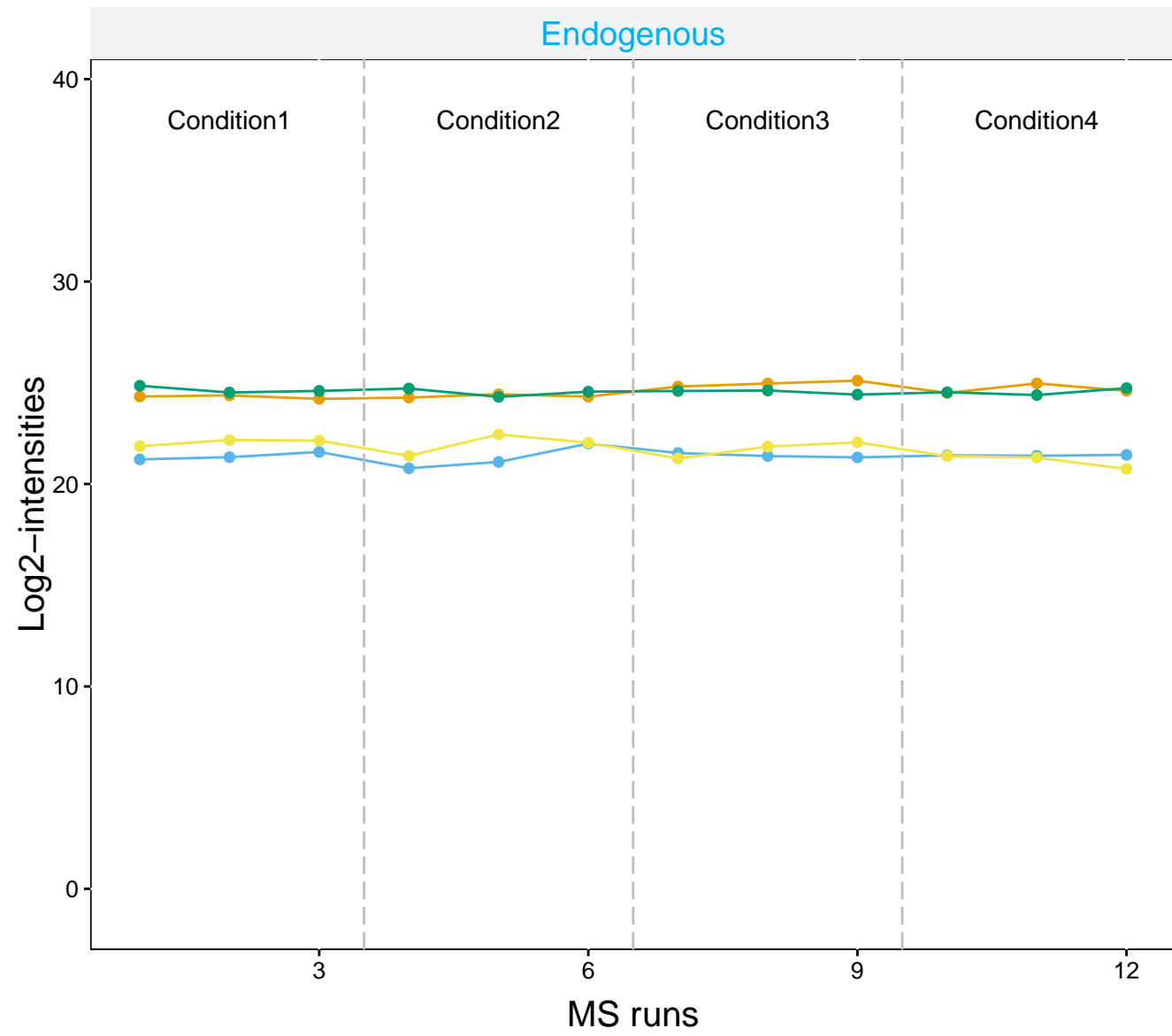
sp|O13585|YP089_YEAST

peptide: 4

DVGEIDLSDPILR_2_sum_NA
HEIPQISQPK_2_sum_NA

ITPVQLLK_2_sum_NA
NVATYKPIDENNK_2_sum_NA

• Detected data



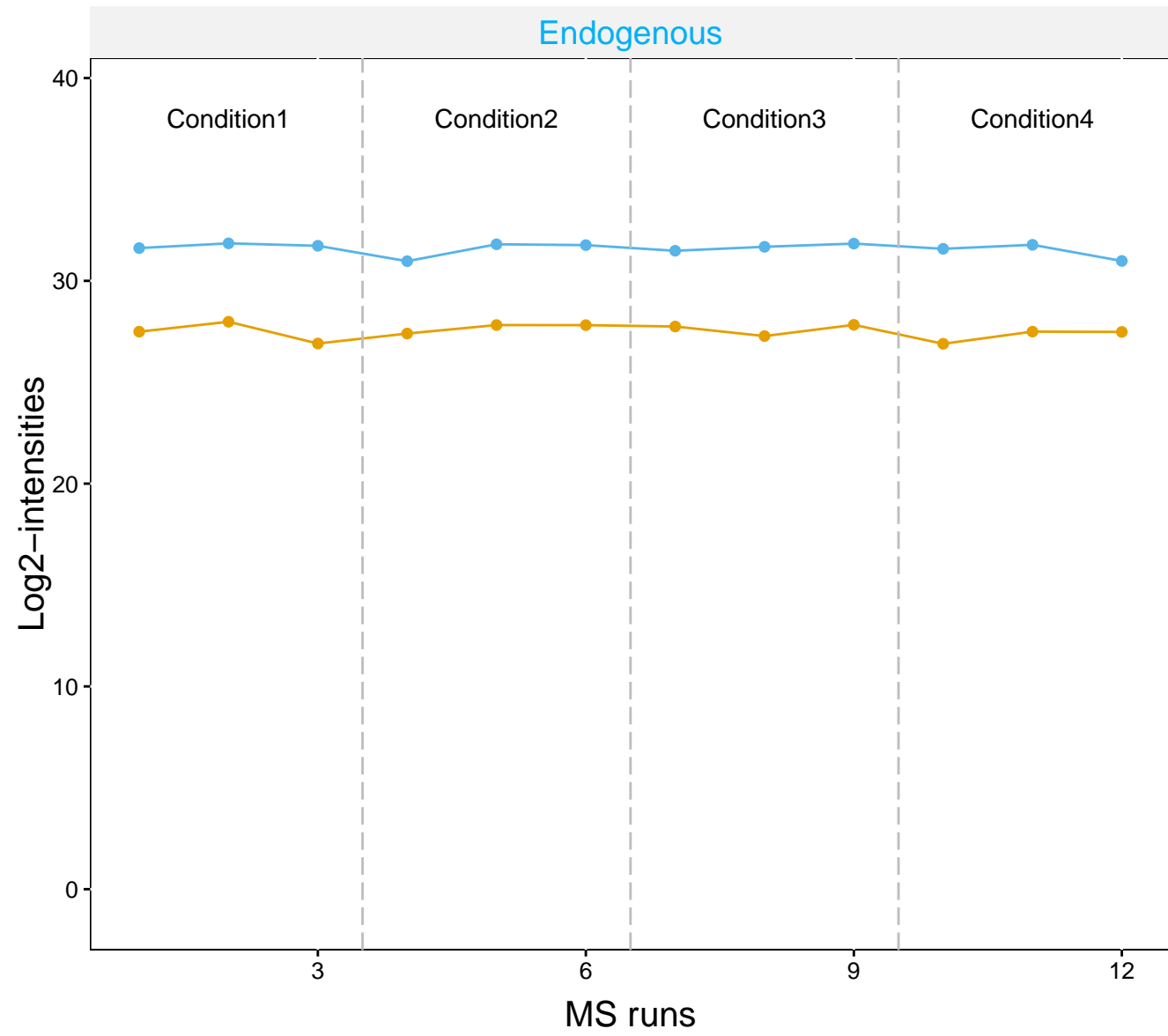
sp|O14455|RL36B_YEAST

peptide: 2

● VTQM[+16]TPAPK_2_sum_NA

● VTQMTPAPK_2_sum_NA

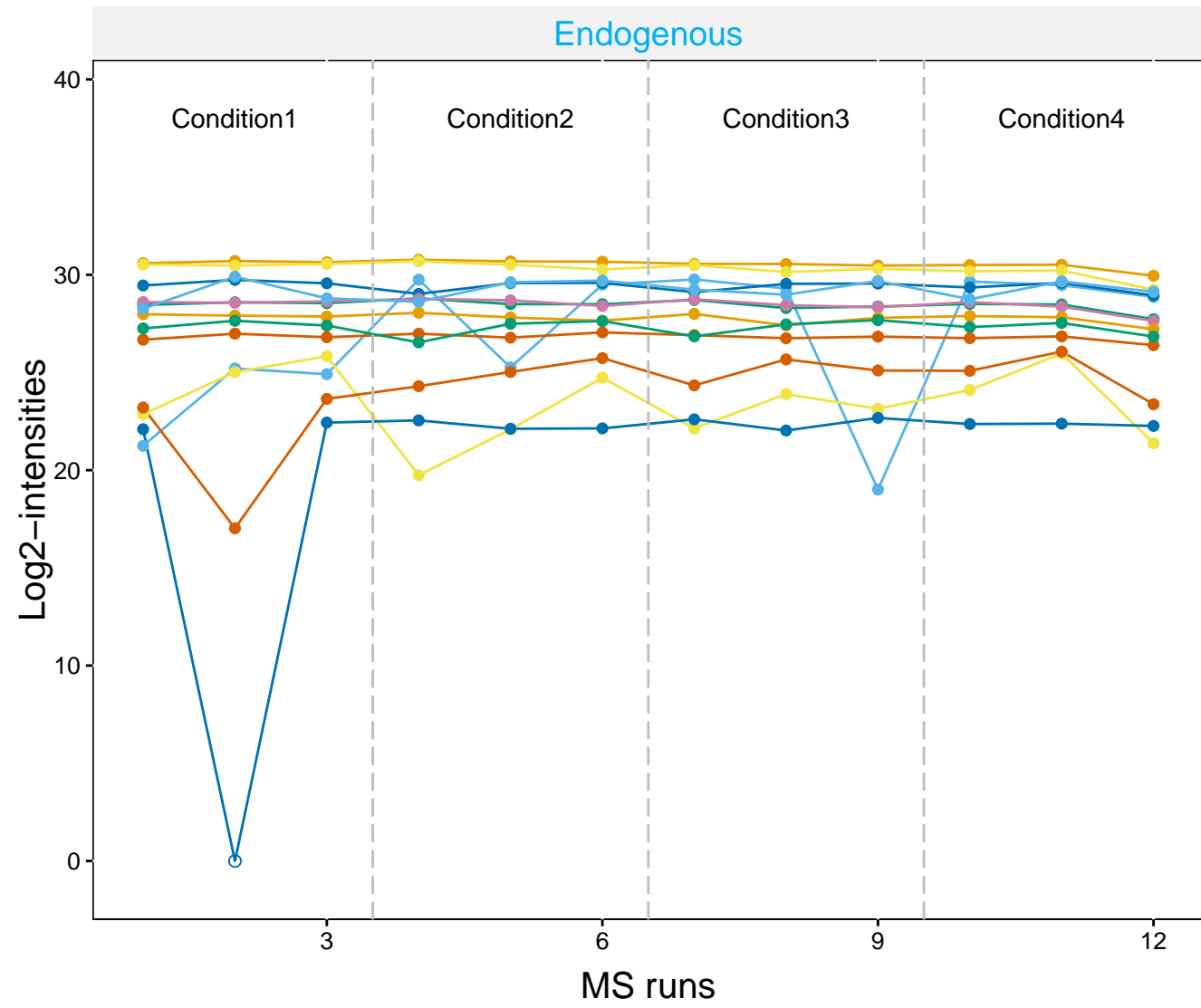
● Detected data



sp|O14467|MBF1_YEAST

● Detected data ○

- AIPNQVLSK_2_sum_NA
- GNNIGSPLGAPK_2_sum_NA
- INEKPTVVNDYEAAR_2_sum_NA
- INEKPTVVNDYEAAR_3_sum_NA
- LDPNVGR_2_sum_NA
- LRGNNIGSPLGAPK_2_sum_NA
- LRGNNIGSPLGAPK_3_sum_NA
- QGLVVSVDK_2_sum_NA
- SQQQINAAR_2_sum_NA
- VDRETDIVKPK_3_sum_NA
- YGSTNTR_2_sum_NA
- YGSTNTRGDNEGQR_2_sum_NA
- YGSTNTRGDNEGQR_3_sum_NA



sp|O14468|YO304_YEAST

○ Censored missing data

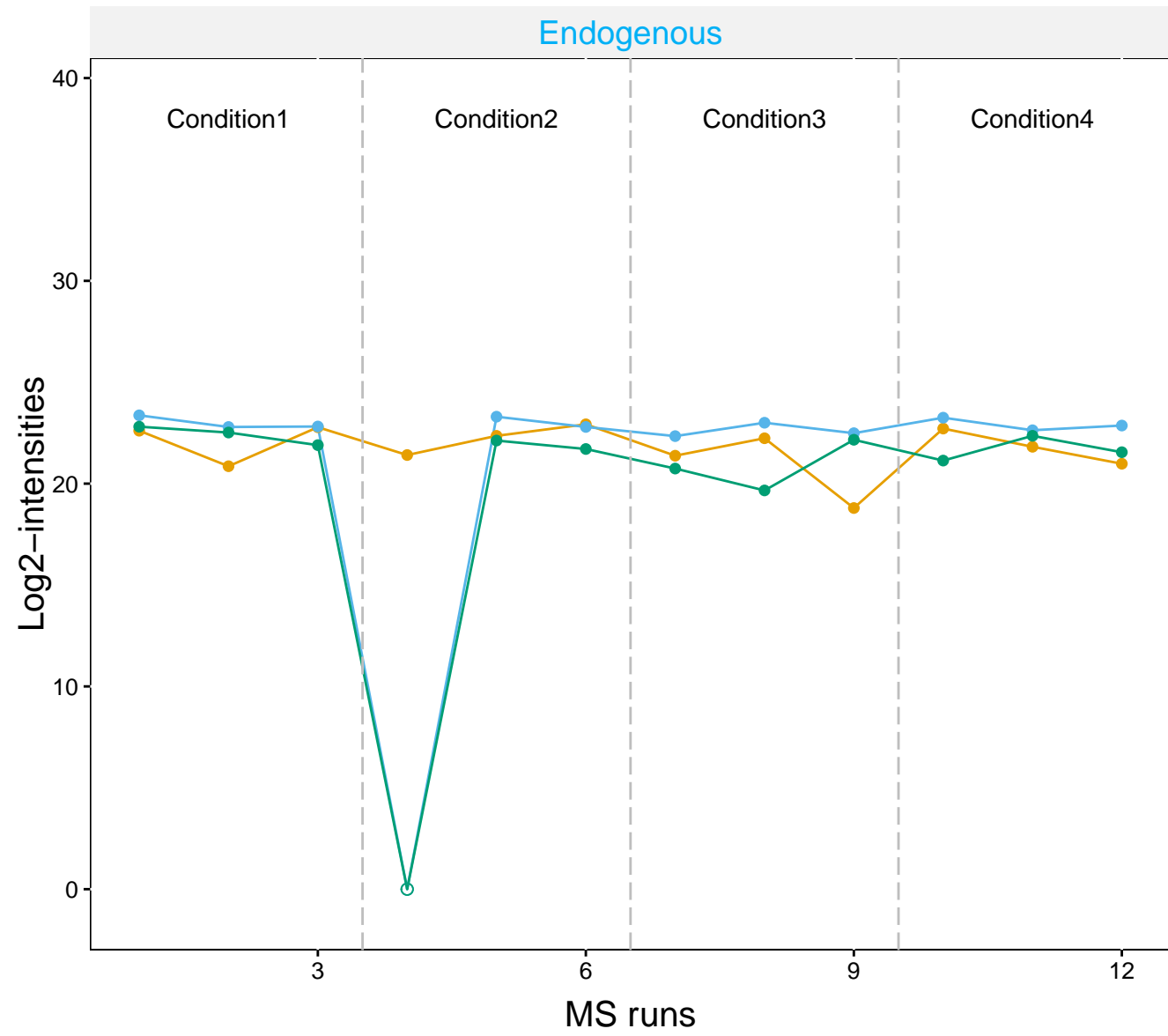
peptide: 3

● ENDDFSR_2_sum_NA

● LEASEEPQAPLANTSETNSIK_2_sum_NA

● LE

Endogenous



sp|O43137|YB085_YEAST

Detected data ○ Censored missing data

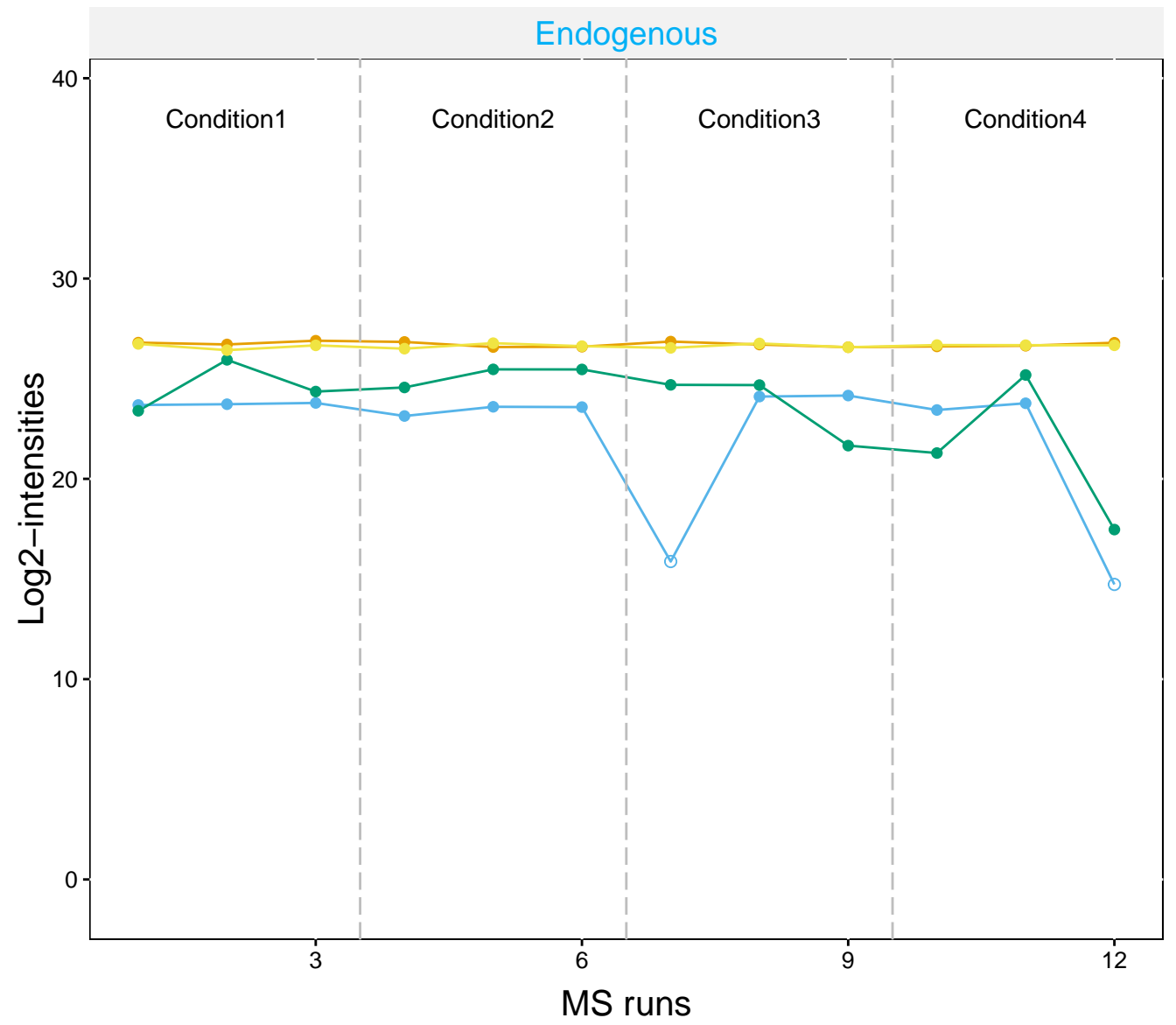
peptide: 4

● DGEVIANIIGEAK_2_sum_NA

● DYLLDLAQQA_2_sum_NA

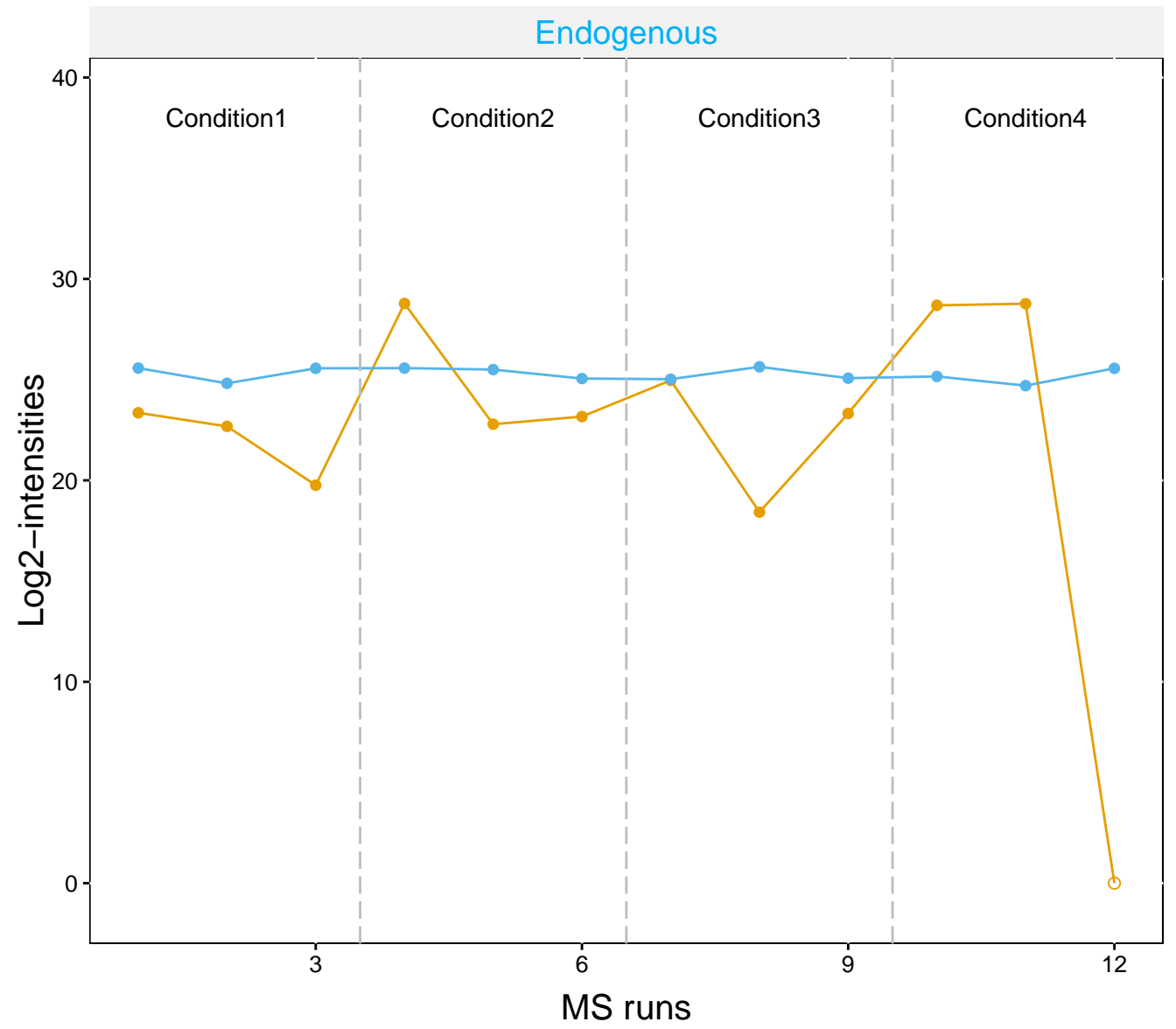
● KNYEIQTVLSEDK_3

● NYEIQTVLSEDK_2_3



sp|O60200|MDM35_YEAST

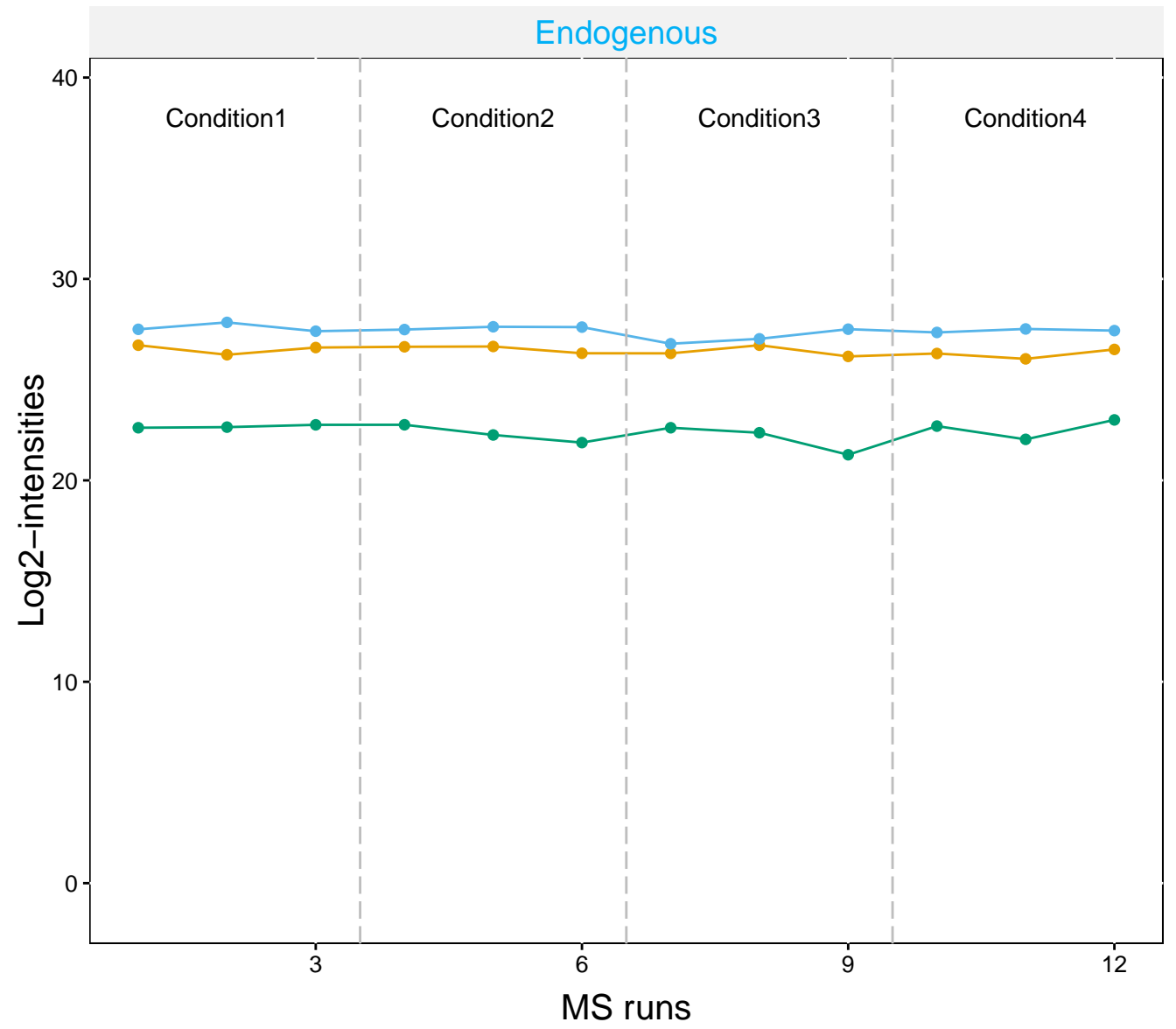
ected data ○ Censored missing data # peptide: 2 ● QWYAYTTC[+57]VNAALVK_2_sum_NA ● YDSC[+57]FNEWY



sp|O74700|TIM9_YEAST

peptide: 3 ● Detected

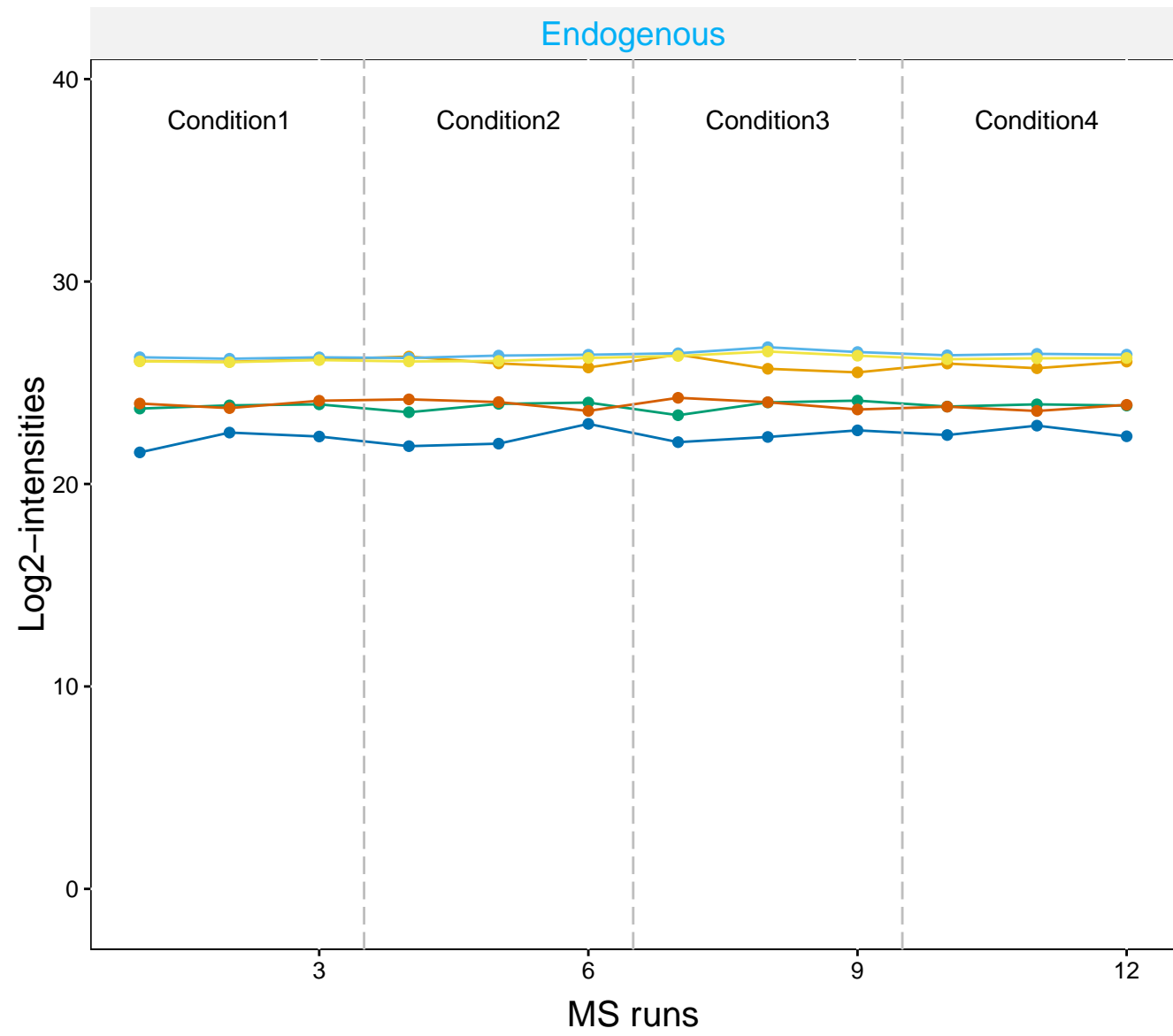
● C[+57]FTDC[+57]VNDFTTSK_2_sum_NA ● FQEQNAALGQGLGR_2_sum_NA ● M[+42]DALNSK_2_sum_NA



sp|P00128|QCR7_YEAST

de: 6

DELDNIEVSK_2_sum_NA FDDLIAEENPIMQTALR_3_sum_NA LC[+57]VPVANQFINLAGYK_3_sum_NA
FDDLIAEENPIMQTALR_2_sum_NA LC[+57]VPVANQFINLAGYK_2_sum_NA LC[+57]VPVANQFINLAGYKK_3_sum_NA

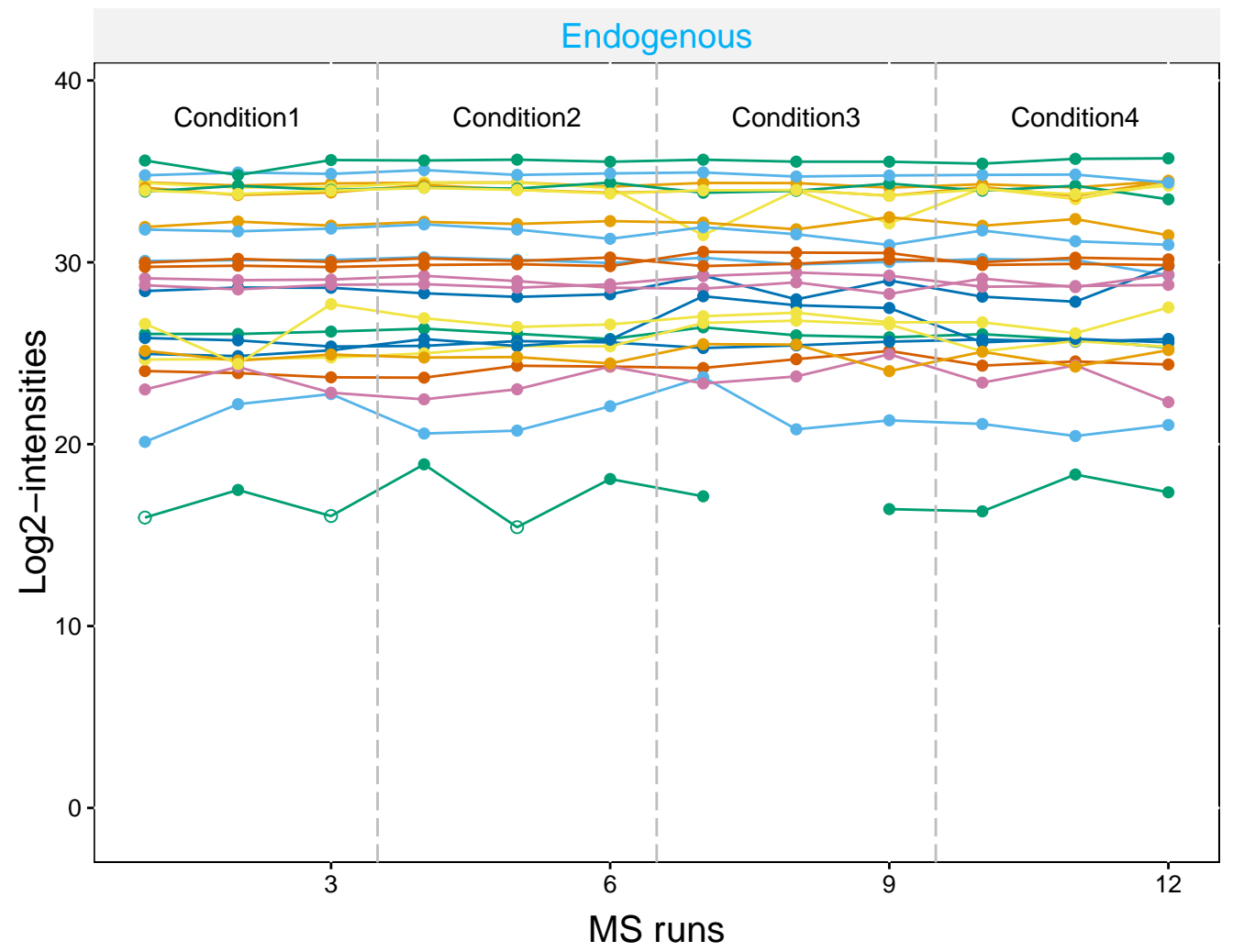


sp|P00330|ADH1_YEAST

Missing data

peptide: 25

- ANELLINVK_2_sum_NA
- ANGTTVLVGM[+16]PAGAK_2_sum_NA
- ANGTTVLVGM[+16]PAGAK_3_sum_NA
- ANGTTVLVGMPAGAK_2_sum_NA
- ANGTTVLVGMPAGAK_3_sum_NA
- ATDGGAHGVINVSVEAAIEASTR_2_sum_NA
- ATDGGAHGVINVSVEAAIEASTR_3_sum_NA
- C[+57]C[+57]SDVFNQVVK_2_sum_NA
- C[+57]C[+57]SDVFNQVVK_3_sum_NA
- DIPVPKPK_2_sum_NA
- DIVGAVLK_2_sum_NA
- E[-18]KDIVGAVLK_2_sum_NA
- EKDIVGAVLK_2_sum_NA
- EKDIVGAVLK_3_sum_NA
- GVIFYESHGK_2_sum_NA
- GVIFYESHGK_3_sum_NA
- MEKGQIVGRYVVDTSK_2_sum_NA
- SANLM[+16]AGHWVAISGAAGGLGSLAVQYAK_3_sum_NA



sp|P00358|G3P2_YEAST

ected data

peptide: 6

● HIIVDGHK_2_sum_NA

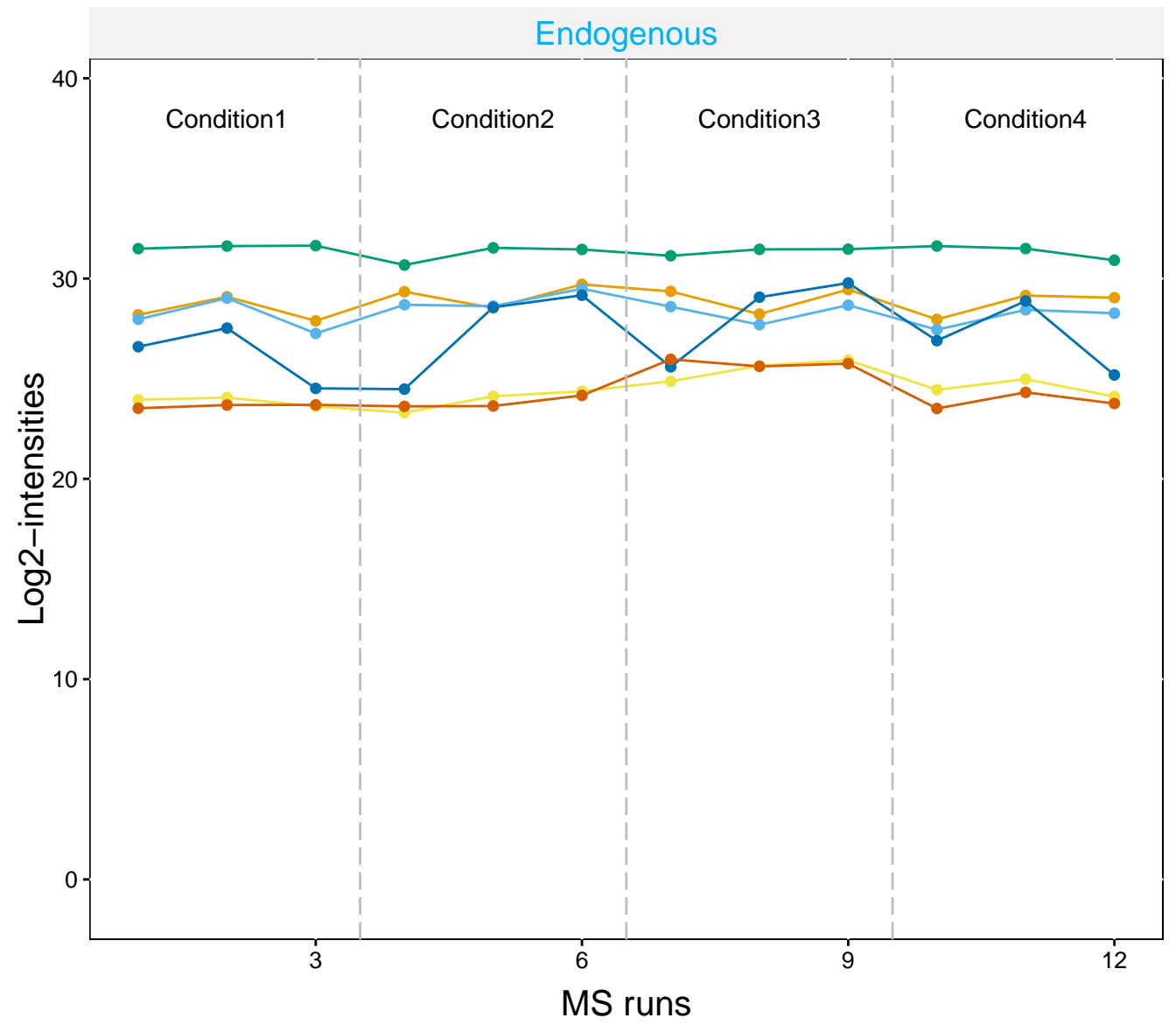
● IATFQER_2_sum_NA

● HIIVDGHK_3_sum_NA

● NVEVVALNDPFISNDYSAYM[+16]FK_2_sum_NA

● NVEVVALNDPFISNDYS

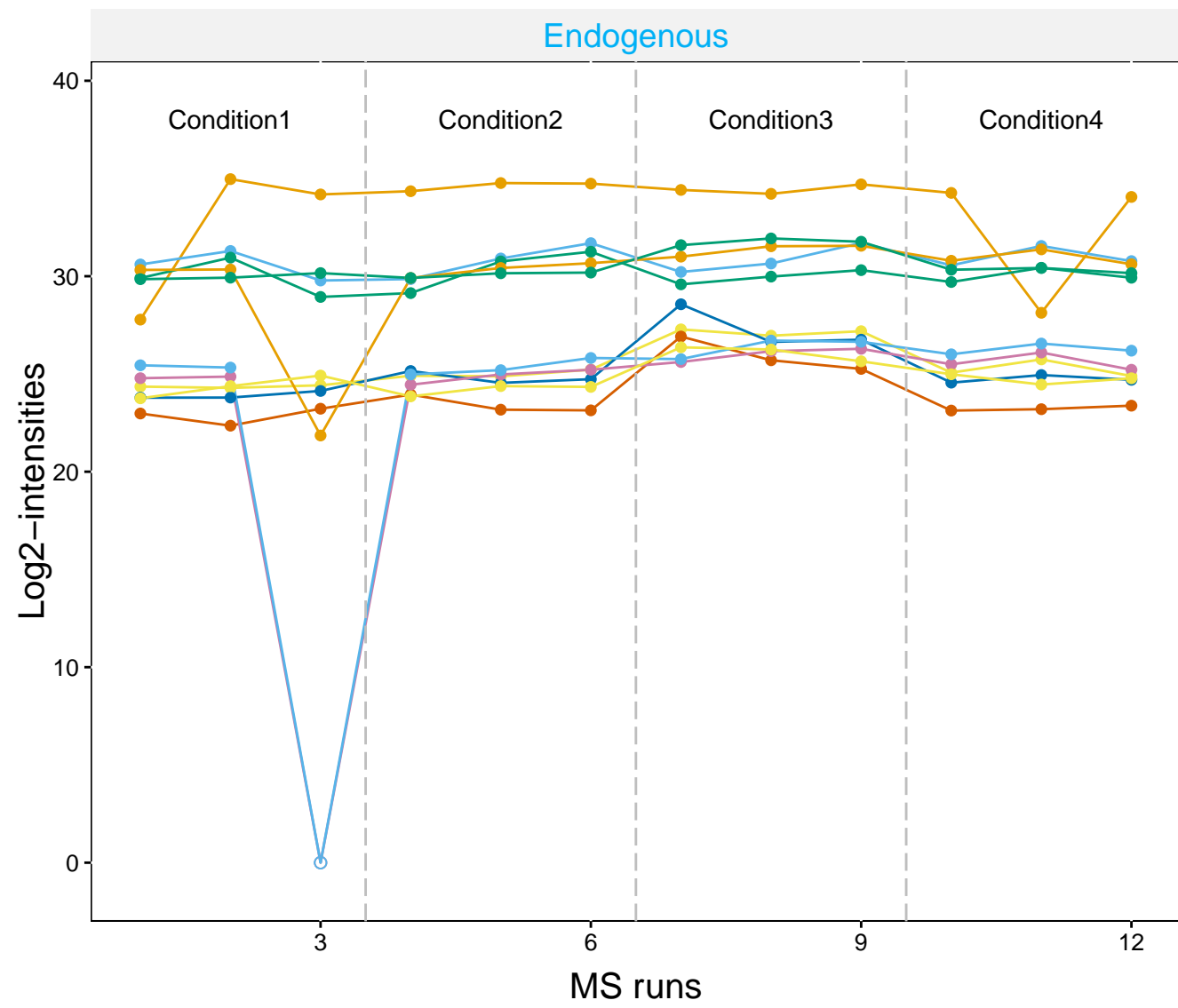
● NVEVVALNDPFISNDYS



sp|P00359|G3P3_YEAST

●

IALSRPNVEVVALNDPFITNDYAYMFK_3_sum_NA
IATYQERDPANLPWGSSNVDIAIDSTGVFK_3_sum_NA
VINDAFGIEEGLM[+16]TTVHSLTATQK_2_sum_NA
VINDAFGIEEGLM[+16]FK_3_sum_NA
VINDAFGIEEGLM[+16]TTVHSLTATQK_3_sum_NA
VINDAFGIEEGLM[+16]TTVHSLTATQK_4_sum_NA
VINDAFGIEEGLM[+16]TTVHSLTATQK_4_sum_NA



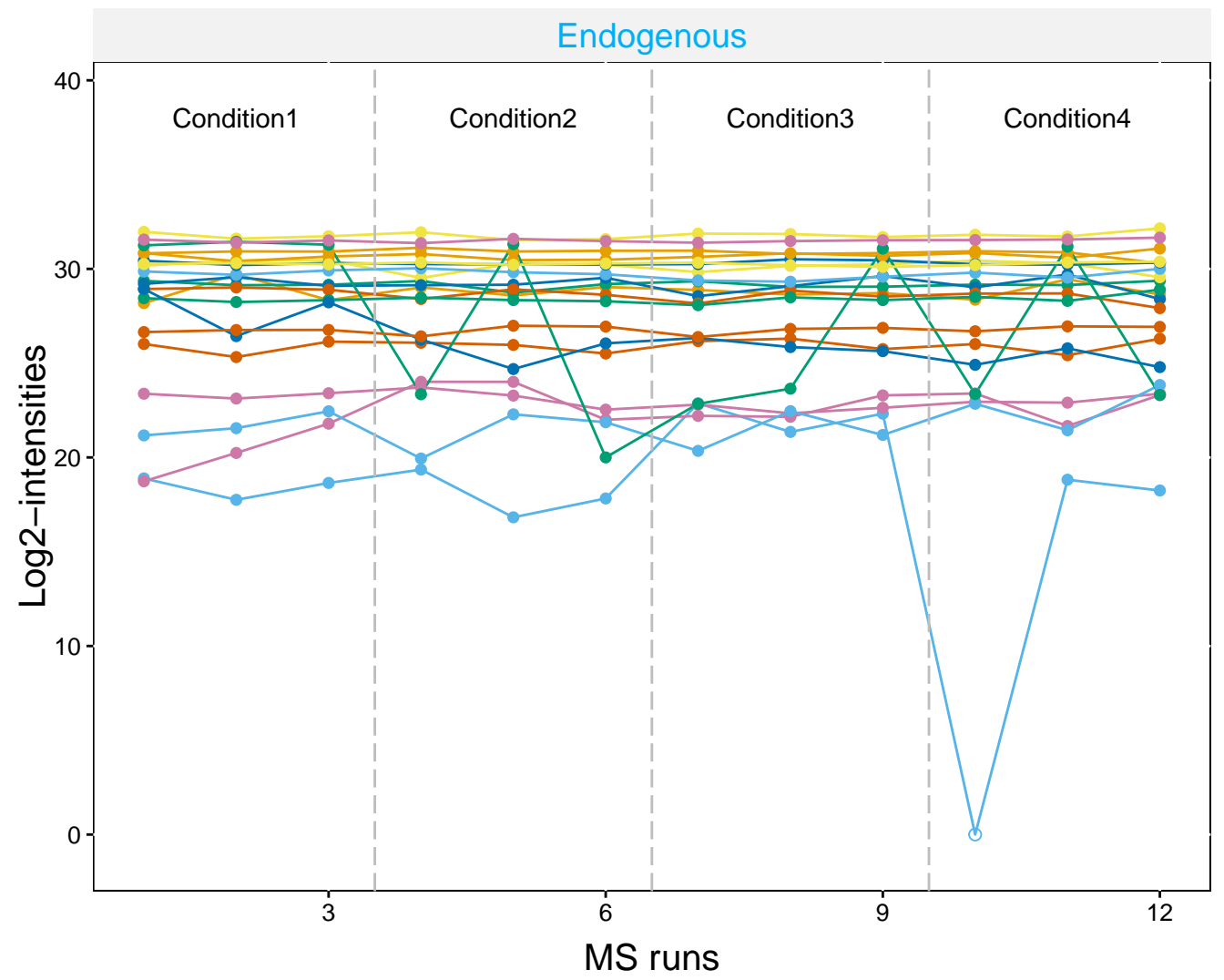
sp|P00360|G3P1_YEAST

ored missing data

peptide: 21

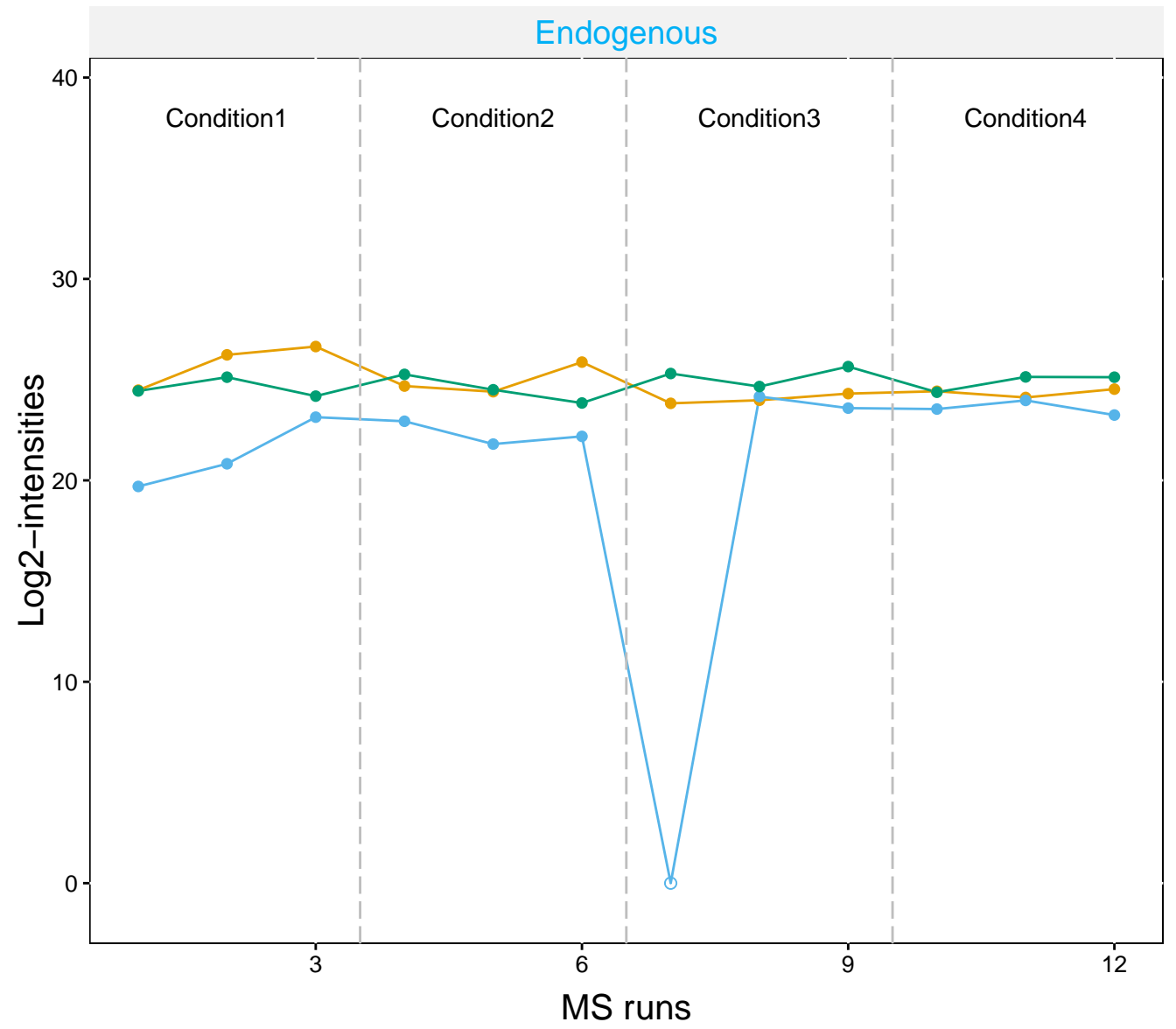
AAAEGPMK_2_sum_NA
DIEVVAVNDPFISNDYAAVMVK_2_sum_NA
DPANLPWGSGLK_2_sum_NA
EATYDQIK_2_sum_NA
EATYDQIKK_2_sum_NA
EATYDQIKK_3_sum_NA
GTVSHDDK_2_sum_NA

HIIDGVK_2_sum_NA
IAINGFGR_2_sum_NA
IATYQERDPANLPWGSGLK_2_sum_NA
IATYQERDPANLPWGSGLK_3_sum_NA
IDVAVDSTGVFK_2_sum_NA
IDVAVDSTGVFKELDTAQK_3_sum_NA
KIVSNASC[+57]TTNC[+57]LAPLAK_3_sum_NA



sp|P00410|COX2_YEAST

3 ● IEAVSLPK_2_sum_NA ● LLDTDTSMVVPVDTHIR_3_sum_NA ● LNQVSALIQR_2_sum_NA ● Detected data ○ Censored



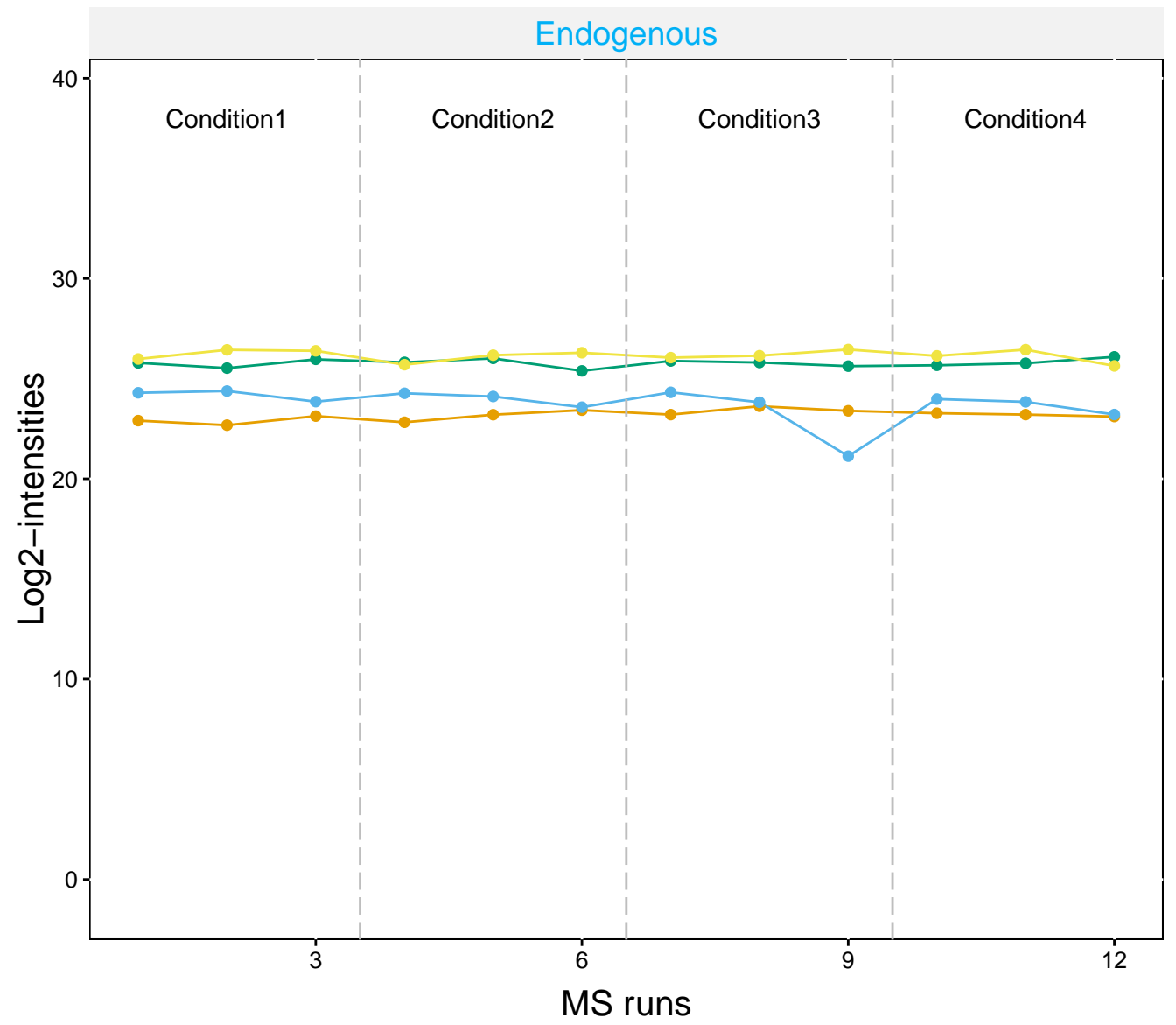
sp|P00425|COX5B_YEAST

peptide: 4

● AAWYISYGEWGPR_2_sum_NA
● ATLTDLPER_2_sum_NA

● ATLTDLPERWENMPNLEQK_3_sum_NA
● TLNNEEIK_2_sum_NA

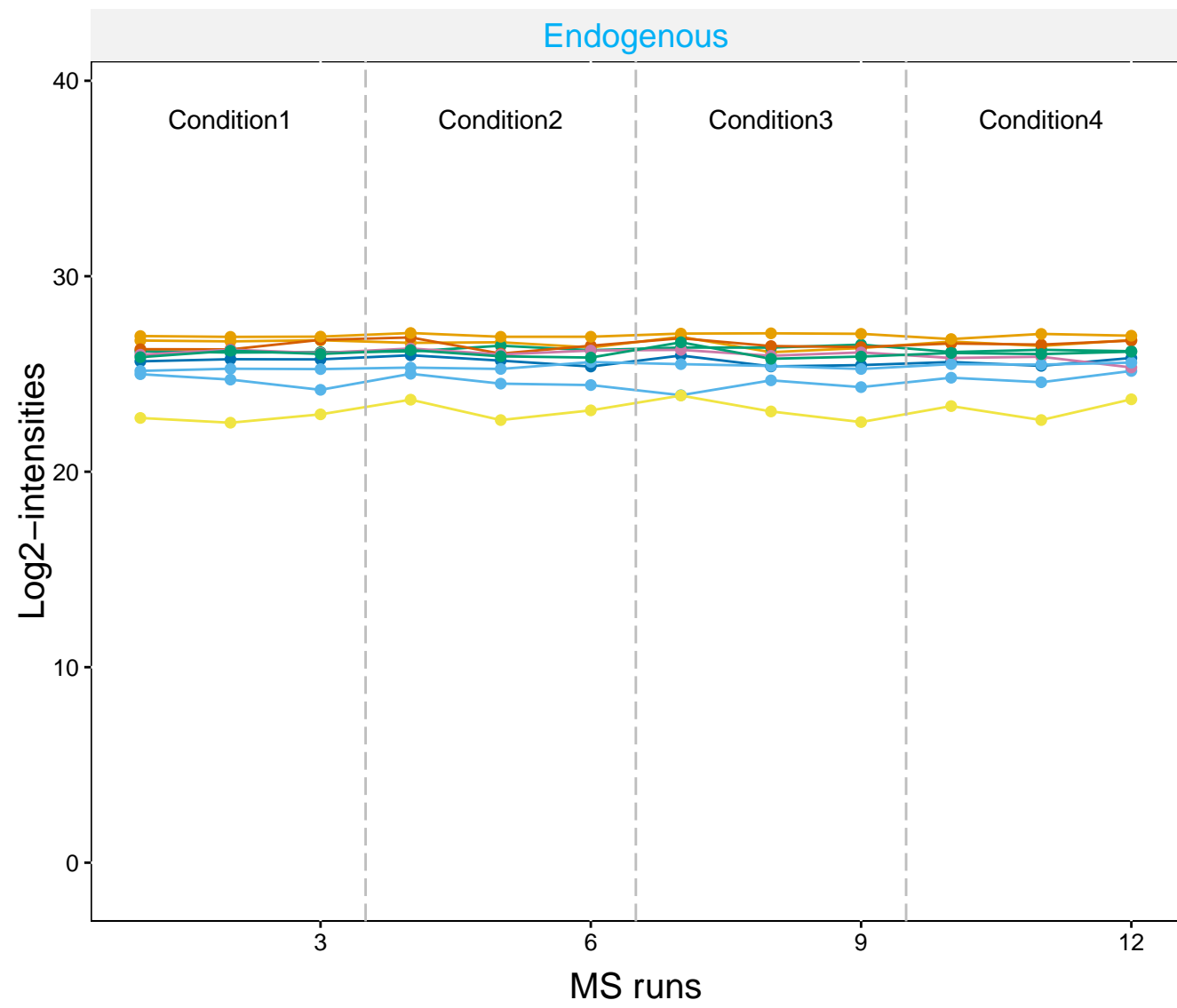
● Detected data



sp|P00427|COX6_YEAST

10

- AYLDELK_2_sum_NA
- AYLDELKDVR_3_sum_NA
- EFDEAYDLFEVQR_2_sum_NA
- Q[-17]ELGVPLKEELFPSSS_2_sum_NA
- QELGVPLK_2_sum_NA
- QELGVPLKEELFPSSS_2_sum_NA
- RVNDLPTAIR_2_sum_NA
- VLNNC[+57]FSYDLVPAPAVIEK_2_sum_NA
- VLNNC[+57]FSYDLVPAPAVIEK_3_sum_NA
- VNDLPTAIR_2_sum_NA

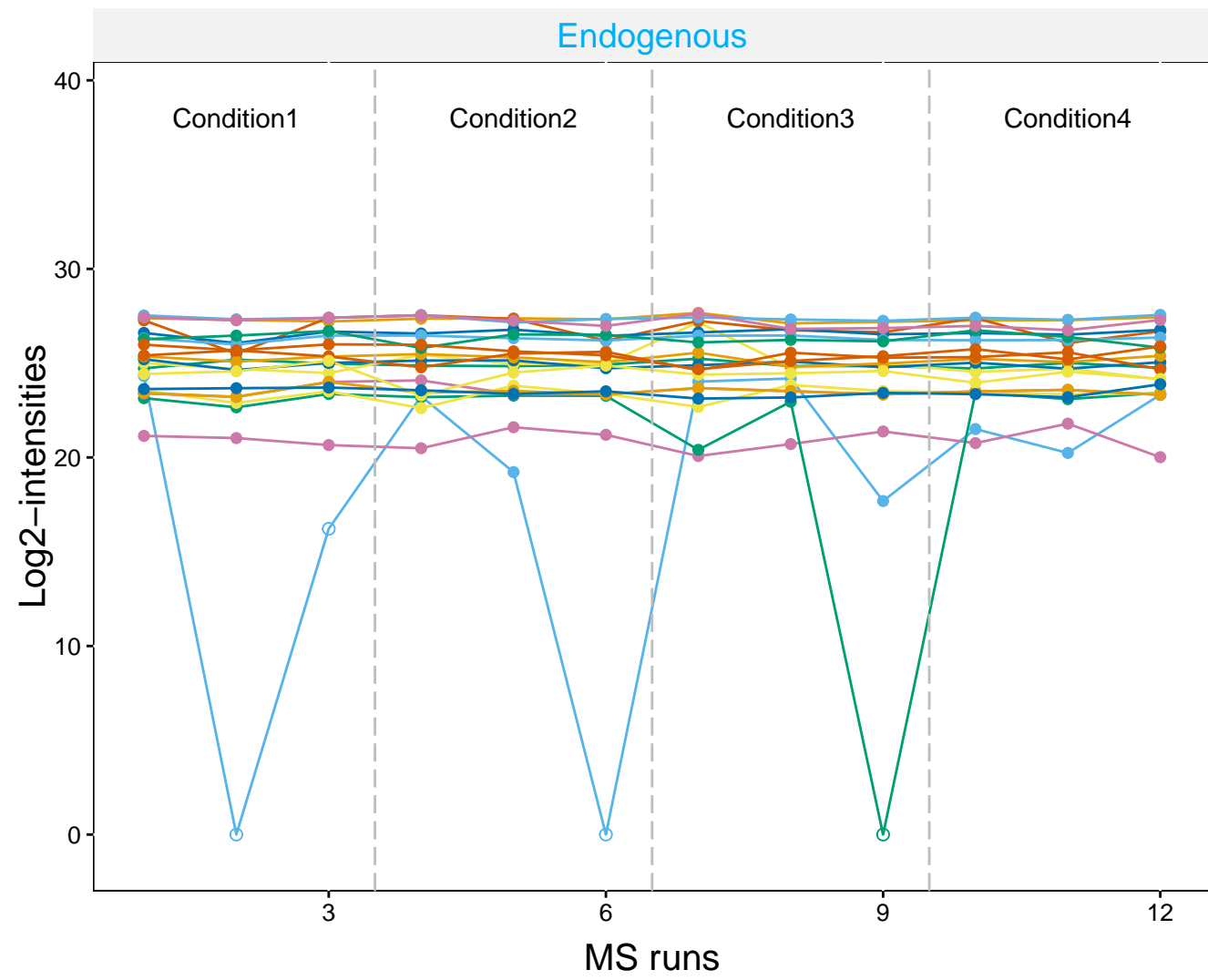


sp|P00431|CCPR_YEAST

Filtered missing data

peptide: 21

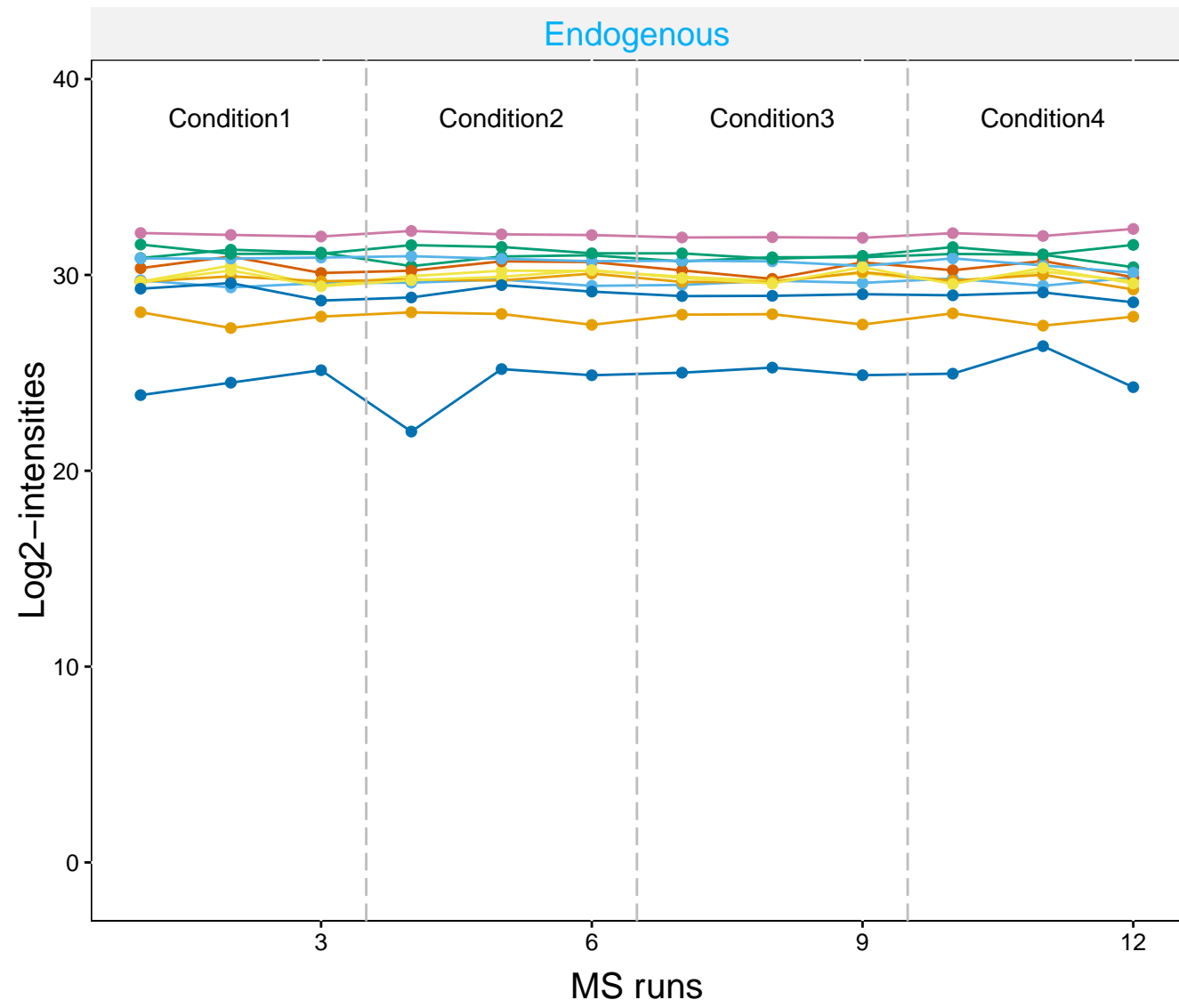
DAPSPFIFK_2_sum_NA
E[-18]VVALM[+16]GAHALGK_3_sum_NA
EFNDPSNAGLQNGFK_2_sum_NA
EVVALMGAHALGK_2_sum_NA
EVVALMGAHALGK_3_sum_NA
FLEPIHK_2_sum_NA
HDNTGGSYGGTYR_2_sum_NA
KEFNDPSNAGLQNGFK_3_sum_NA
LLENGITFPK_2_sum_NA
LNMNDREVVALMGAHALGK_4_sum_NA
LREDDEYDNYIGYGPVLVR_2_sum_NA
LREDDEYDNYIGYGPVLVR_3_sum_NA
NDANNEQWDSK_2_sum_NA
SGYM[+16]MLPTDYSLIQDPK_2_sum_NA



sp|P00445|SODC_YEAST

e: 12

- FEQASESEPTTVSYEIAGNSPNAER_2_sum_NA
- FEQASESEPTTVSYEIAGNSPNAER_3_sum_NA
- GDAGVSGVVK_2_sum_NA
- GDTEESLK_2_sum_NA
- HVGDM(+16)GNVK_2_sum_NA
- HVGDMGNVK_2_sum_NA
- LIGPTSVVGR_2_sum_NA
- SVVIHAGQDDLKG_2_sum_NA
- SVVIHAGQDDLKG_3_sum_NA
- TGNAGPRPAC(+57)GVIGLTN_2_sum_NA
- THGAPTDEVR_2_sum_NA
- THGAPTDEVR_3_sum_NA



sp|P00447|SODM_YEAST

● Detected data

peptide: 6

● AIWNVVNWK_2_sum_NA

● GGLSLLSTTAR_2_sum_NA

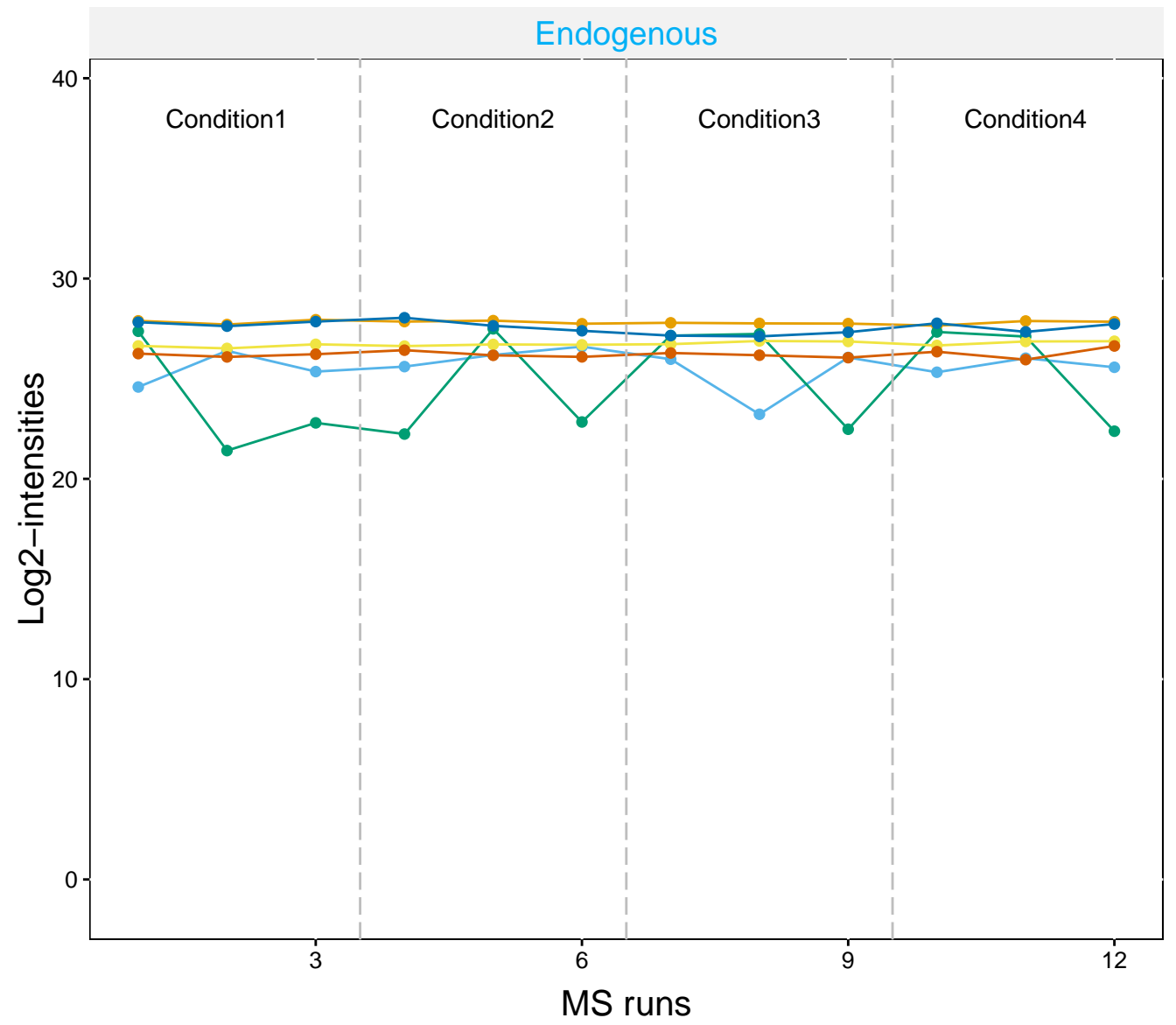
● MIAIQQNIK_2_sum_NA

● EPSPANAR_2_sum_NA

● LAGVQGSGWAFIVK_2_sum_NA

● VTLPDLK_2_sum_NA

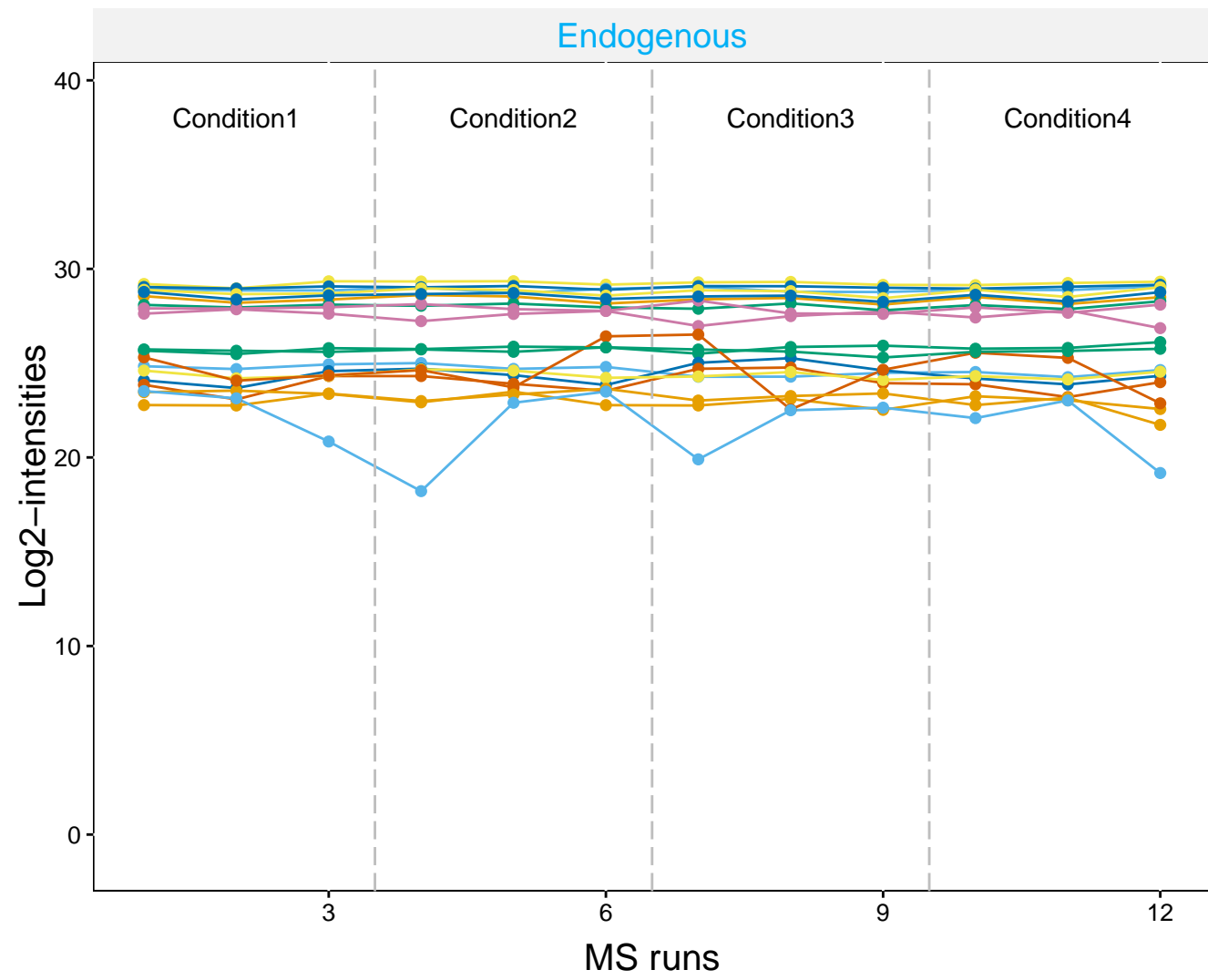
Endogenous



sp|P00498|HIS1_YEAST

de: 19

- C[+57]DLGITGVDQVR_2_sum_NA
- FVSC[+57]IYNAPEDK_2_sum_NA
- FVSC[+57]IYNAPEDKLPPELLK_2_sum_NA
- FVSC[+57]IYNAPEDKLPPELLK_3_sum_NA
- IDDEGWVAVSSMIER_2_sum_NA
- IDDEGWVAVSSMIER_3_sum_NA
- KPEQLIGK_2_sum_NA
- KPEQLIGK_3_sum_NA
- LLFAIPK_2_sum_NA
- LQVQVPVNGEYK_2_sum_NA
- LQVQVPVNGEYKKPEQLIGK_4_sum_NA
- M[+42]DLVNHILTDR_2_sum_NA
- RAPTISK_2_sum_NA
- SLIATIK_2_sum_NA
- SRIEGVMTAQR_2_sum_NA
- SVSILNGADITFHR_2_sum_NA
- SVSILNGADITFHR_3_sum_NA
- TIVTSFVK_2_sum_NA
- YFADLEGTTVEK_2_sum_NA

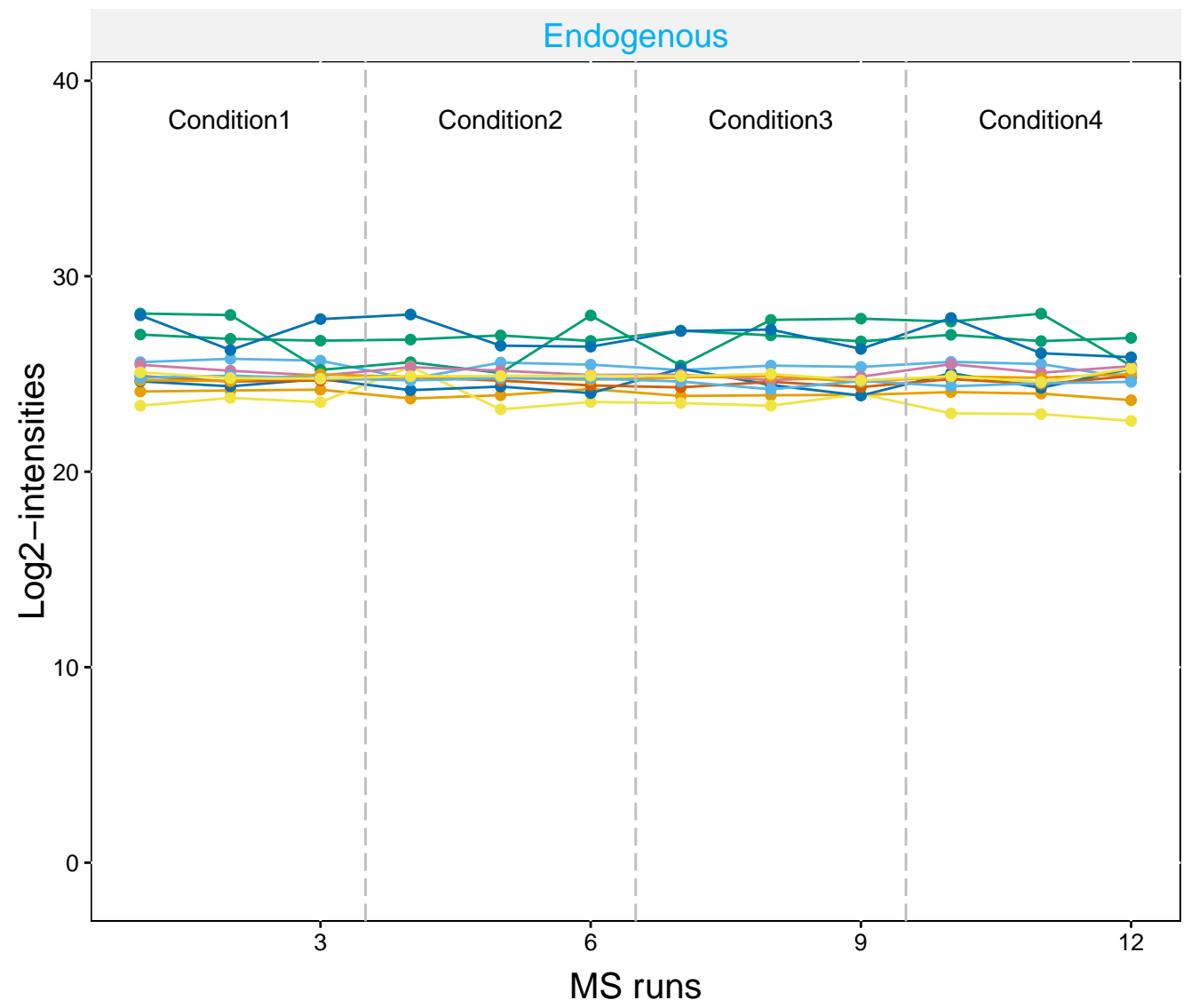


sp|P00546|CDK1_YEAST

peptide: 12

● Detected

● ALDLRPGQGQR_2_sum_NA	● FMMQLC[+57]K_2_sum_NA	● LESEDEGVPSTAIR_2_sum_NA
● ALDLRPGQGQR_3_sum_NA	● GIDLLDK_2_sum_NA	● LLAYDPINR_2_sum_NA
● DLSQVPSLDPR_2_sum_NA	● IRLESEDEGVPSTAIR_3_sum_NA	● RYMEGIPK_2_sum_NA
● DQPLGADIVKK_2_sum_NA	● KPIFGDSEIDQIFK_3_sum_NA	● VGEGTYGVVYK_2_sum_NA

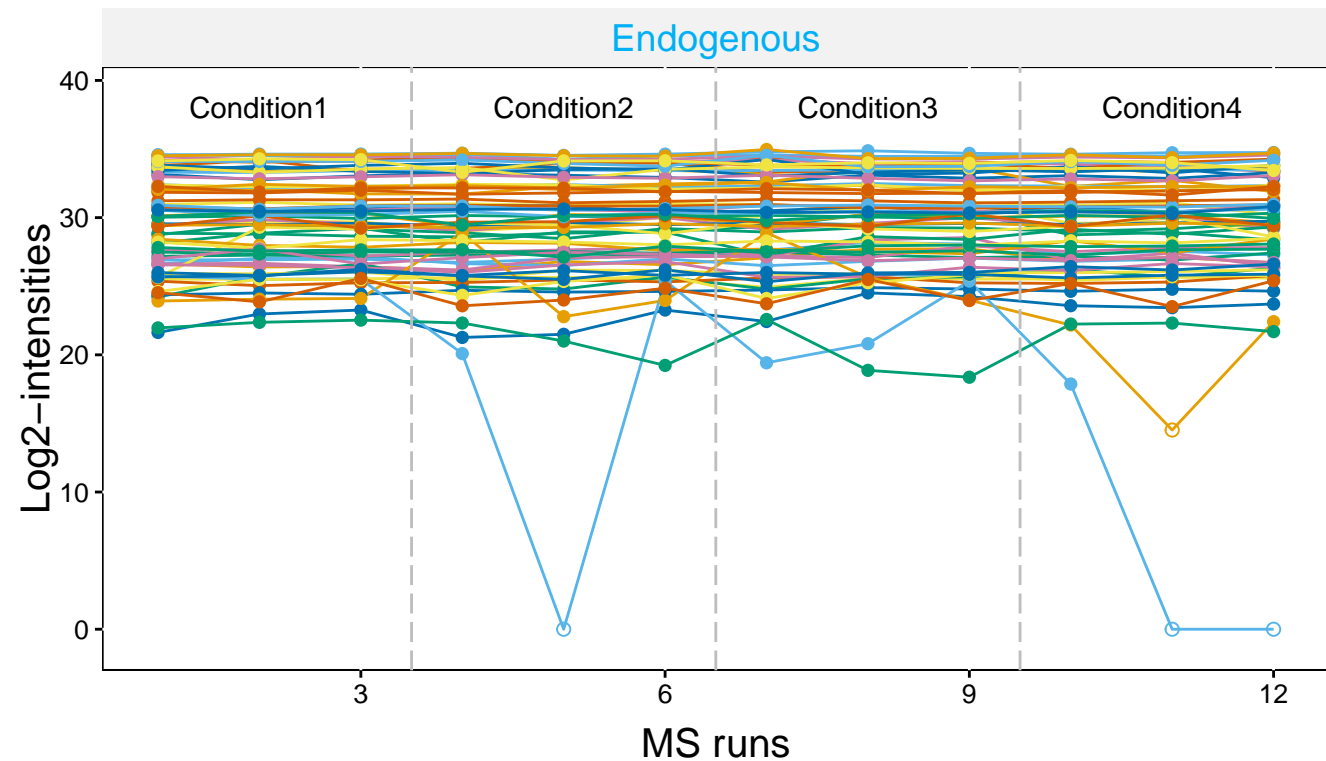


sp|P00549|KPYK1_YEAST

ing data

peptide: 69

● AEVSDVGNAILDGADC[+57]VM[+16]LSGETAK_2_sum_NA
● AEVSDVGNAILDGADC[+57]VM[+16]LSGETAK_3_sum_NA
● AEVSDVGNAILDGADC[+57]VMLSGETAK_2_sum_NA
● AEVSDVGNAILDGADC[+57]VMLSGETAK_3_sum_NA
● AEVSDVGNAILDGADC[+57]VMLSGETAK_4_sum_NA
● AGAGHSNTLQVSTV_2_sum_NA
● AGLNIVR_2_sum_NA
● AIIVLSTSGTTTPR_2_sum_NA
● AIIVLSTSGTTTPR_3_sum_NA
● E[-18]FGILKK_2_sum_NA
● EFGILKK_2_sum_NA
● EPVSDWTDDVEAR_2_sum_NA
● EVLGEQGG_2_sum_NA
● EVLGEQGGKDVK_2_sum_NA
● EVLGEQGGKDVK_3_sum_NA
● GDLGIEIPAEVLAVQK_2_sum_NA
● GDLGIEIPAEVLAVQK_3_sum_NA
● GDLGIEIPAEVLAVQKK_3_sum_NA
● GDTYYSIQGFK_2_sum_NA
● GVFPFVFEK_2_sum_NA
● GVFPFVFEKEPVSQWTDDEAR_2_sum_NA
● GVFPFVFEKEPVSQWTDDEAR_3_sum_NA
● GVNLPGTDVDPALSEK_2_sum_NA
● GVNLPGTDVDPALSEK_3_sum_NA
● GVNLPGTDVDPALSEKDK_2_sum_NA
● GVNLPGTDVDPALSEKDKEDLR_2_sum_NA
● GVNLPGTDVDPALSEKDKEDLR_3_sum_NA
● GVNLPGTDVDPALSEKDKEDLR_4_sum_NA
● IENQQGVNNFDEILK_2_sum_NA
● IENQQGVNNFDEILK_3_sum_NA
● IM[+16]YVDYK_2_sum_NA
● IMYVDYK_2_sum_NA
● INFGIEK_2_sum_NA
● KGDTYYSIQGFK_2_sum_NA
● KGDTYYSIQGFK_3_sum_NA
● LTSLNVVAGSDLR_2_sum_NA
● LTSLNVVAGSDLR_2_sum_NA
● LTSLNVVAGSDLR_3_sum_NA
● LTSLNVVAGSDLR_3_sum_NA
● NC[+57]TPKPTSTTETVAASAVAAVFEQK_2
● NC[+57]TPKPTSTTETVAASAVAAVFEQK_3
● NC[+57]TPKPTSTTETVAASAVAAVFEQK_4
● NGVHVM[+16]VFASFIR_2_sum_NA
● NGVHVM[+16]VFASFIR_3_sum_NA
● NGVHVM[+16]VFASFIR_3_sum_NA
● NGVHVM[+16]VFASFIR_3_sum_NA



sorted missing data

peptide: 58

AAGFLLEK_2_sum_NA
AGAEIVPK_2_sum_NA
AHSSM[+16]VGFDLPQR_2_sum_NA
AHSSM[+16]VGFDLPQR_3_sum_NA
AHSSMVGFDLPQR_2_sum_NA
AHSSMVGFDLPQR_3_sum_NA
ALENPTRPFLAILGGAK_2_sum_NA

