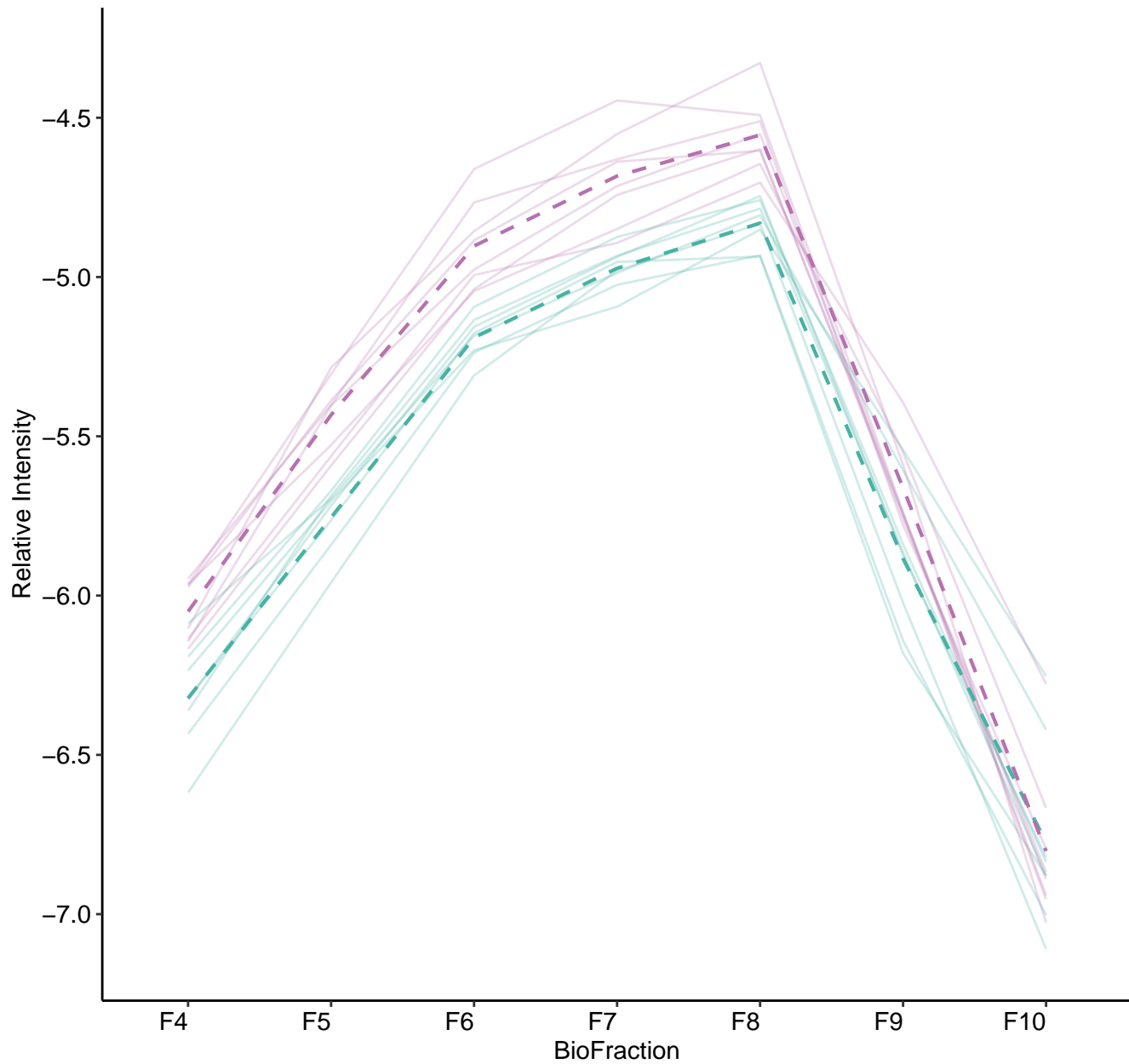
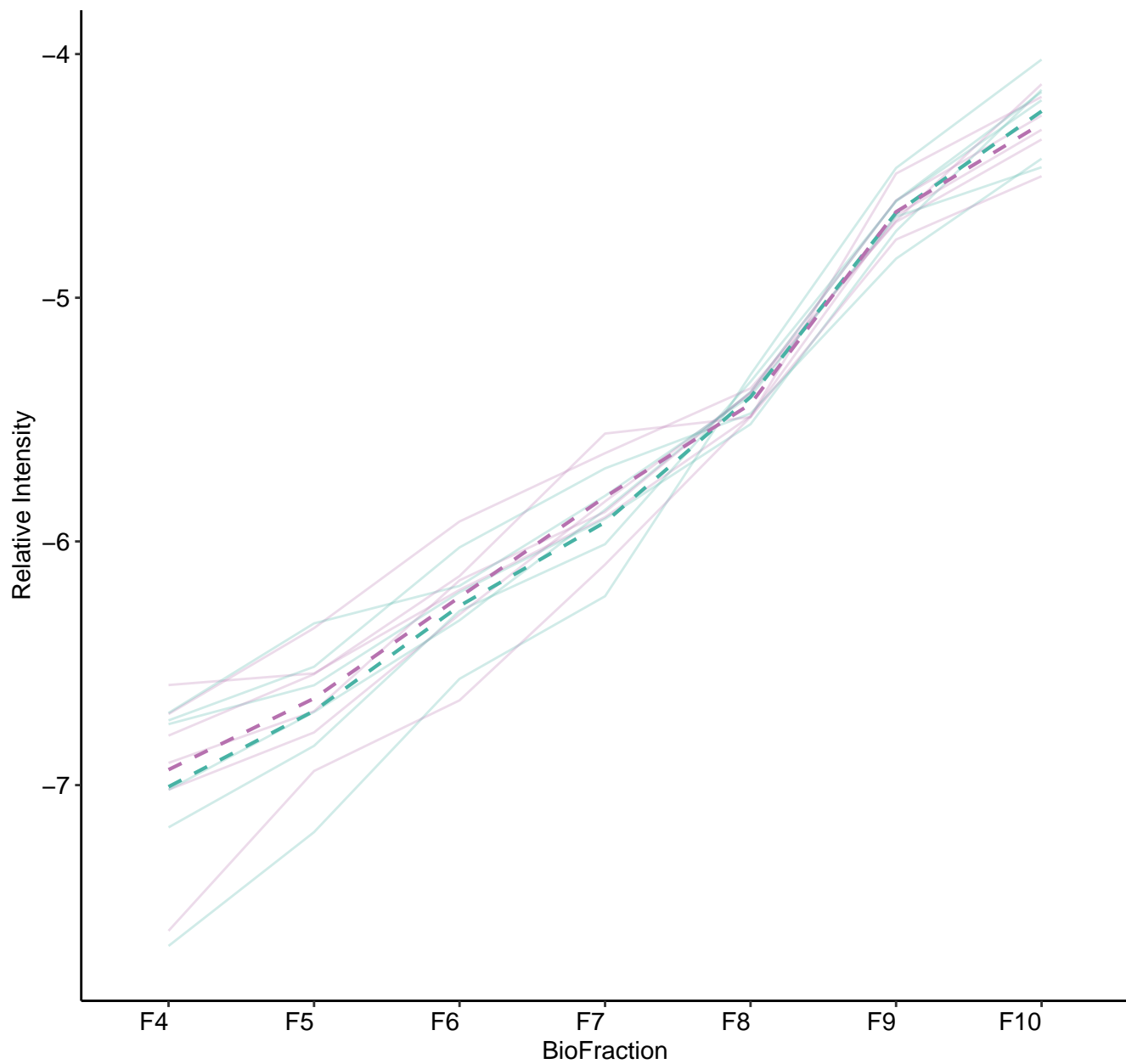


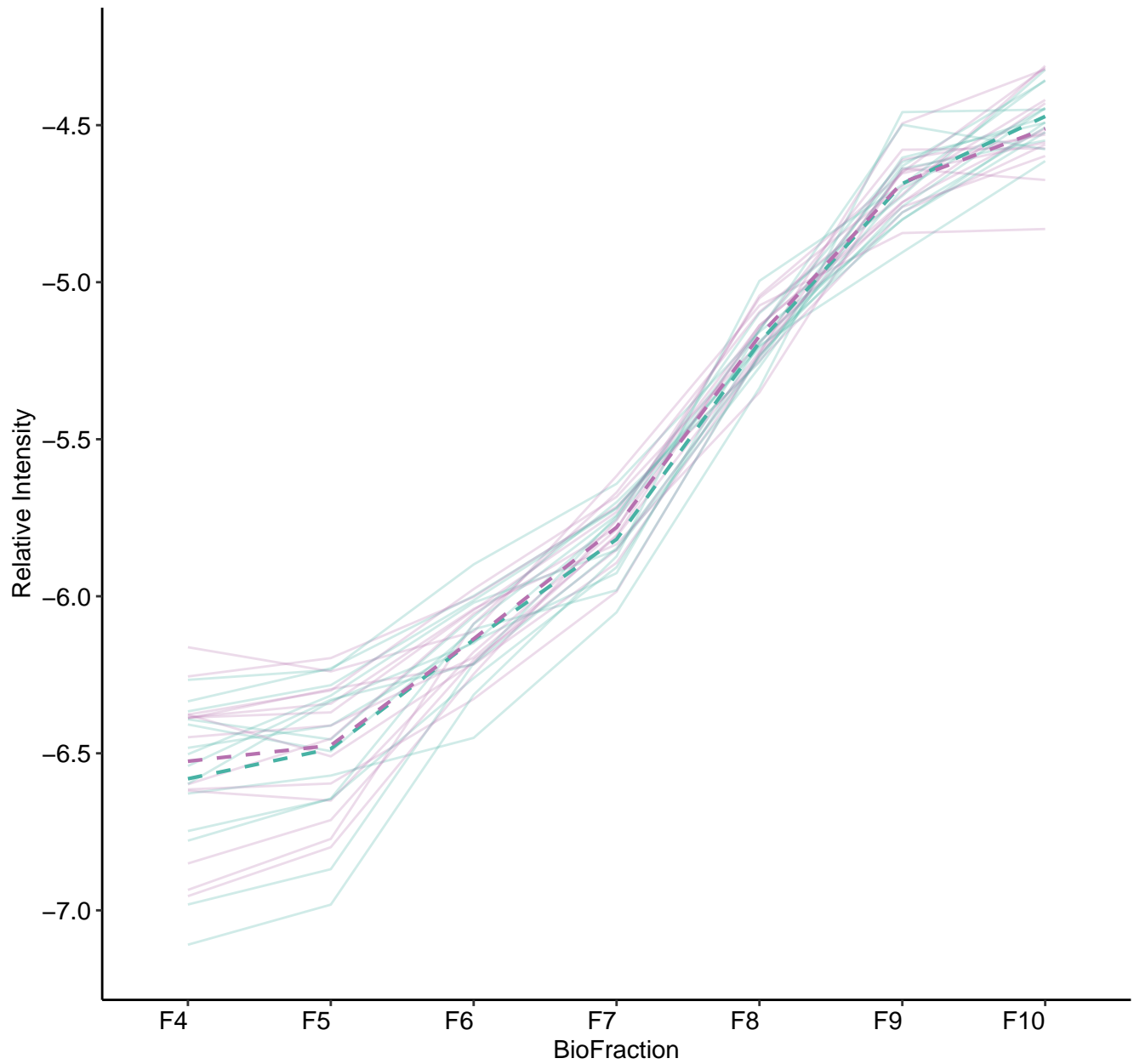
(ER)-localized multiprotein complex, Ig heavy chains associated (n = 8)
(R2.Fixef = 0.956)



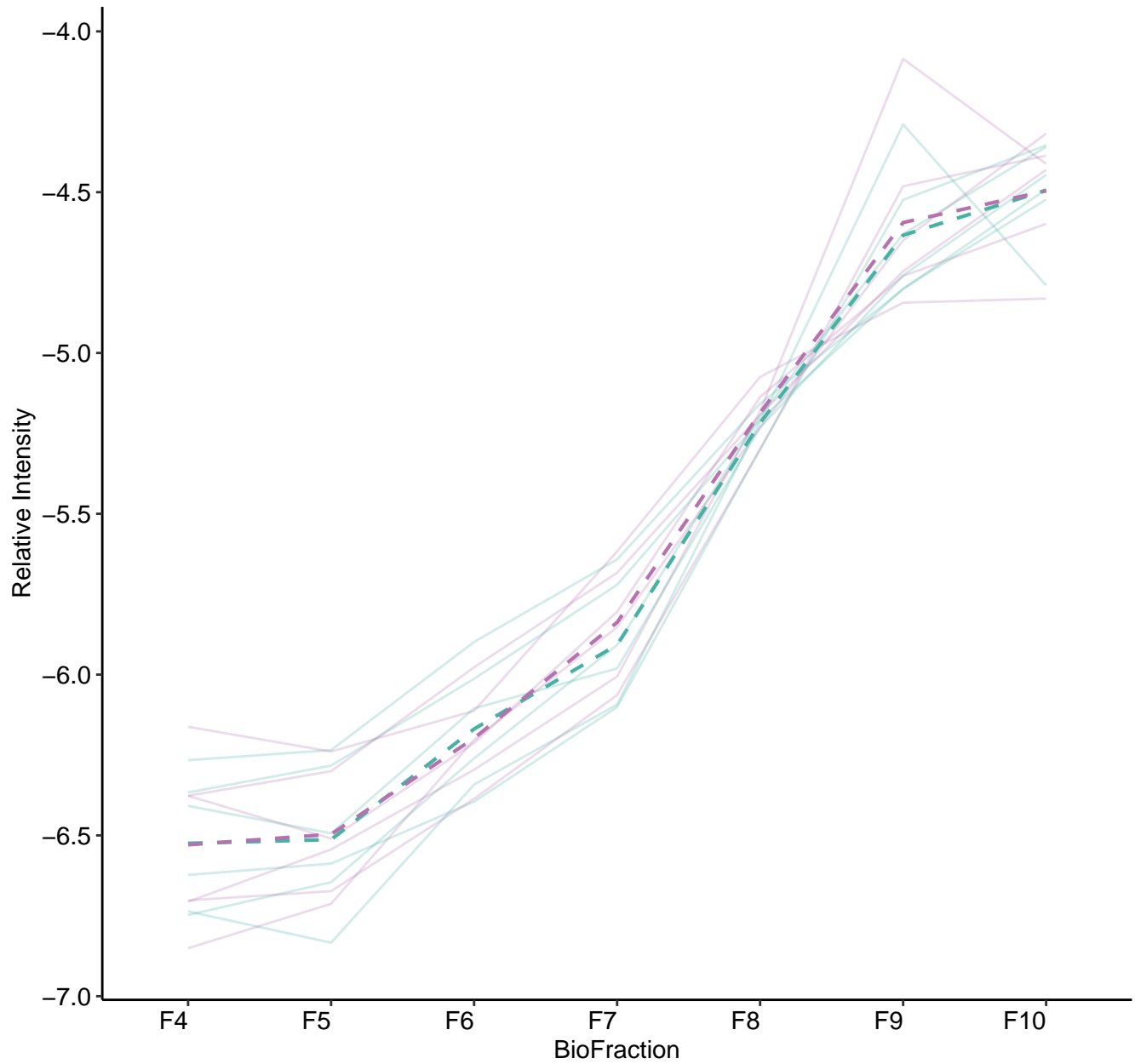
12S U11 snRNP (n = 6)
(R2.Fixef = 0.951)



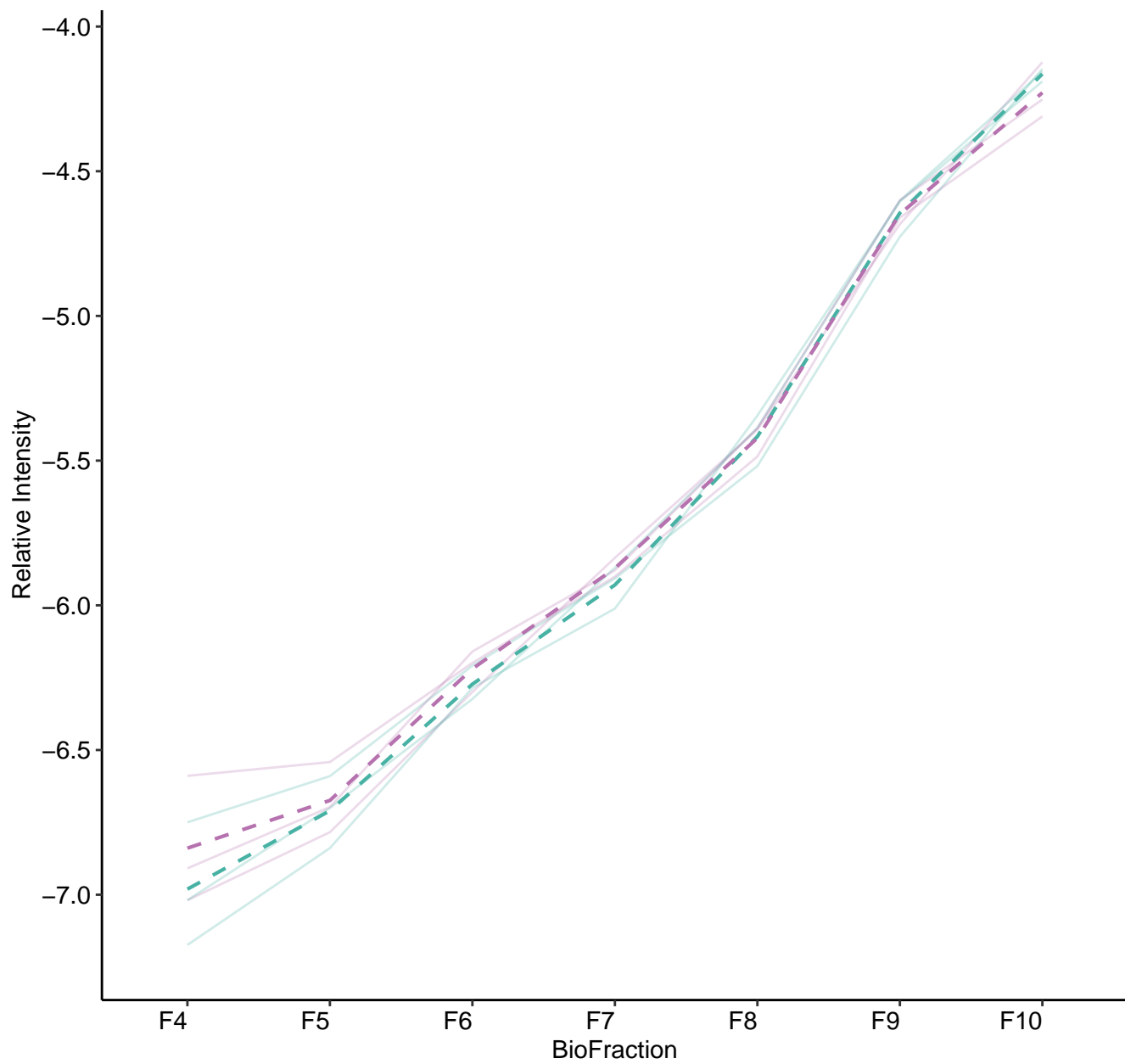
17S U2 snRNP (n = 14)
(R2.Fixef = 0.962)



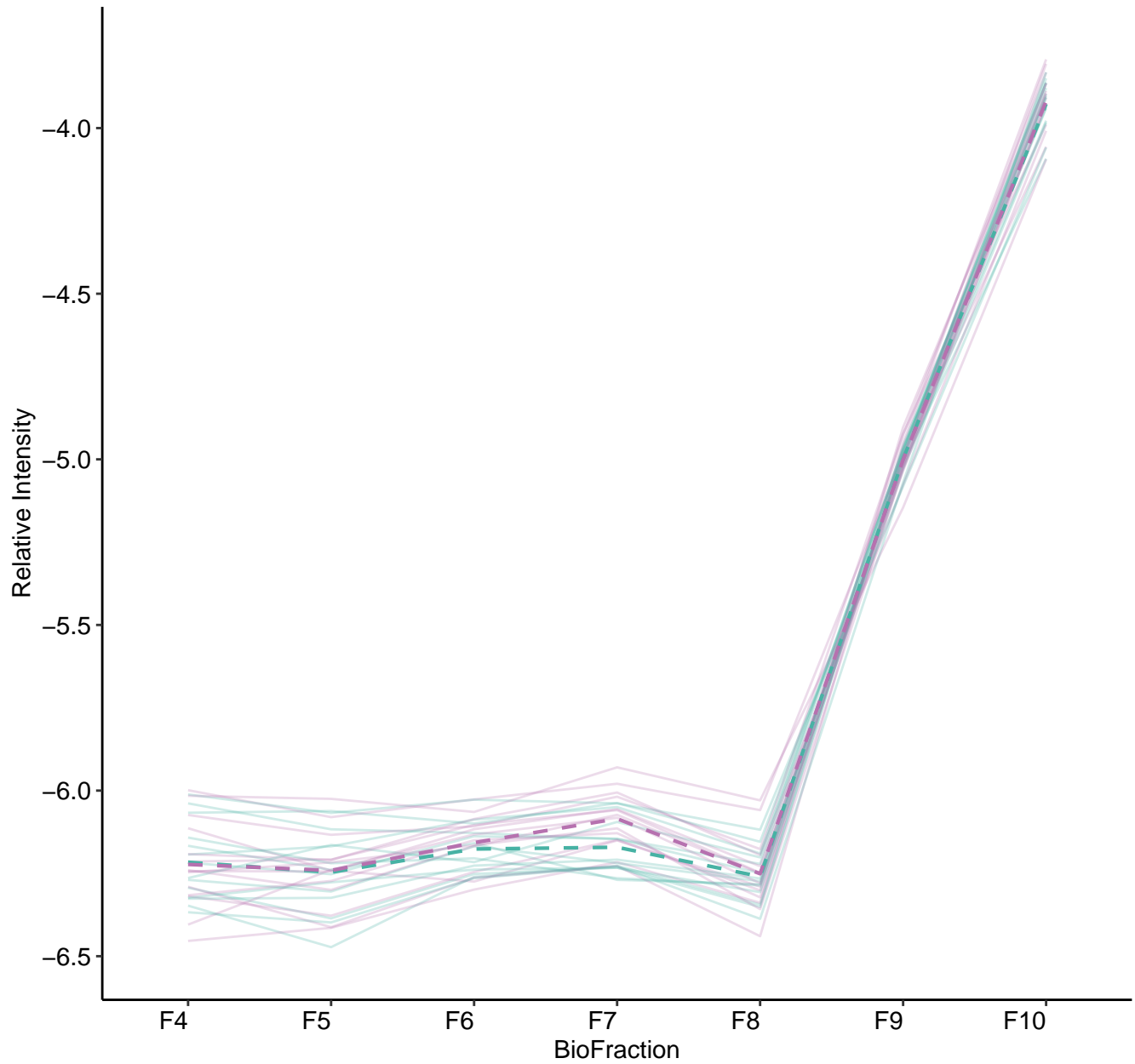
18S U11/U12 snRNP (n = 6)
(R2.Fixef = 0.946)



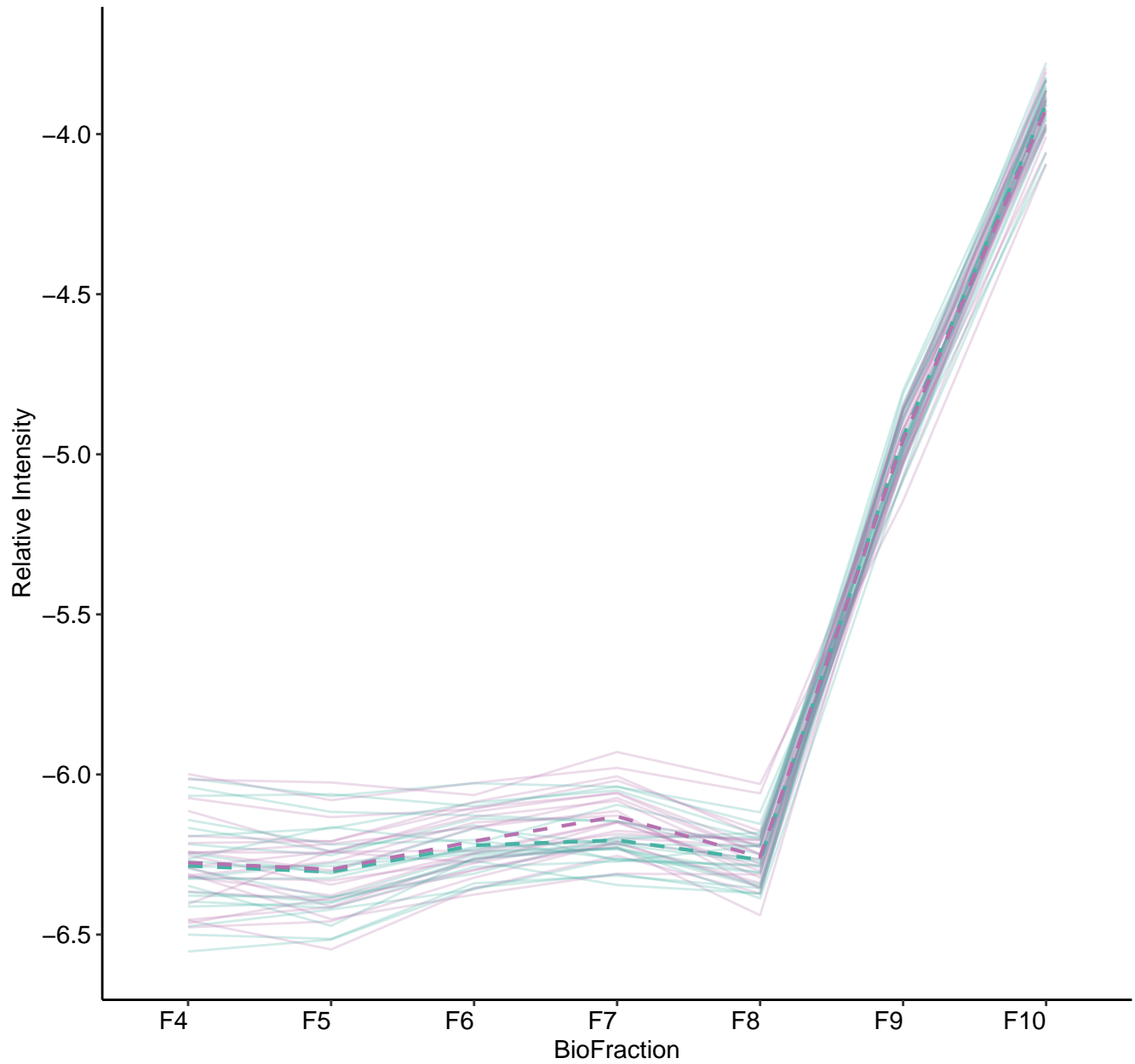
20S methylosome and RG-containing Sm protein complex (n = 3)
(R2.Fixef = 0.987)



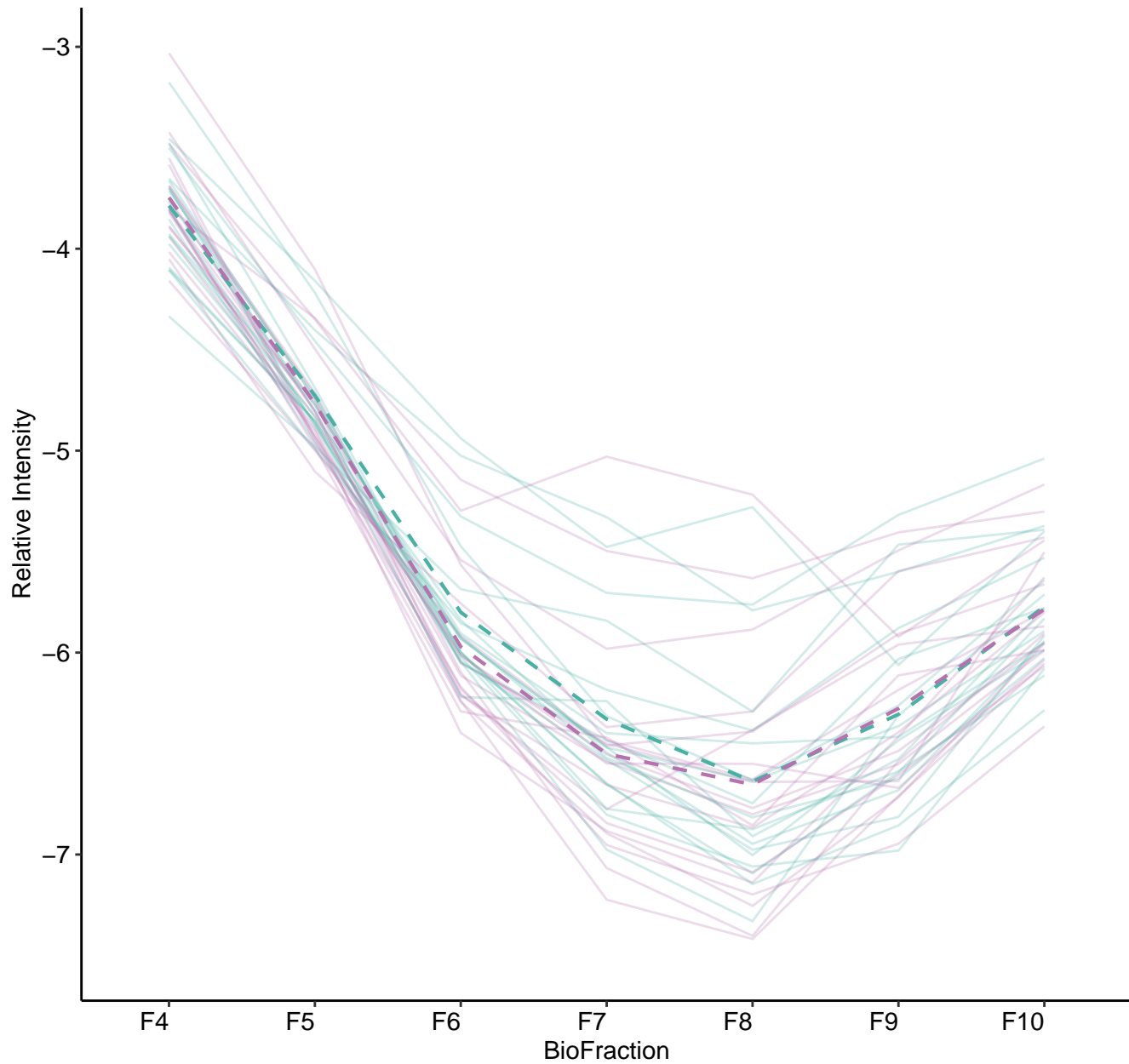
20S proteasome (n = 14)
(R2.Fixef = 0.987)



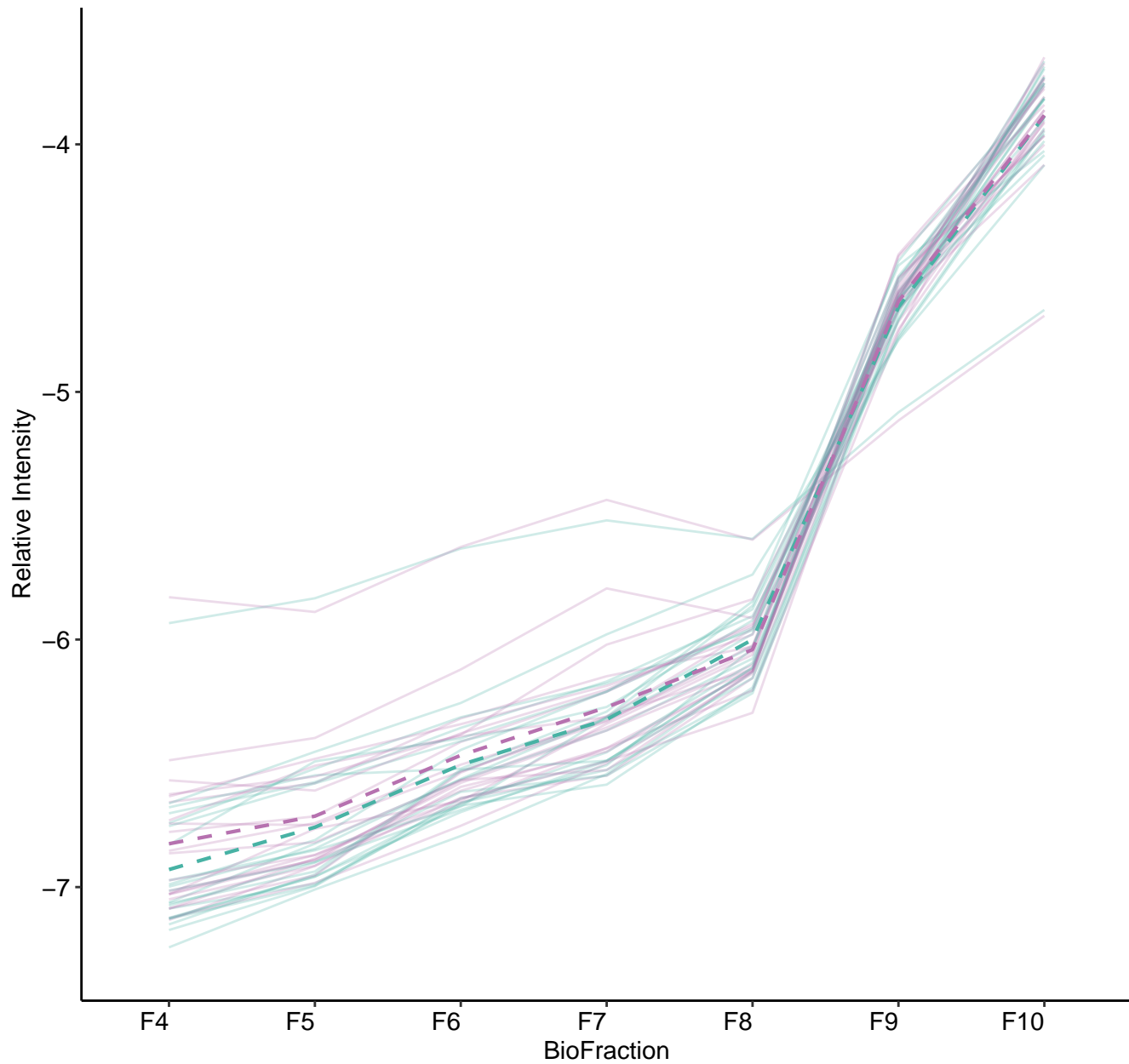
26S proteasome (n = 22)
(R2.Fixef = 0.986)



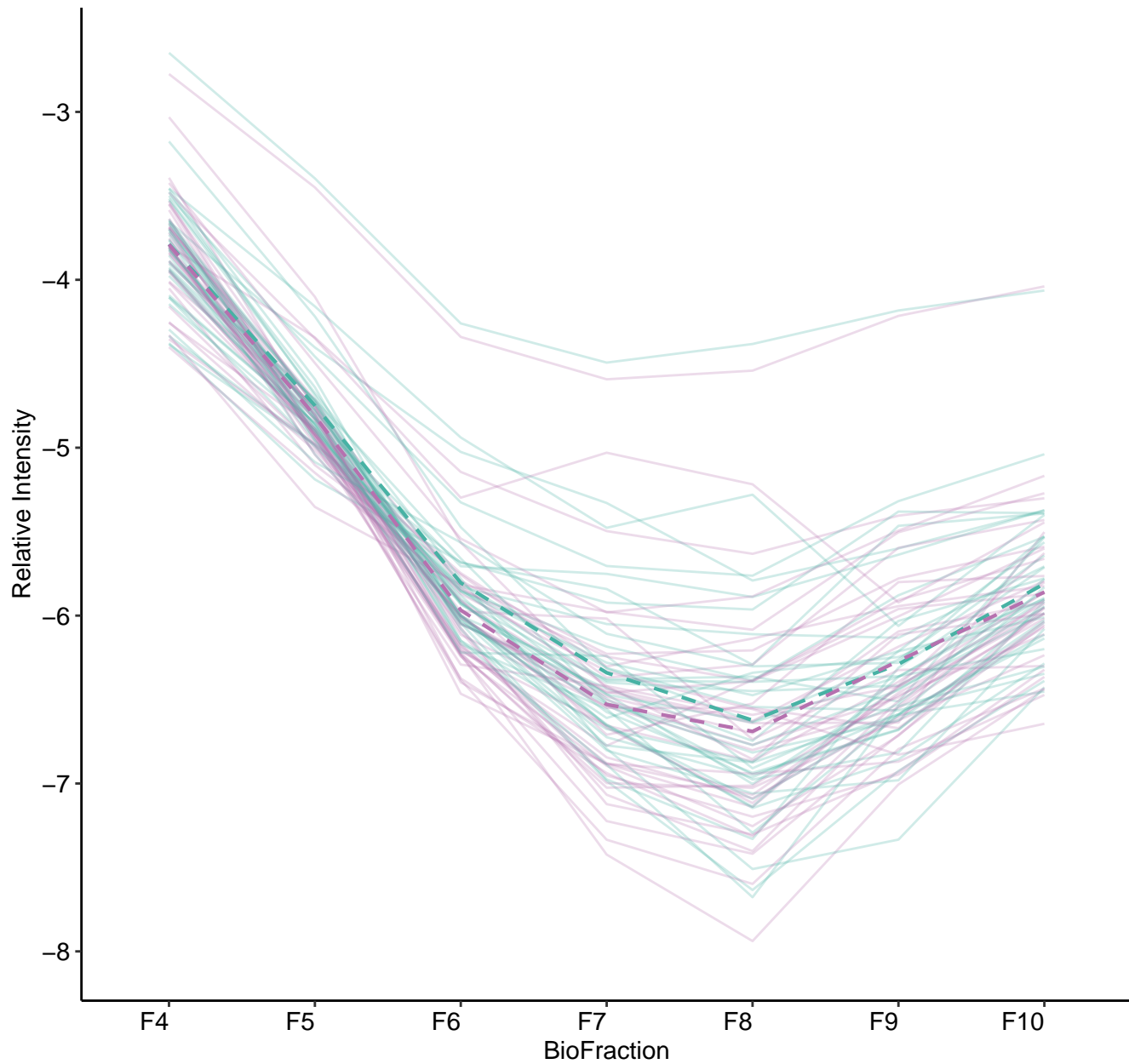
28S ribosomal subunit, mitochondrial (n = 19)
(R2.Fixef = 0.851)



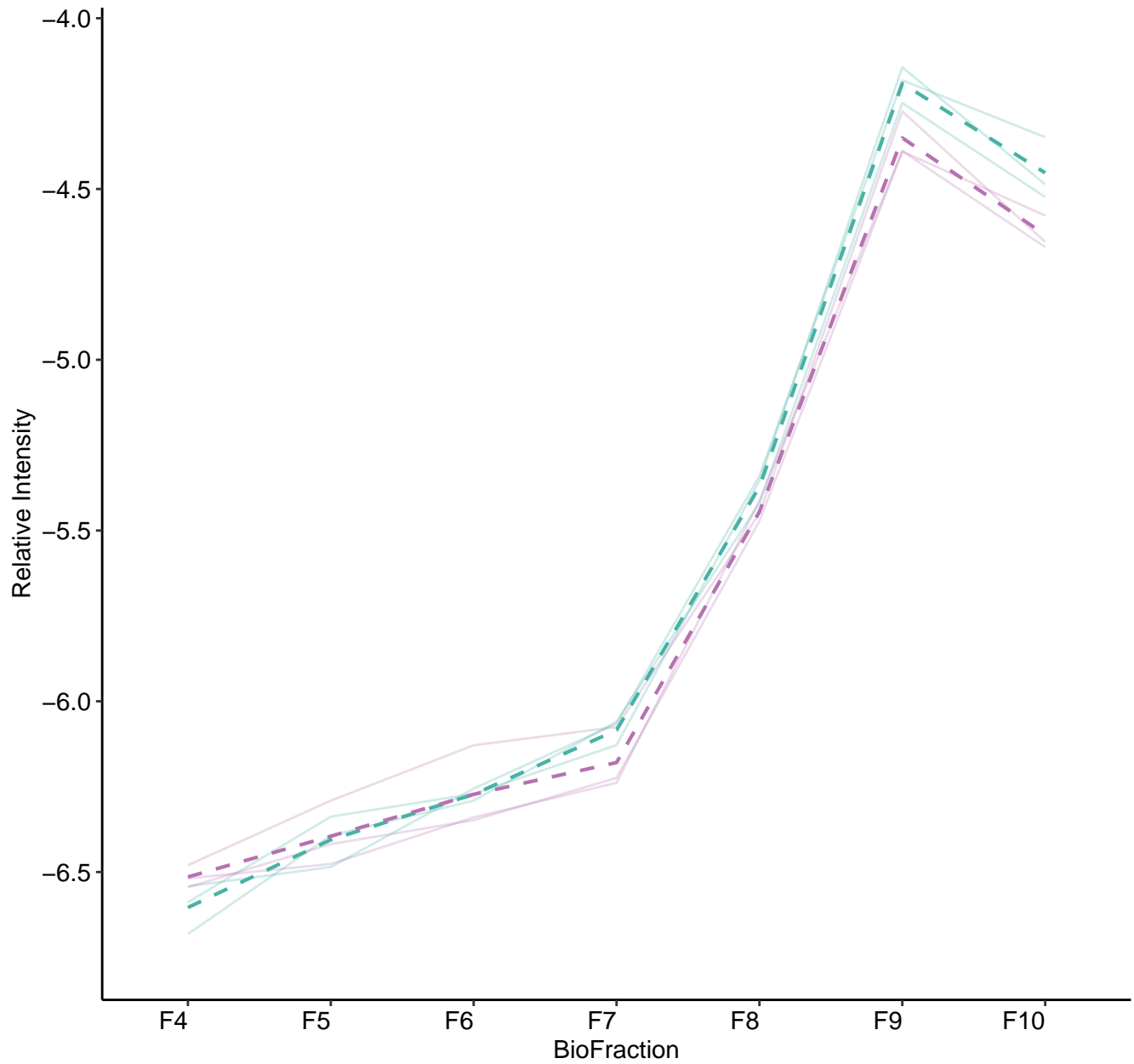
40S ribosomal subunit, cytoplasmic (n = 21)
(R2.Fixef = 0.954)



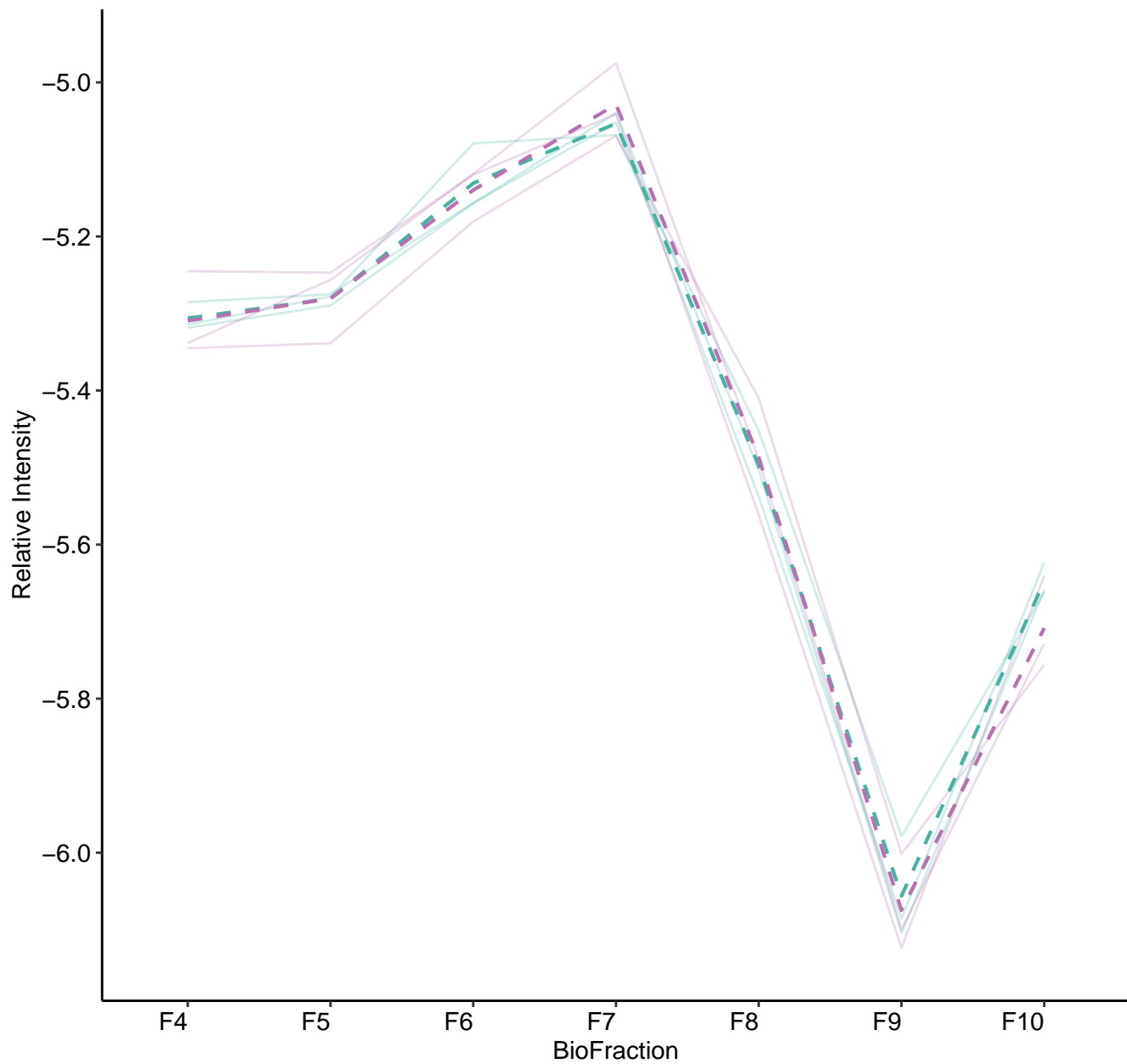
55S ribosome, mitochondrial (n = 38)
(R2.Fixef = 0.799)



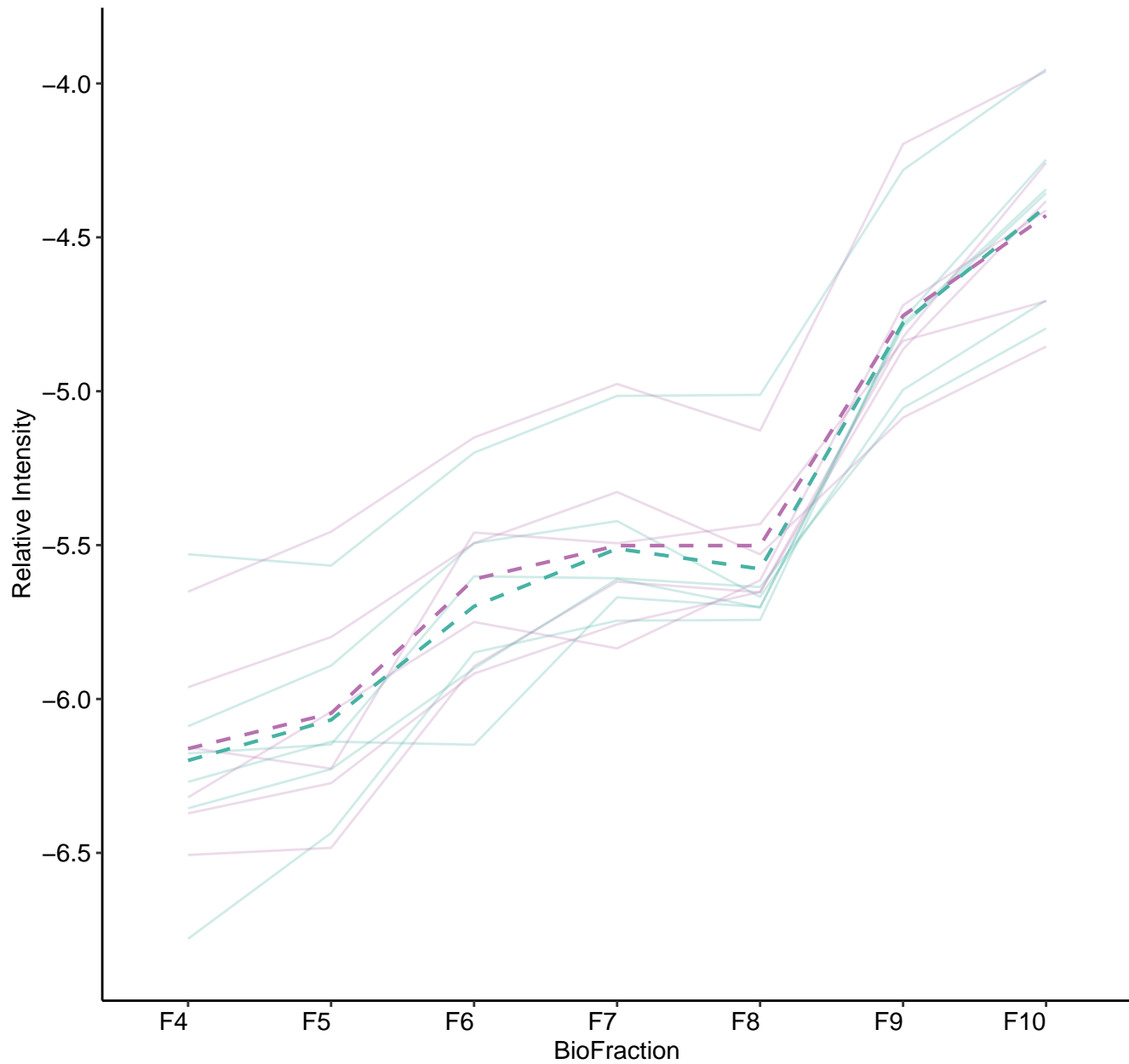
5FMC (Friends of Methylated Chtop) complex (n = 3)
(R2.Fixef = 0.994)



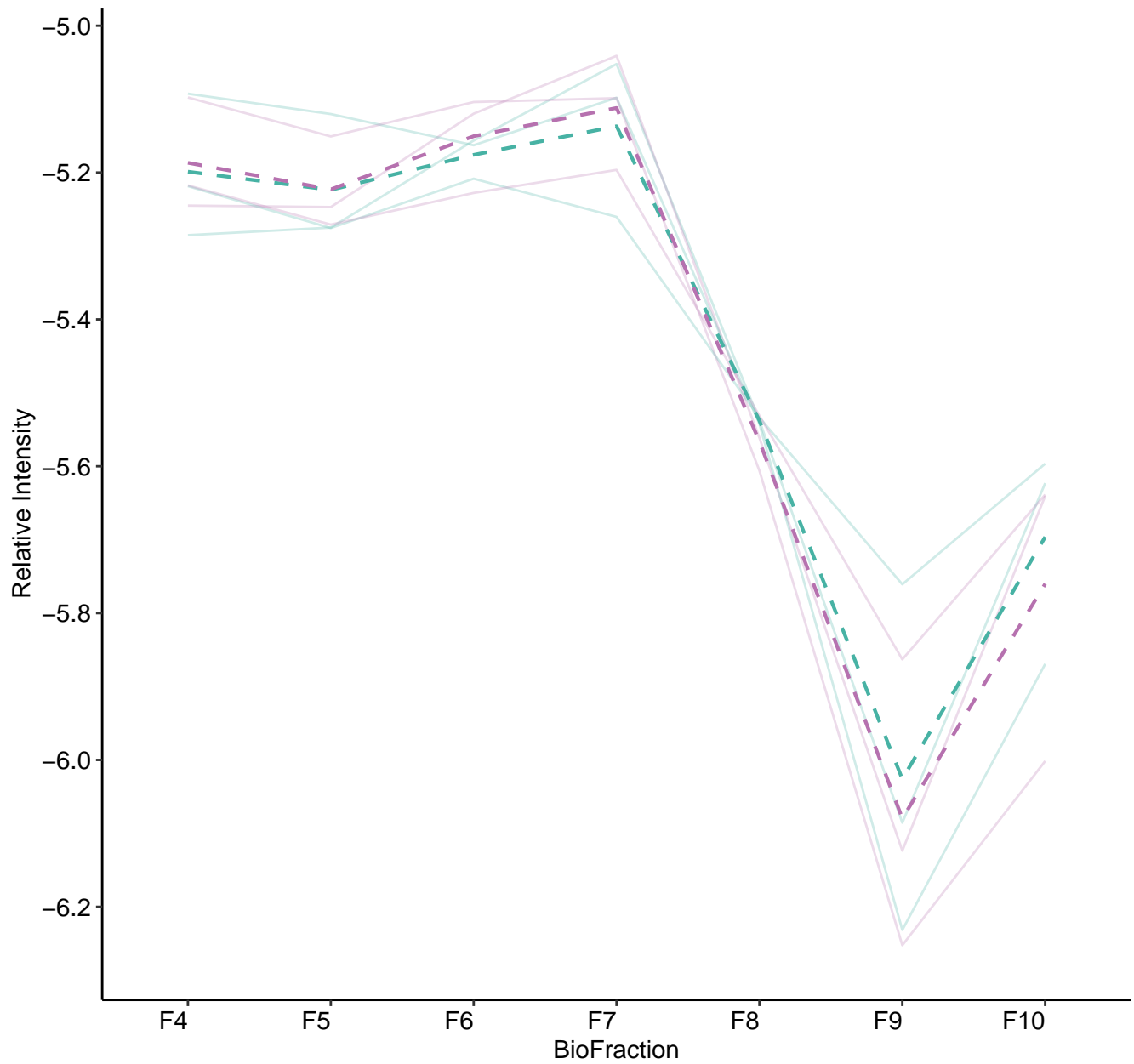
9S–cytosolic aryl hydrocarbon (Ah) receptor non–ligand activated complex (n = 3)
(R2.Fixef = 0.979)



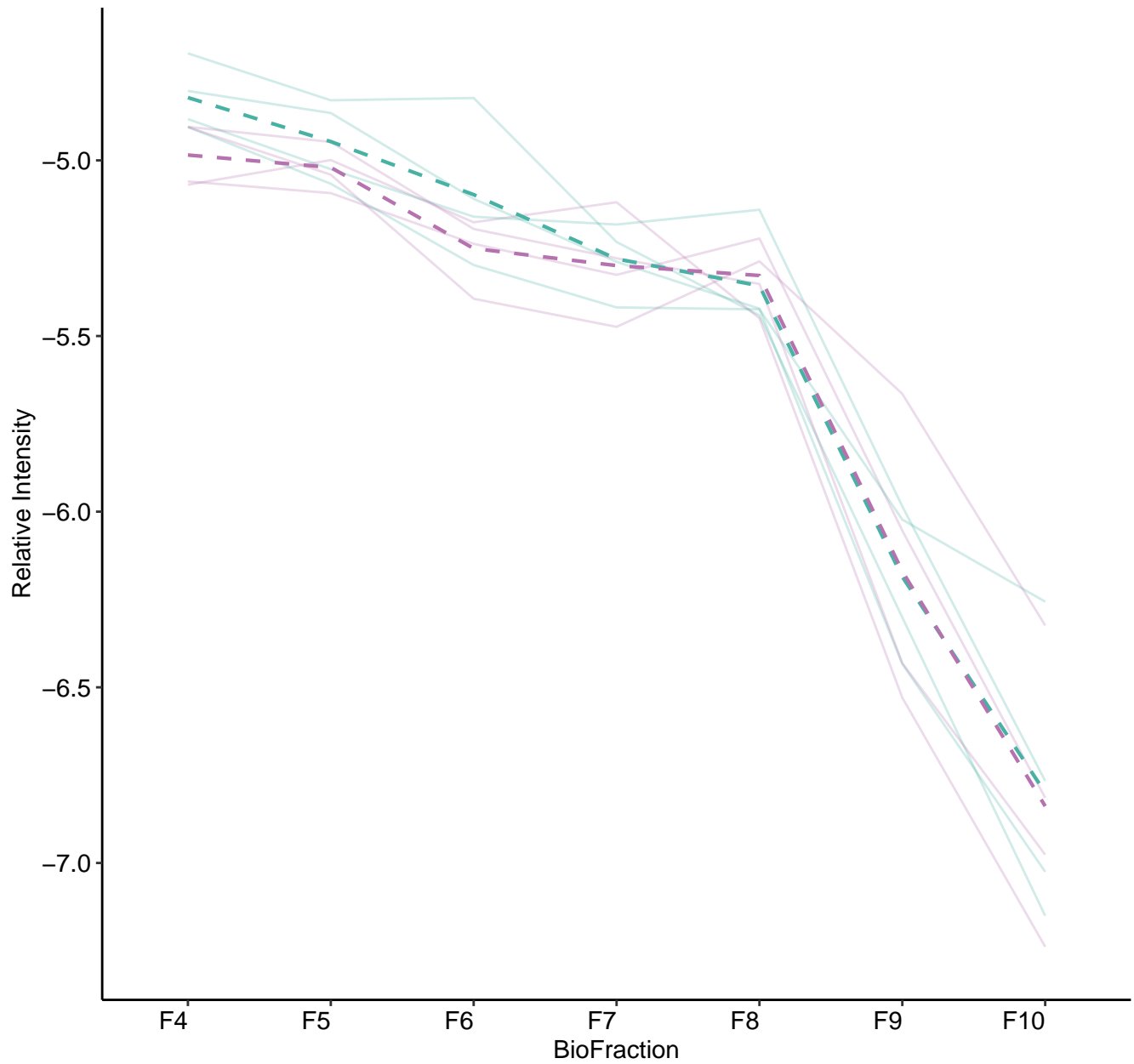
ALL-1 supercomplex (n = 6)
(R2.Fixef = 0.79)



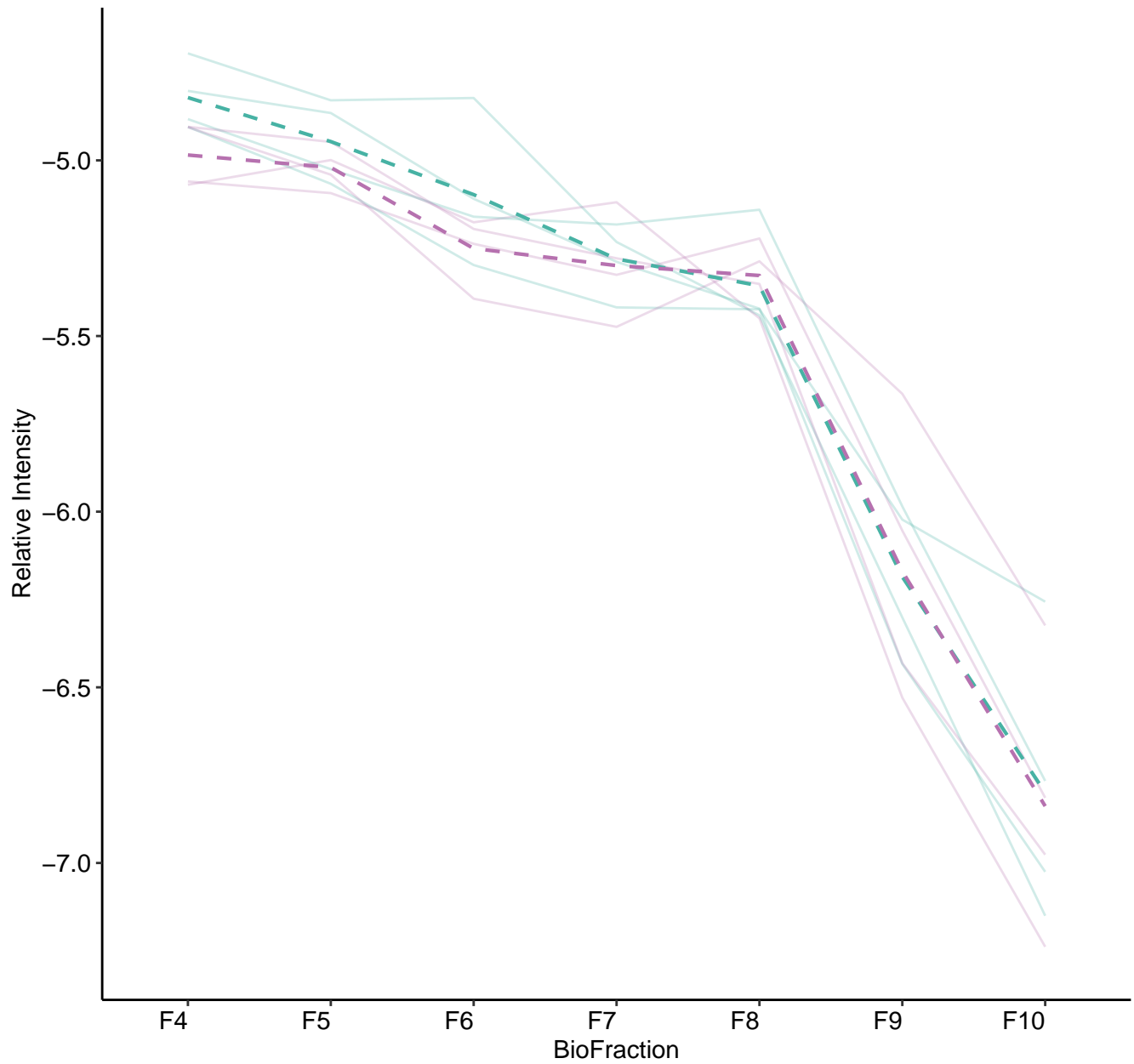
Alpha-GDI-Hsp90 chaperone complex, ATP dependent (n = 3)
(R2.Fixef = 0.878)



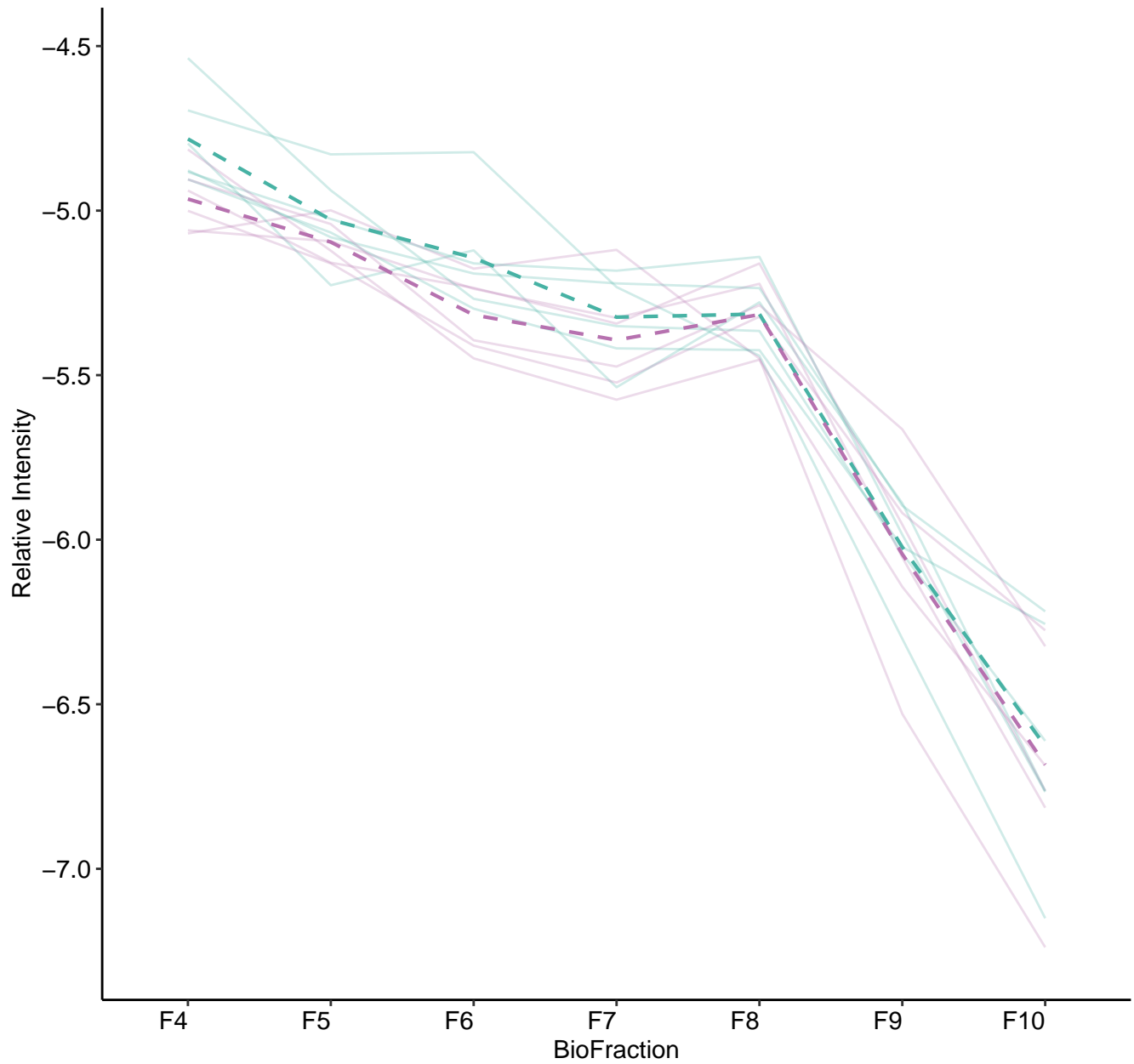
AMPA receptor complex (anti-GluA1-a) (n = 4)
(R2.Fixef = 0.903)



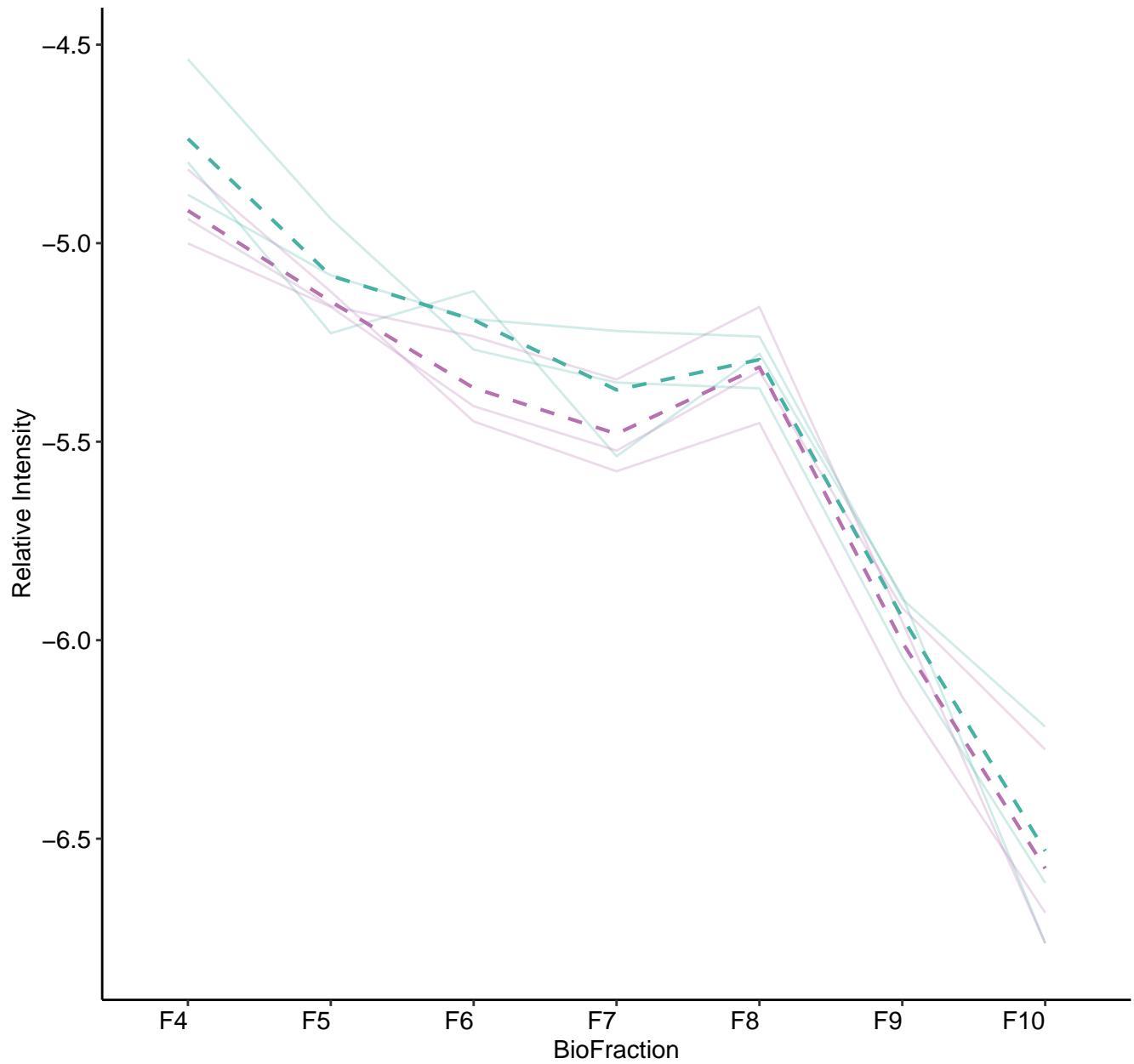
AMPA receptor complex (anti-GluA1-b) (n = 4)
(R2.Fixef = 0.903)



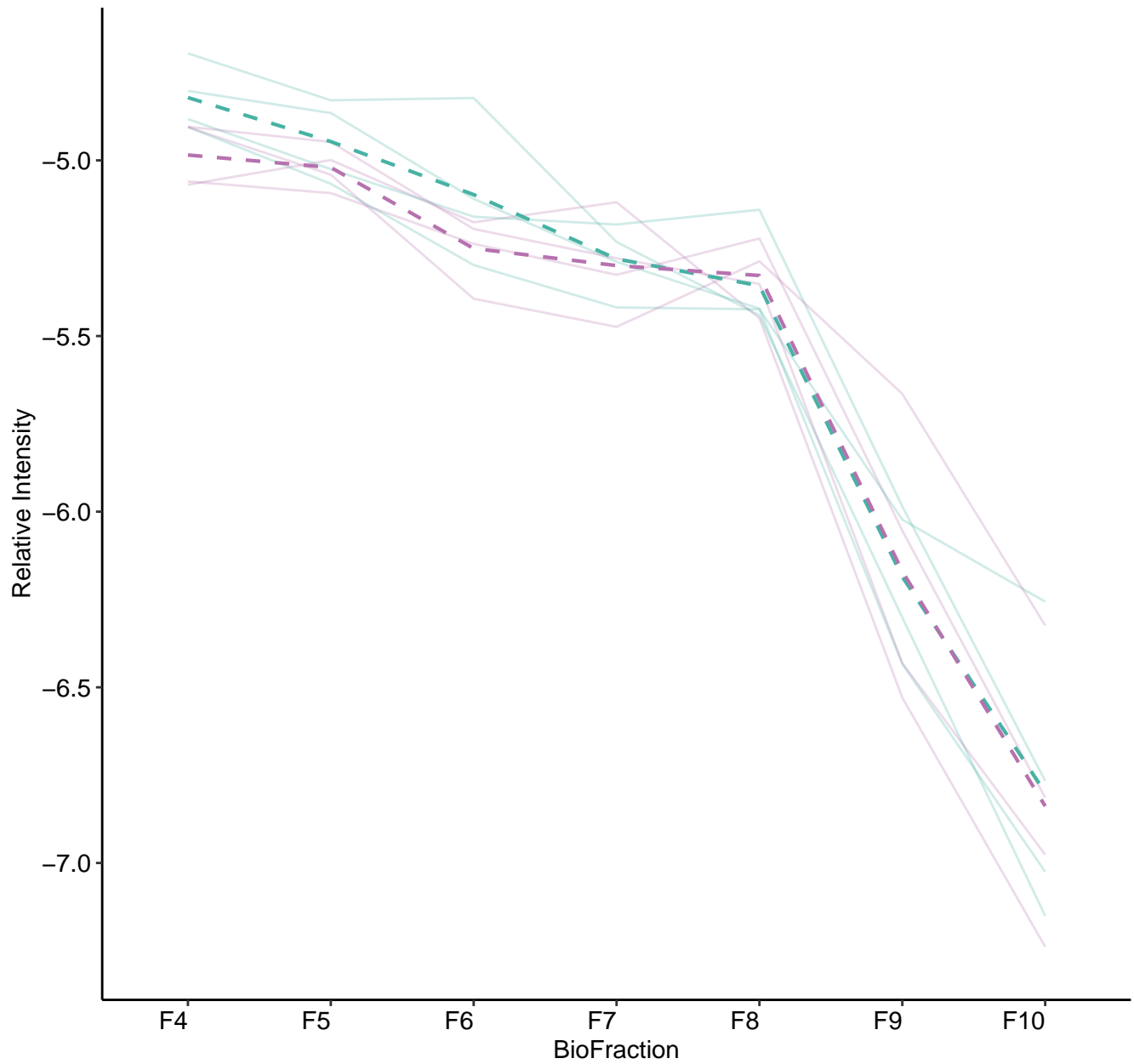
AMPA receptor complex (anti-GluA1-c) (n = 6)
(R2.Fixef = 0.901)



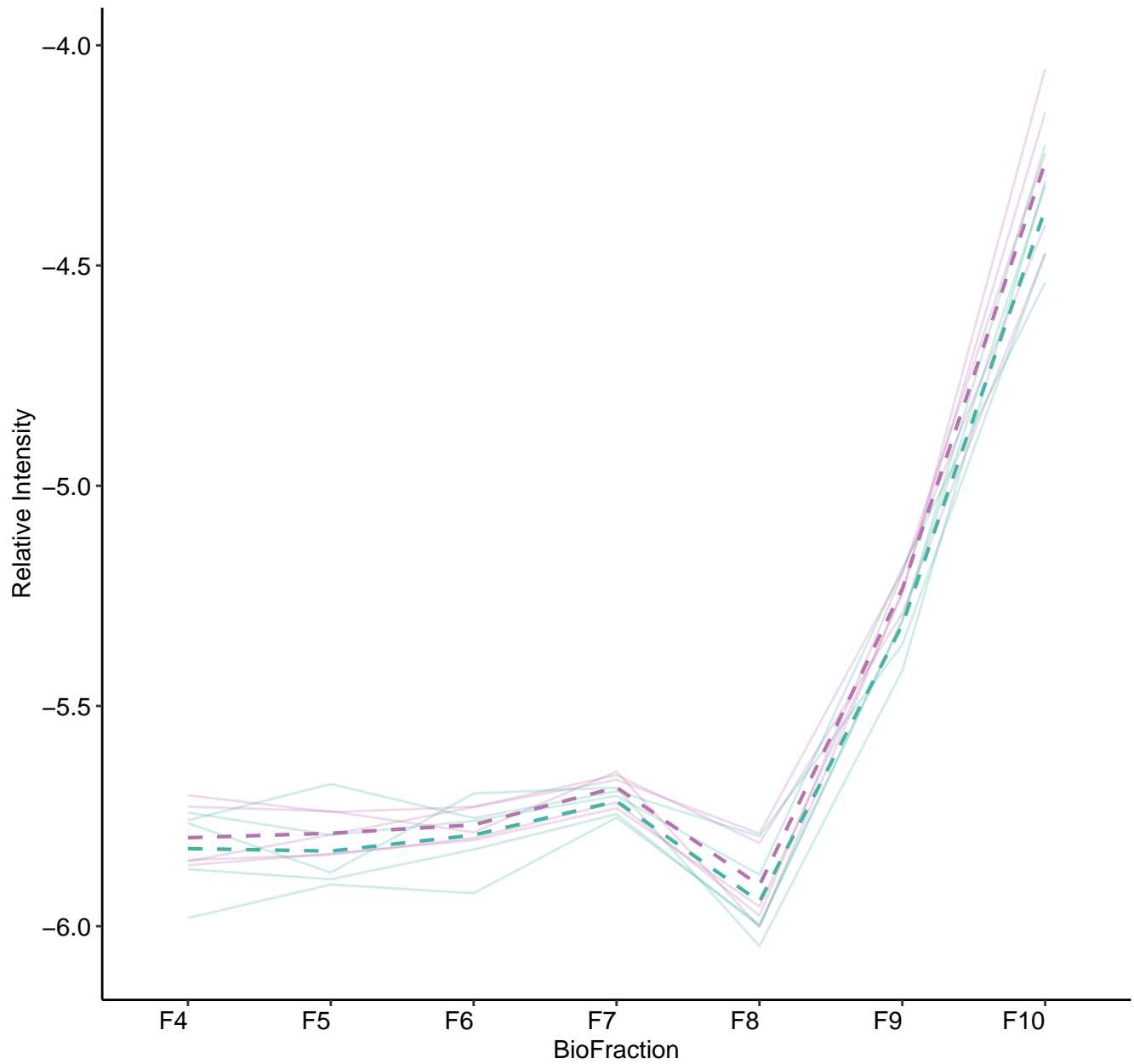
AMPA receptor complex (anti-GluA2-a) (n = 3)
(R2.Fixef = 0.93)



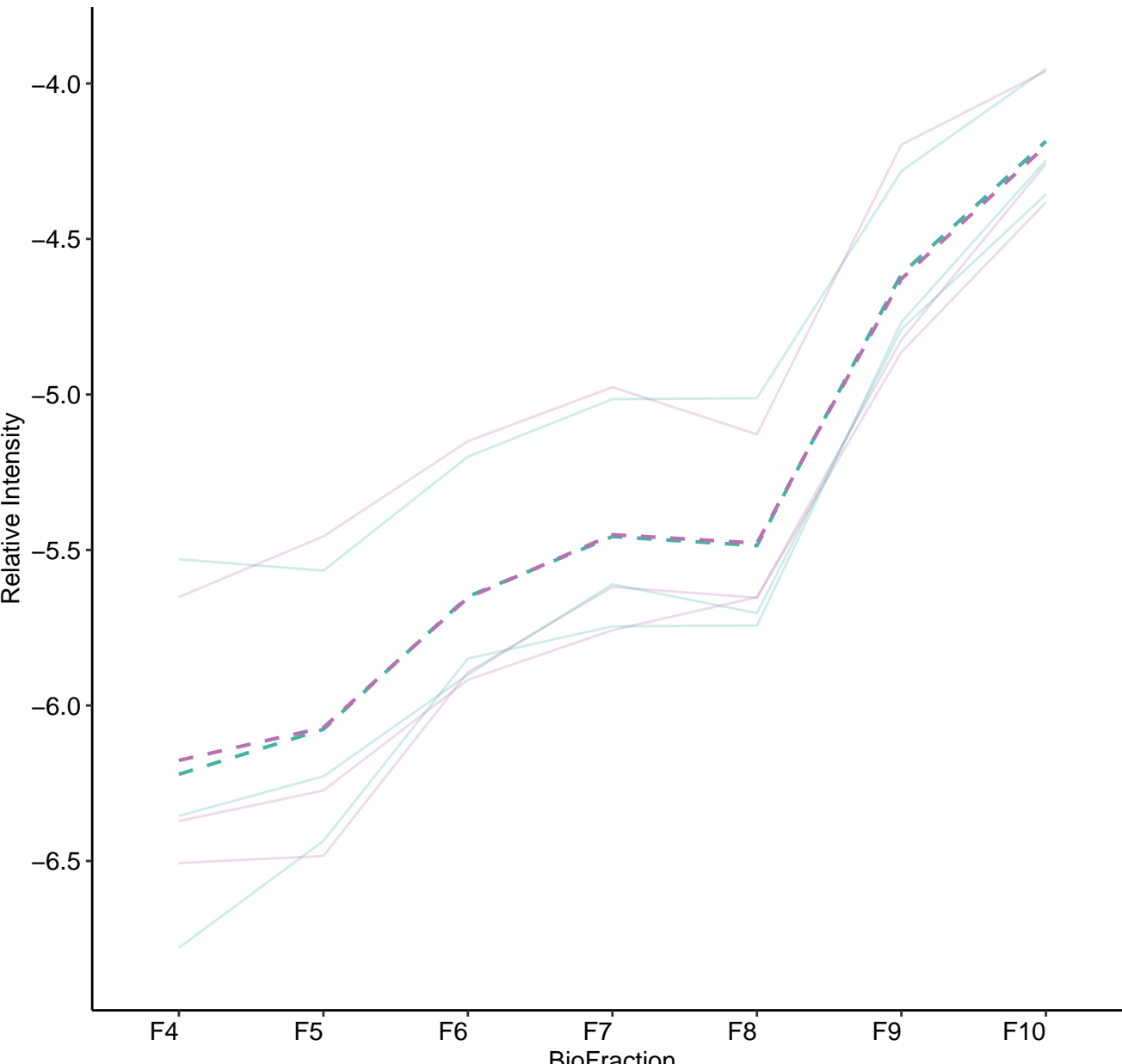
AMPA receptor complex (anti-GluA2-b) (n = 4)
(R2.Fixef = 0.903)



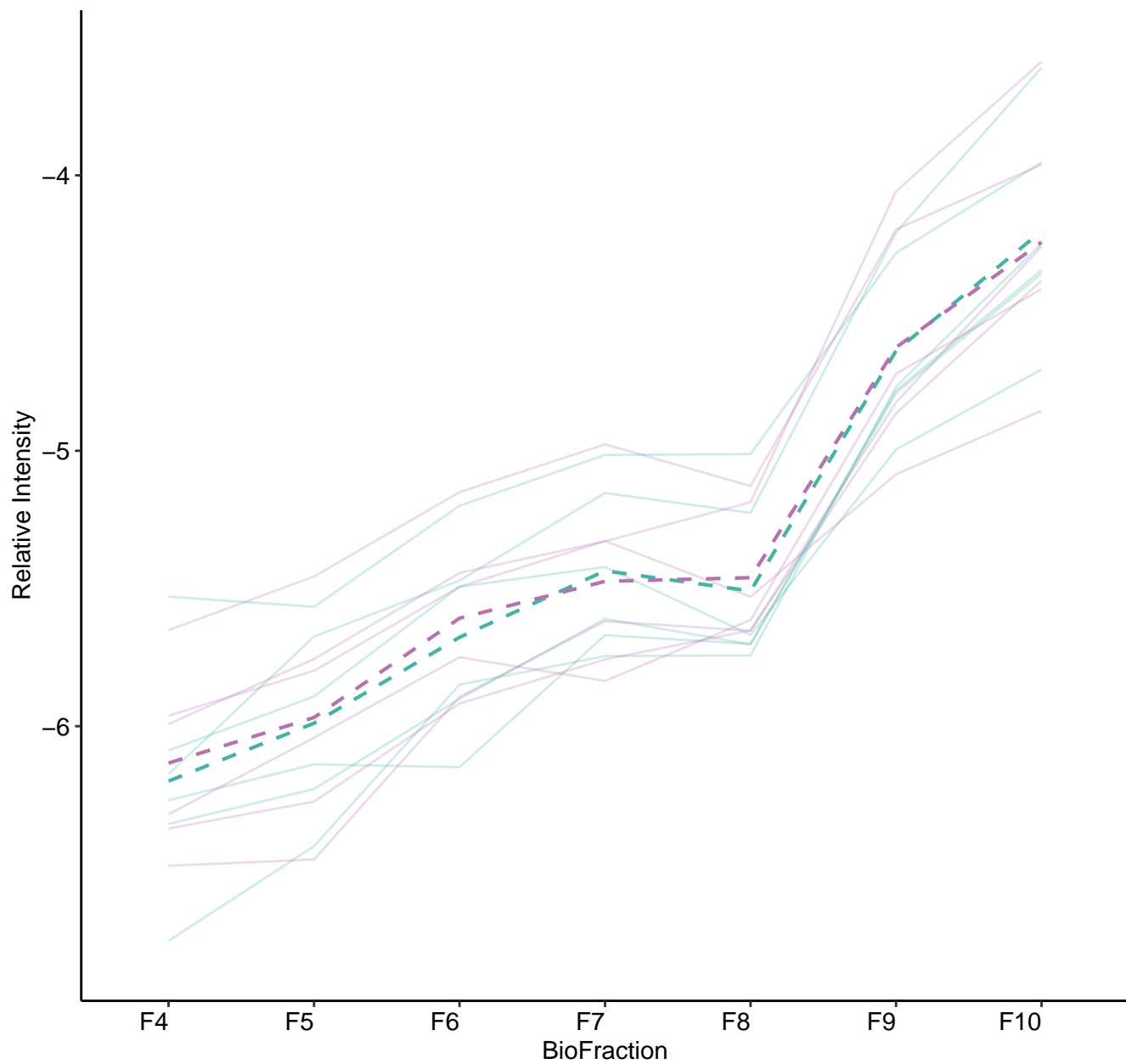
Anaphase-promoting complex (n = 5)
(R2.Fixef = 0.972)



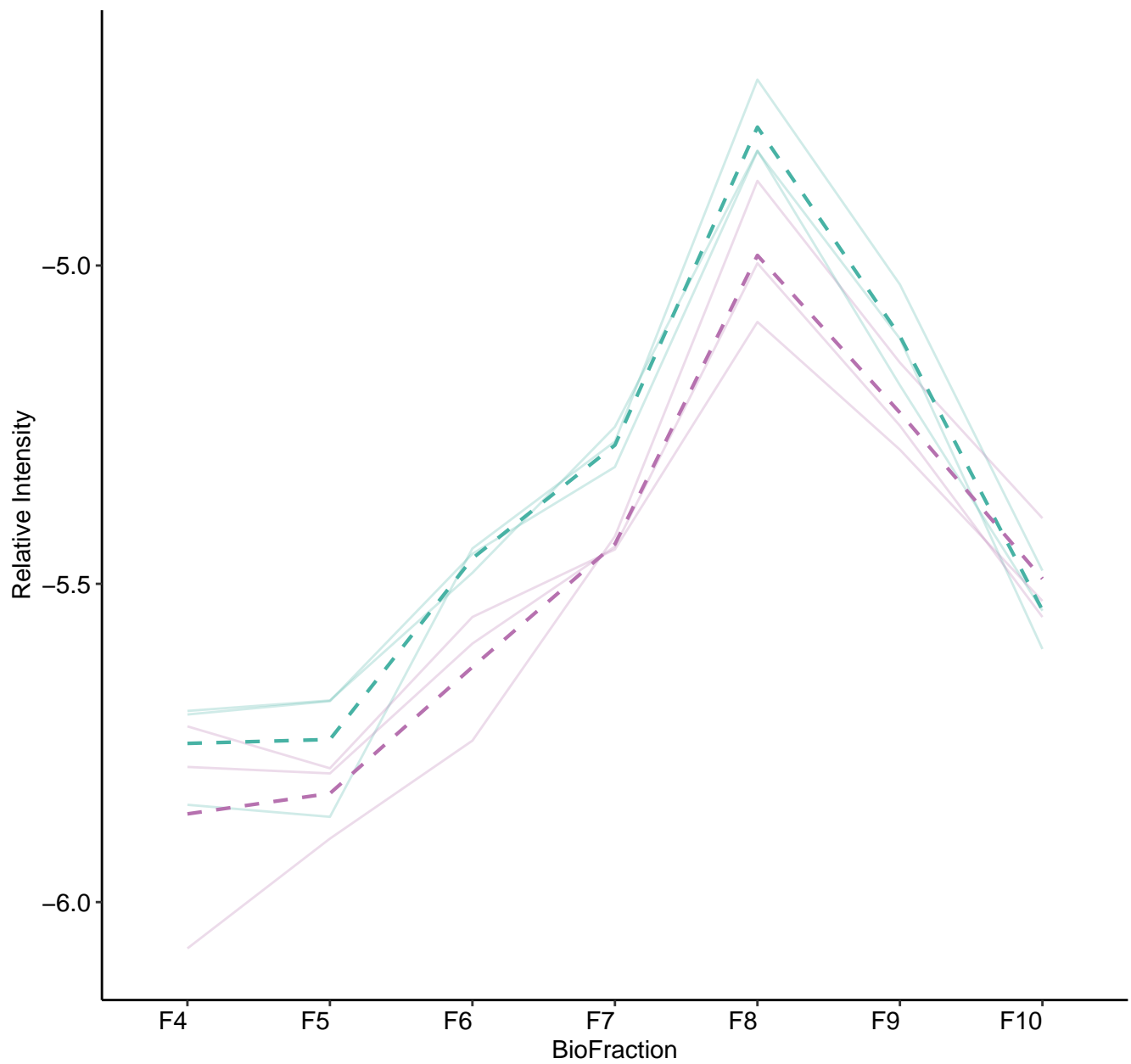
anti-BHC110 complex (n = 3)
(R2.Fixef = 0.737)



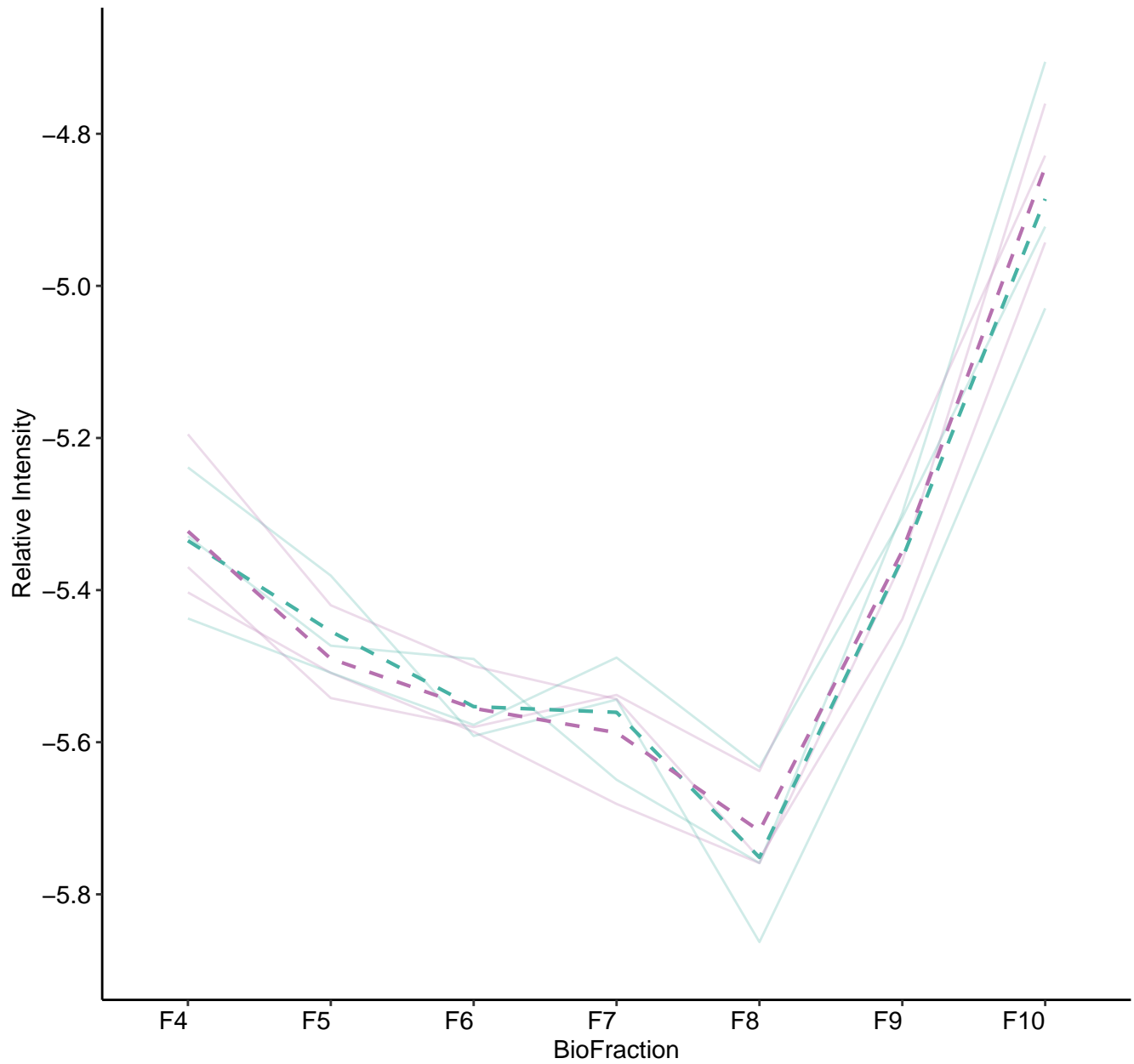
Anti-HDAC2 complex (n = 6)
(R2.Fixef = 0.782)



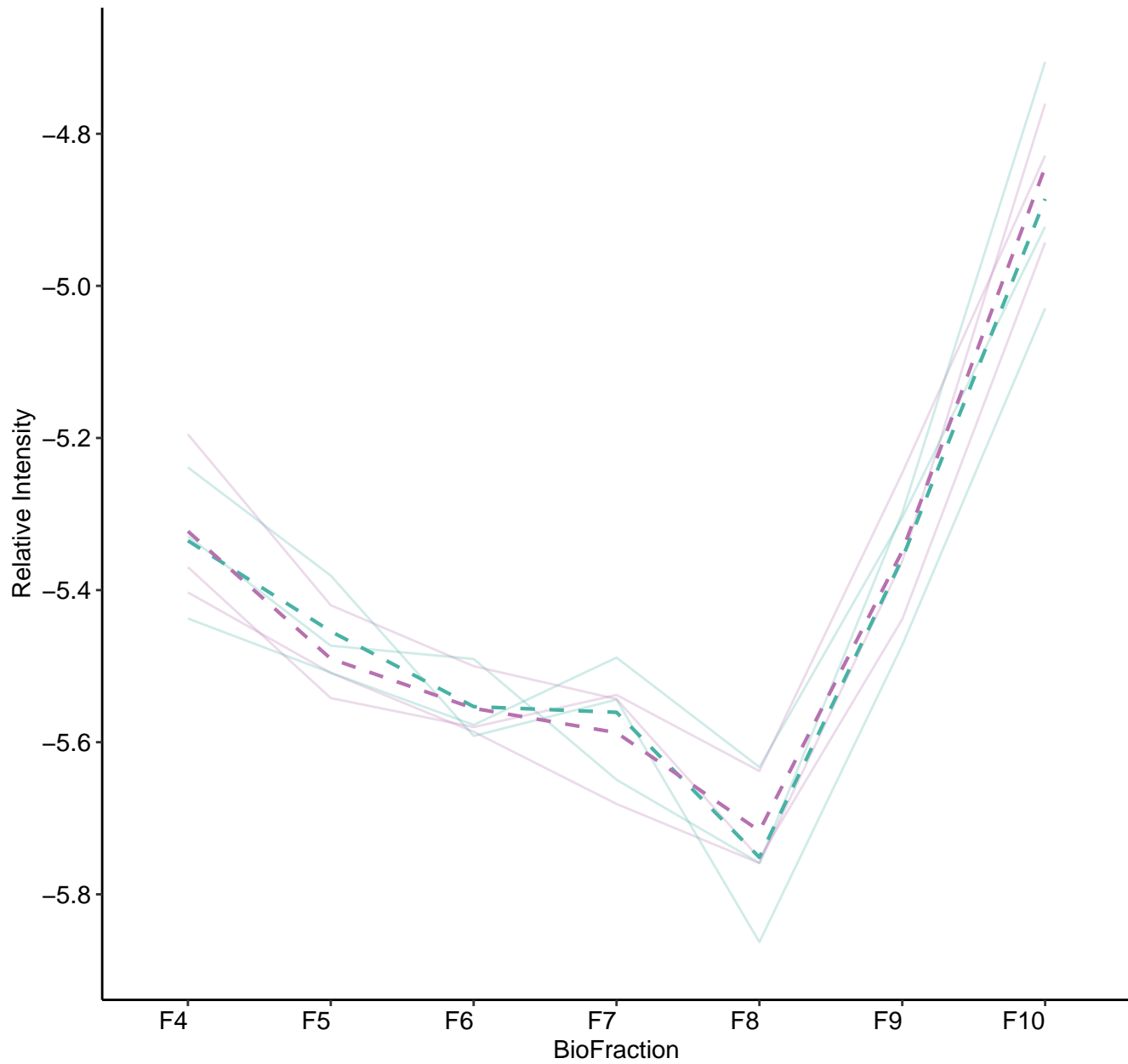
AP1 adaptor complex (n = 3)
(R2.Fixef = 0.93)



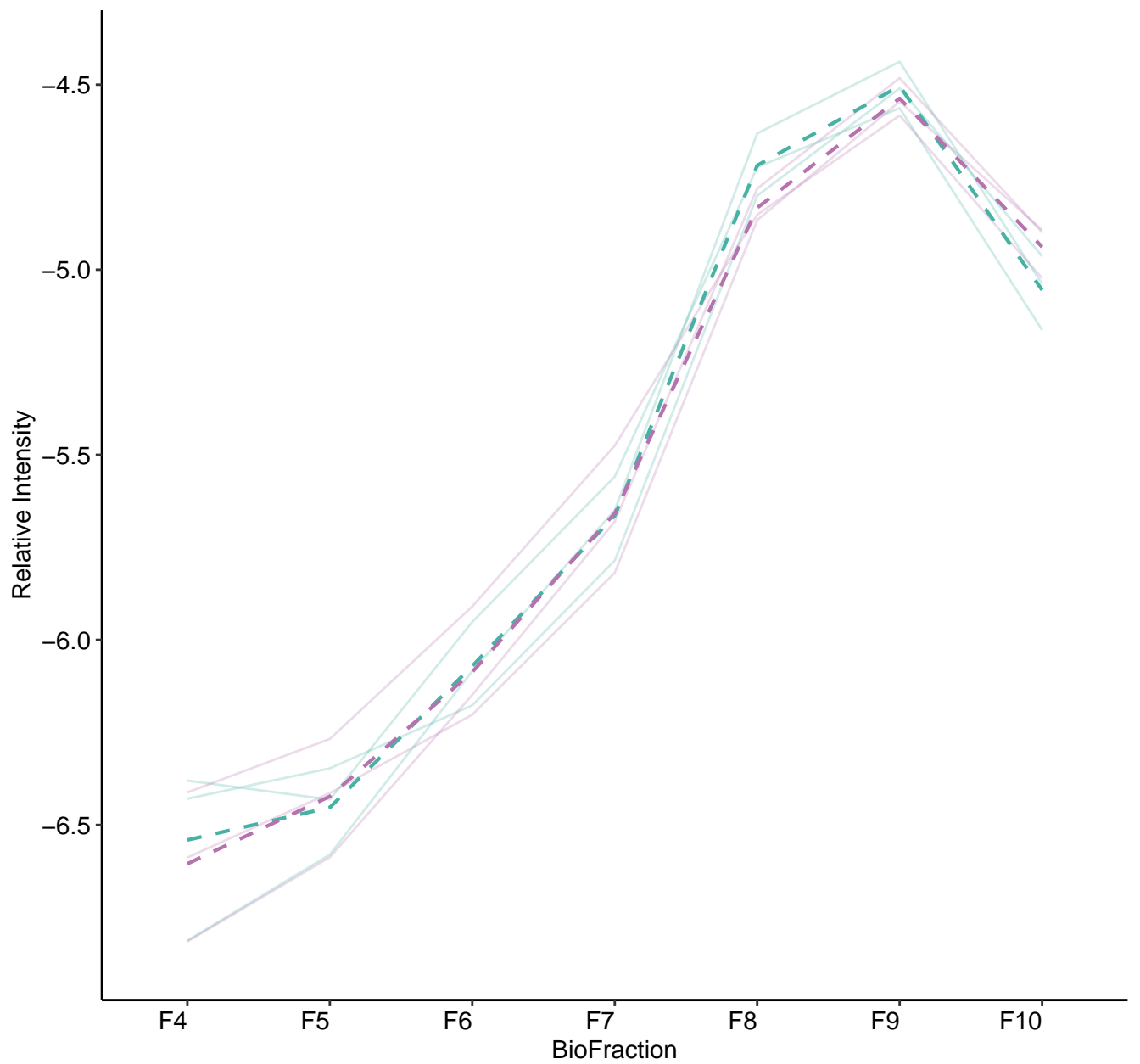
AP3 adaptor complex (n = 3)
(R2.Fixef = 0.886)



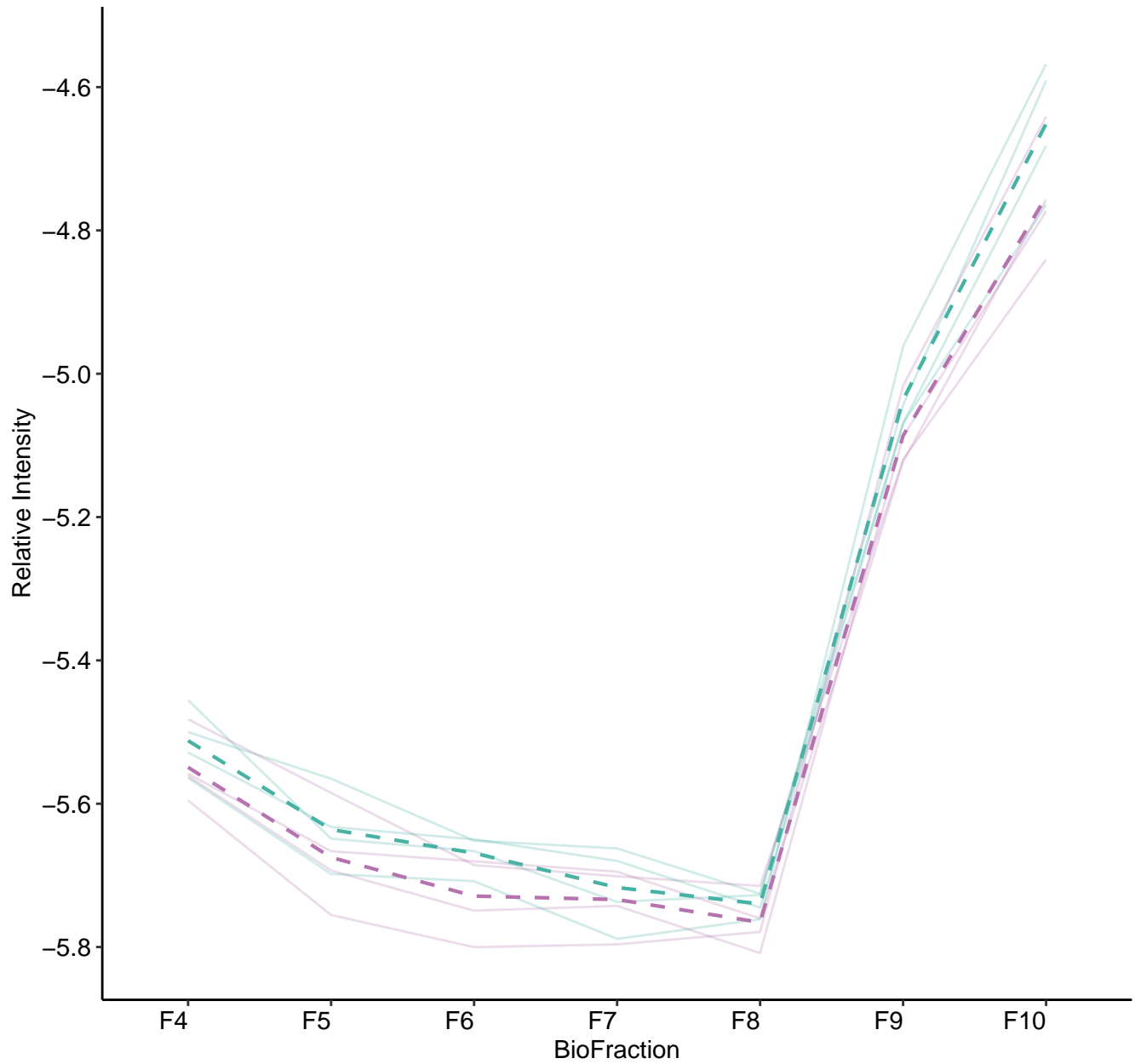
AP3-BLOC1 complex (n = 3)
(R2.Fixef = 0.886)



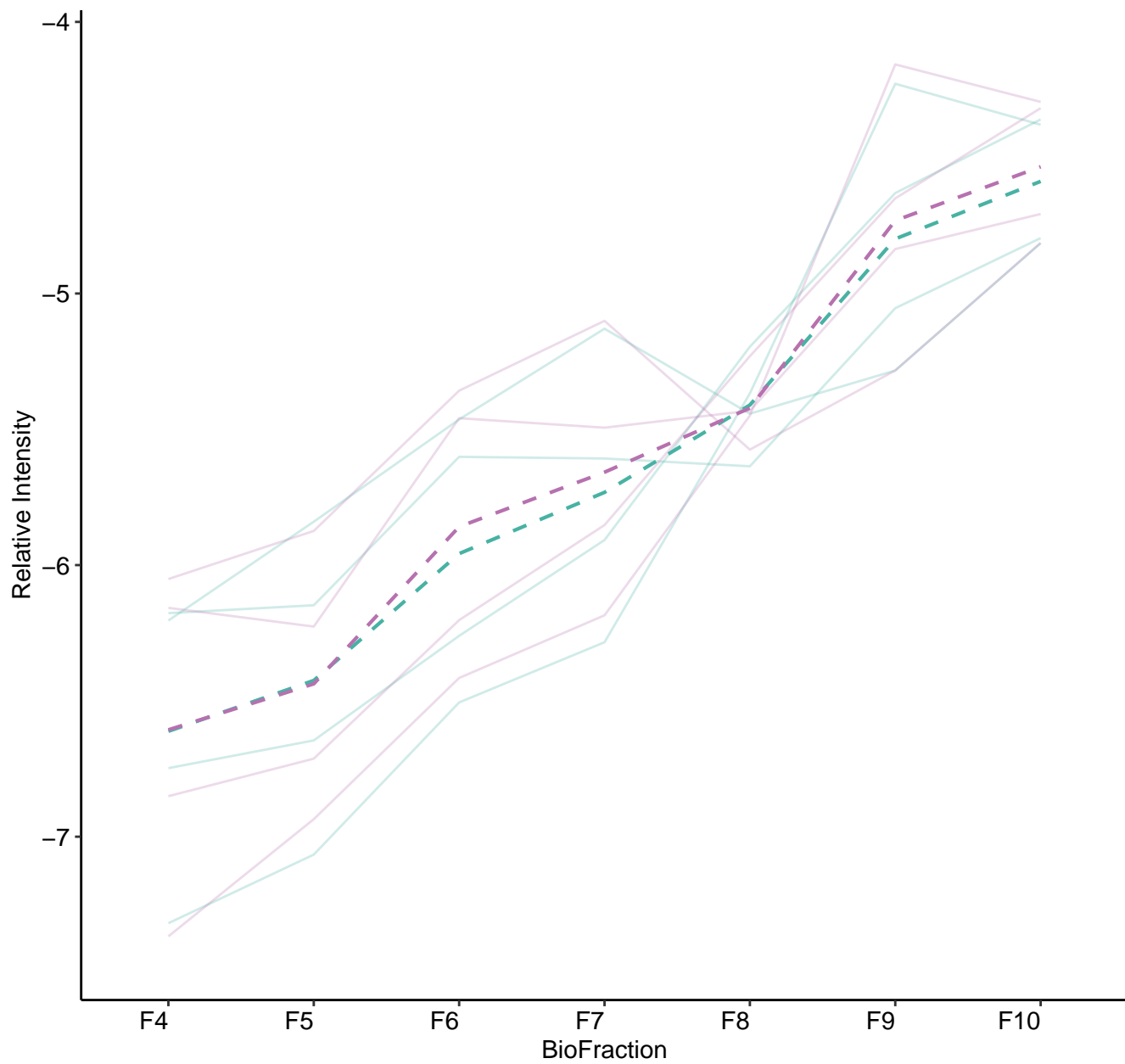
Apoptosis- and splicing-associated protein complex (ASAP-L), SAP18, RNPS1, Acinus-L (n = 3)
(R2.Fixef = 0.971)



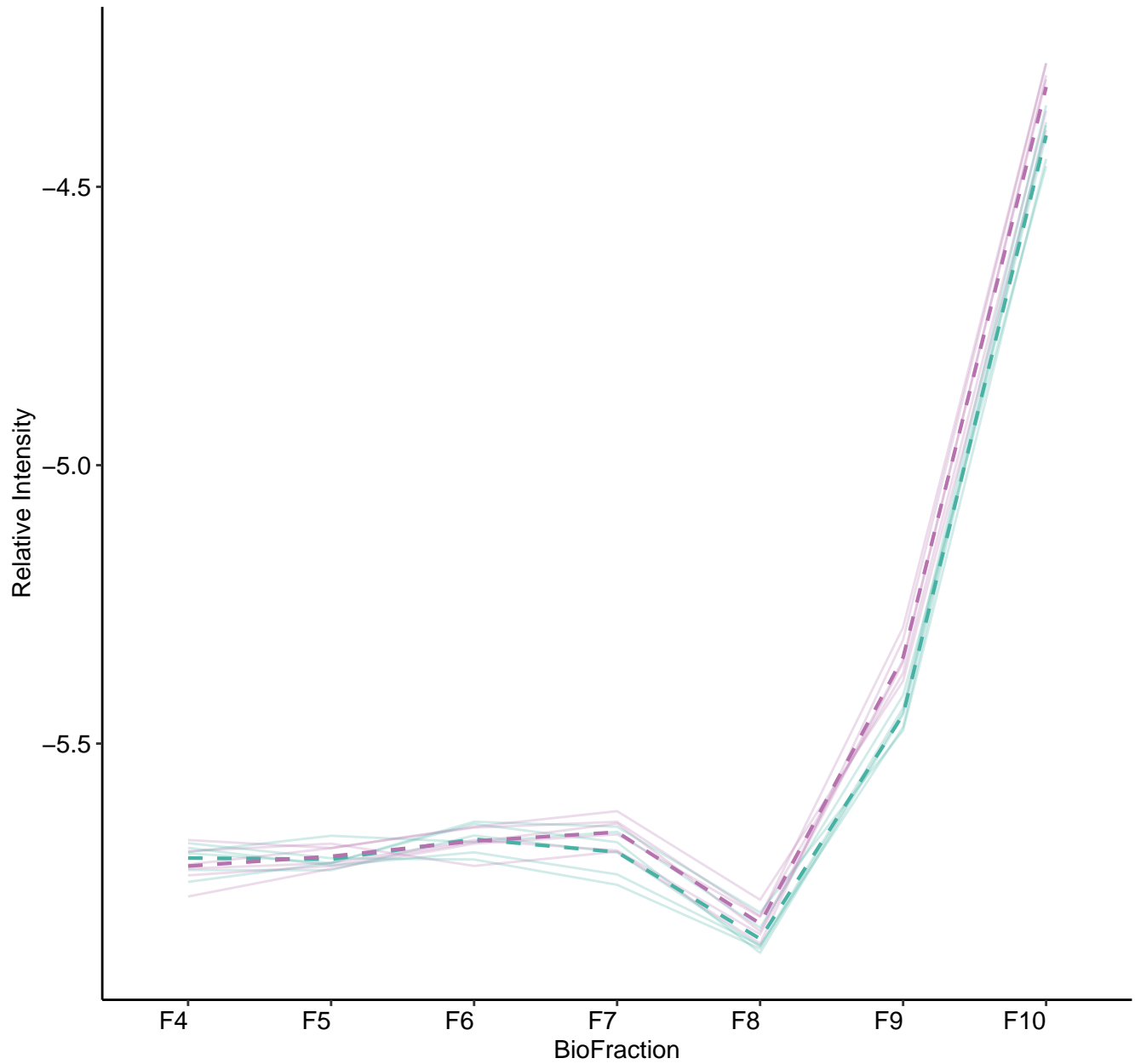
Arp2/3 protein complex (n = 4)
(R2.Fixef = 0.979)



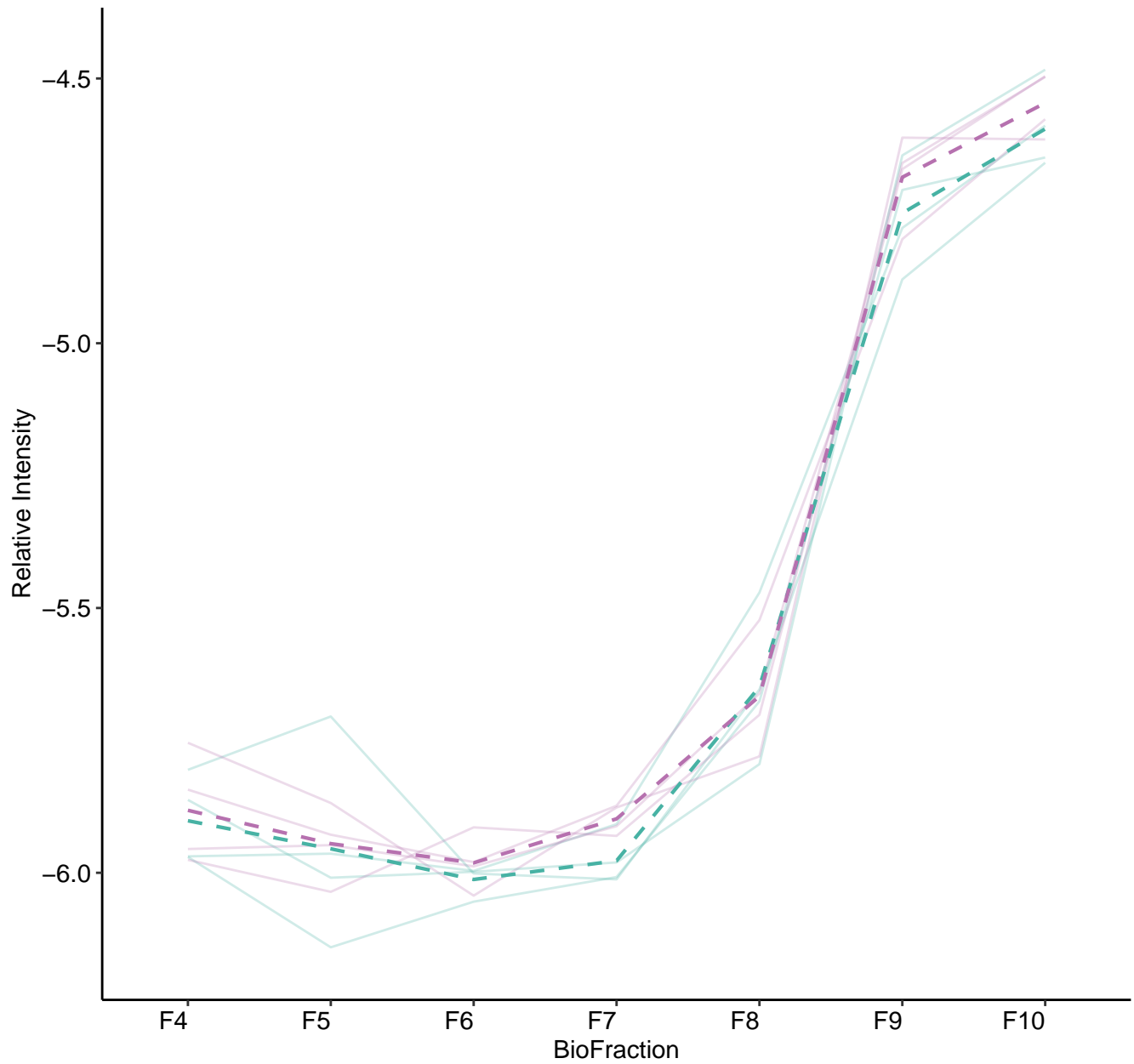
B-WICH complex (n = 4)
(R2.Fixef = 0.724)



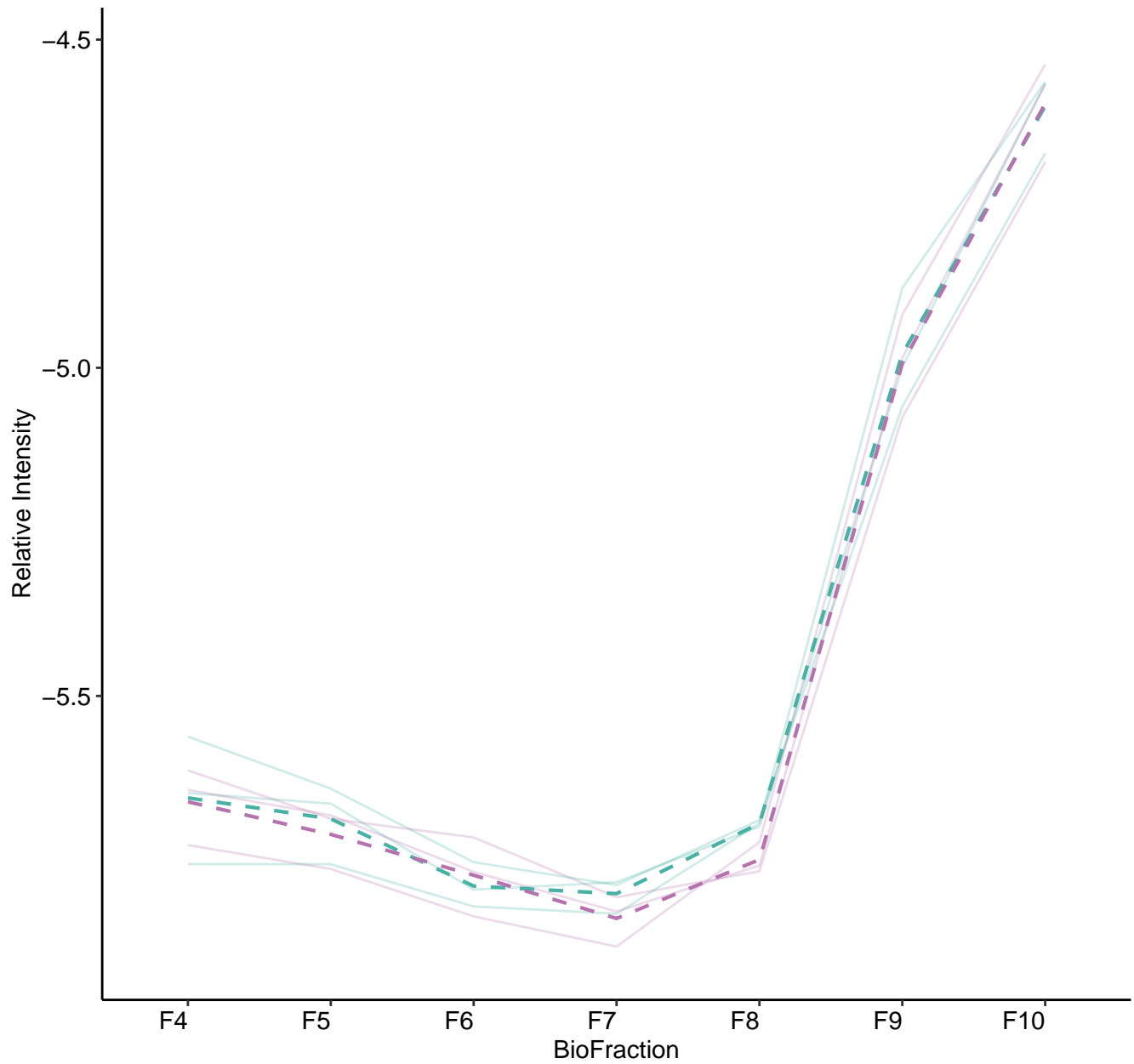
BBS-chaperonin complex (n = 6)
(R2.Fixef = 0.995)



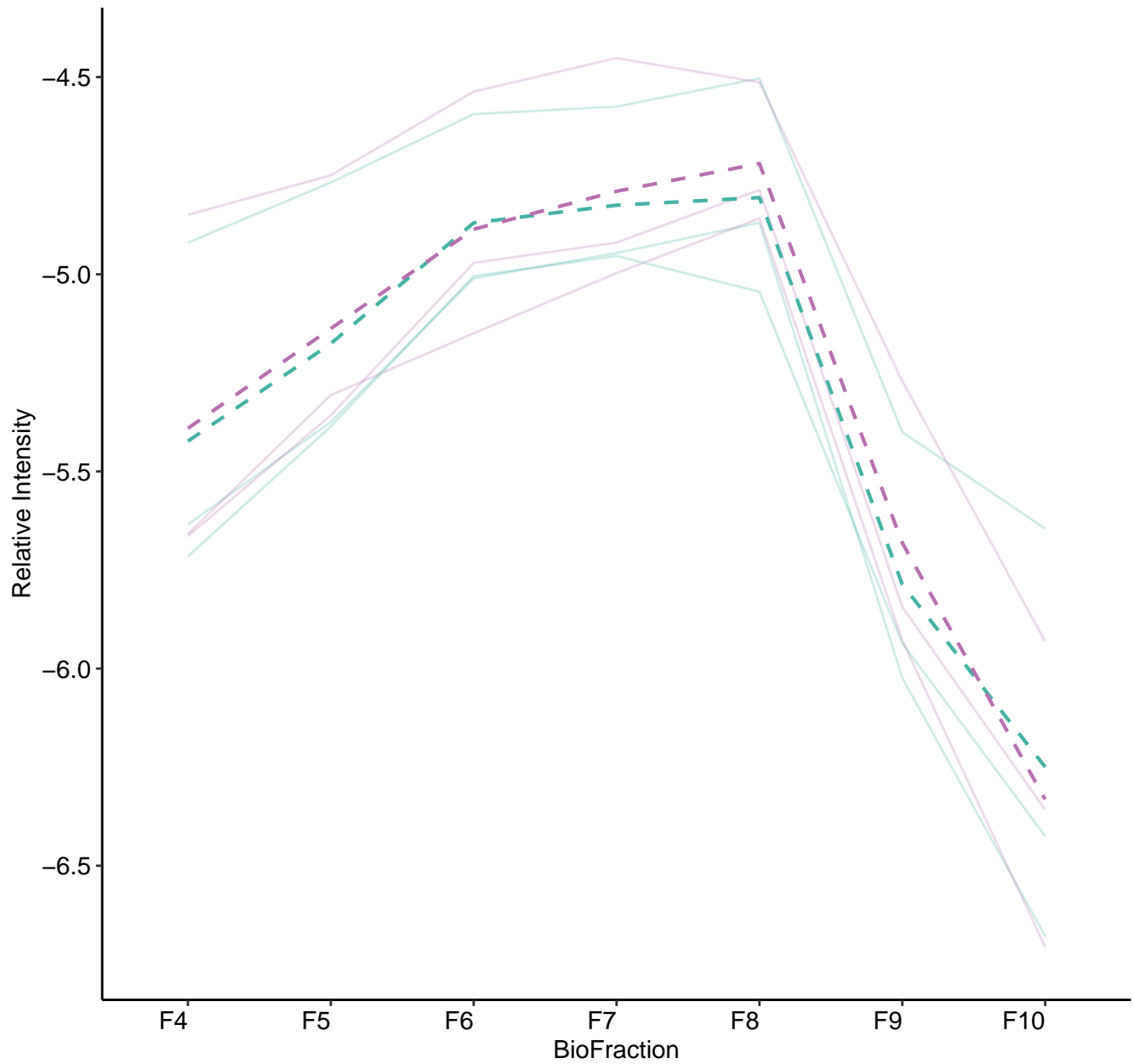
BBSome (n = 4)
(R2.Fixef = 0.975)



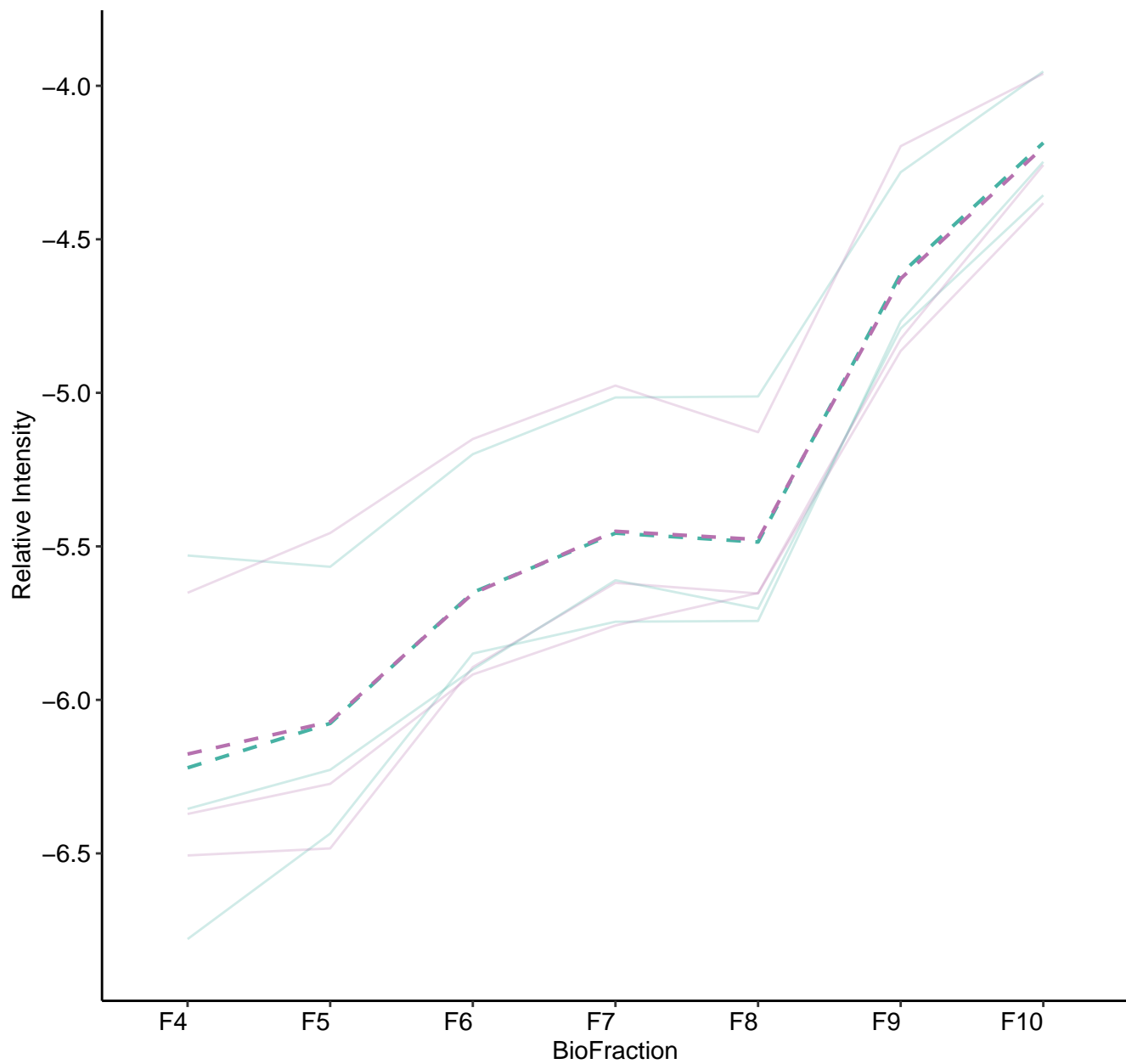
Bcl-2-Beclin1-UVRAG-PI(3)KC3 complex (n = 3)
(R2.Fixef = 0.982)



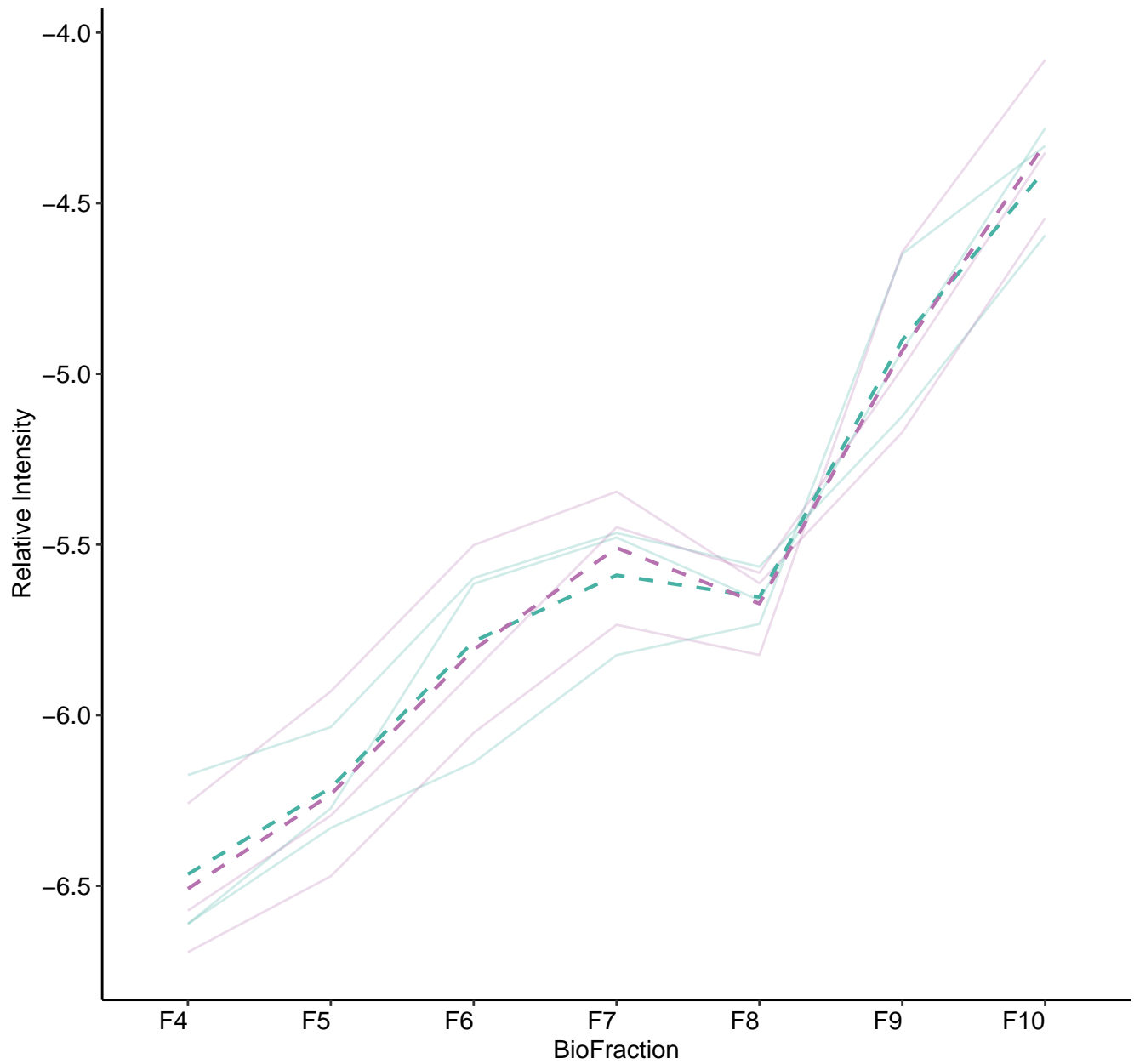
BORC complex (n = 3)
(R2.Fixef = 0.694)



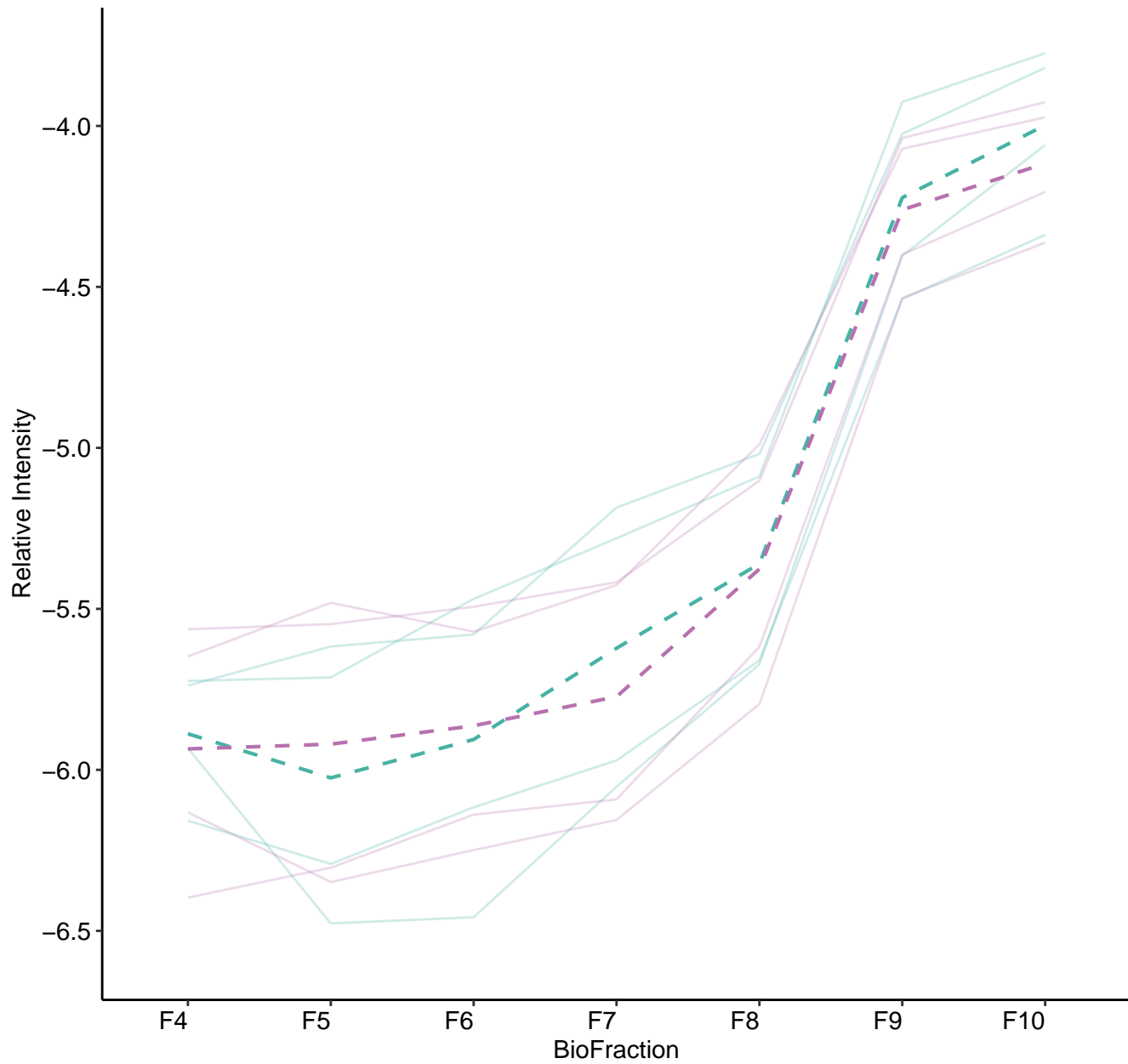
BRAF53-BRCA2 complex (n = 3)
(R2.Fixef = 0.737)



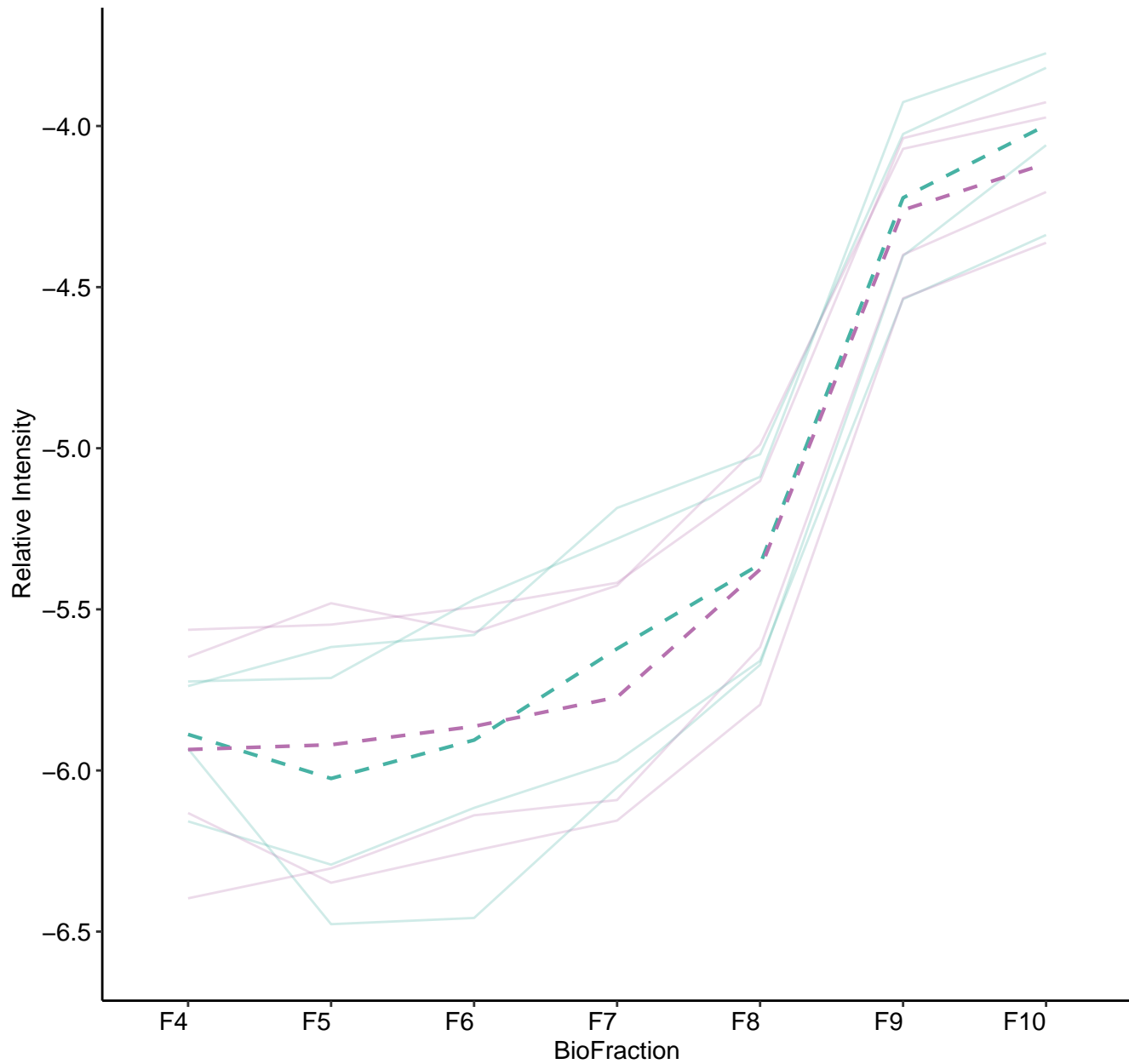
BRCA1–RNA polymerase II complex (n = 3)
(R2.Fixef = 0.903)



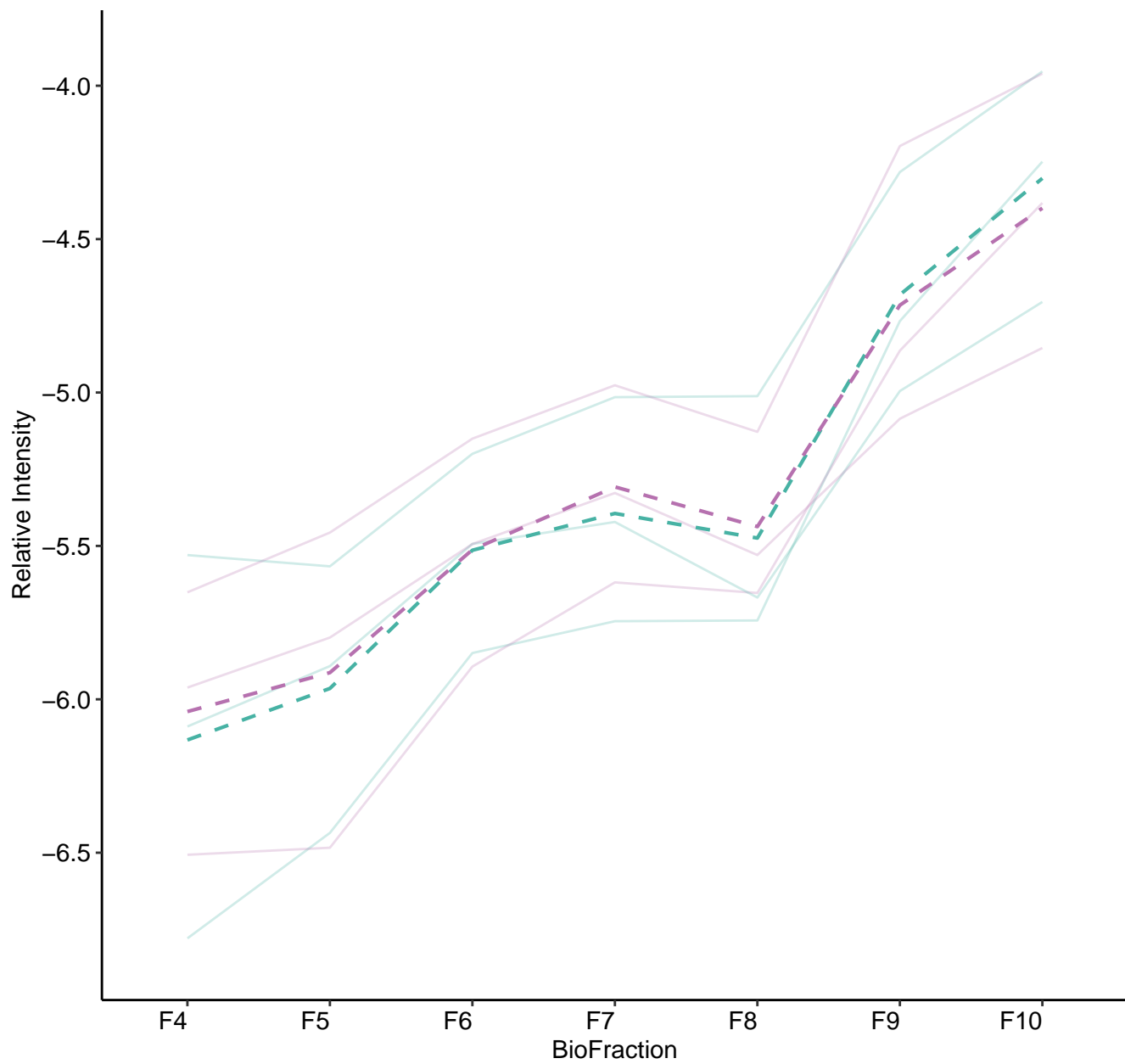
BRG1-SIN3A complex (n = 4)
(R2.Fixef = 0.814)



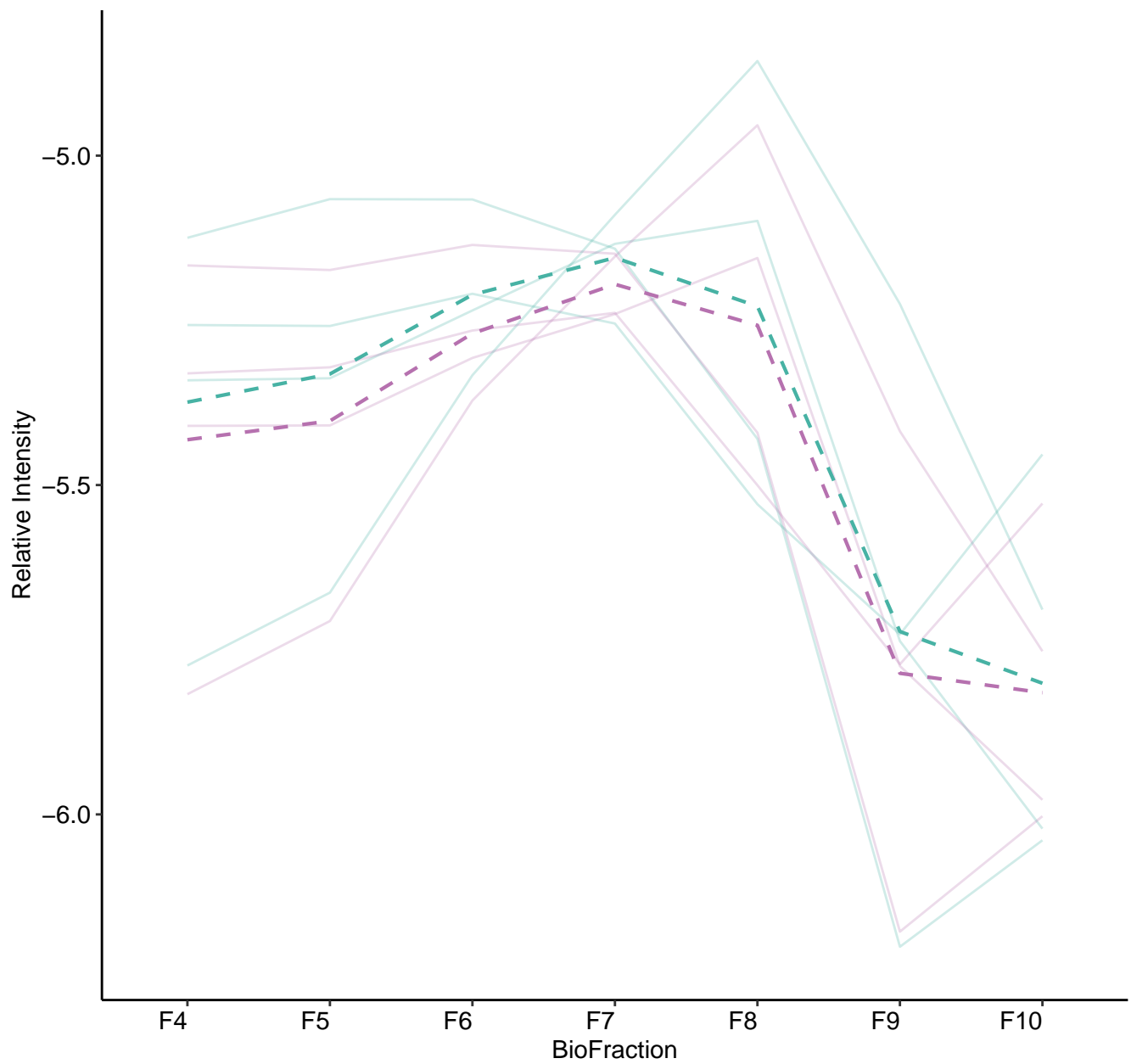
BRG1–SIN3A–HDAC containing SWI/SNF remodeling complex I (n = 4)
(R2.Fixef = 0.814)



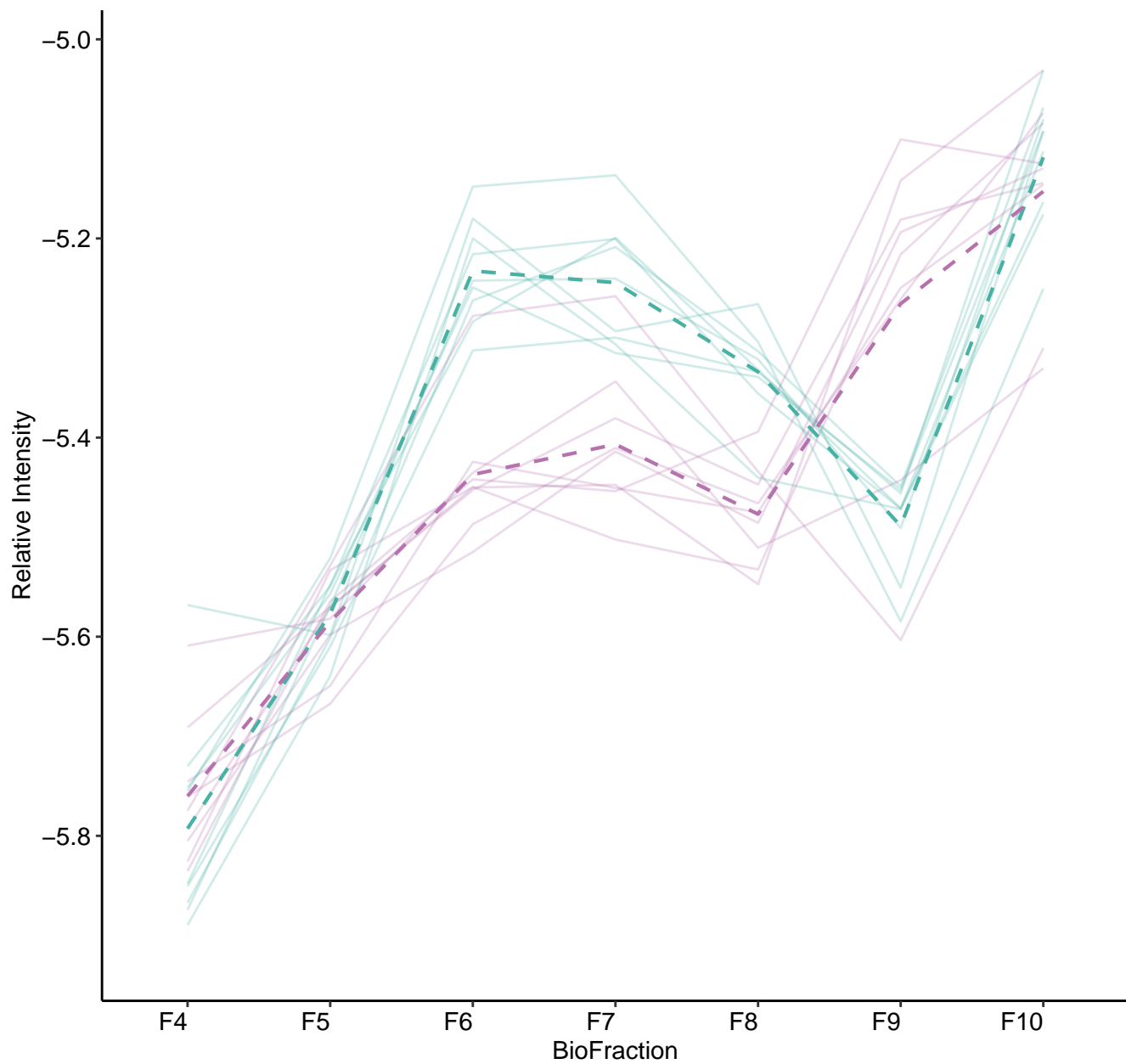
BRMS1–SIN3–HDAC complex (n = 3)
(R2.Fixef = 0.663)



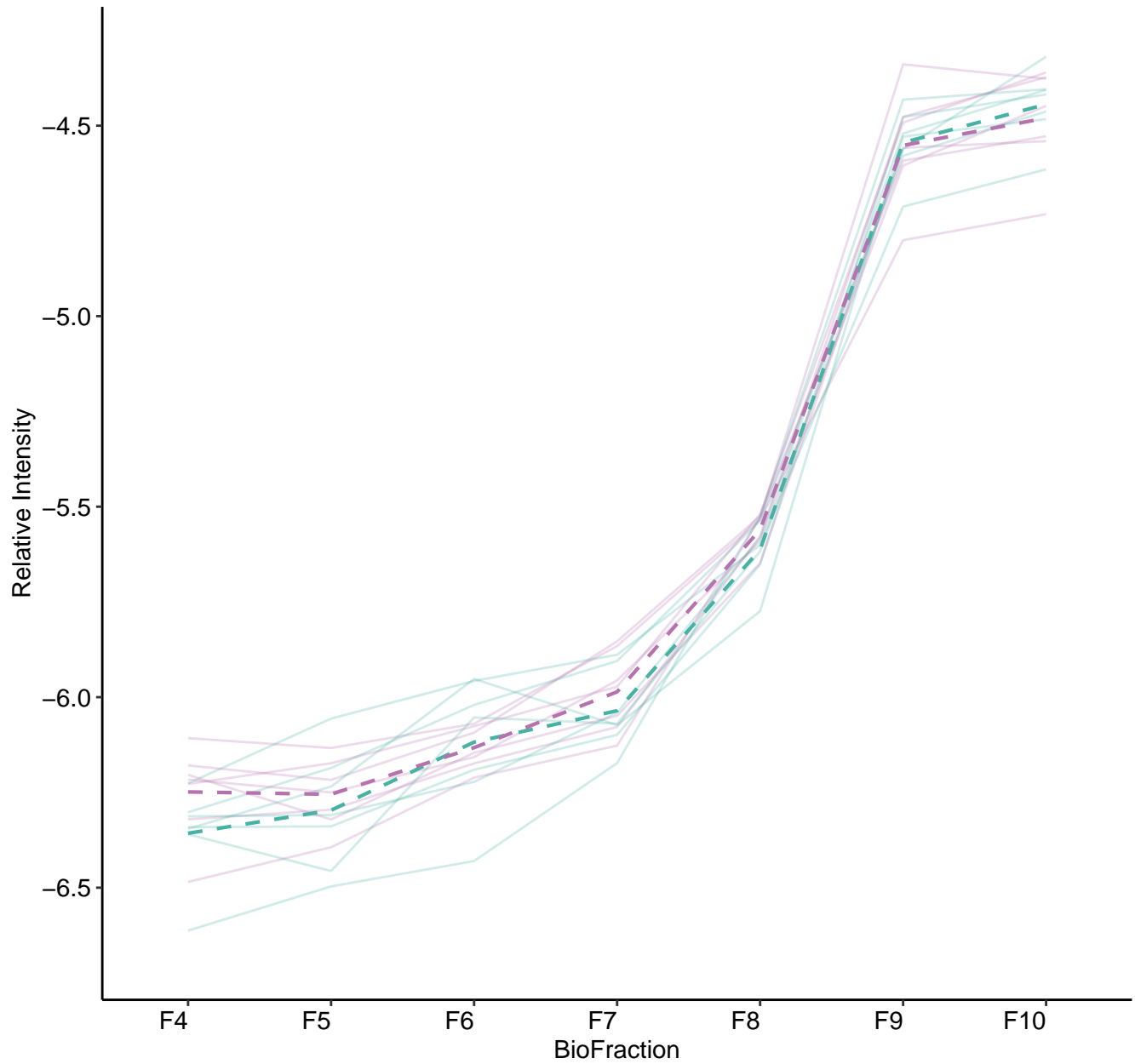
c-Src immune complex (n = 4)
(R2.Fixef = 0.488)



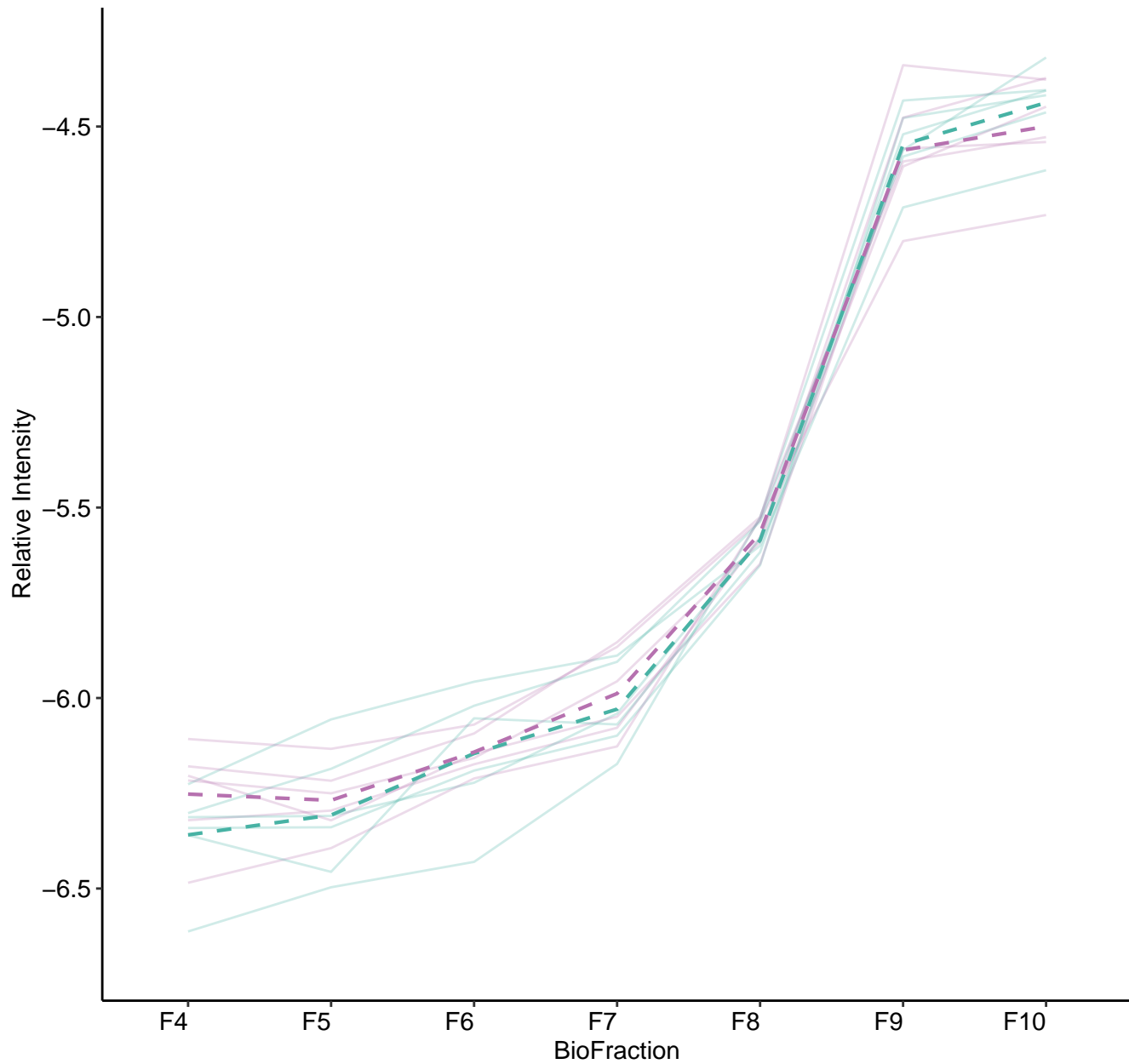
CCC-Wash (WASH1, FAM21C) complex (n = 9)
(R2.Fixef = 0.875)



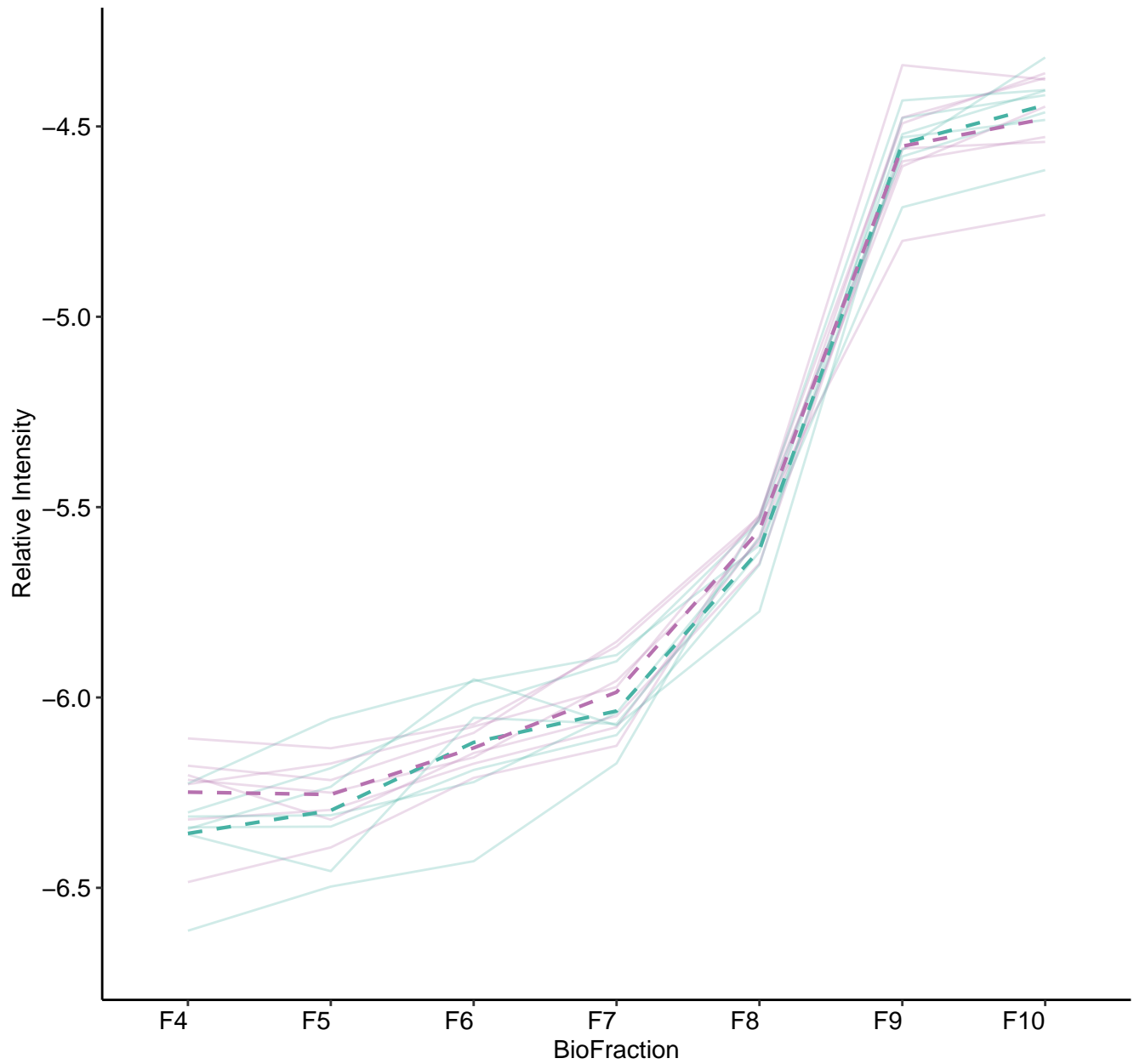
CCR4-NOT complex (n = 7)
(R2.Fixef = 0.977)



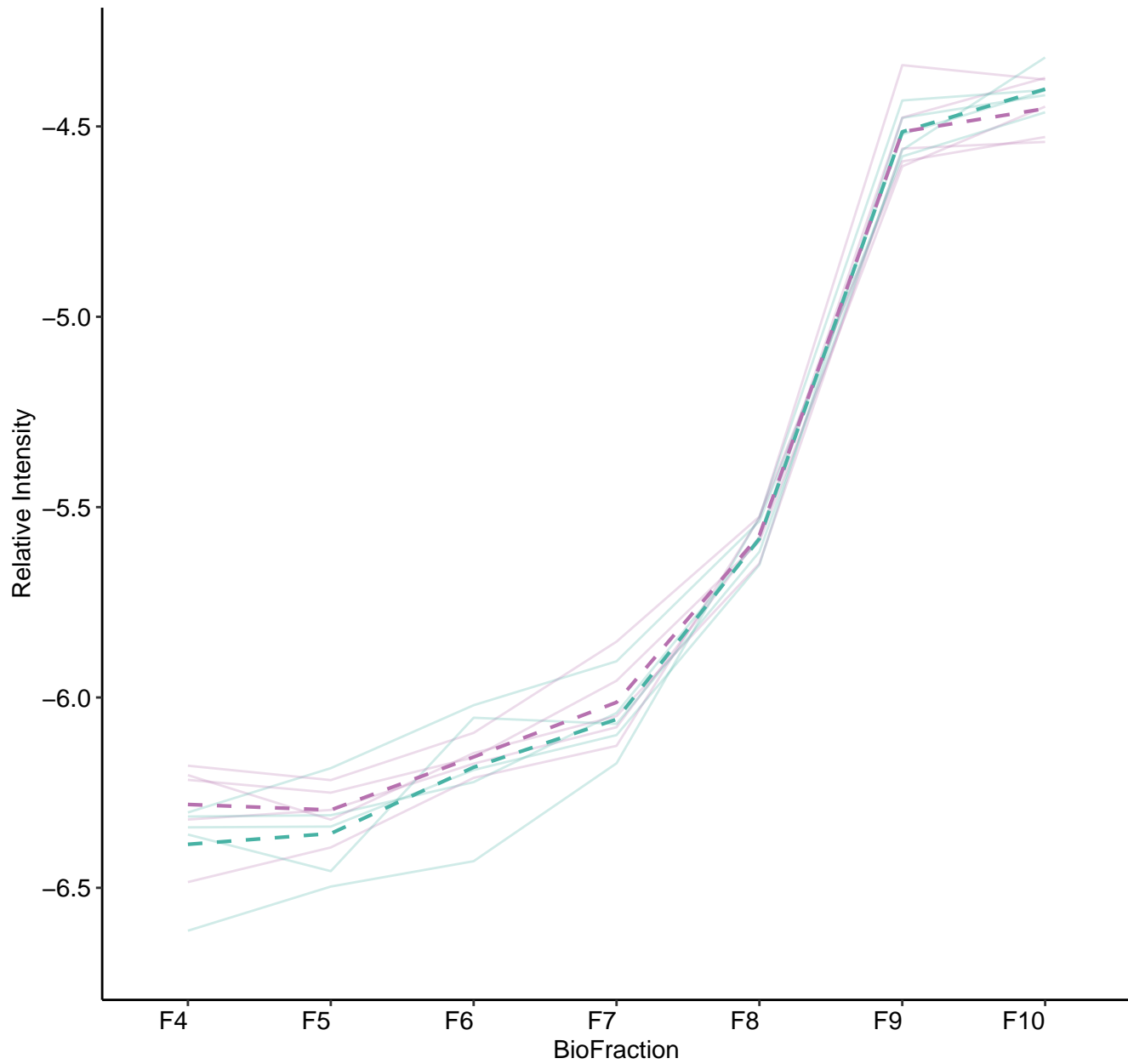
CCR4–NOT–CNOT7–CNOT6 complex (n = 6)
(R2.Fixef = 0.975)



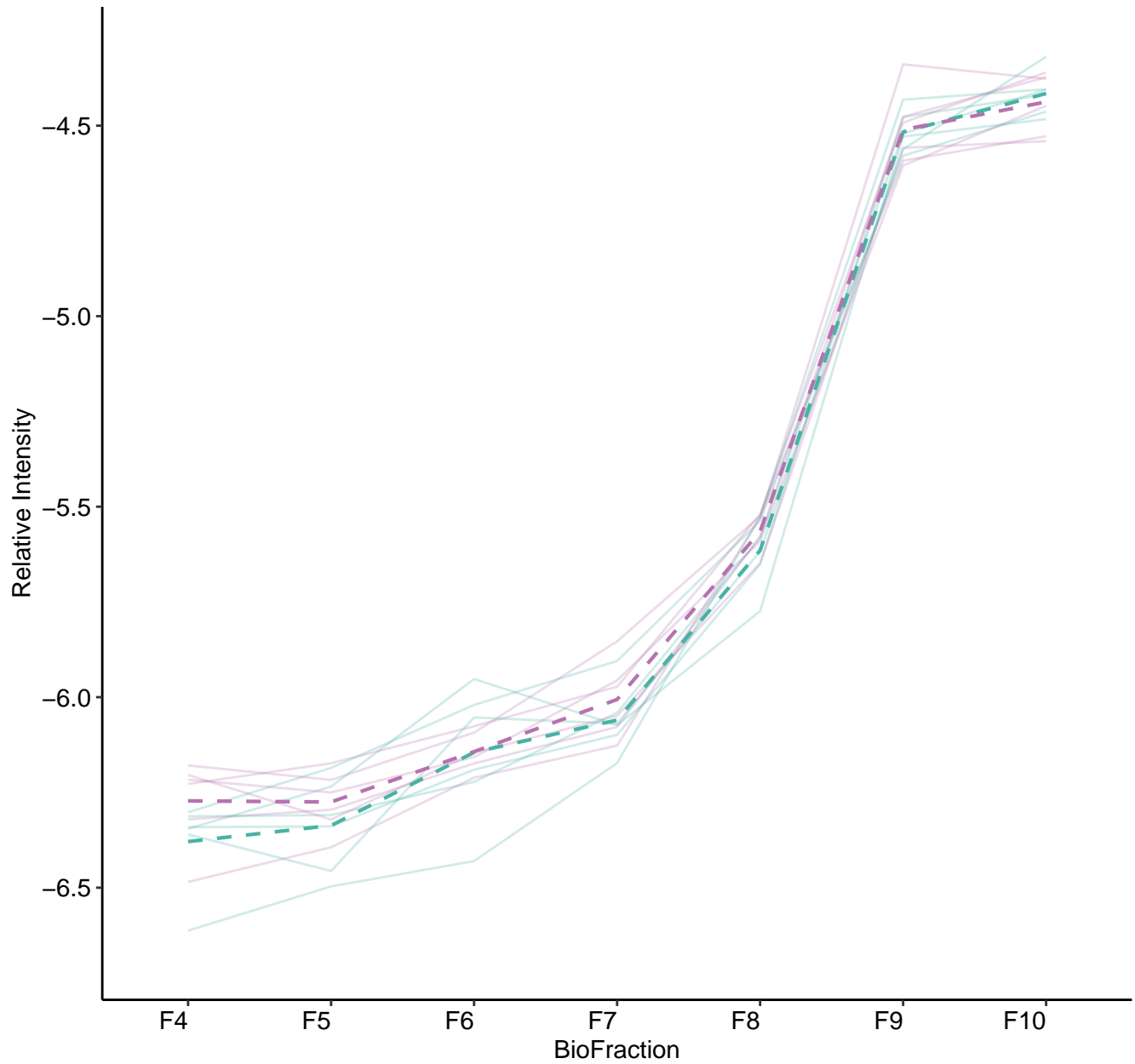
CCR4-NOT-CNOT7-CNOT6L complex (n = 7)
(R2.Fixef = 0.977)



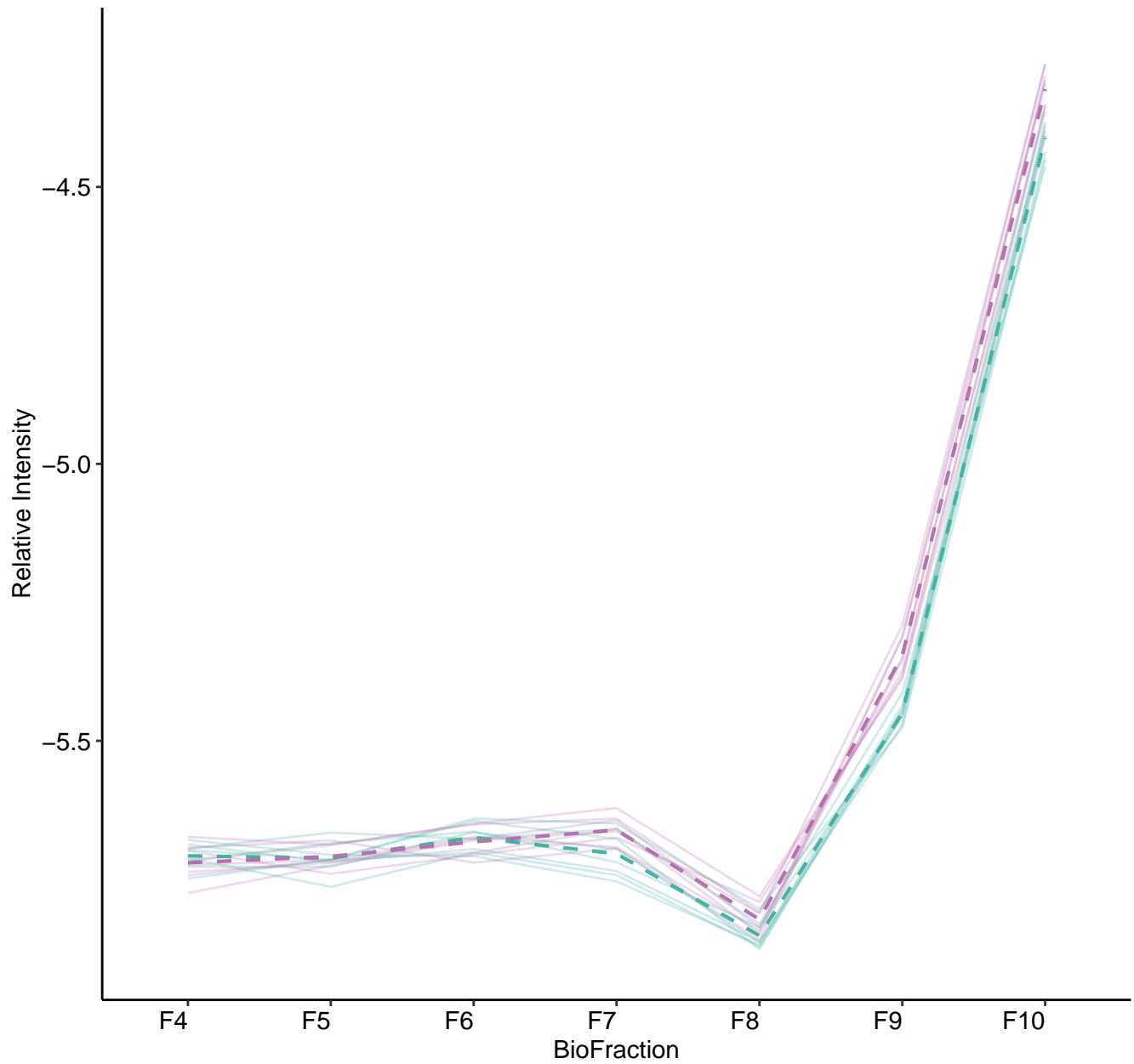
CCR4–NOT–CNOT8–CNOT6 complex (n = 5)
(R2.Fixef = 0.984)



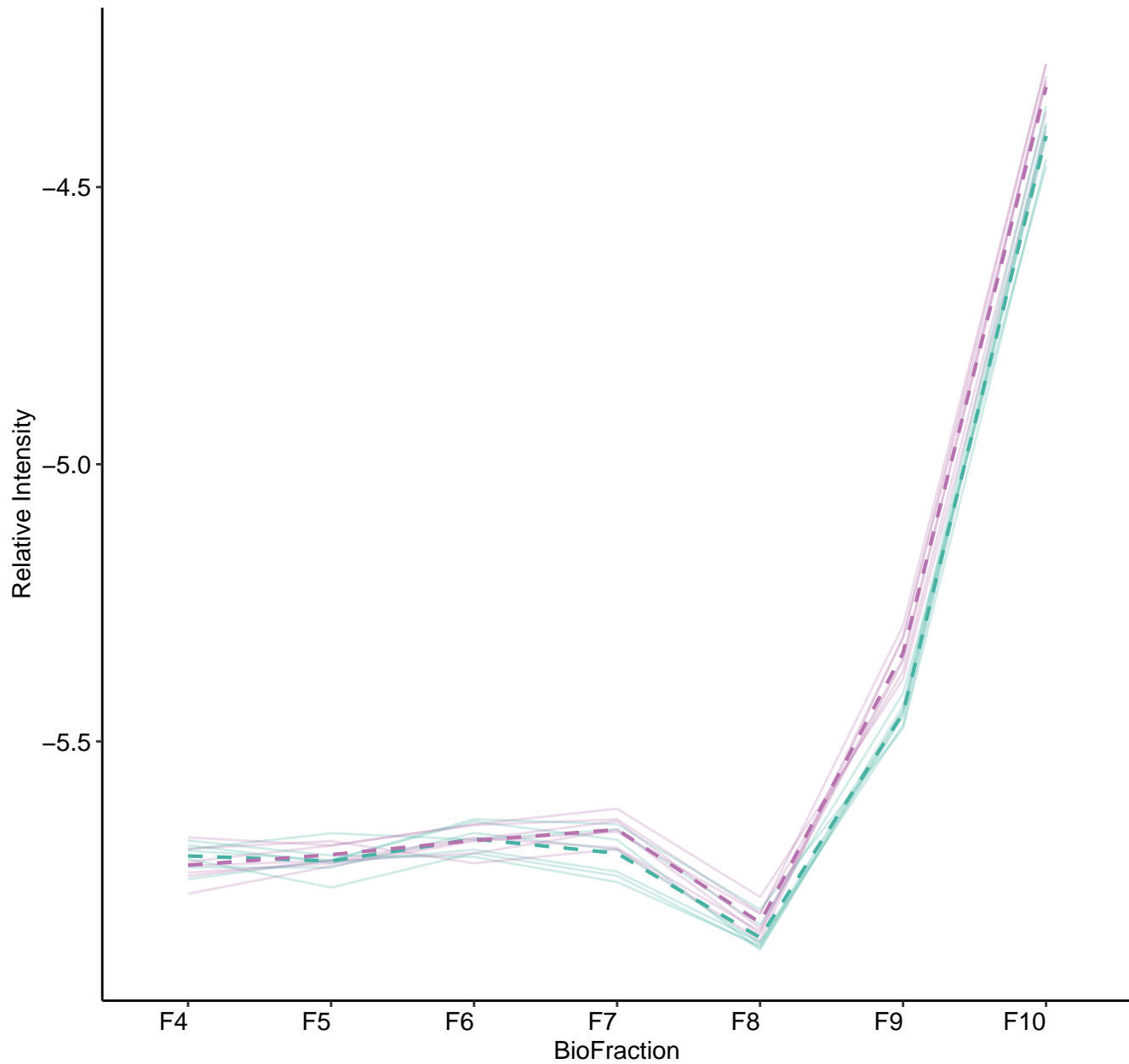
CCR4-NOT-CNOT8-CNOT6L complex (n = 6)
(R2.Fixef = 0.984)



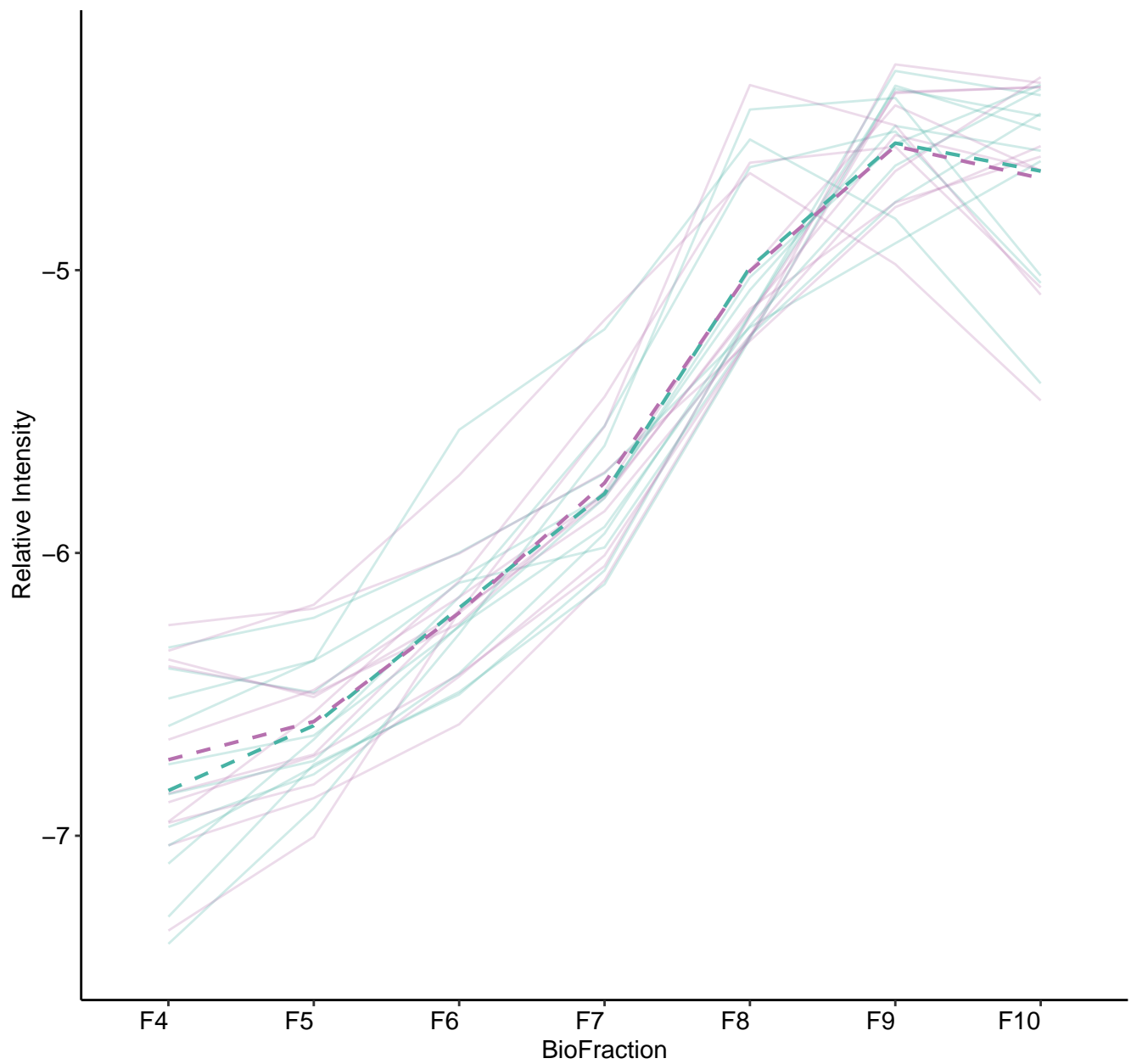
CCT complex (chaperonin containing TCP1 complex) (n = 8)
(R2.Fixef = 0.996)



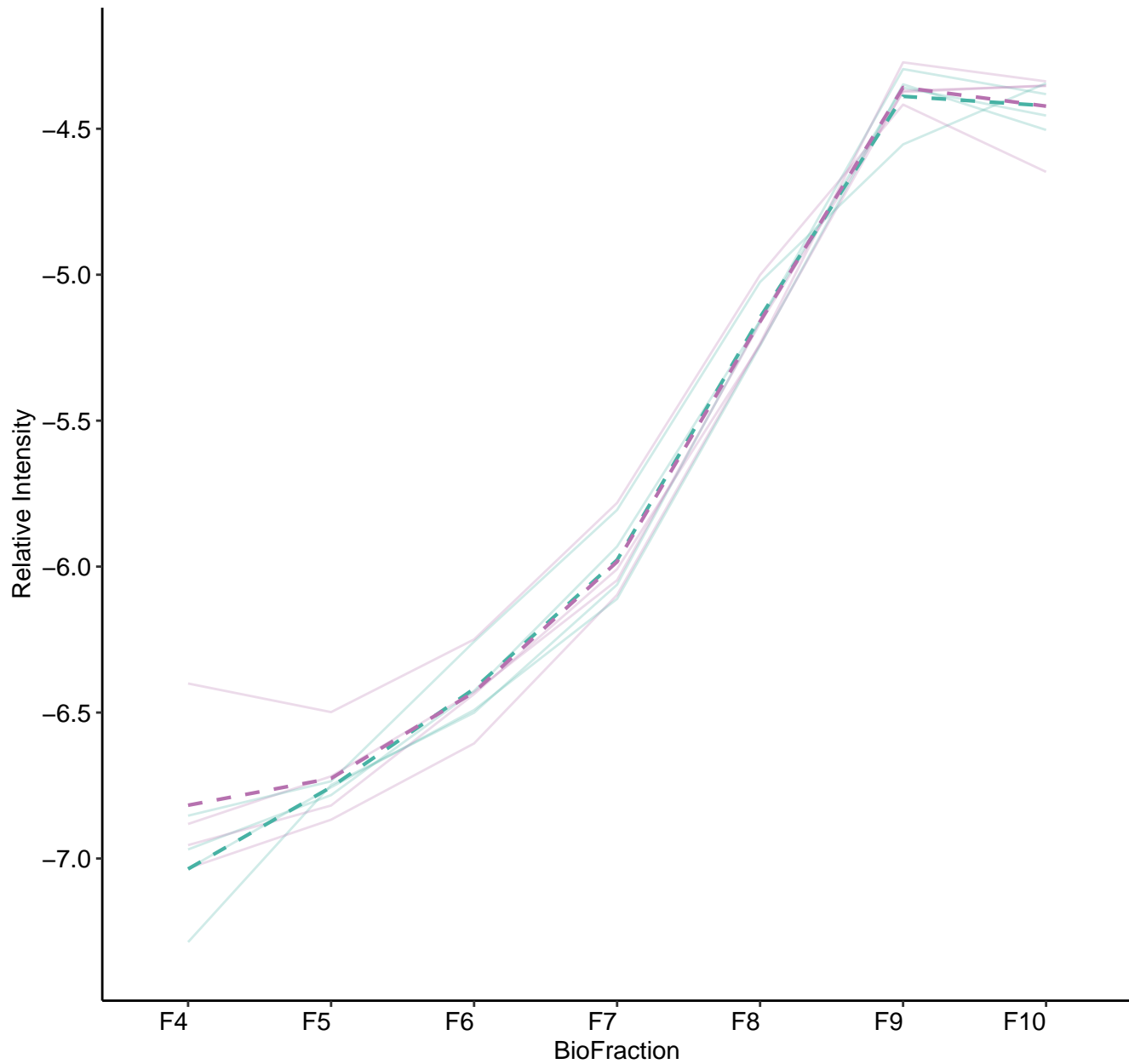
CCT complex (chaperonin containing TCP1 complex), testis specific (n = 7)
(R2.Fixef = 0.996)



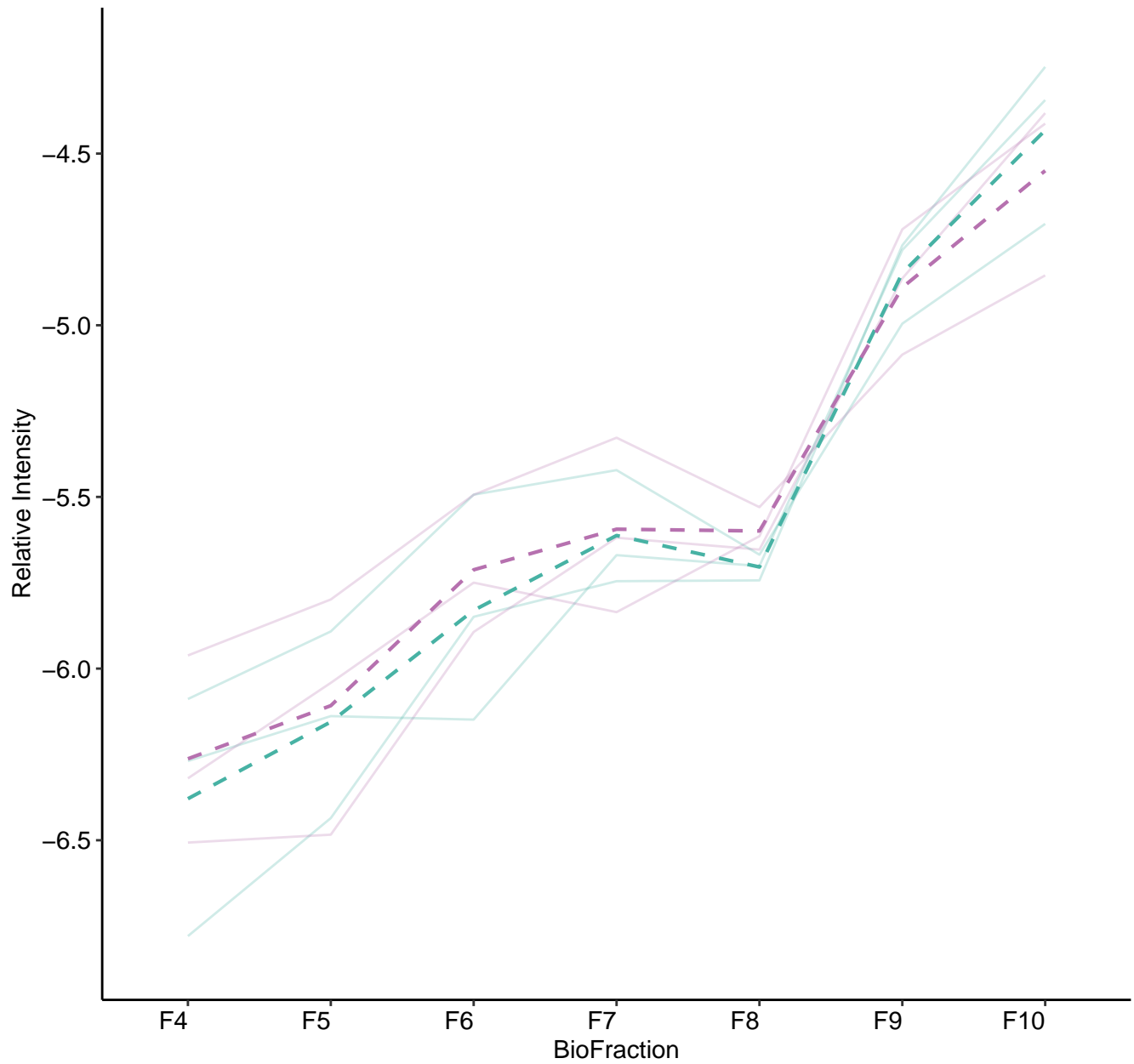
CDC5L complex (n = 11)
(R2.Fixef = 0.899)



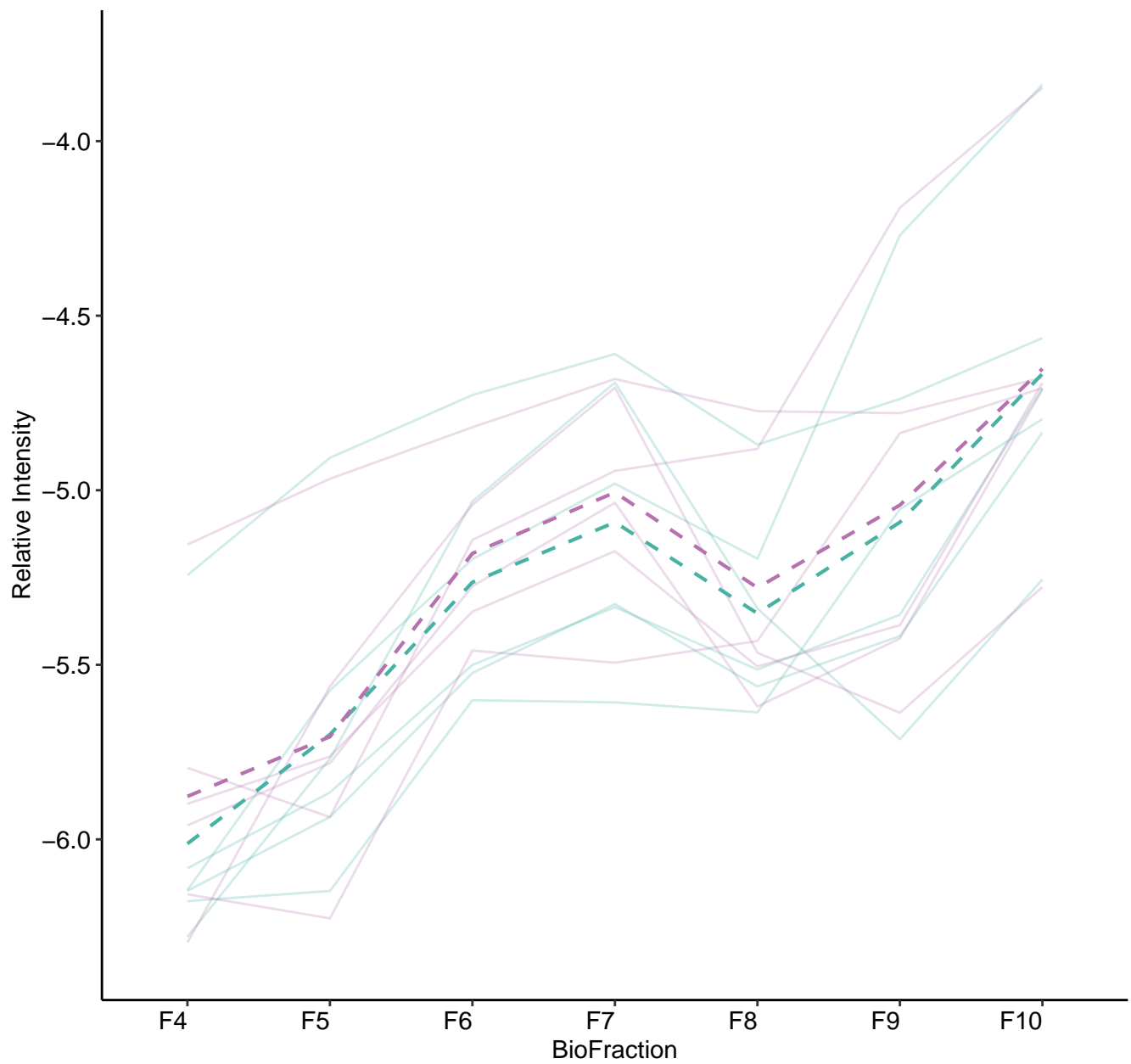
CDC5L core complex (n = 4)
(R2.Fixef = 0.981)



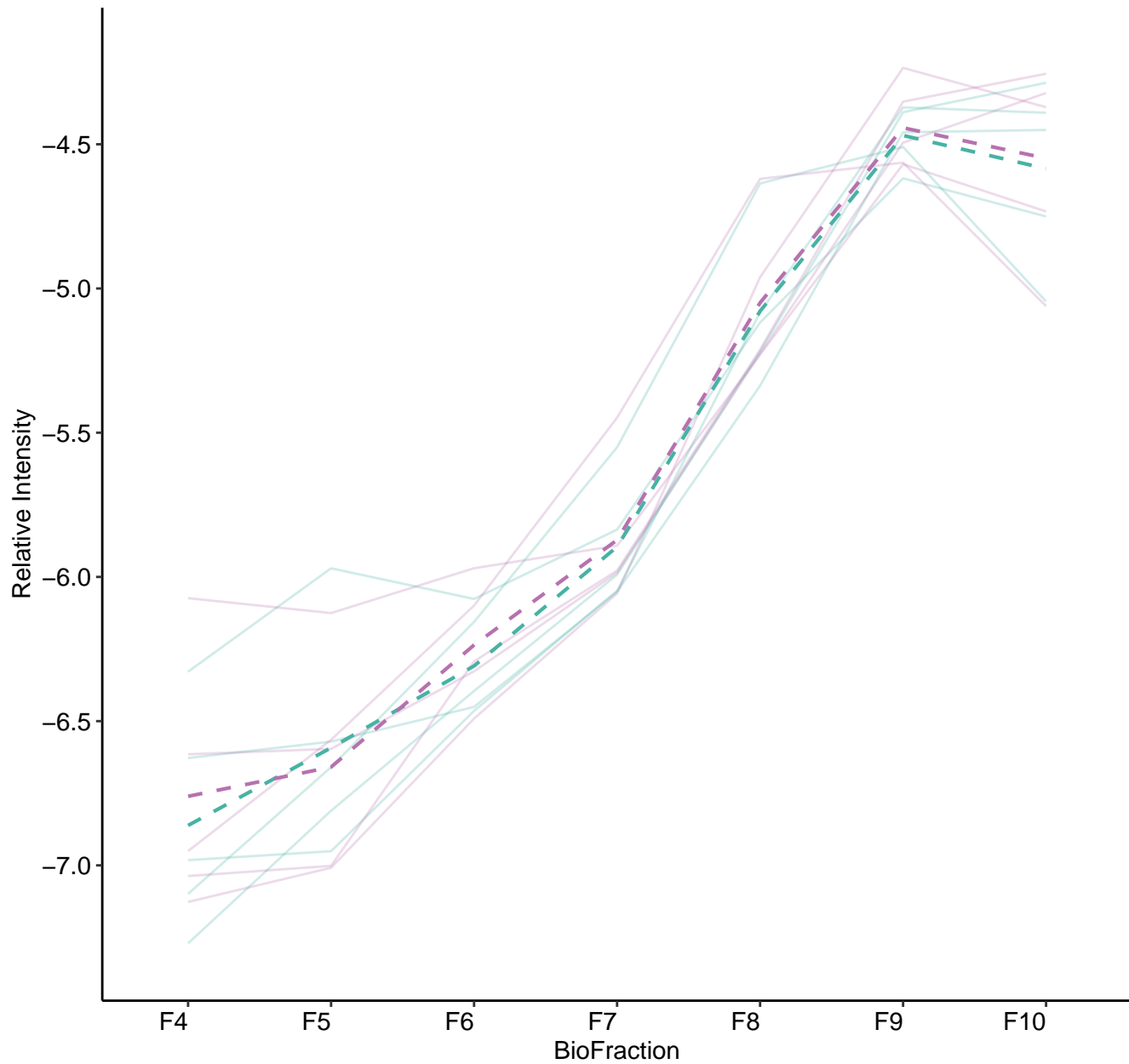
CDH4–HDAC2–MTA2–RBBP7–TWIST1 complex (n = 3)
(R2.Fixef = 0.866)



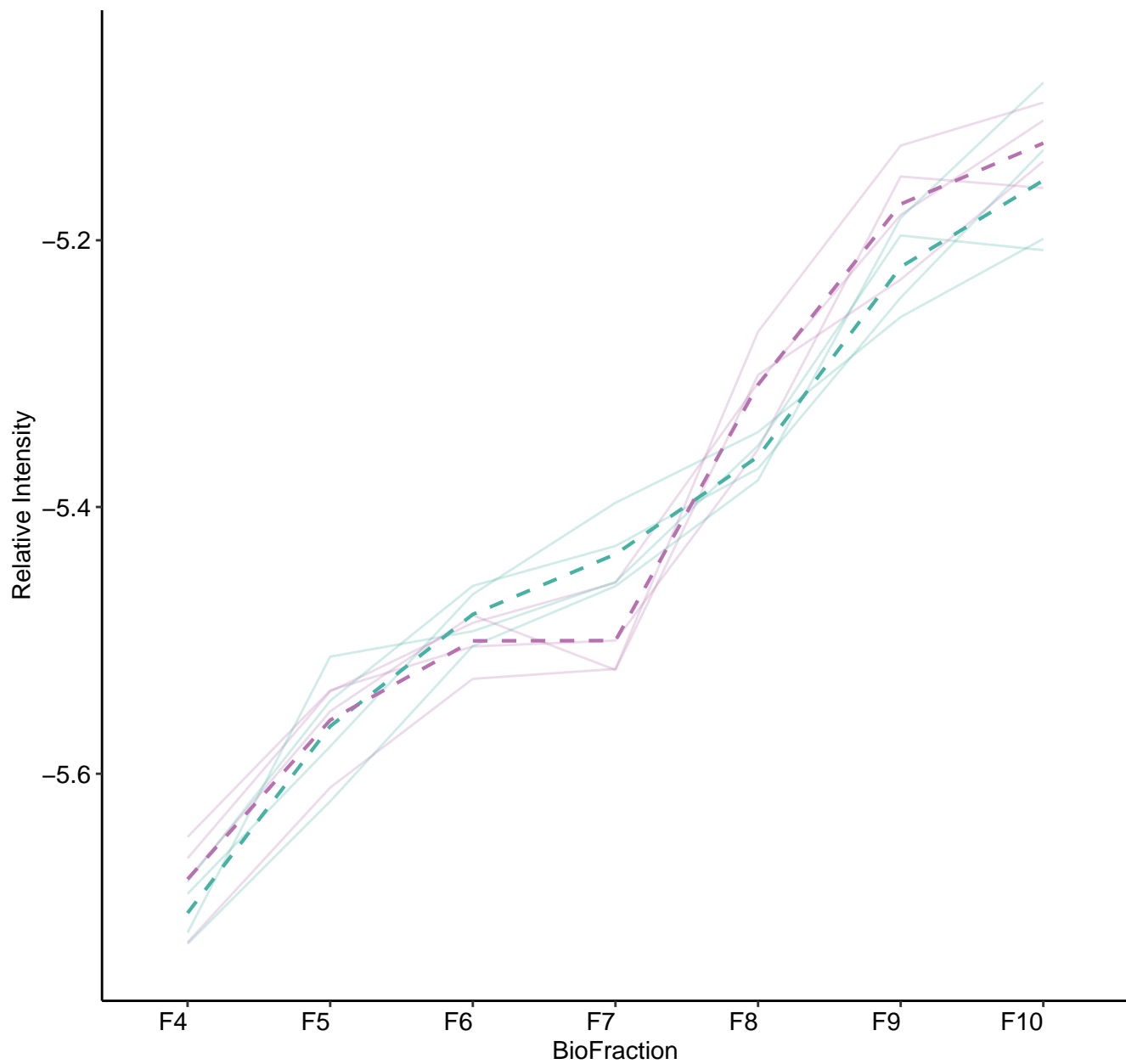
CEN complex (n = 6)
(R2.Fixef = 0.497)



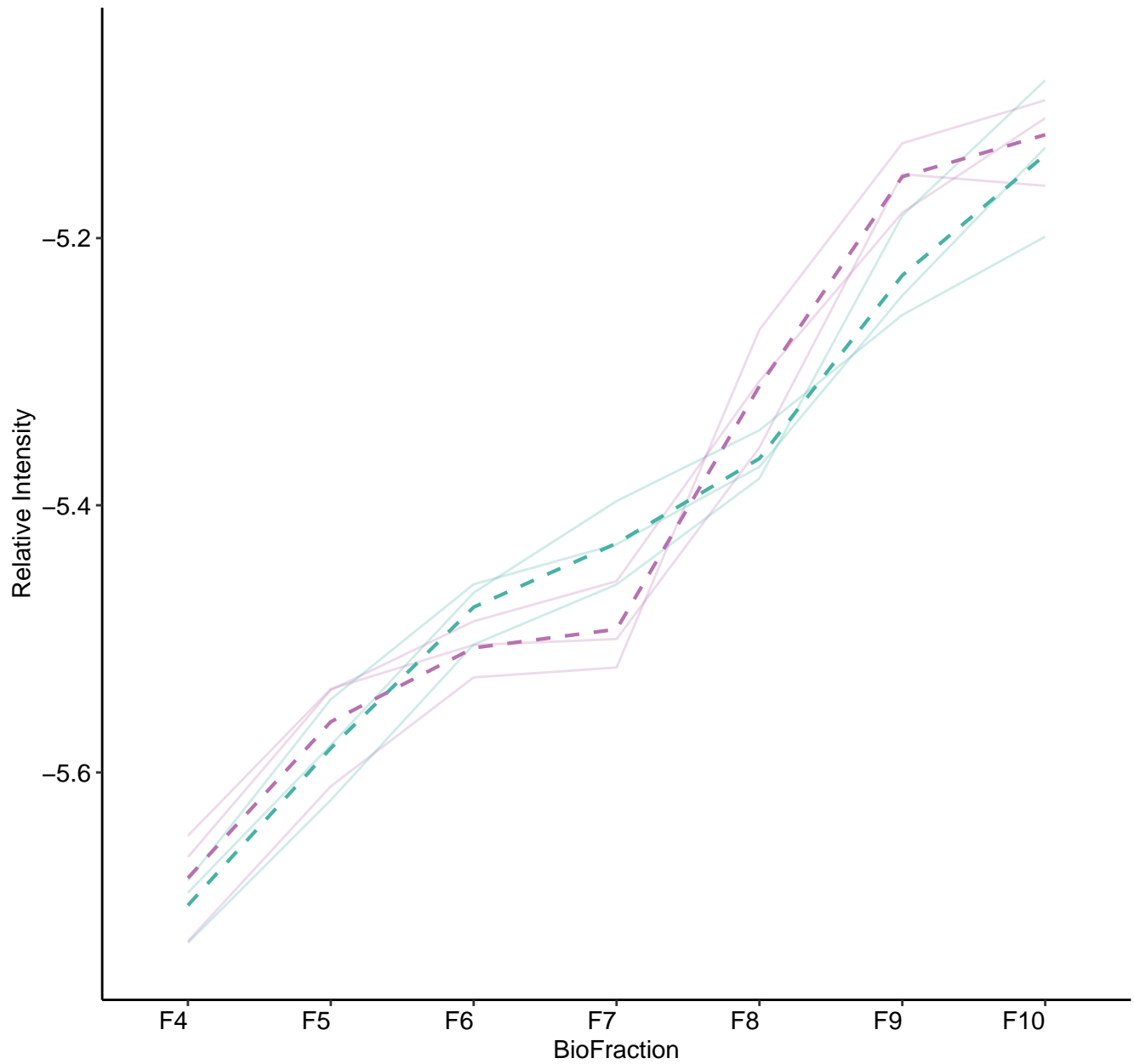
CF IIAm complex (Cleavage factor IIAm complex) (n = 5)
(R2.Fixef = 0.908)



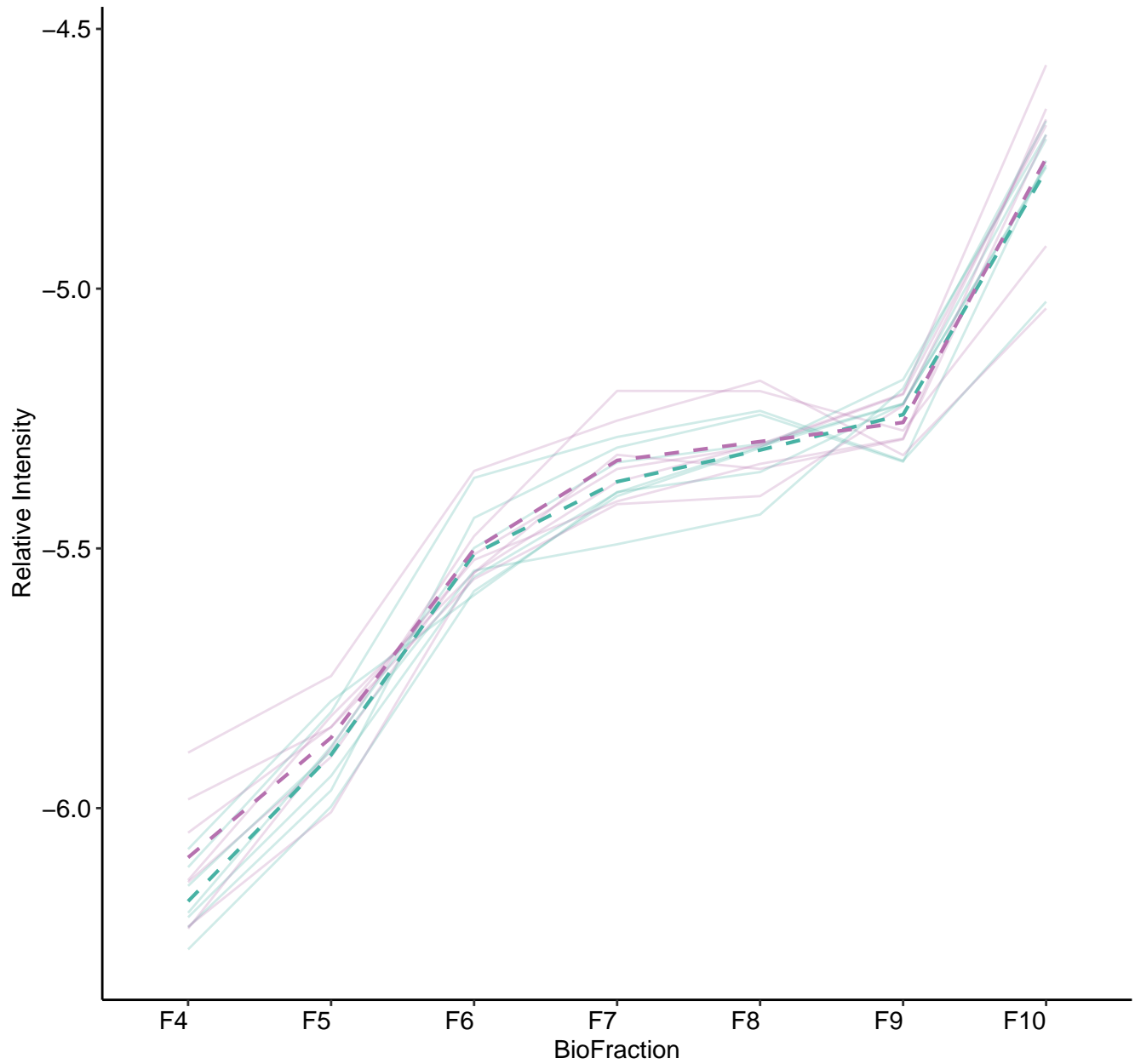
Class C Vps complex (hVPS11, hVPS18, hVPS16, rVPS33a) (n = 4)
(R2.Fixef = 0.967)



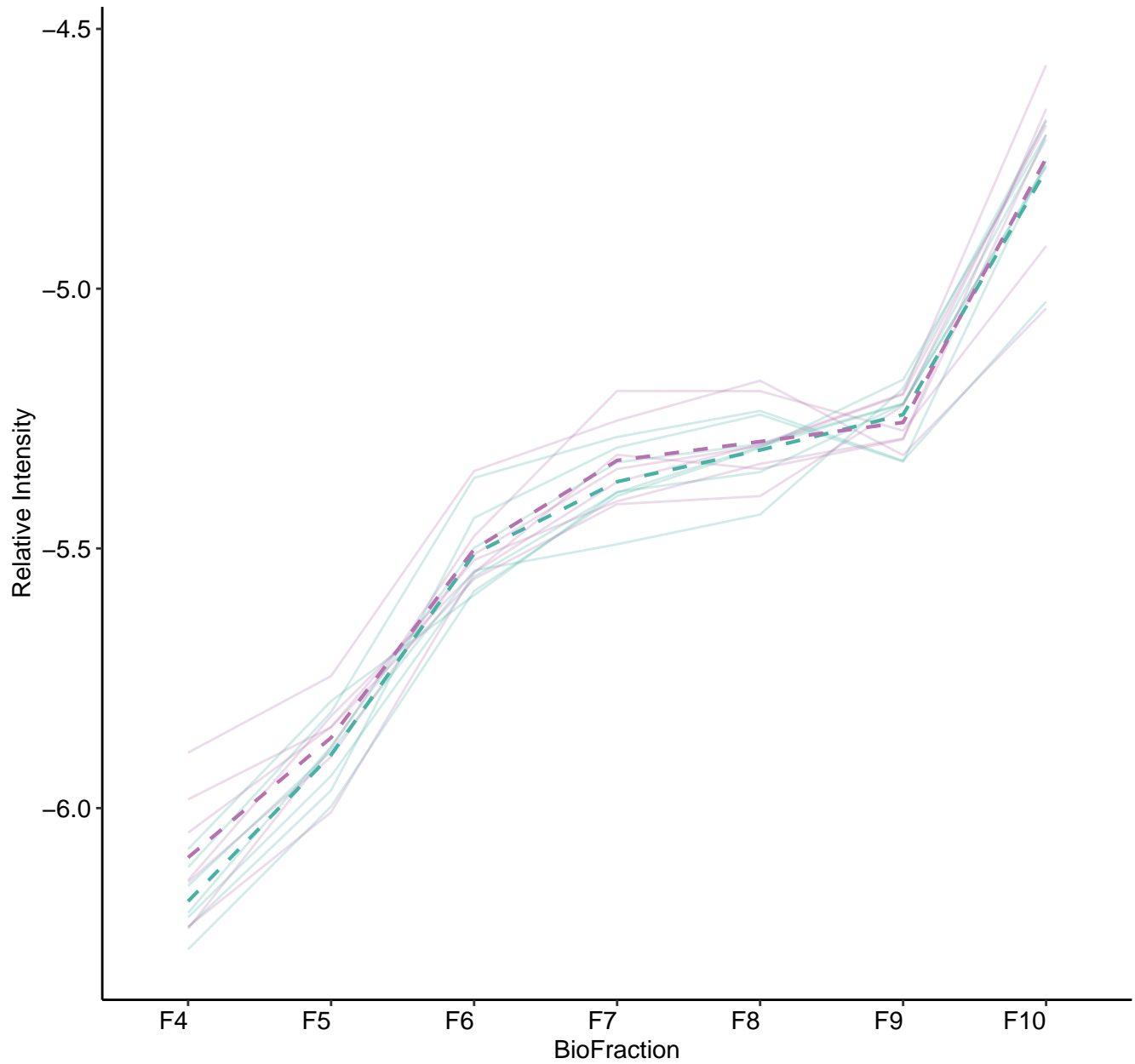
Class C VPS/HOPS complex (n = 3)
(R2.Fixef = 0.966)



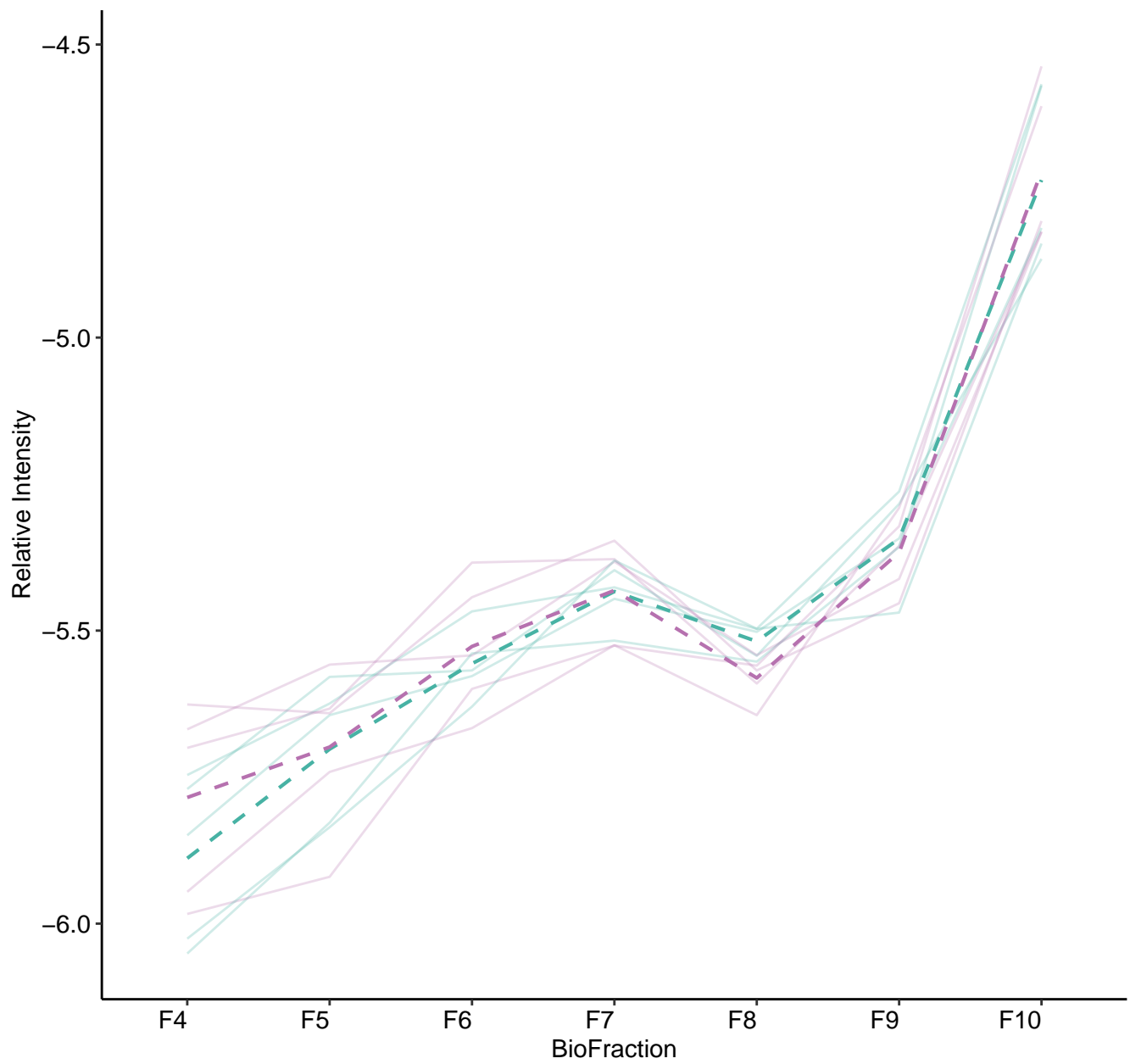
Coatmer complex (n = 7)
(R2.Fixef = 0.955)



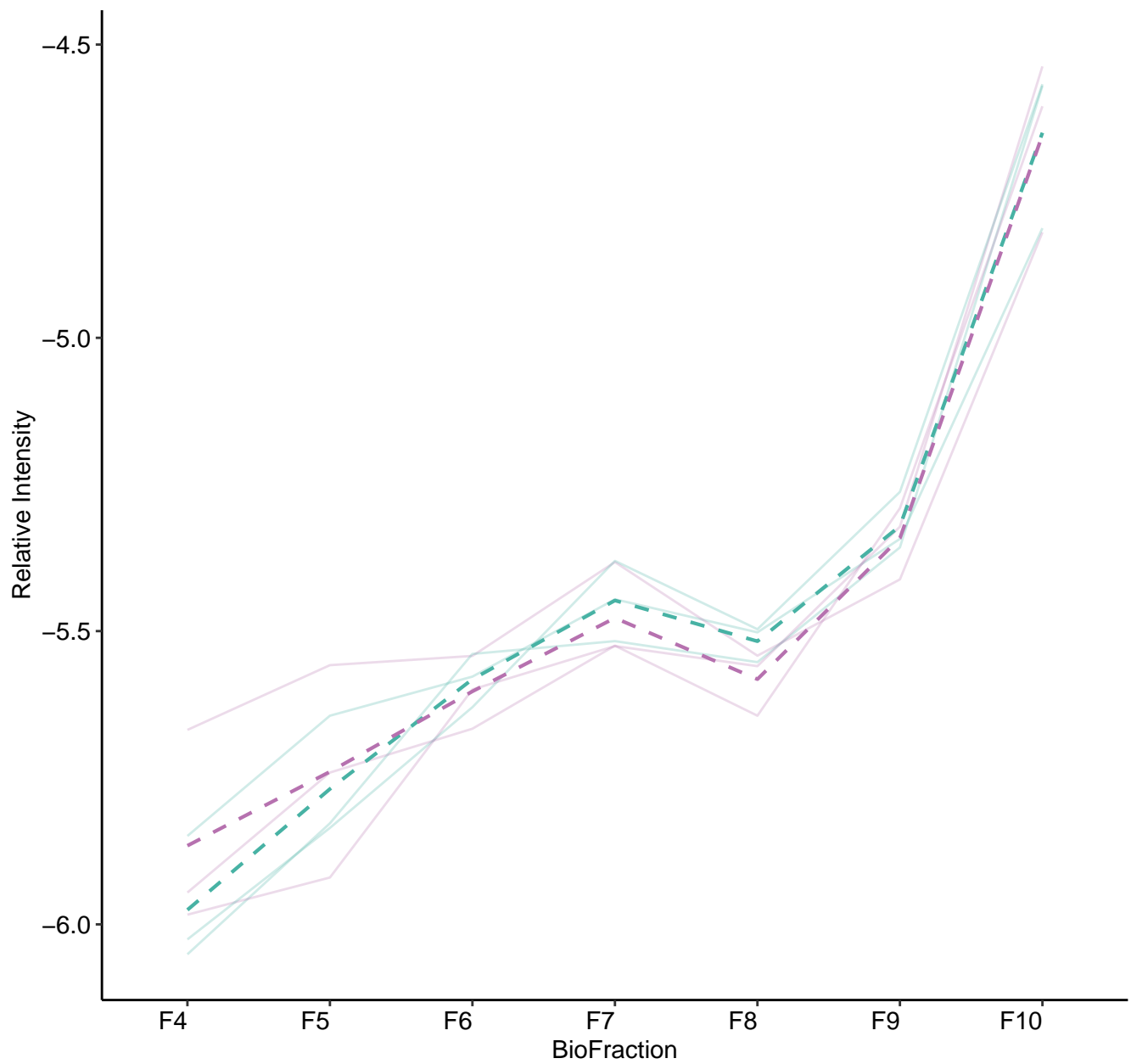
Coatomer-Arf1 complex (n = 7)
(R2.Fixef = 0.955)



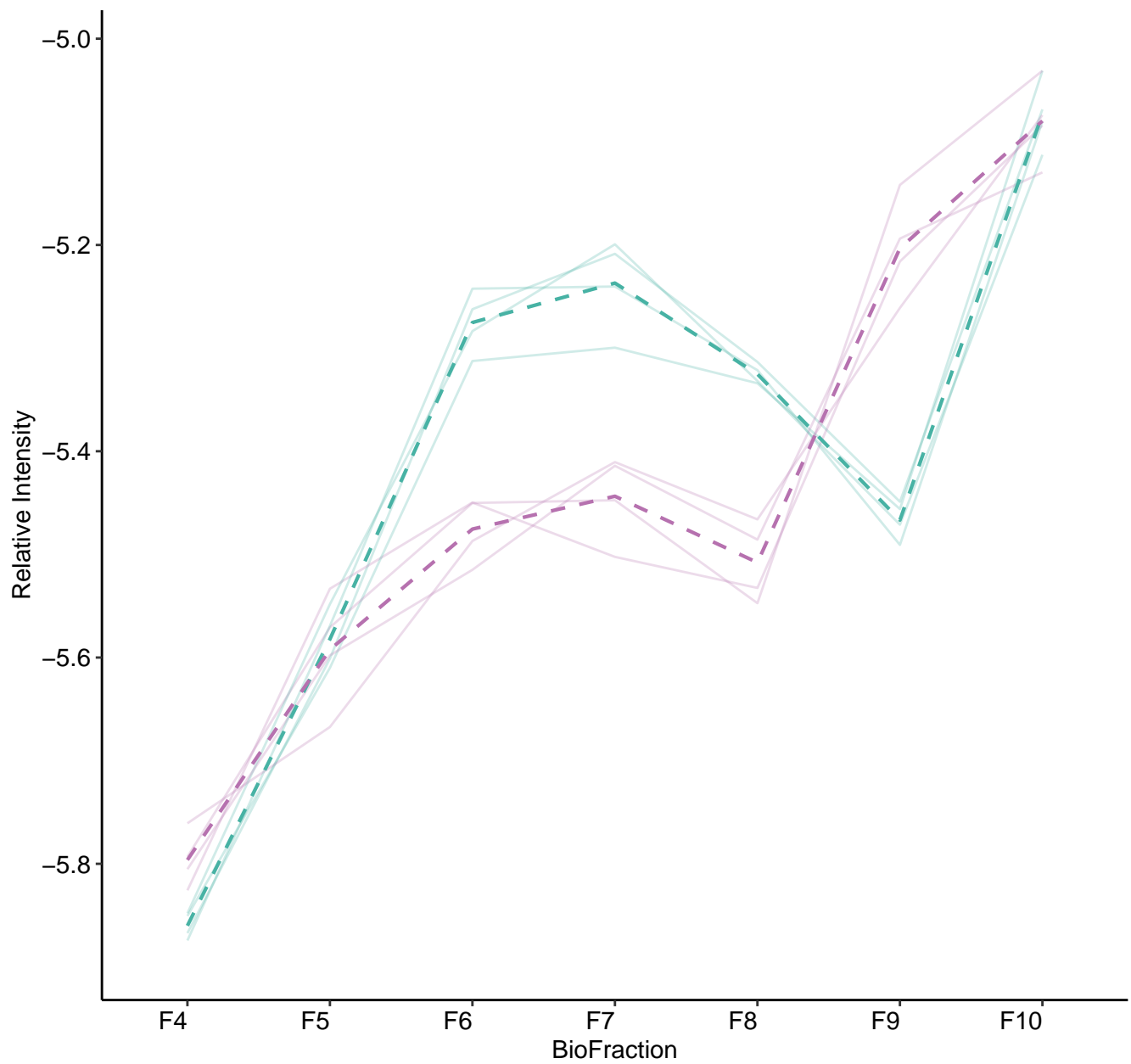
COG complex (n = 5)
(R2.Fixef = 0.906)



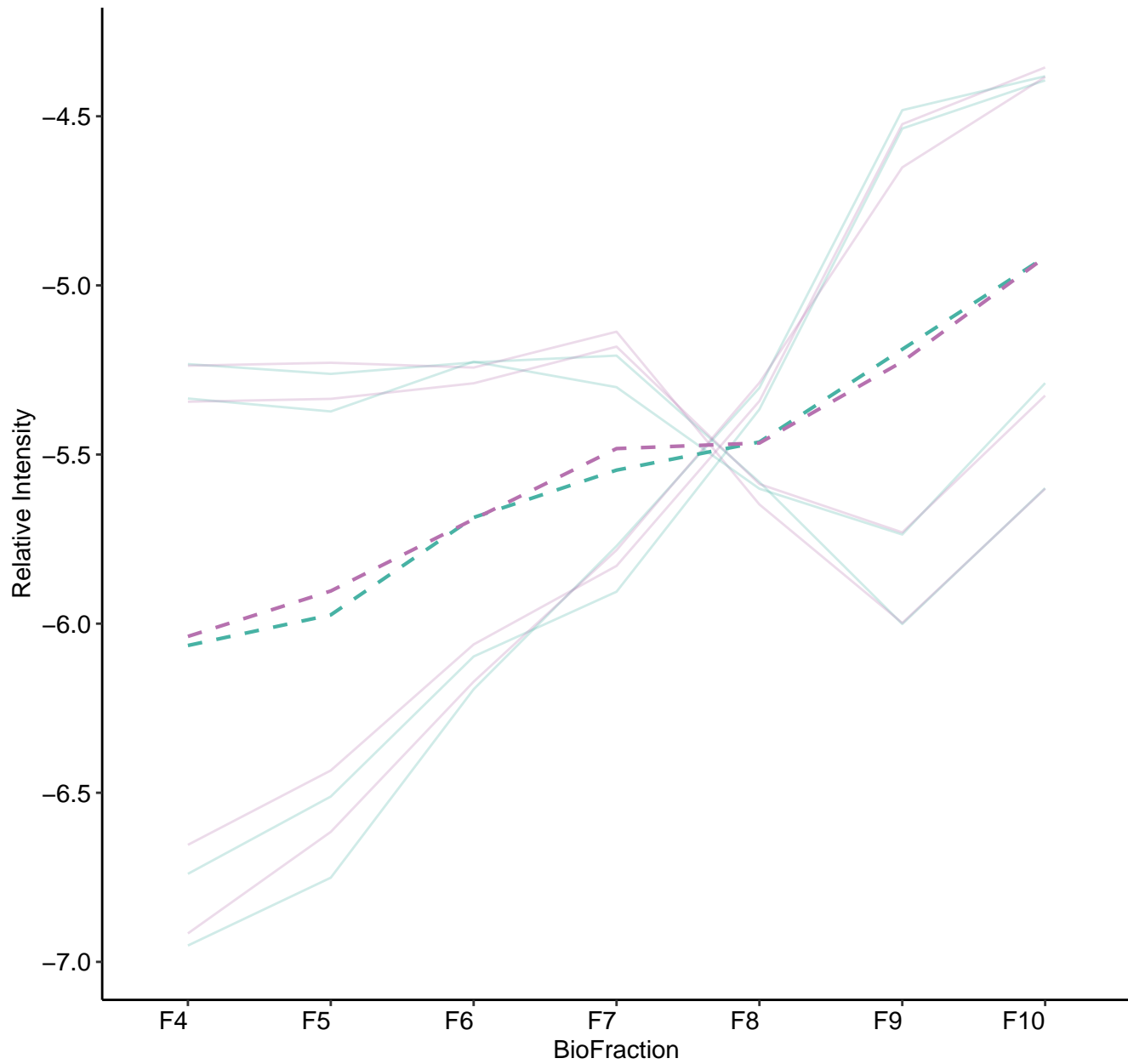
COG1-COG8-COG2-COG3-COG4 subcomplex (n = 3)
(R2.Fixef = 0.929)



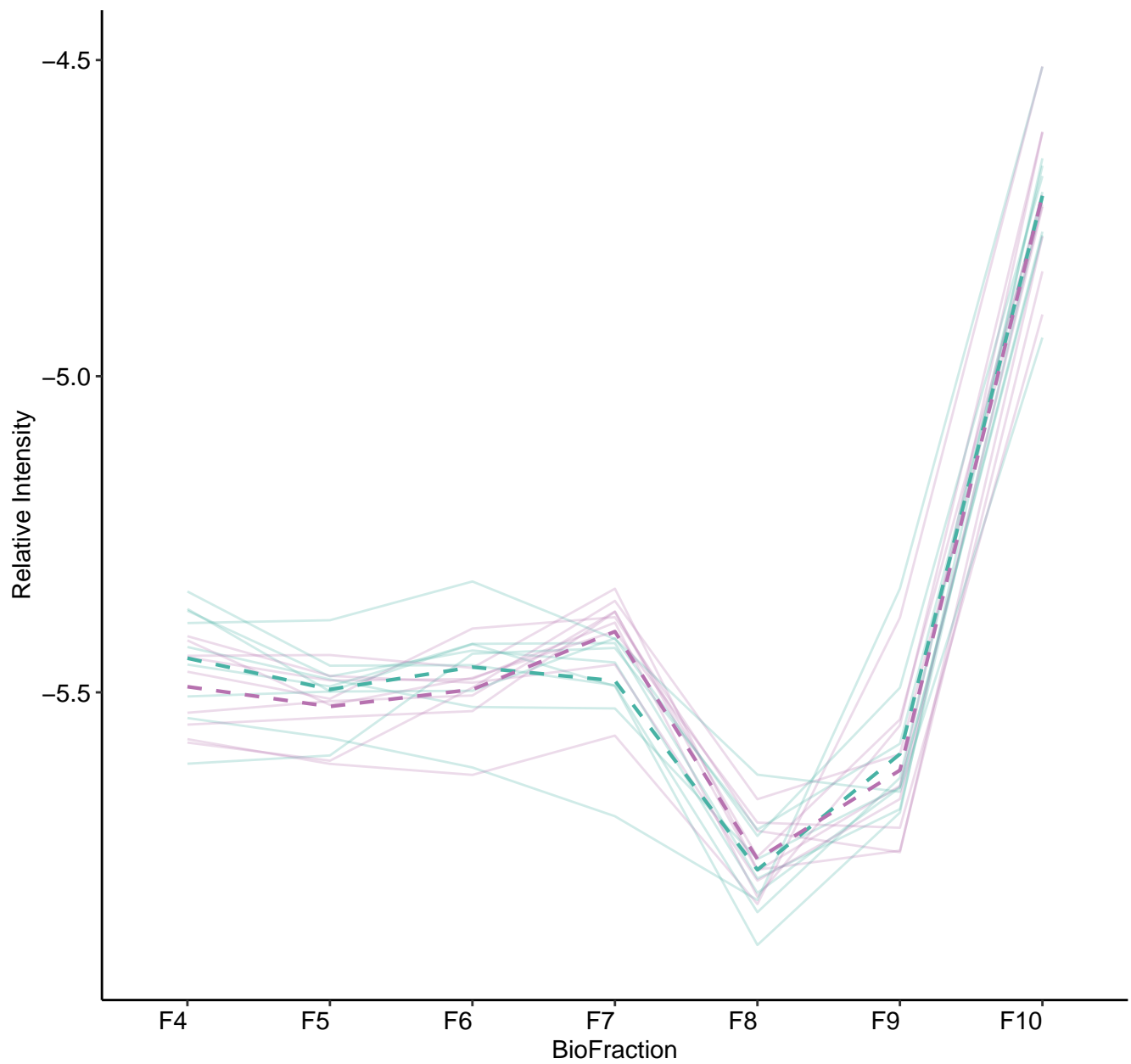
COMMD1-CCDC22-CCDC93-C16orf62 complex (n = 4)
(R2.Fixef = 0.977)



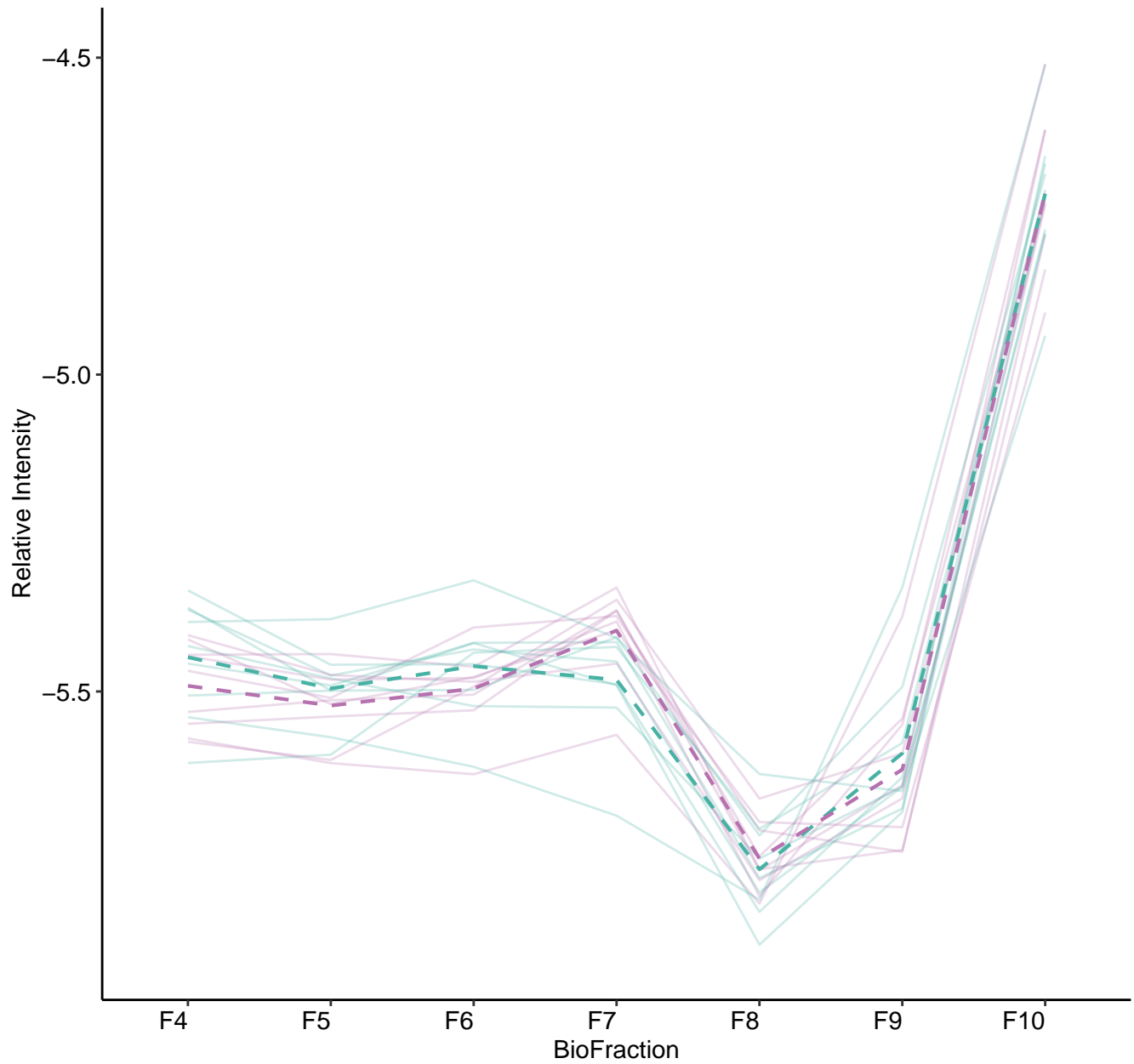
CRM1-RAN-PHAX-CBC complex (cap binding complex) (n = 4)
(R2.Fixef = 0.26)



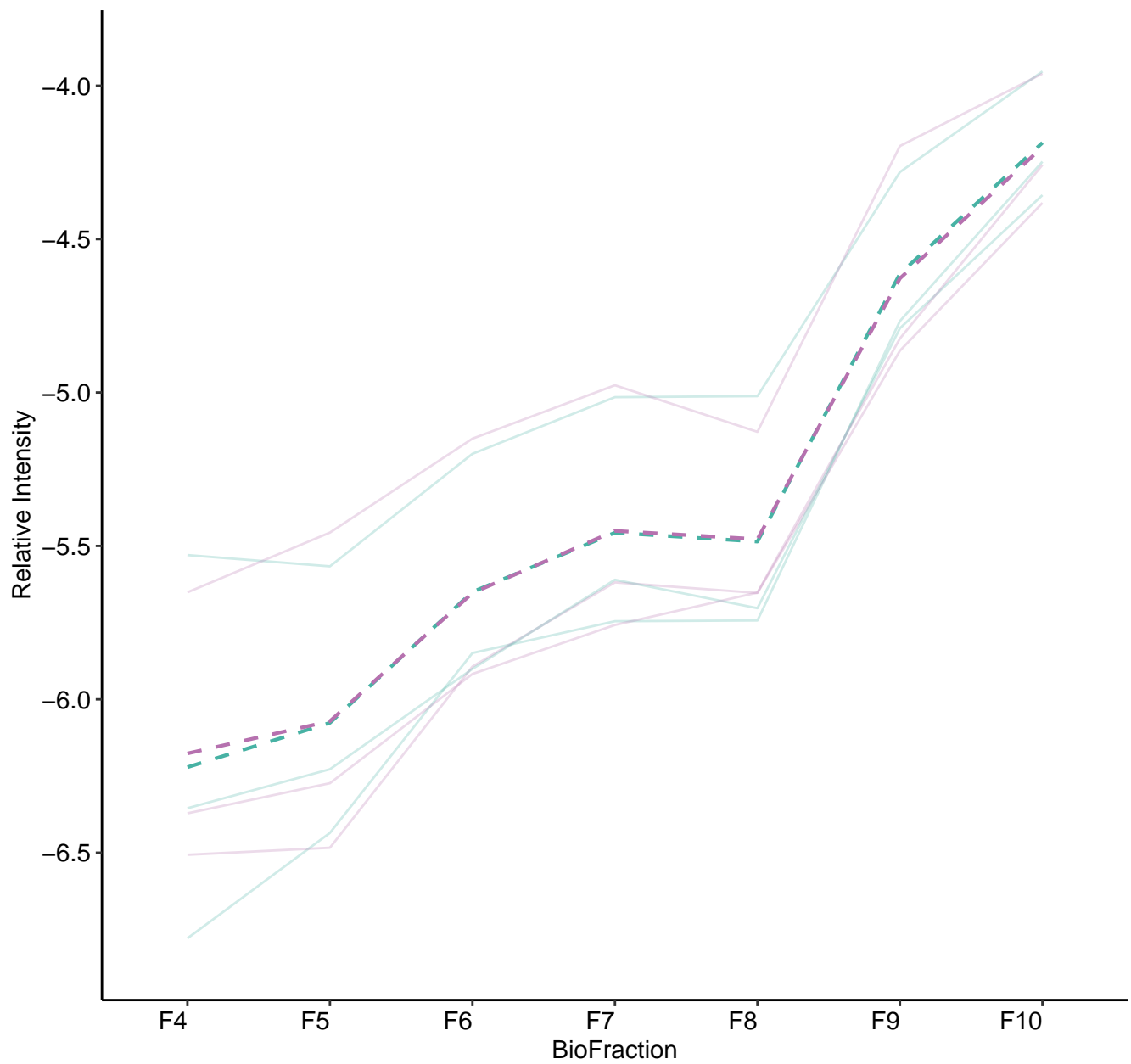
CSA complex (n = 9)
(R2.Fixef = 0.926)



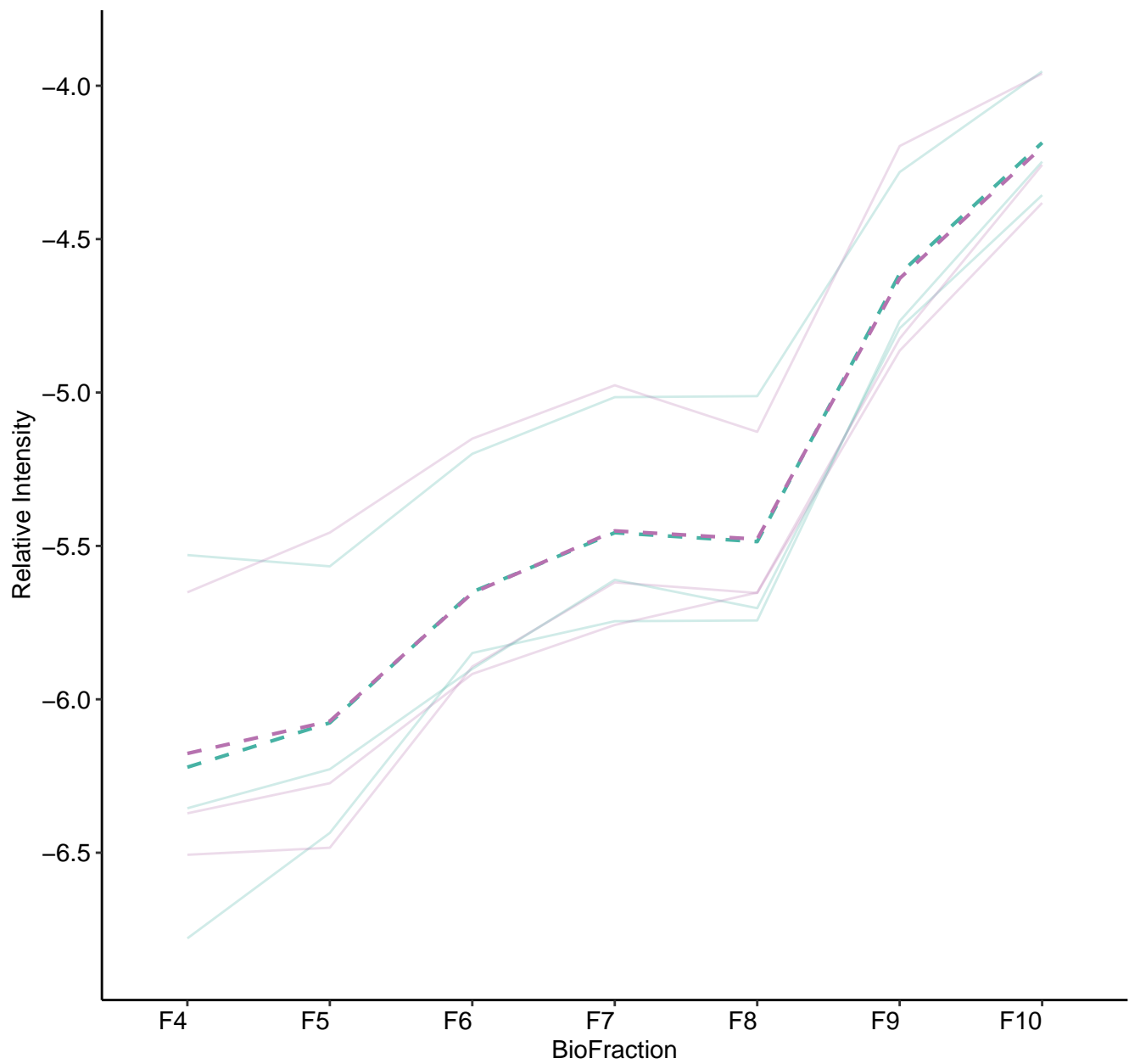
CSA-POLIIa complex (n = 9)
(R2.Fixef = 0.926)



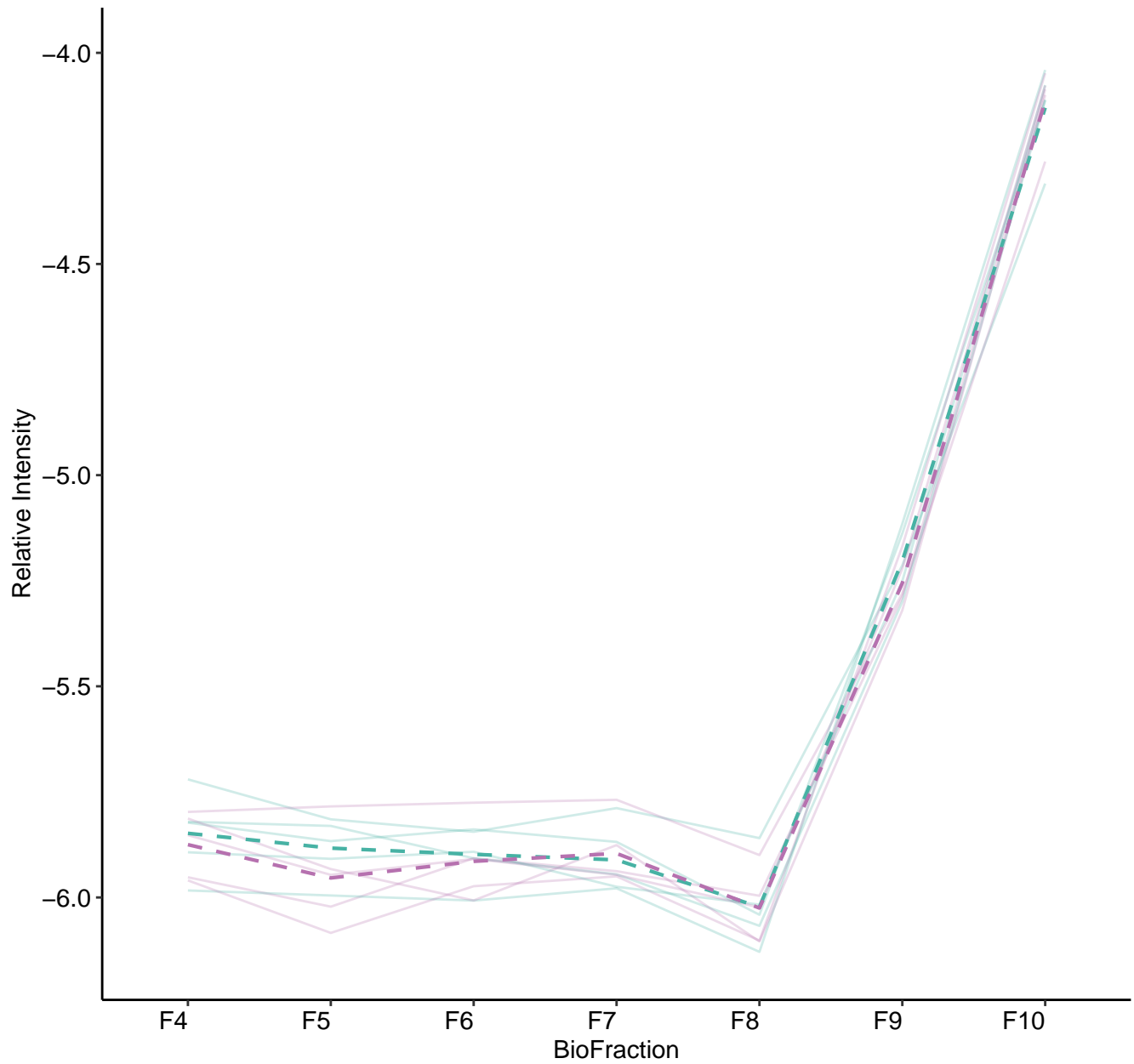
CtBP complex (n = 3)
(R2.Fixef = 0.737)



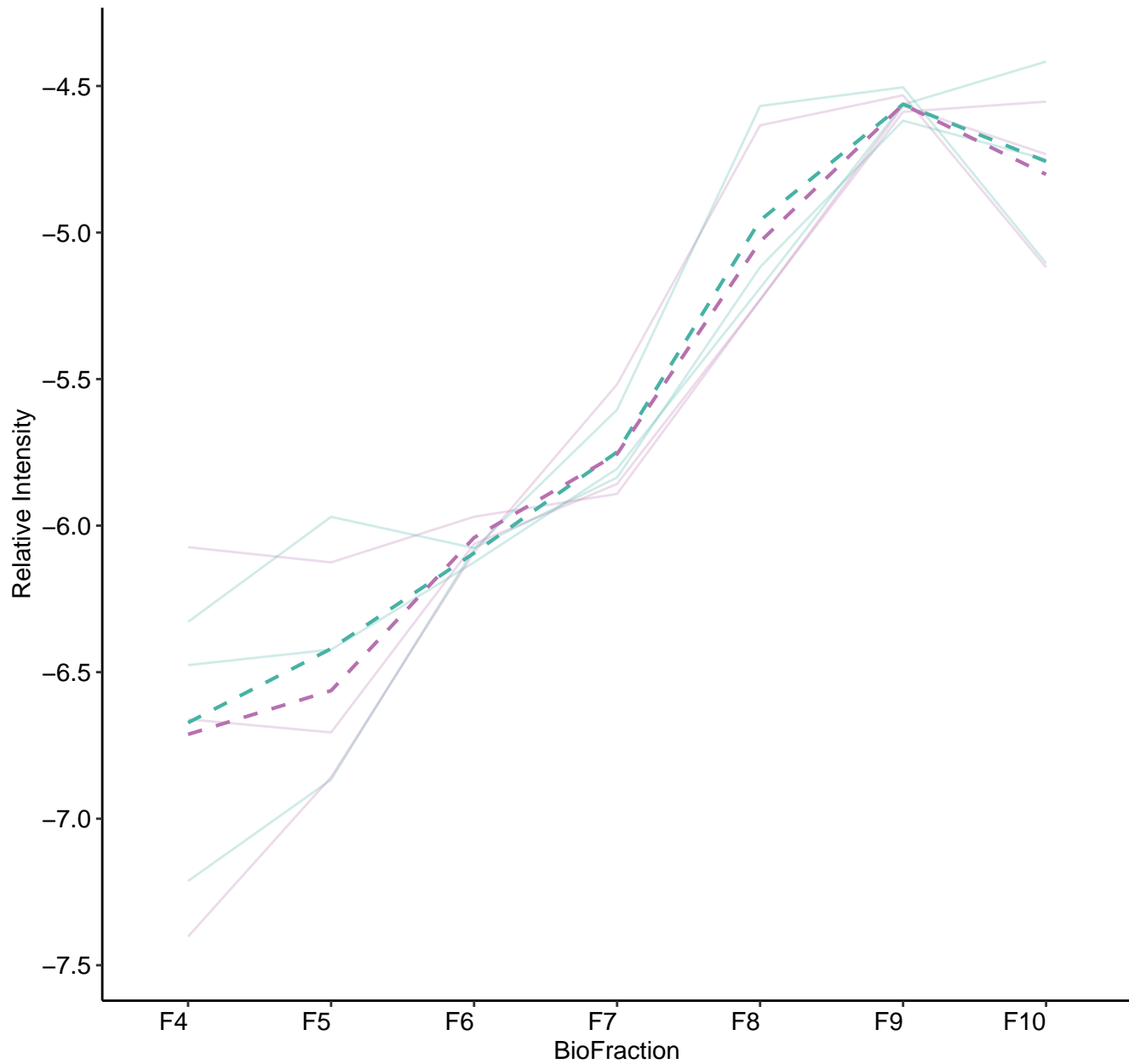
CtBP core complex (n = 3)
(R2.Fixef = 0.737)



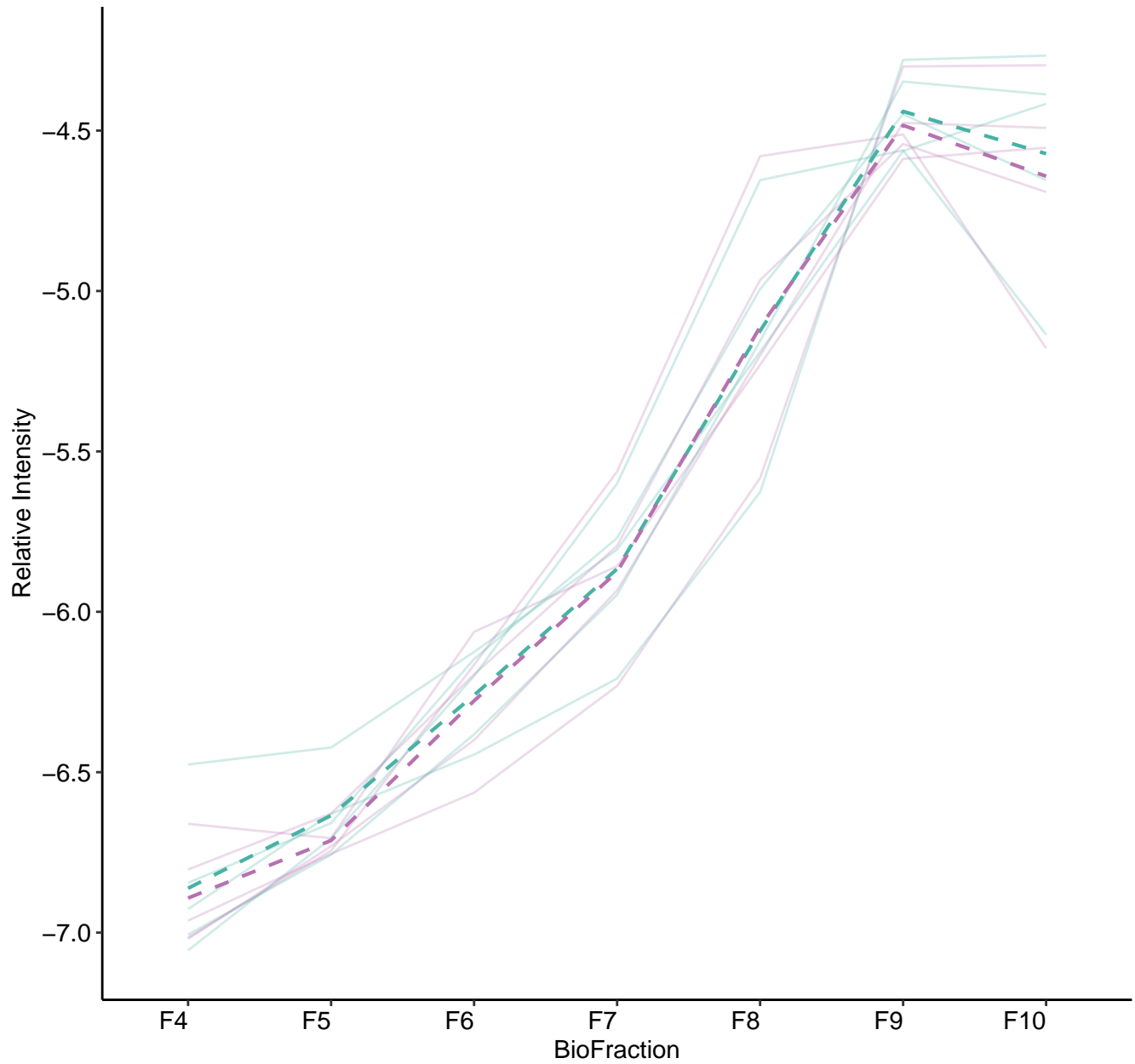
CTLH complex (n = 5)
(R2.Fixef = 0.983)



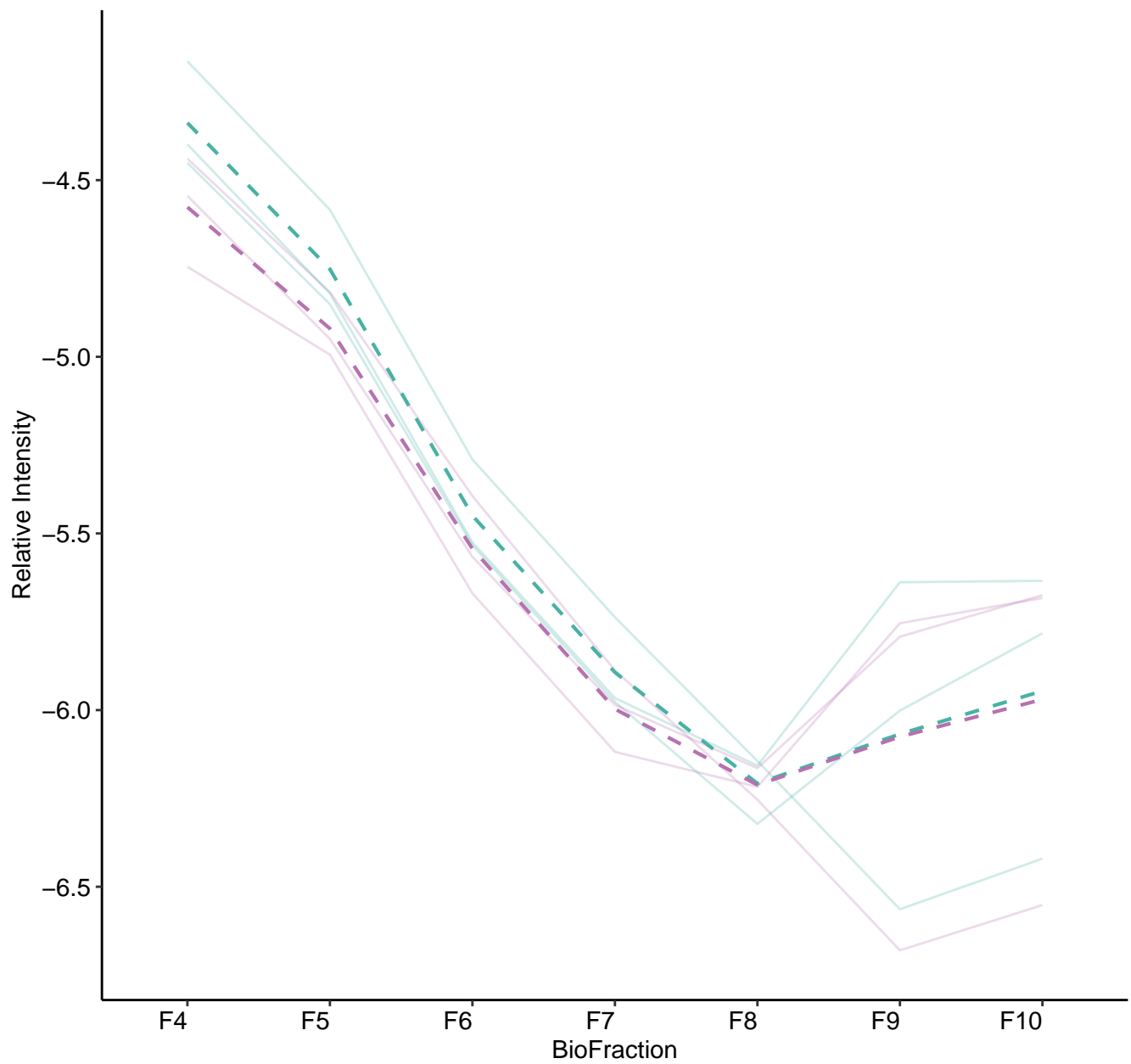
DCS complex (Ptbp1, Ptbp2, Hnrph1, Hnrpf) (n = 3)
(R2.Fixef = 0.854)



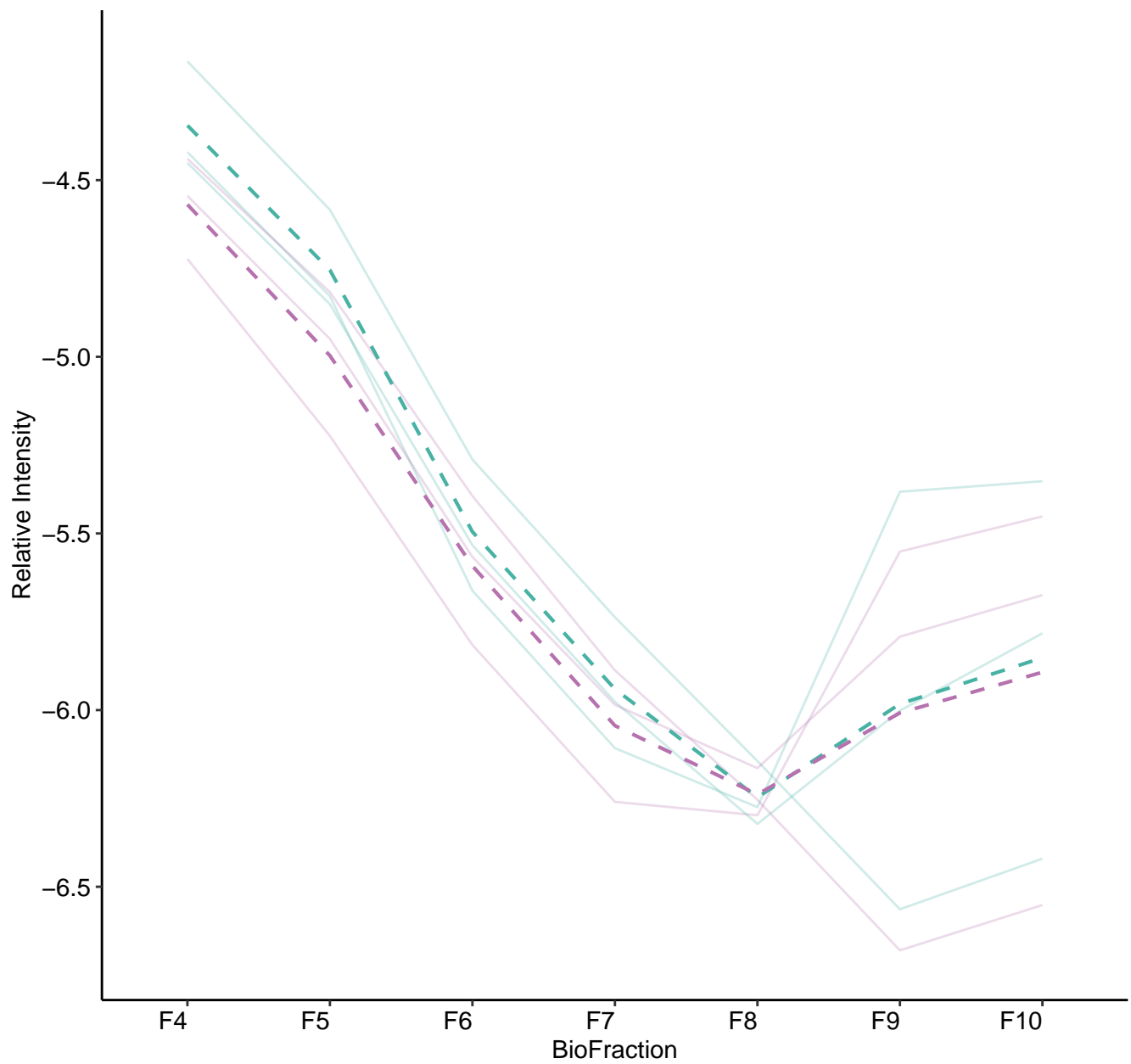
DGCR8 multiprotein complex (n = 5)
(R2.Fixef = 0.937)



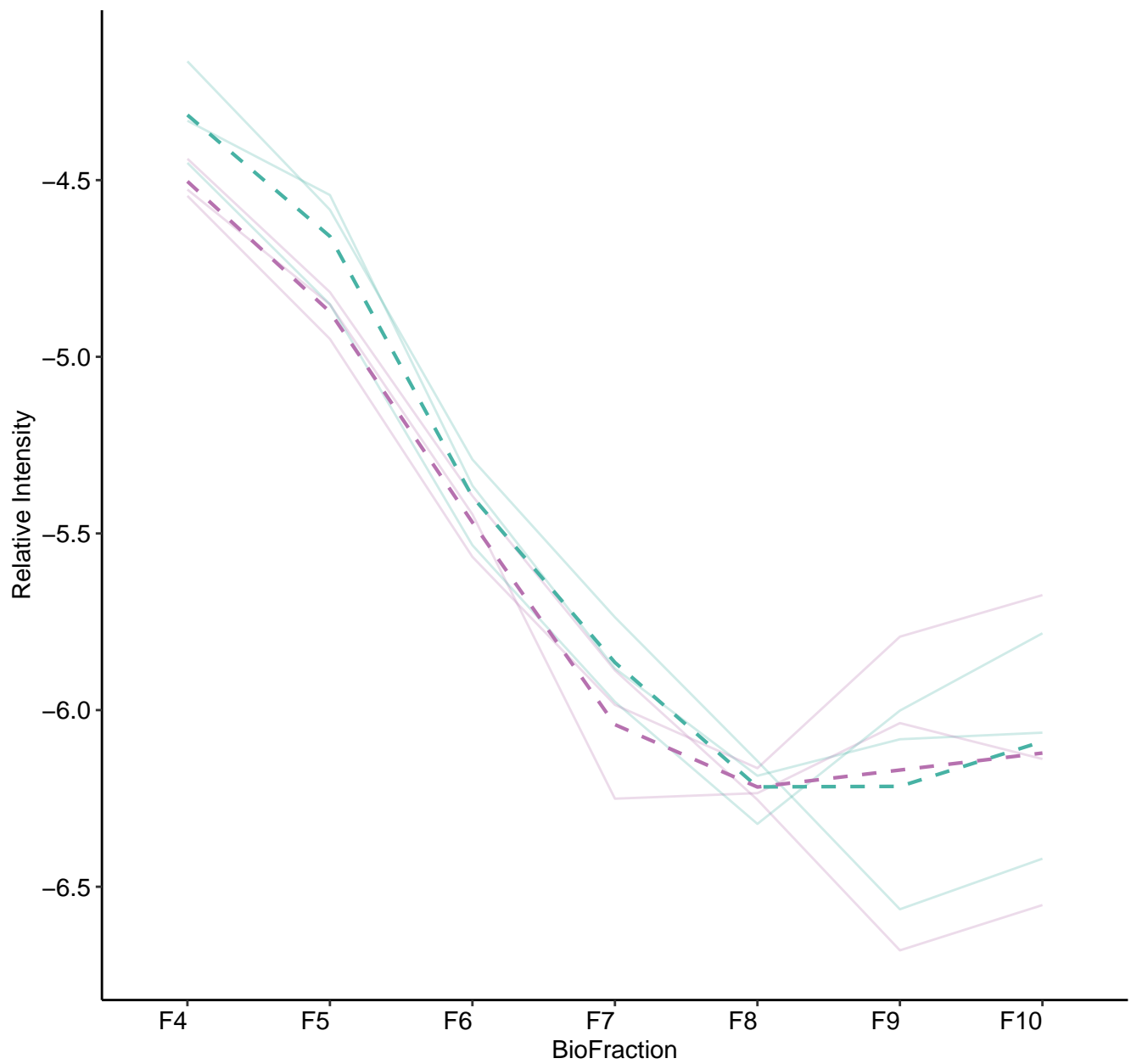
DLG4-DLGAP1-SHANK2 complex (n = 3)
(R2.Fixef = 0.84)



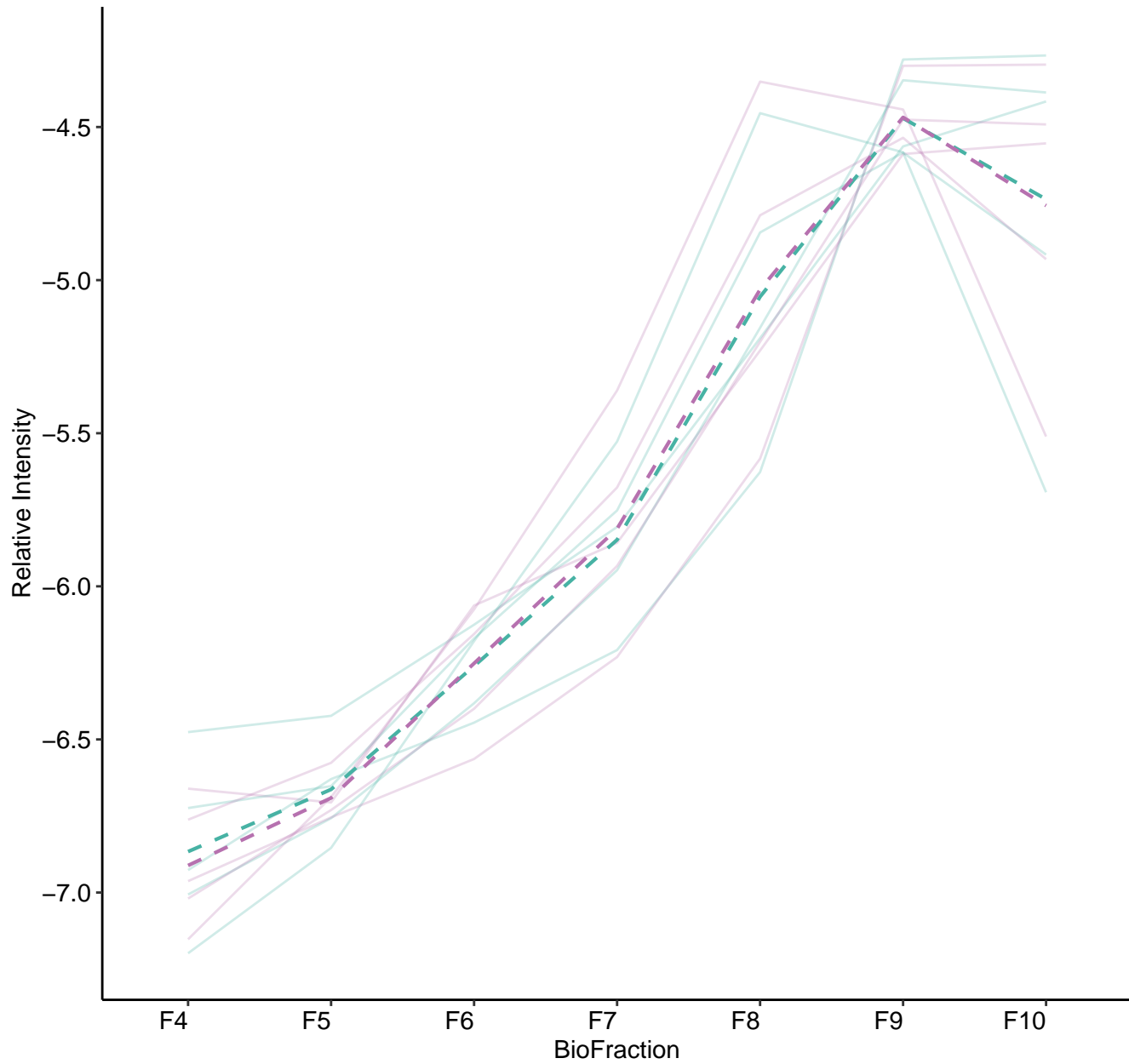
DLG4-DLGAP1-SHANK3 complex (n = 3)
(R2.Fixef = 0.772)



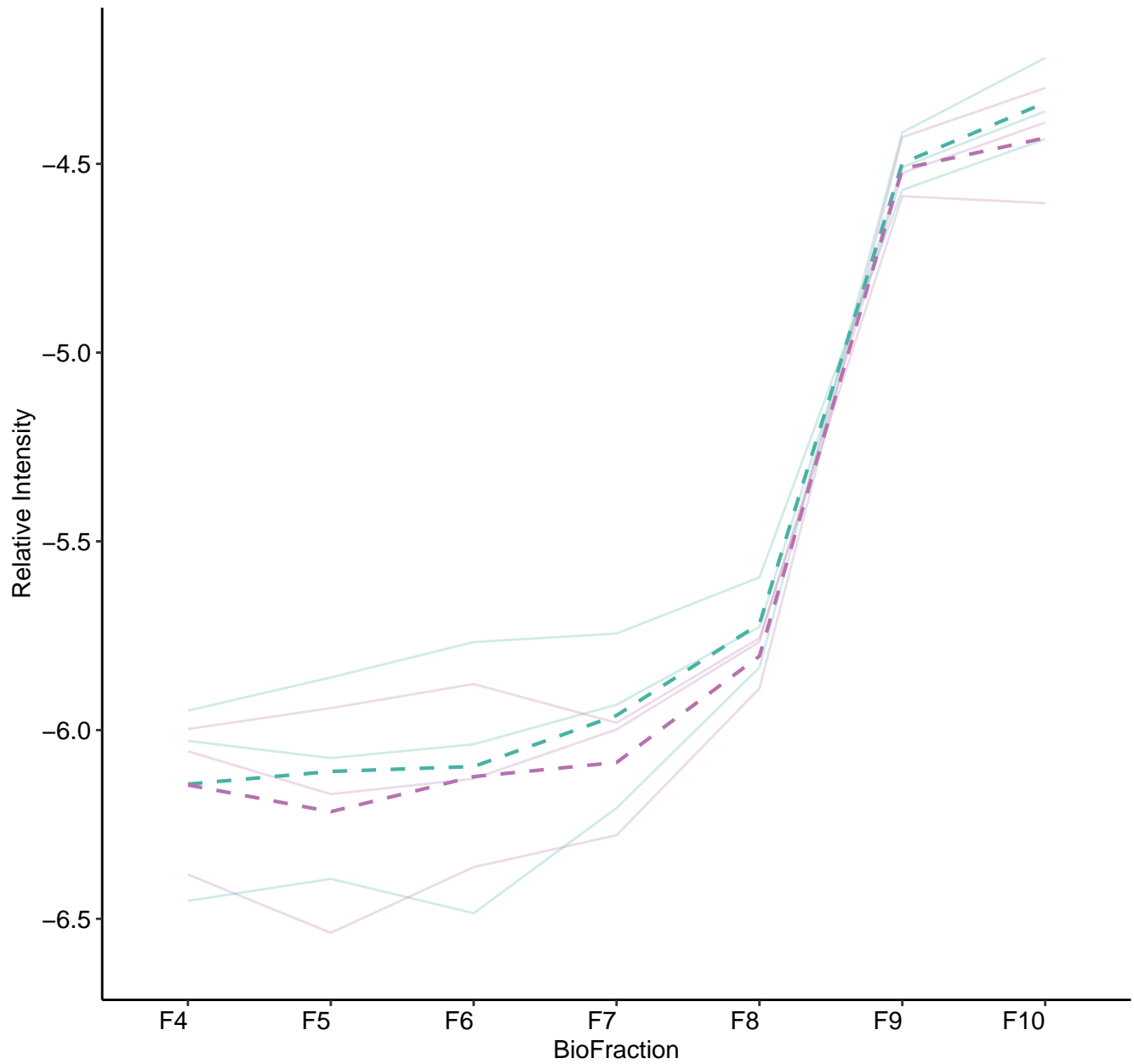
Dlgap1-Dlg4-Begain complex (n = 3)
(R2.Fixef = 0.901)



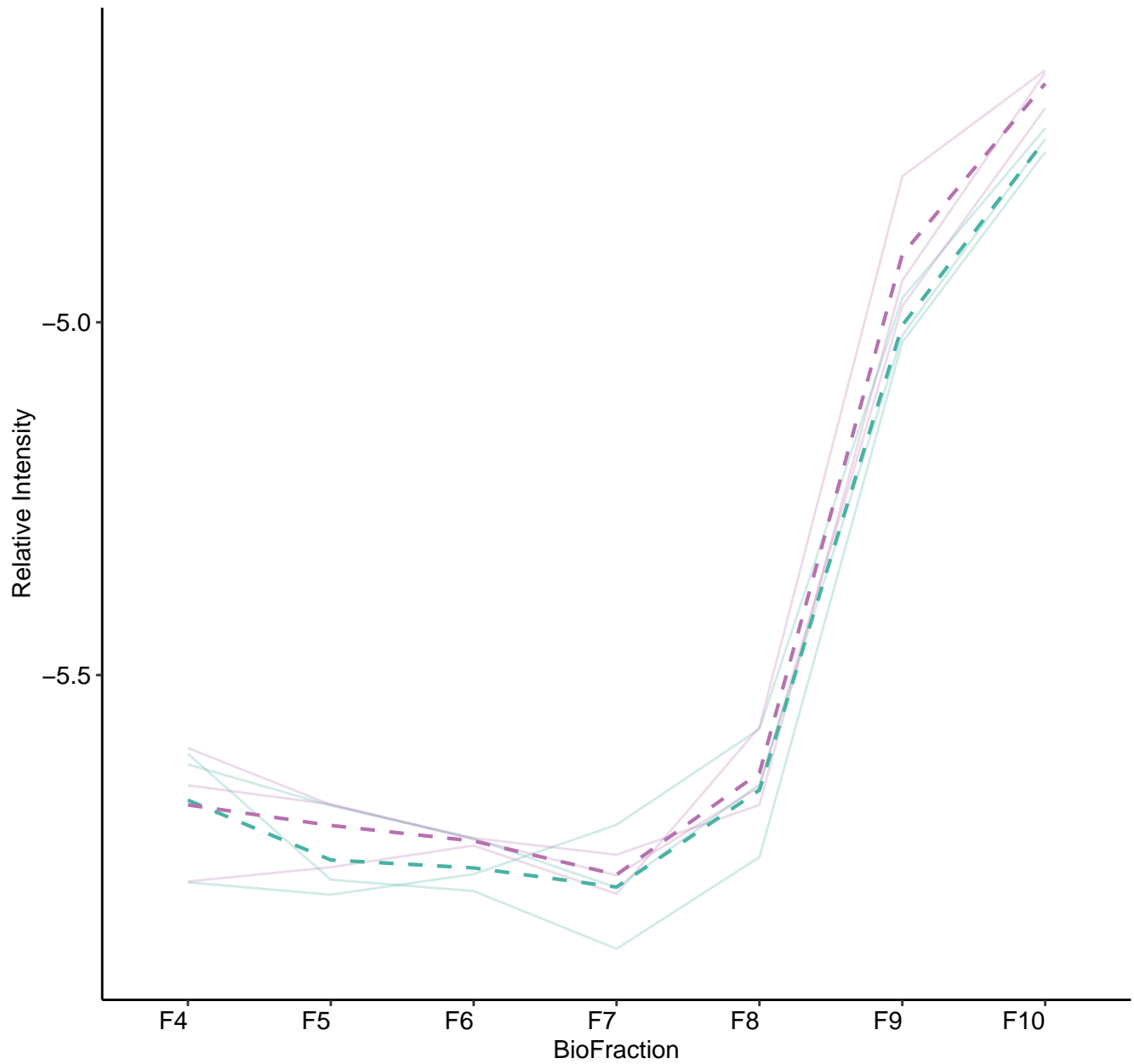
Drosha complex (n = 5)
(R2.Fixef = 0.889)



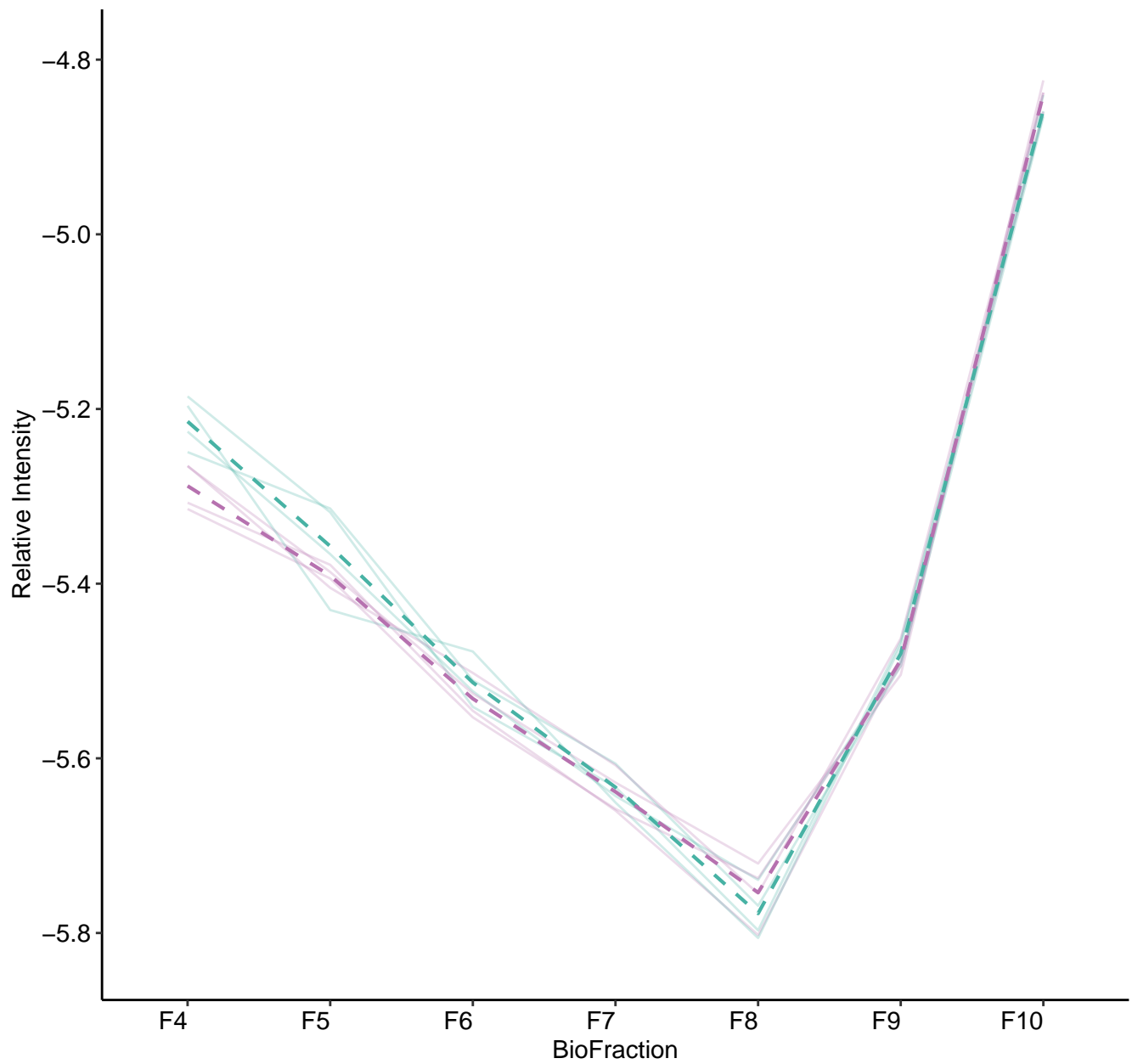
DSS1 complex (n = 3)
(R2.Fixef = 0.926)



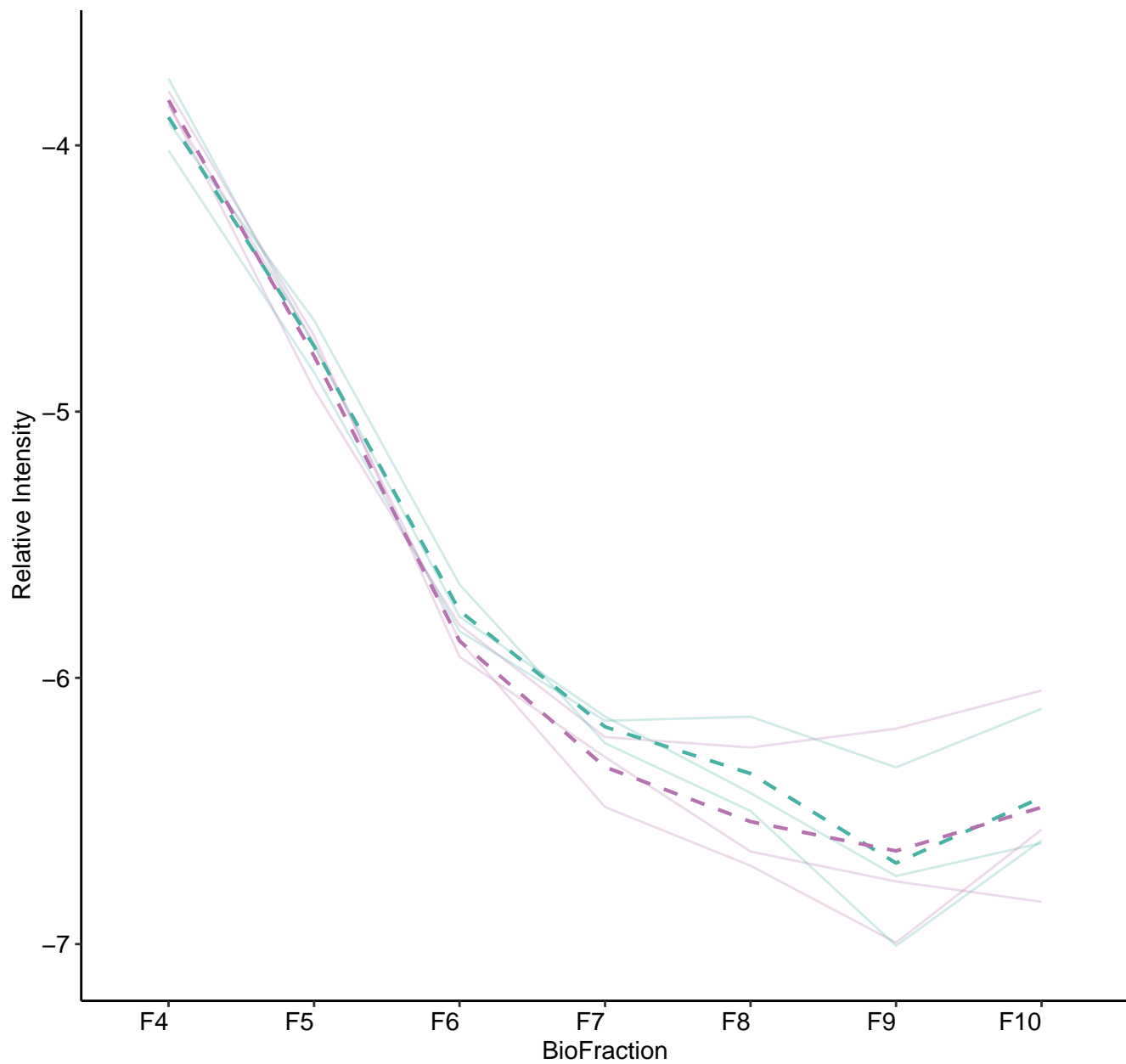
Dynein complex (n = 3)
(R2.Fixef = 0.976)



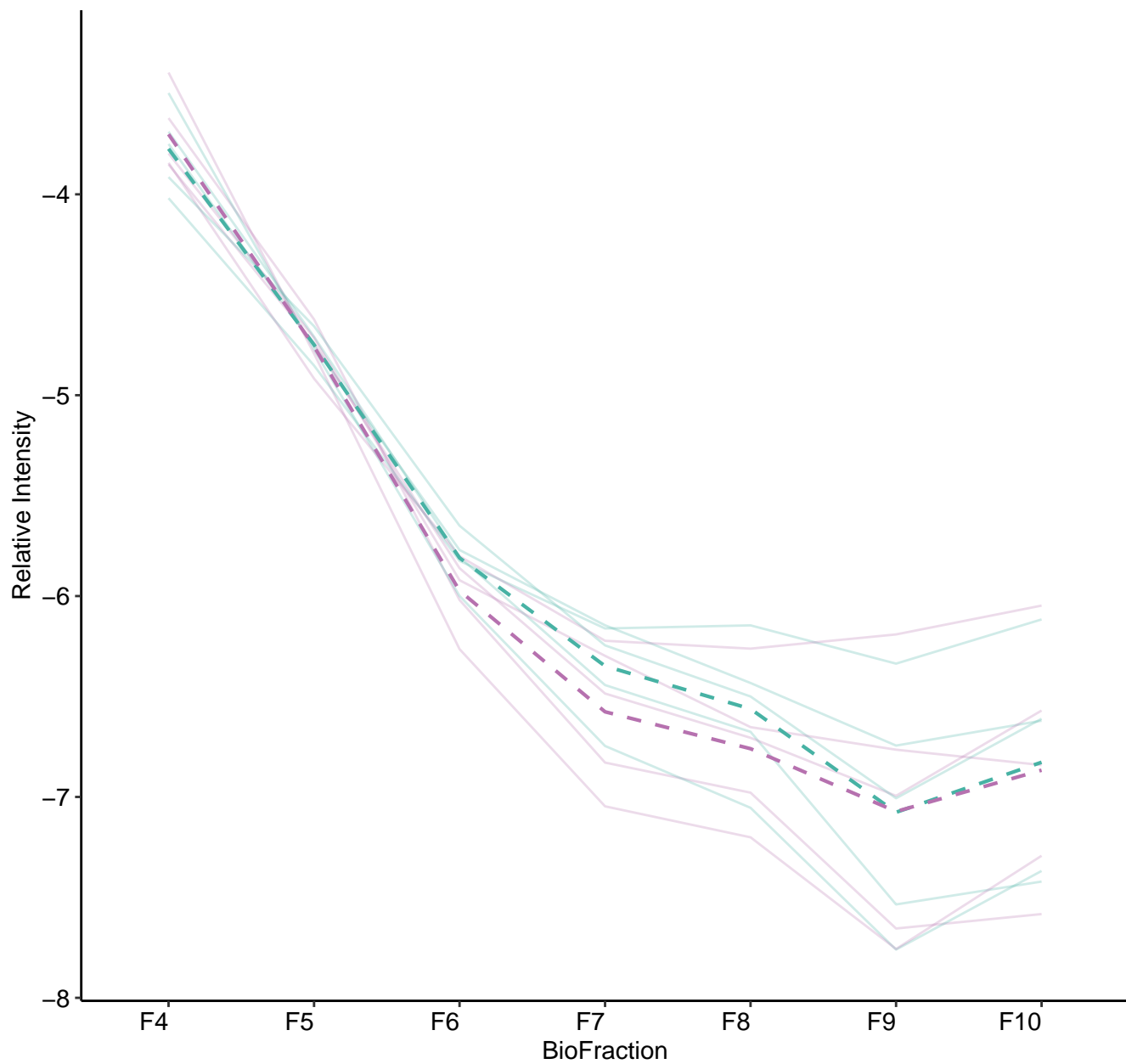
EARP complex (n = 4)
(R2.Fixef = 0.991)



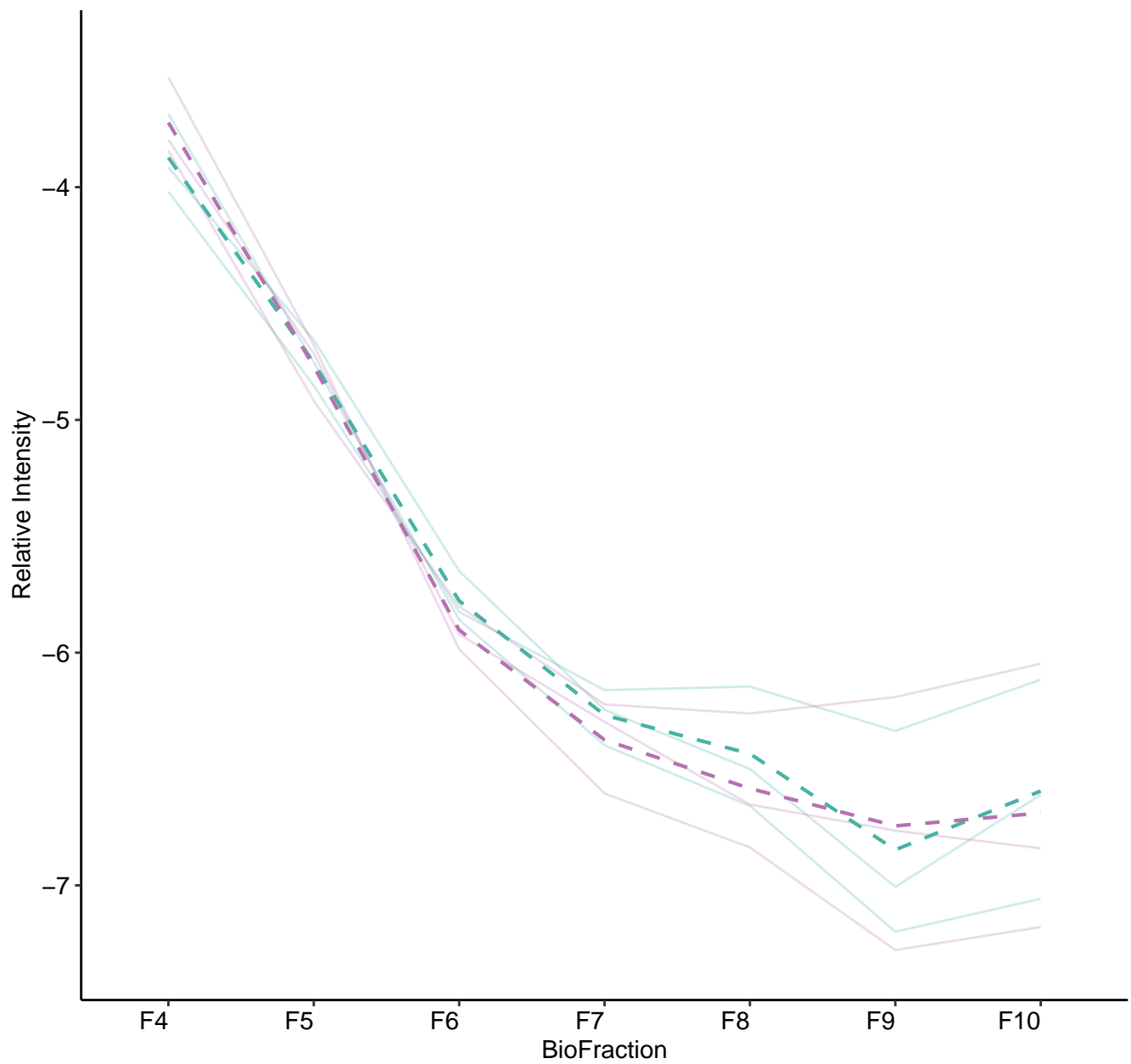
Ecsit complex (ECSIT, MT-CO2, GAPDH, TRAF6, NDUFAF1) (n = 3)
(R2.Fixef = 0.951)



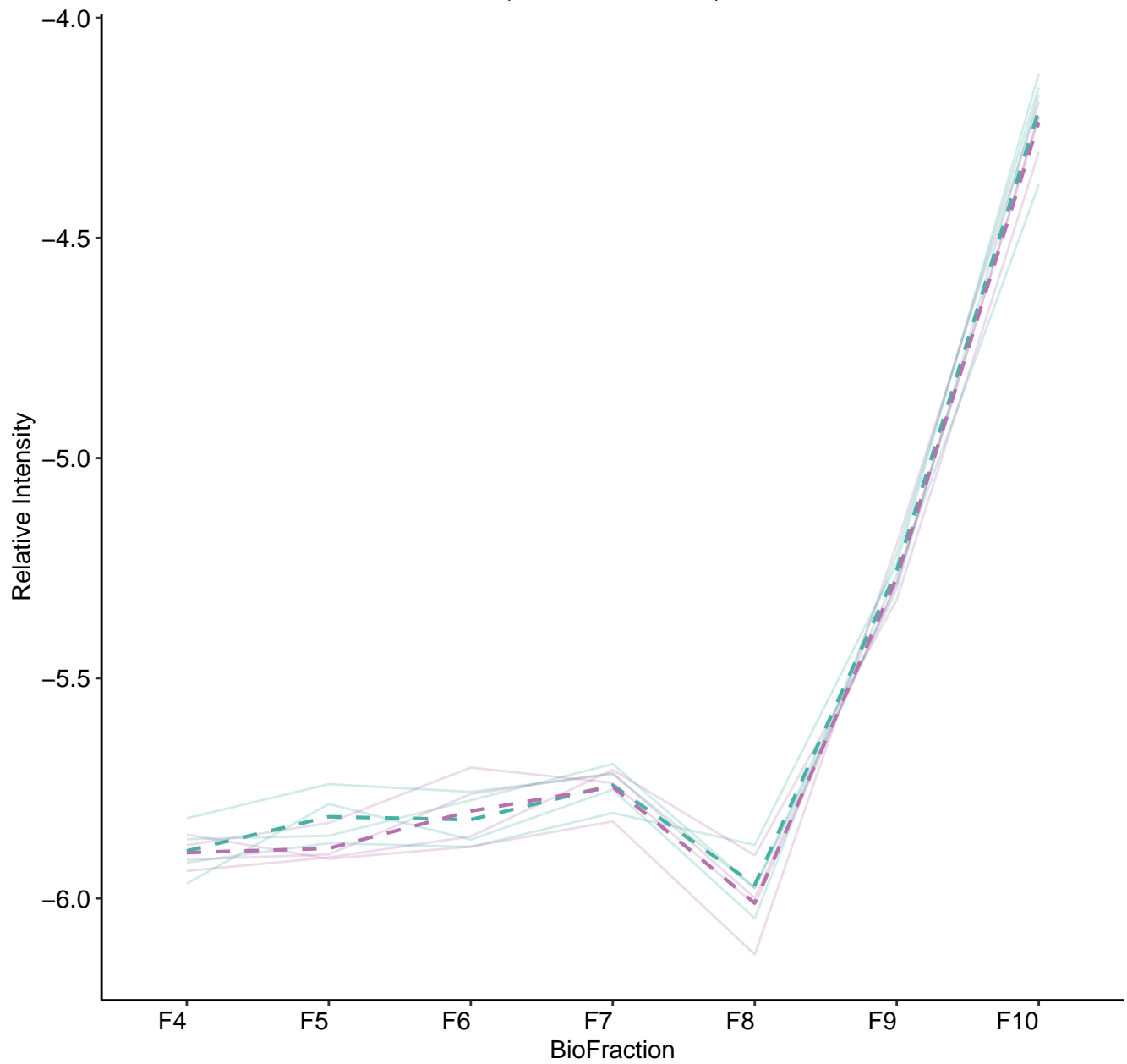
Ecsit complex (ECSIT, MT-CO2, NDUFA1, MT-ND1, TRAF6, NDUFAF1) (n = 5)
(R2.Fixef = 0.903)



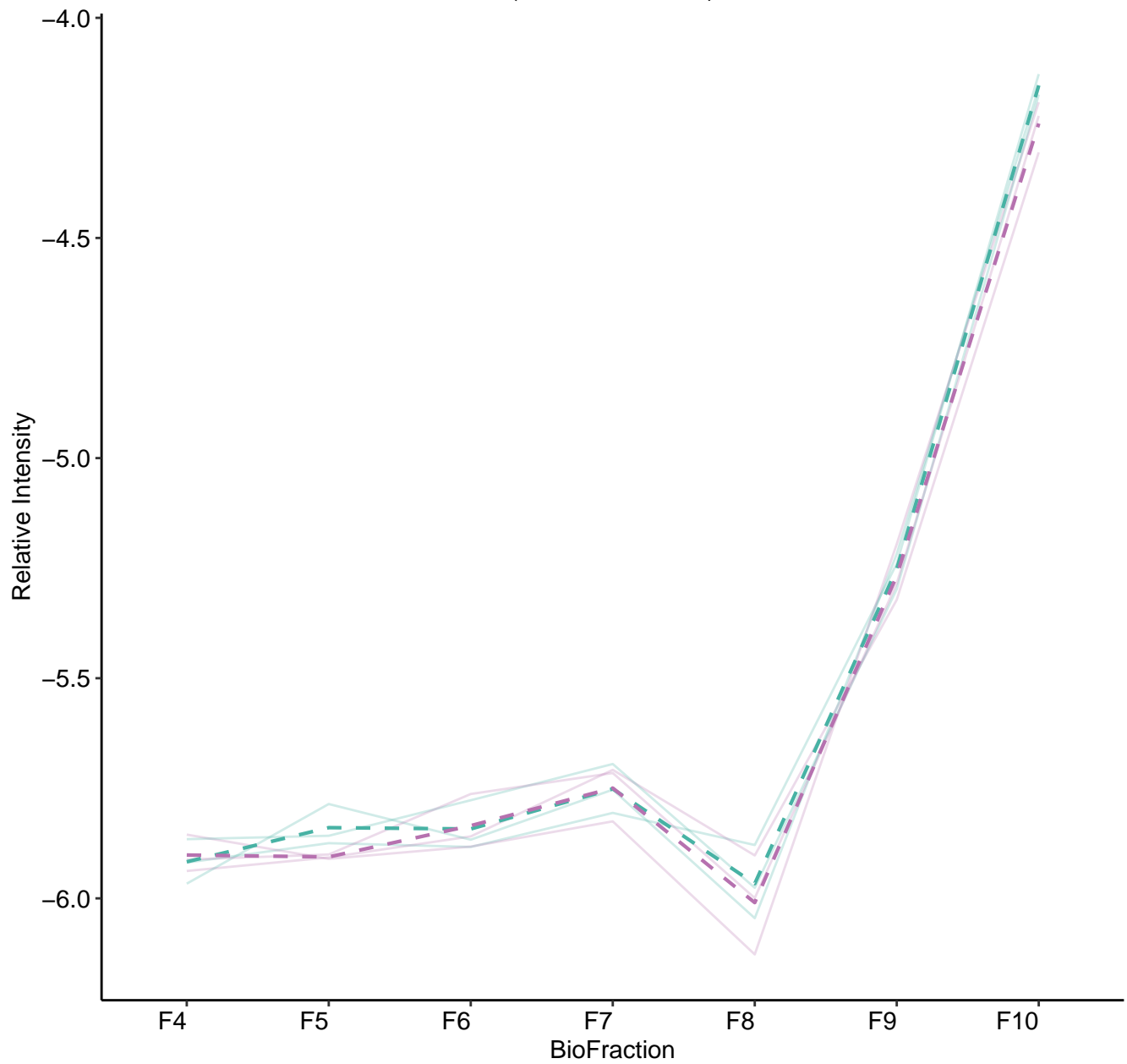
Ecsit complex (ECSIT, NDUFS3, NDUFAF1) (n = 3)
(R2.Fixef = 0.919)



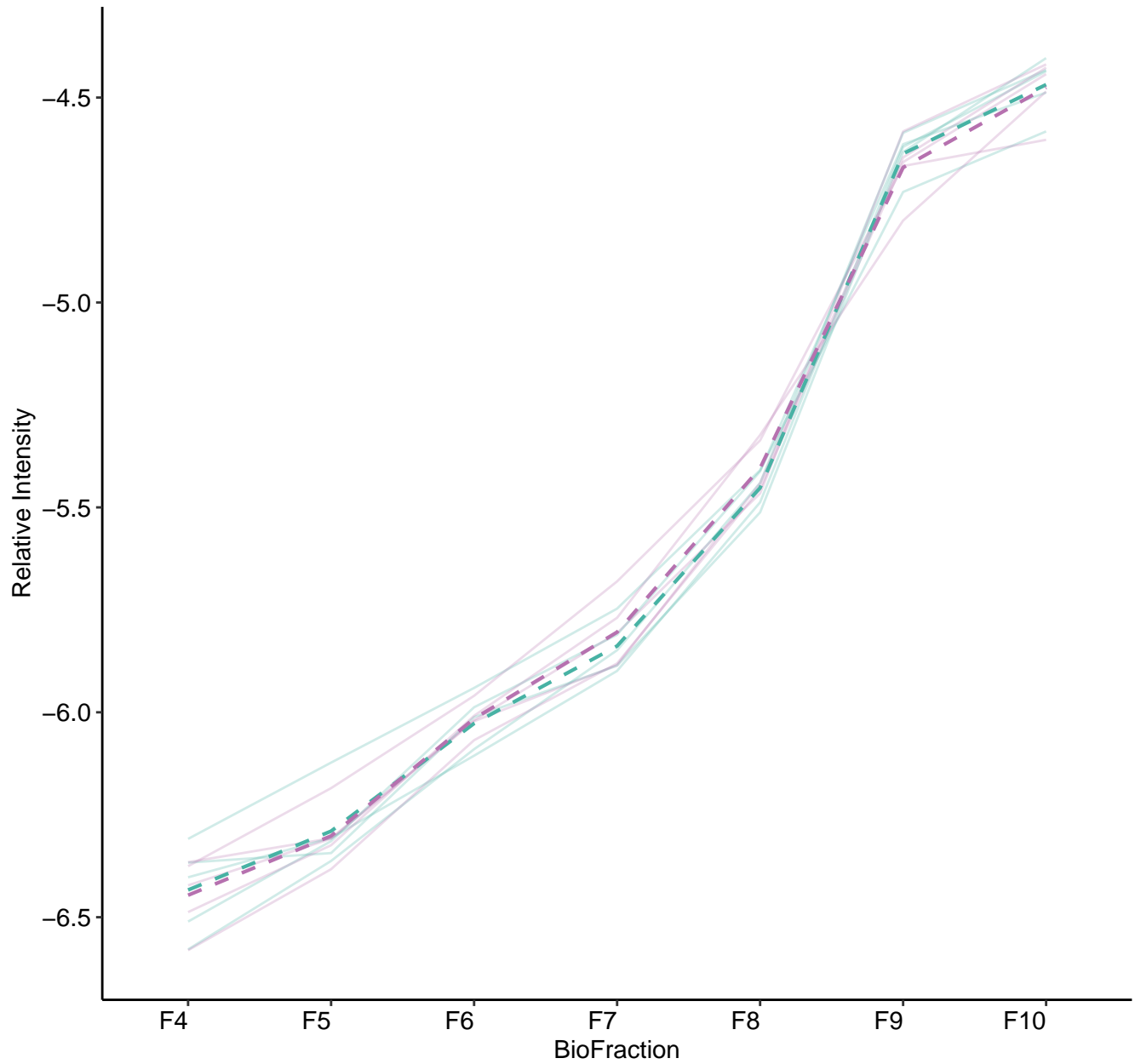
EIF2B1–EIF2B2–EIF2B3–EIF2B4–EIF2B5 complex (n = 4)
(R2.Fixef = 0.988)



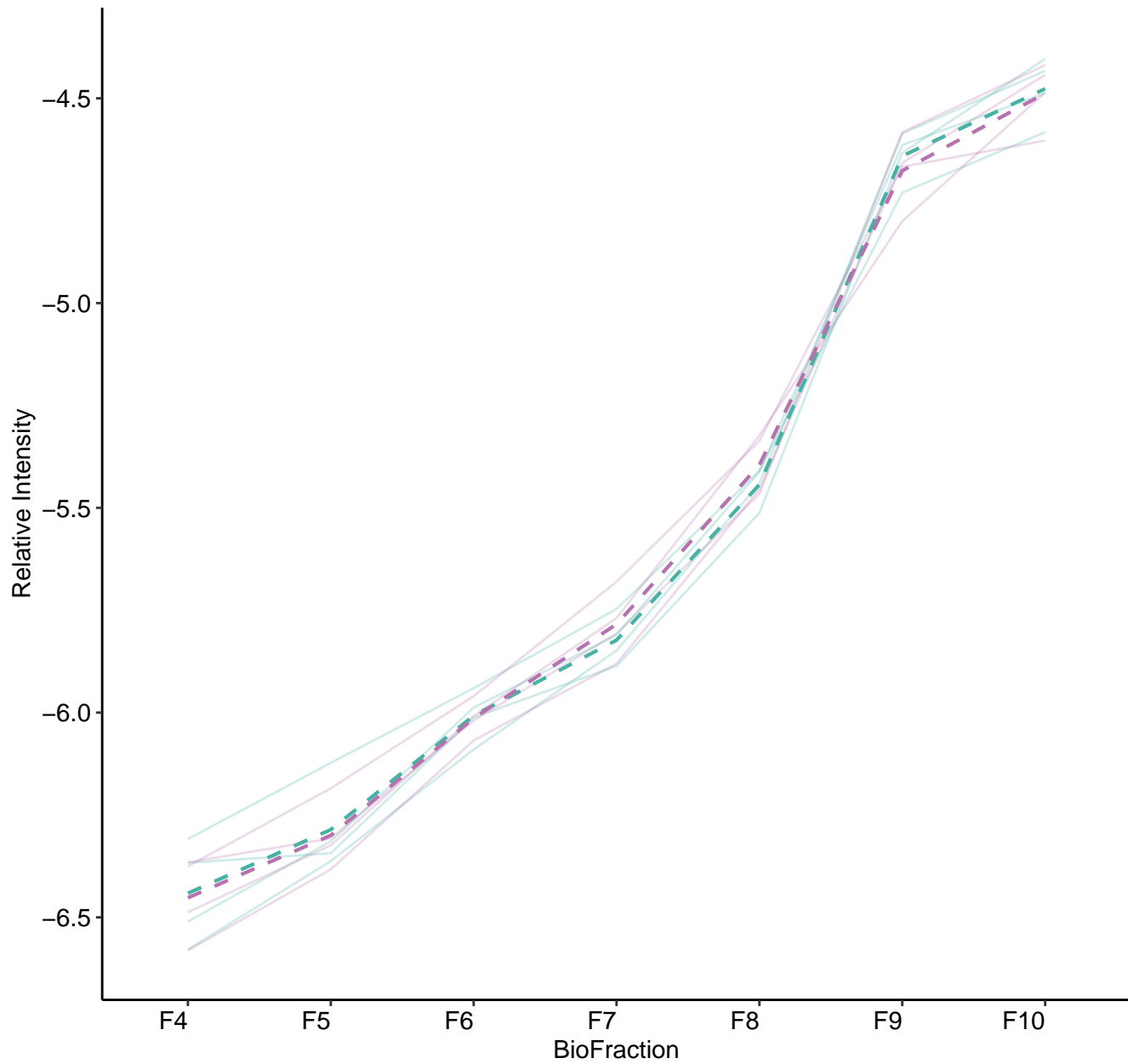
EIF2B2–EIF2B3–EIF2B4–EIF2B5 complex (n = 3)
(R2.Fixef = 0.99)



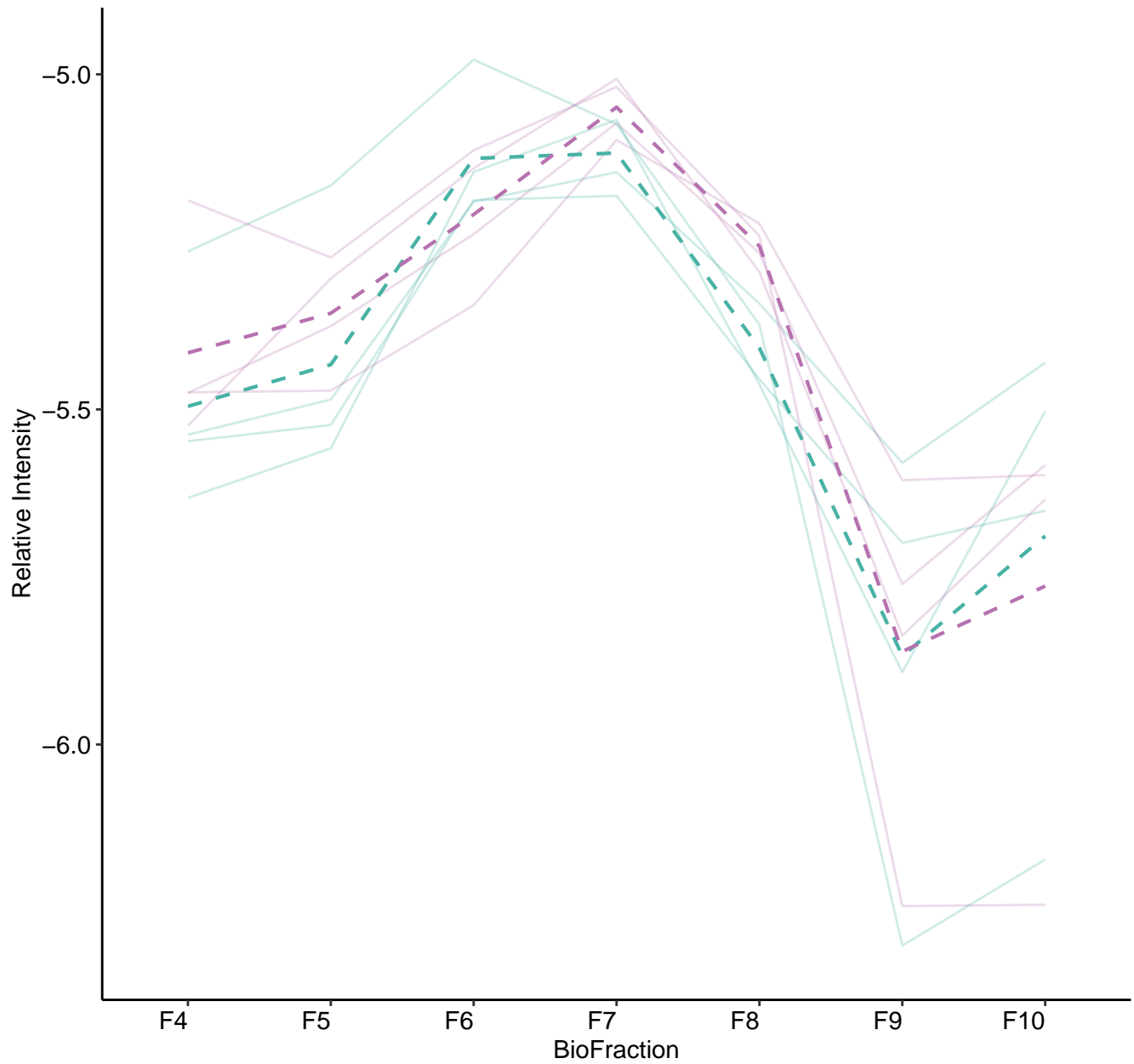
EIF3 complex (EIF3A, EIF3B, EIF3G, EIF3I, EIF3C) (n = 5)
(R2.Fixef = 0.989)



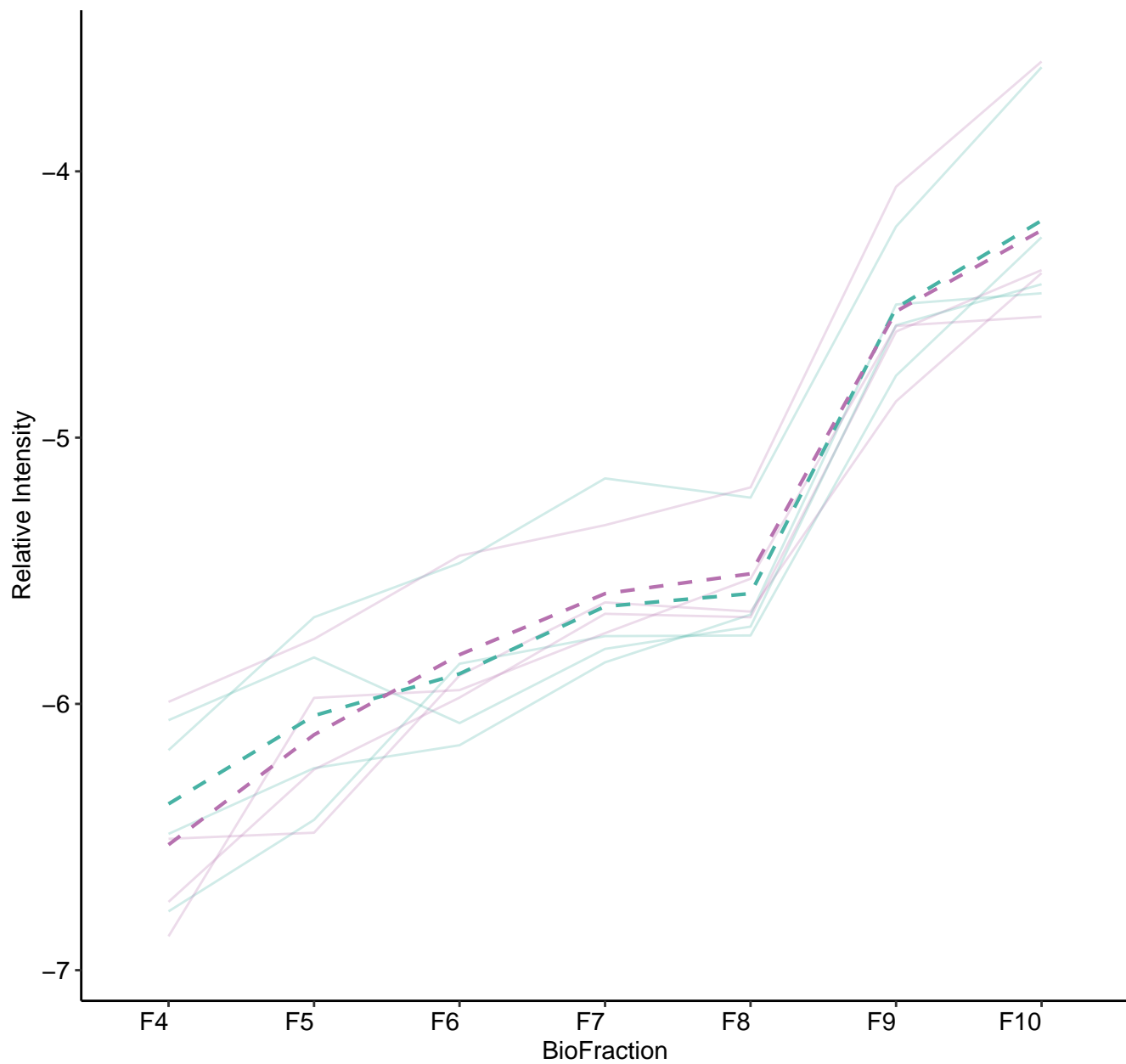
EIF3 complex (EIF3A, EIF3B, EIF3G, EIF3I, EIF3J) (n = 4)
(R2.Fixef = 0.987)



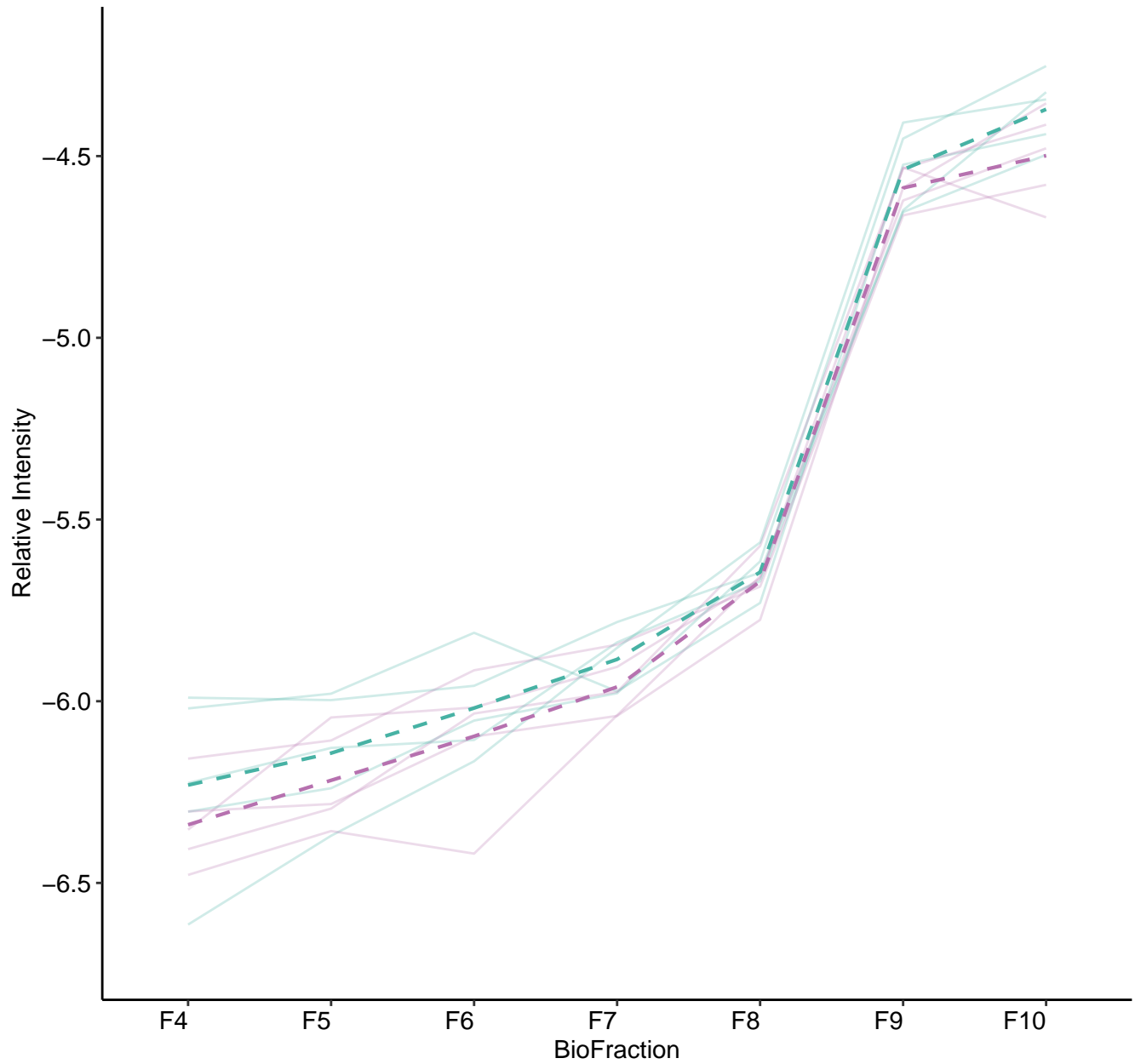
ESCRT-III complex (n = 4)
(R2.Fixef = 0.664)



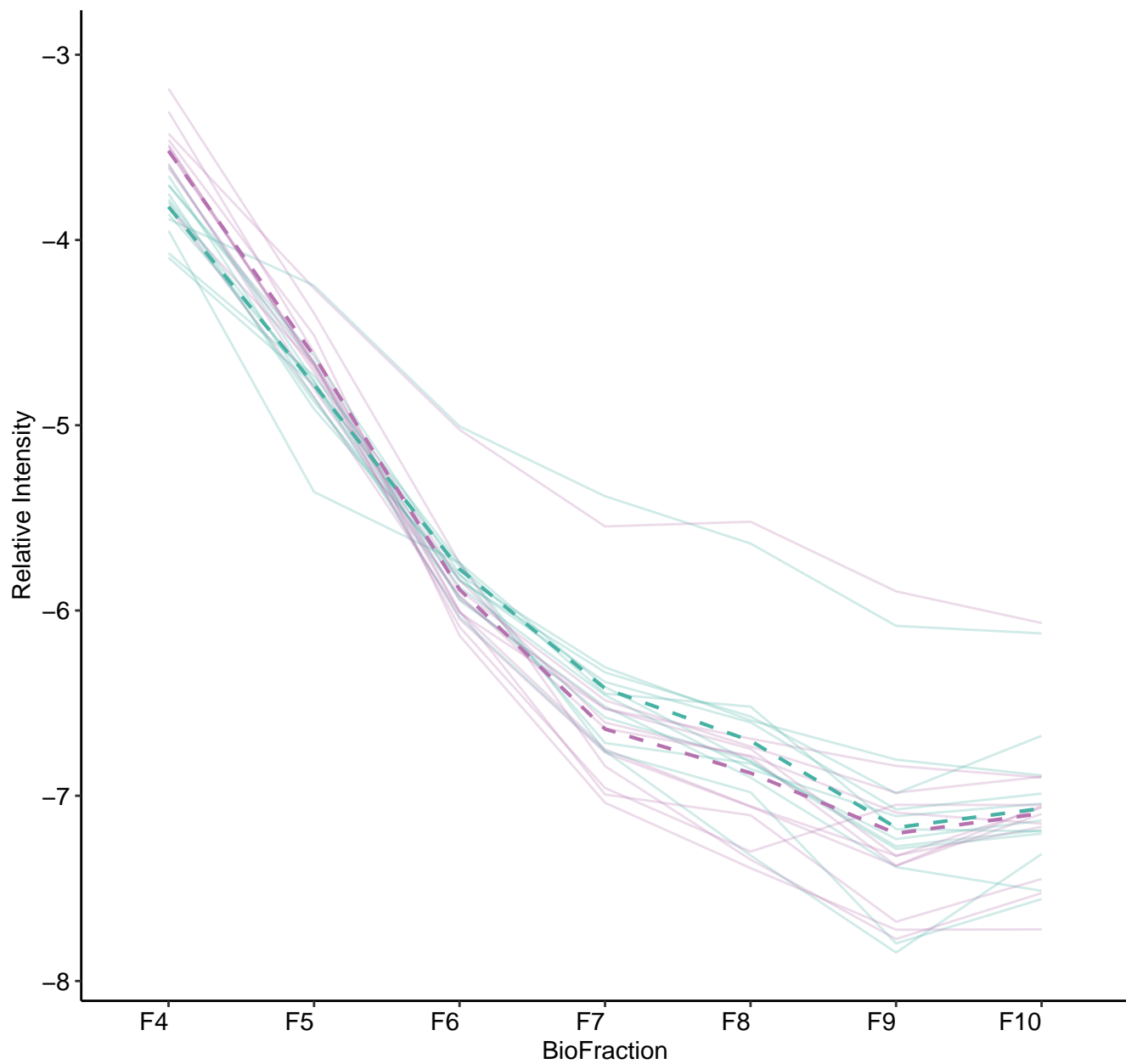
ESR1-CDK7-CCNH-MNAT1-MTA1-HDAC2 complex (n = 4)
(R2.Fixef = 0.855)



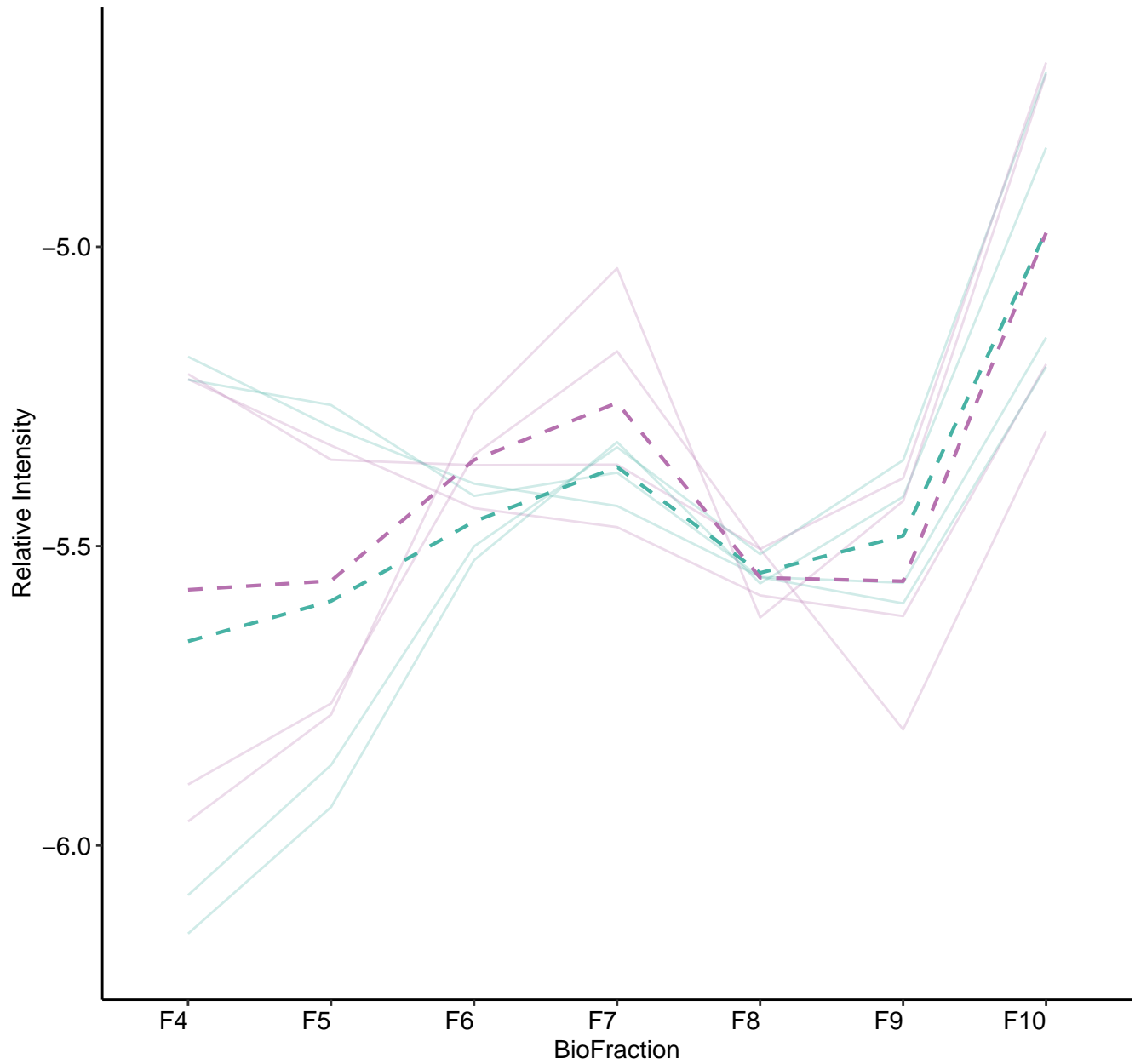
Exosome (n = 5)
(R2.Fixef = 0.967)



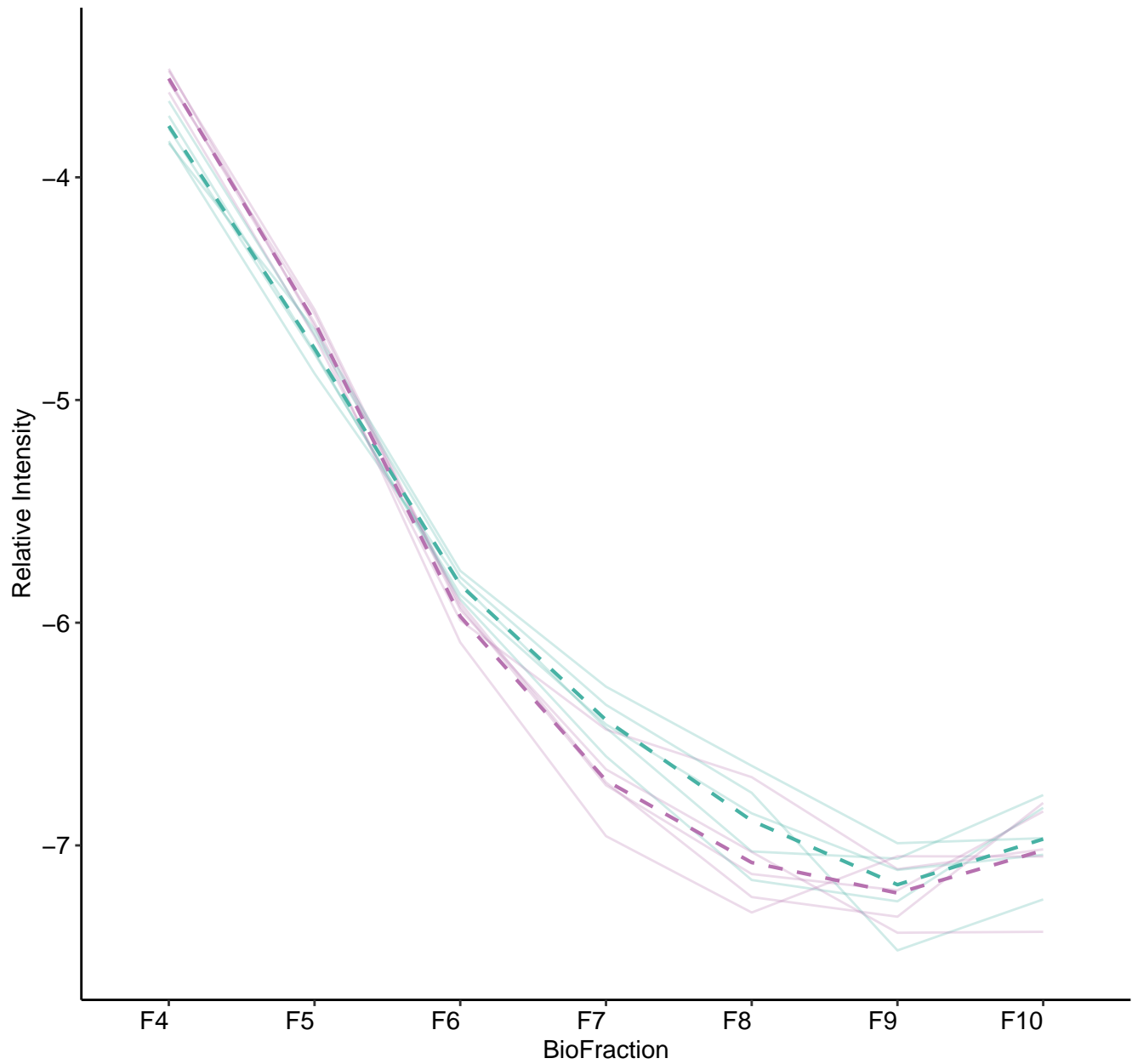
F0F1 ATP synthase, mitochondrial (n = 12)
(R2.Fixef = 0.924)



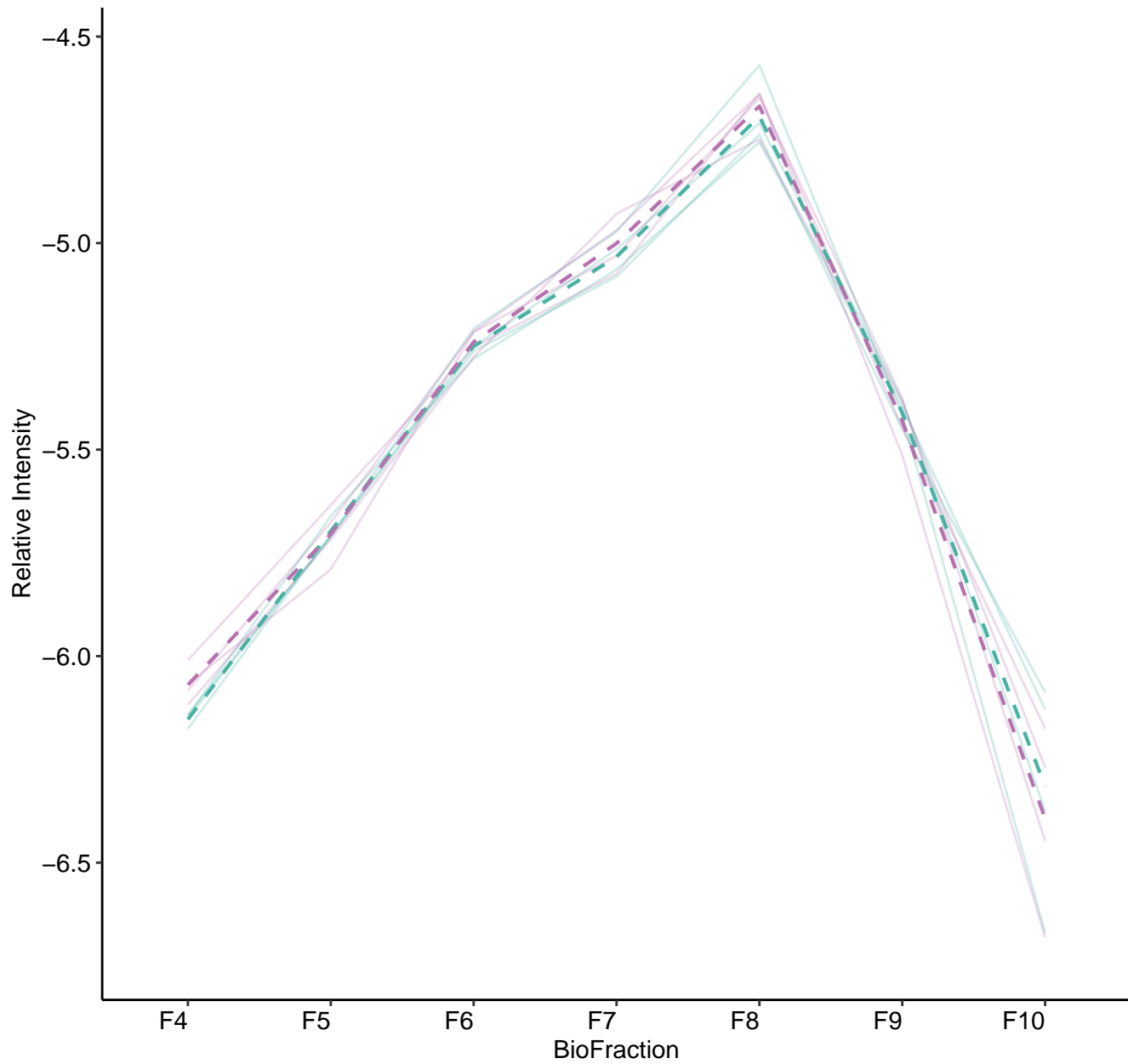
FACT complex, UV-activated (n = 4)
(R2.Fixef = 0.41)



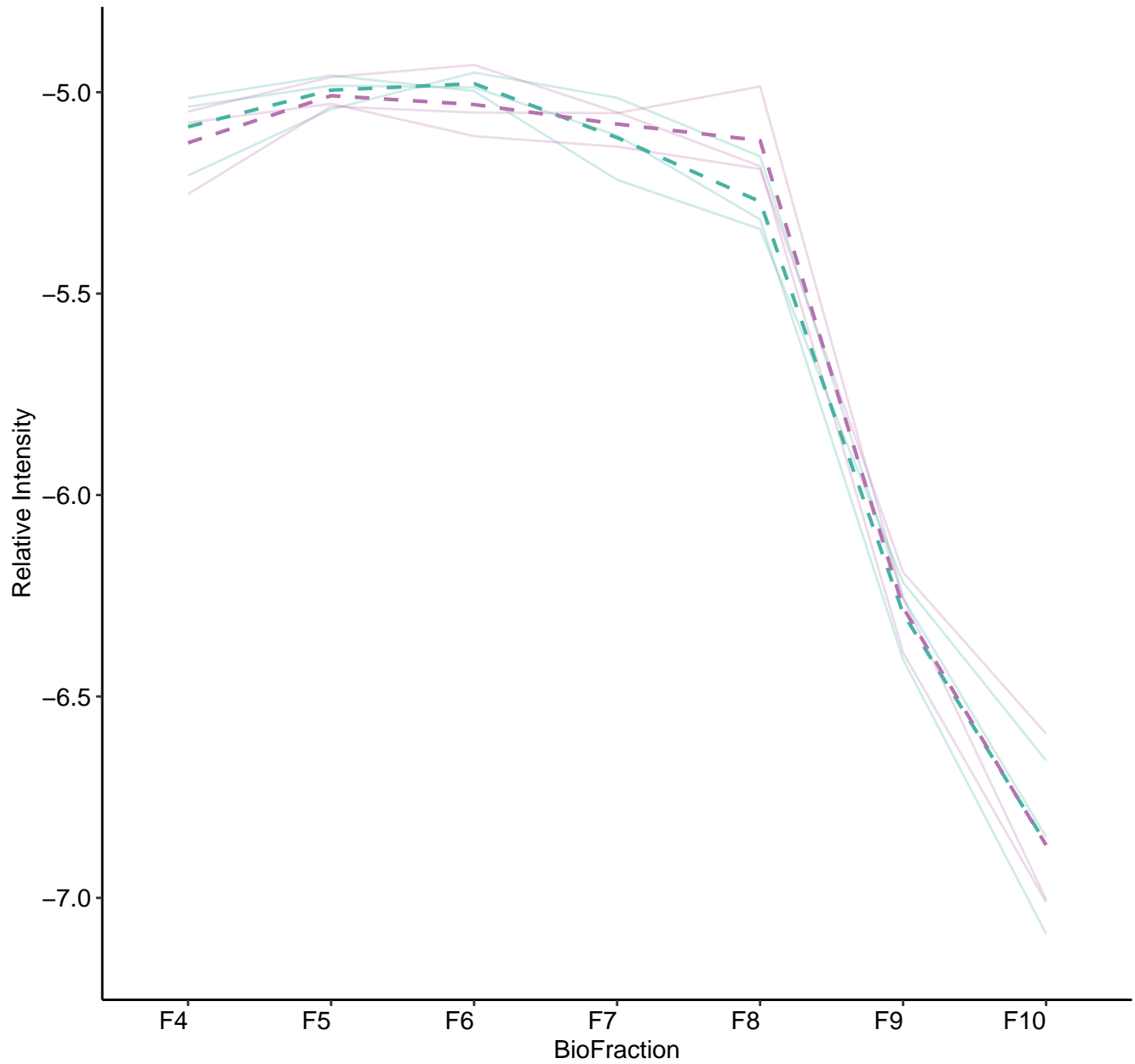
Frataxin complex (n = 5)
(R2.Fixef = 0.986)



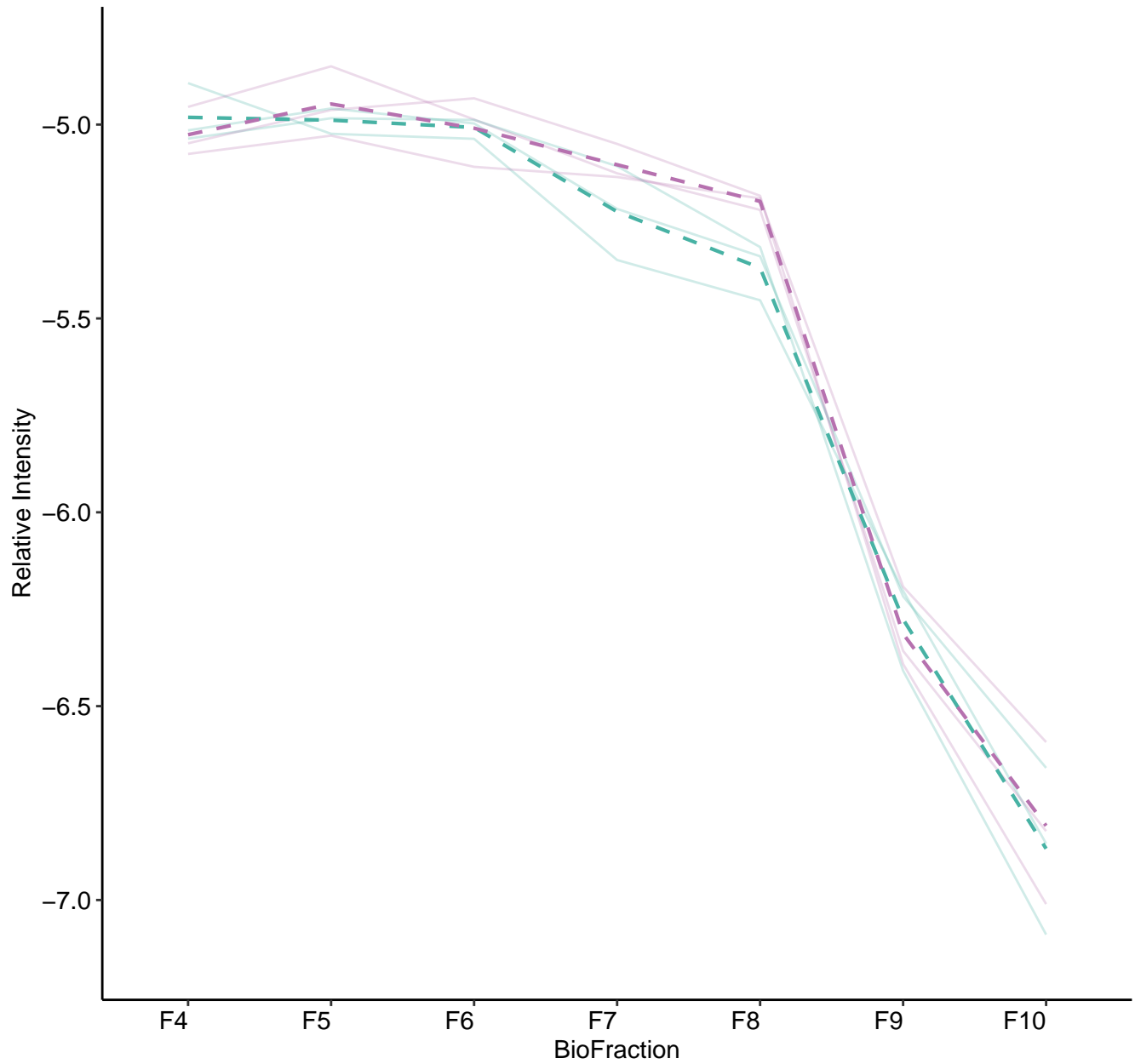
GAA1-GPI8-PIGT-PIG-PIGS complex (n = 4)
(R2.Fixef = 0.966)



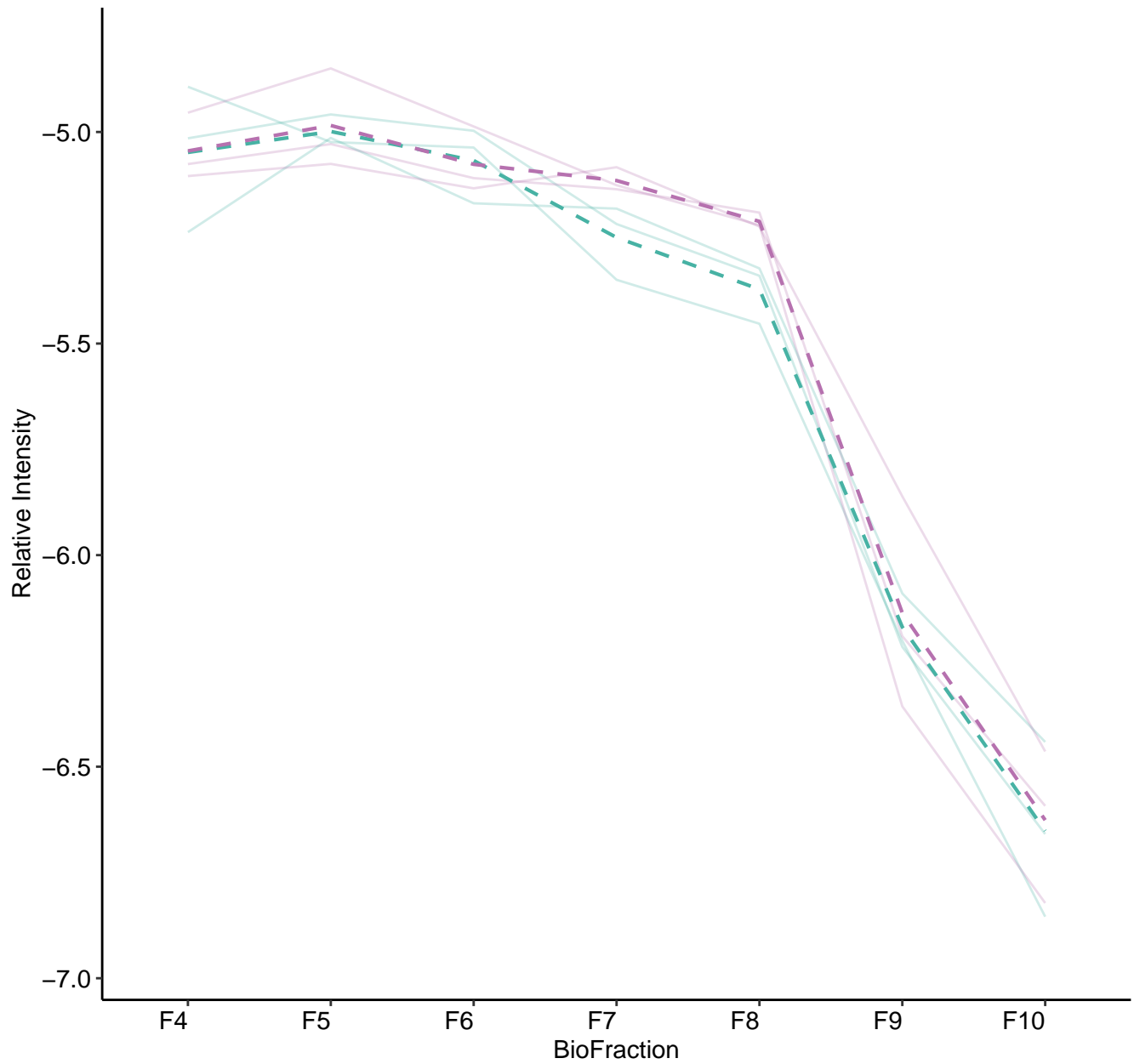
GABA-A receptor (GABRA1, GABRB2, GABRD) (n = 3)
(R2.Fixef = 0.973)



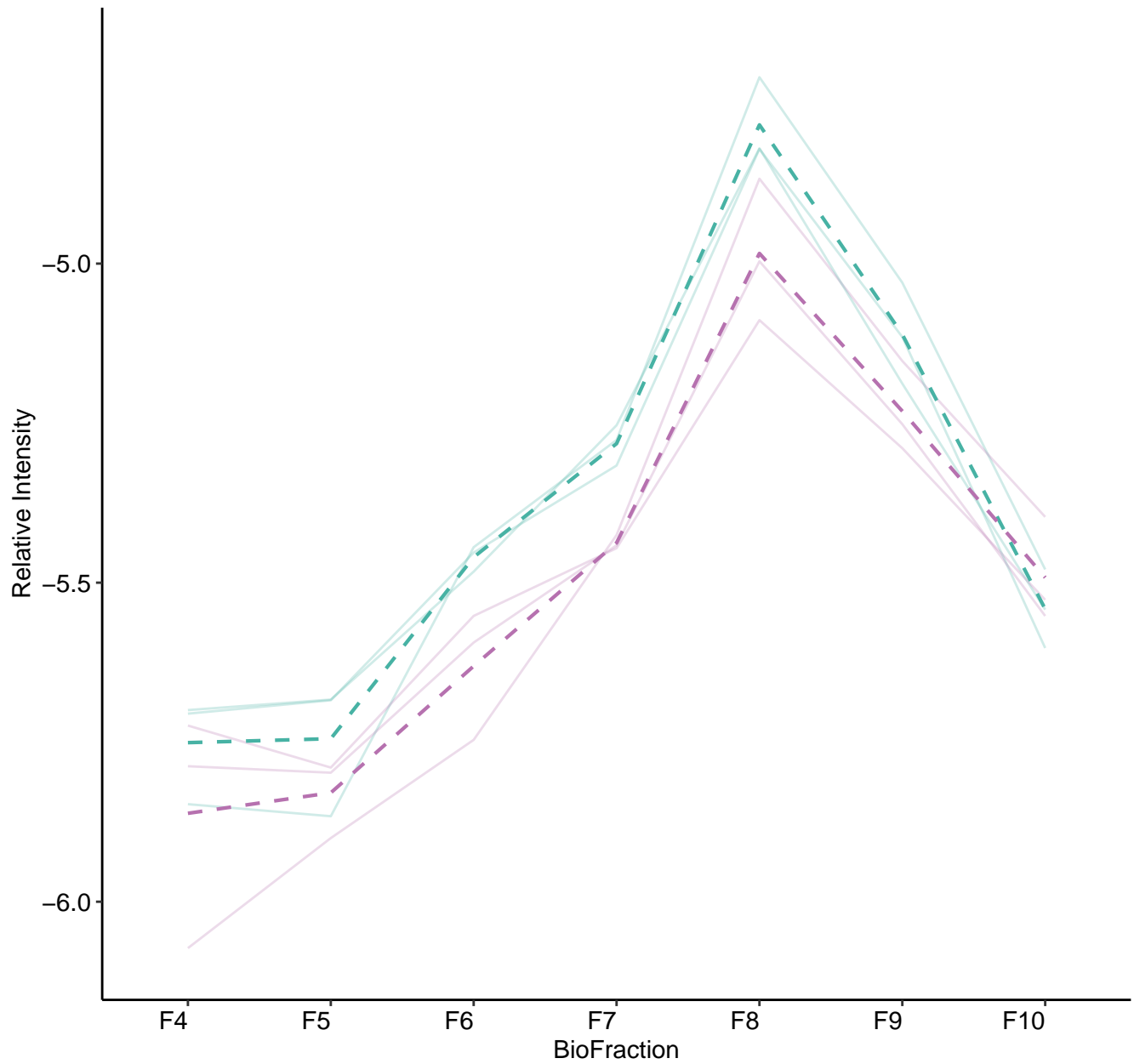
GABA-A receptor (GABRA1, GABRB2, GABRG2) (n = 3)
(R2.Fixef = 0.977)



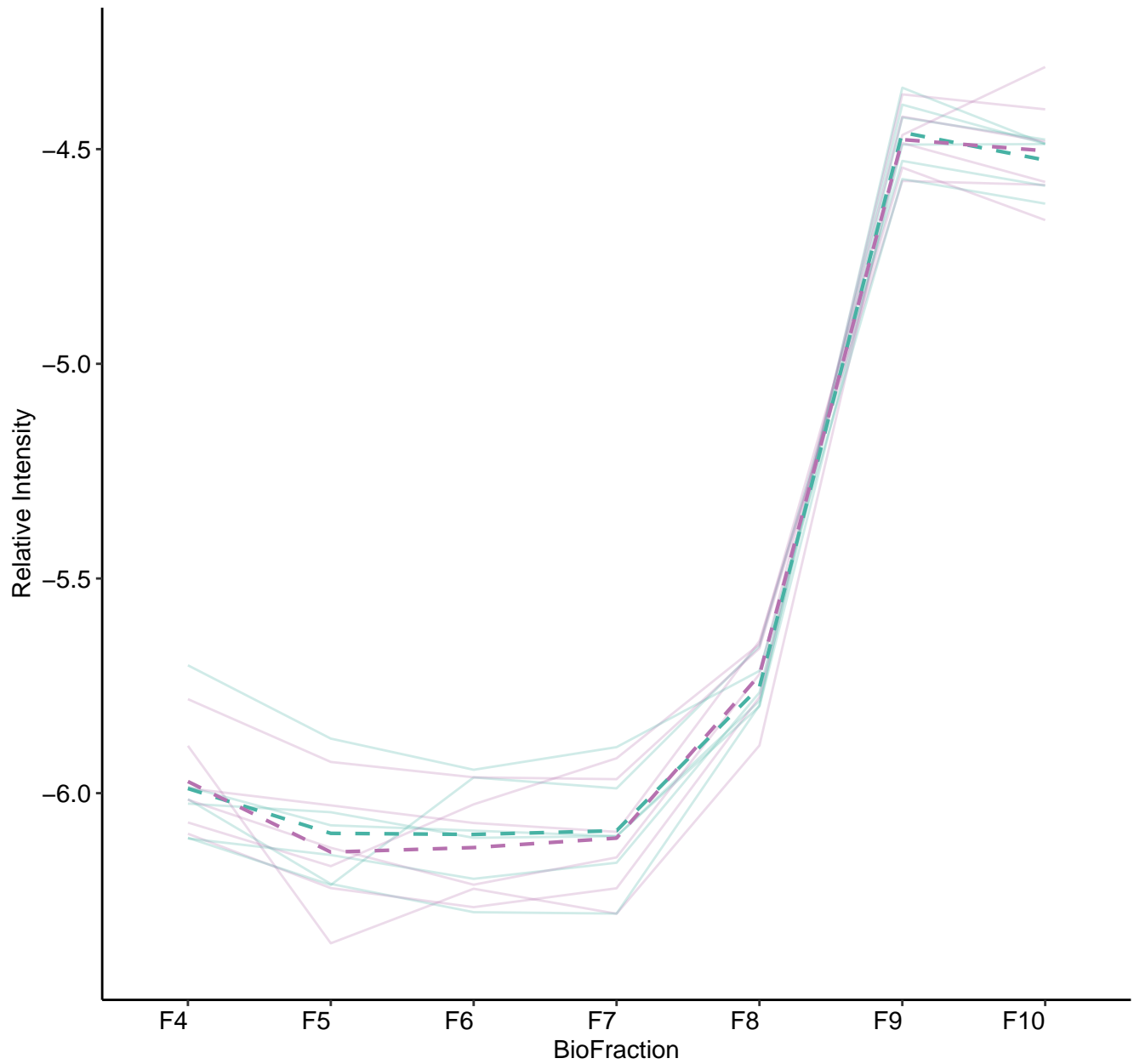
GABA-A receptor (Gabra5, Gabrb2, Gabrg2) (n = 3)
(R2.Fixef = 0.958)



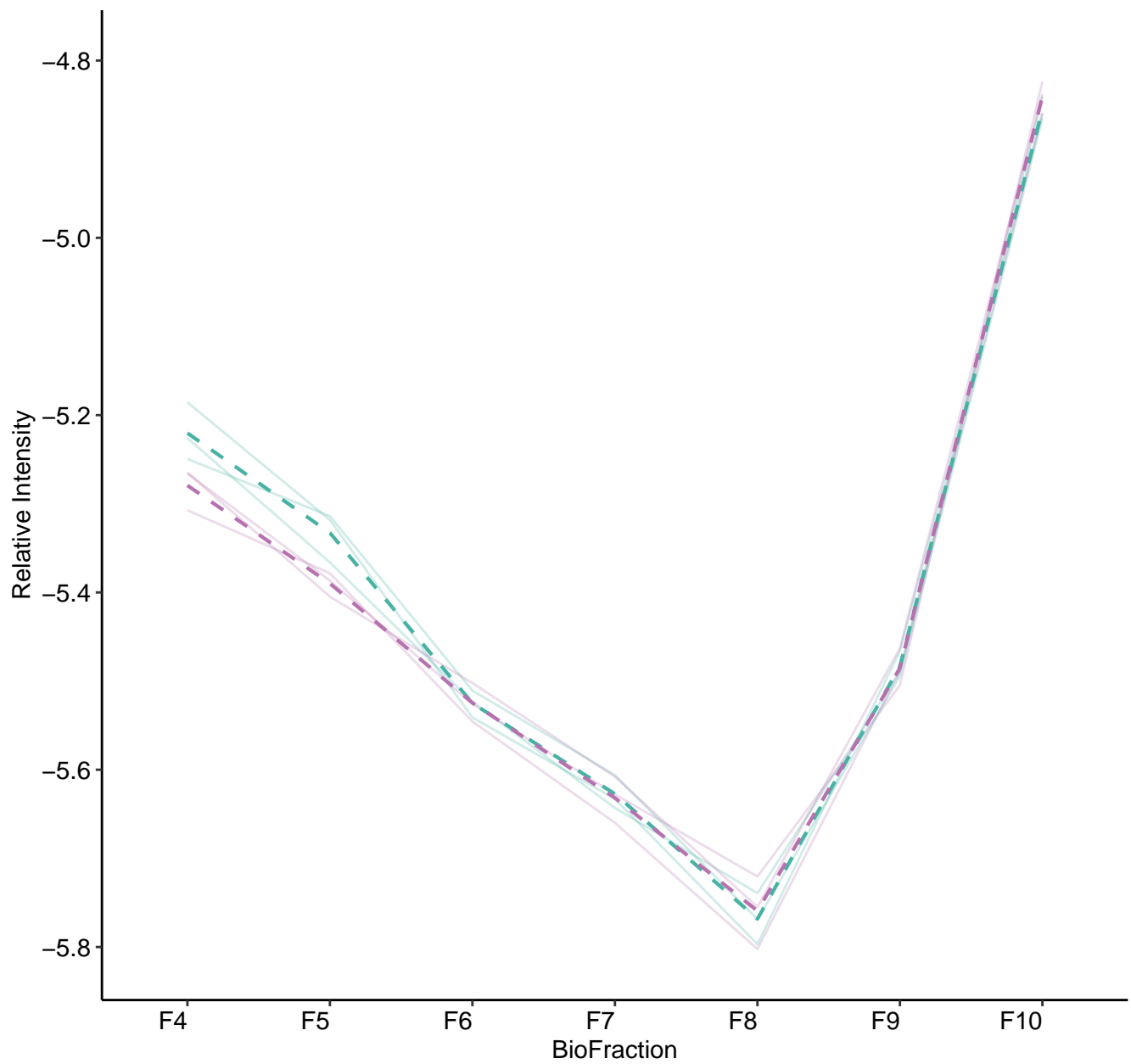
Gamma-BAR-AP1 complex (n = 3)
(R2.Fixef = 0.93)



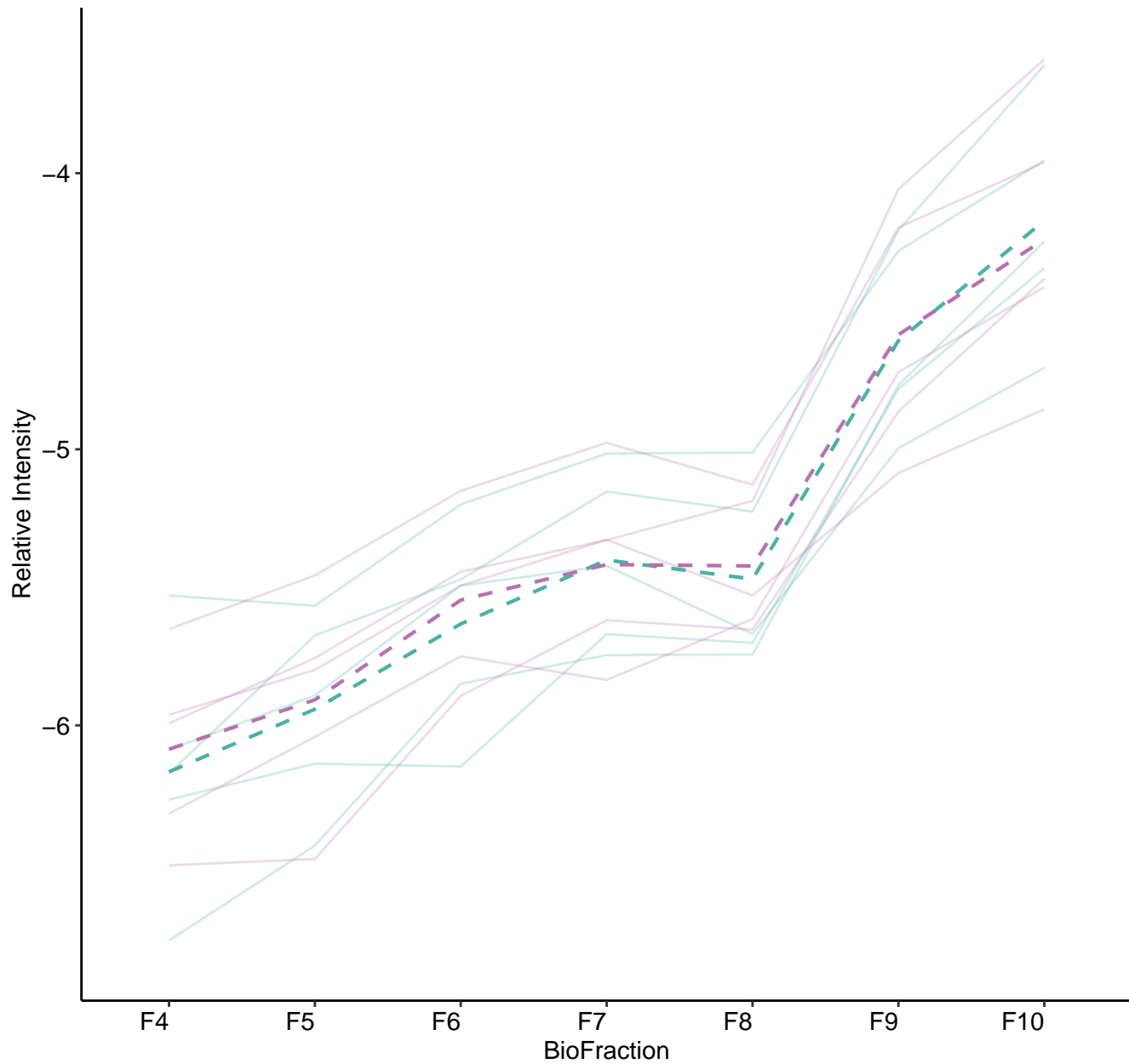
Gamma-tubulin complex (n = 6)
(R2.Fixef = 0.973)



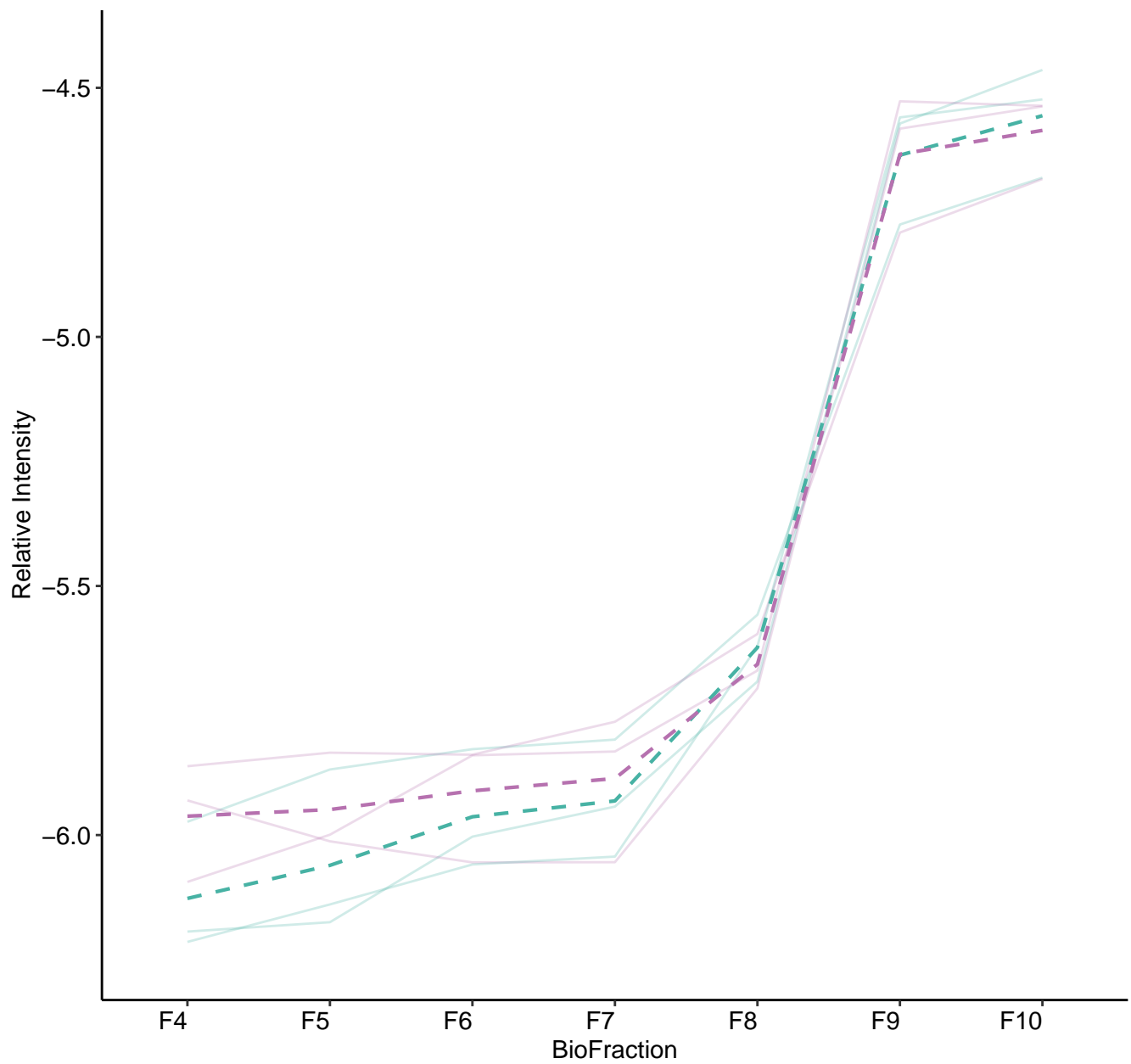
GARP complex (n = 3)
(R2.Fixef = 0.993)



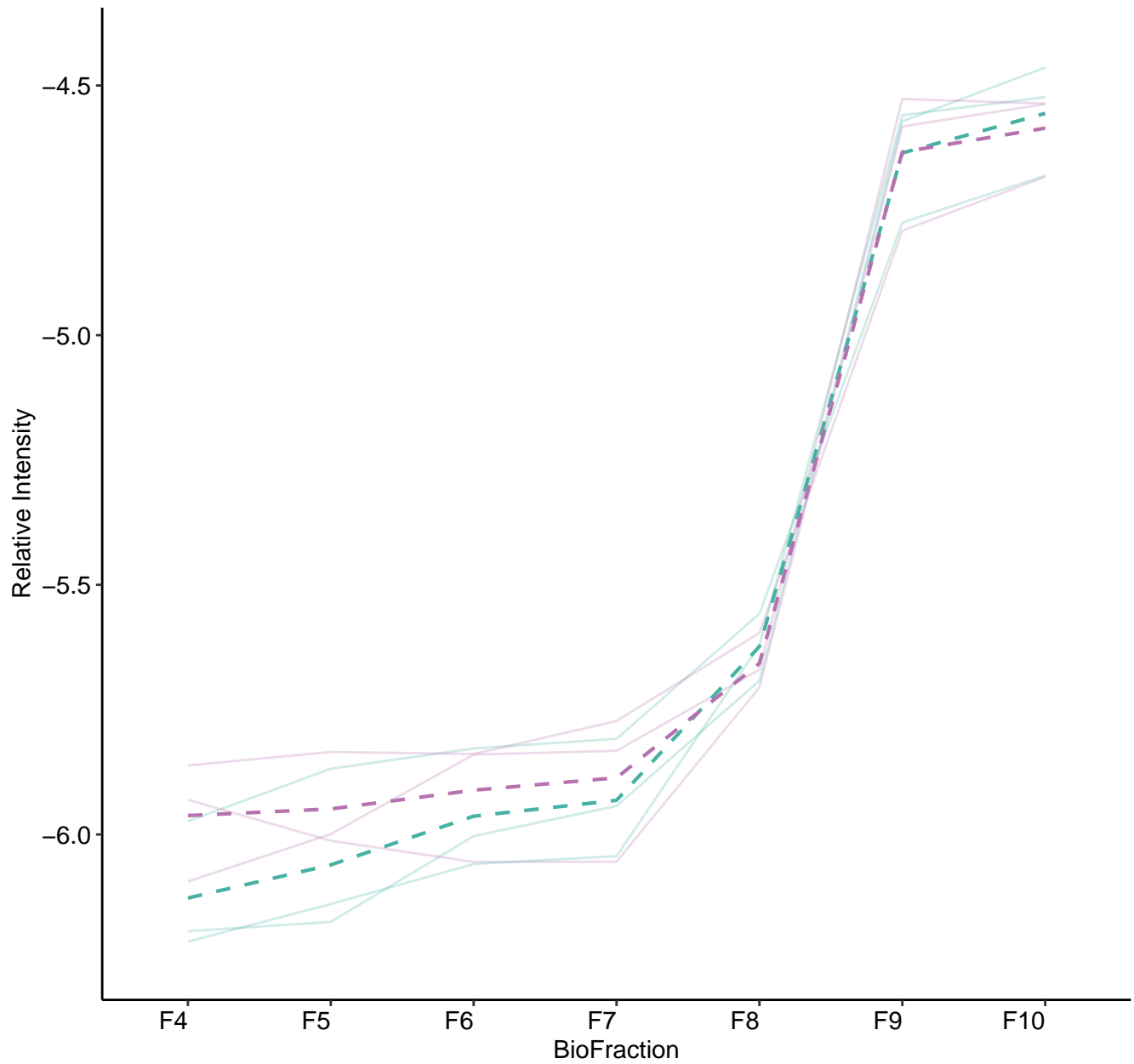
Gata1–Fog1–MeCP1 complex (n = 5)
(R2.Fixef = 0.756)



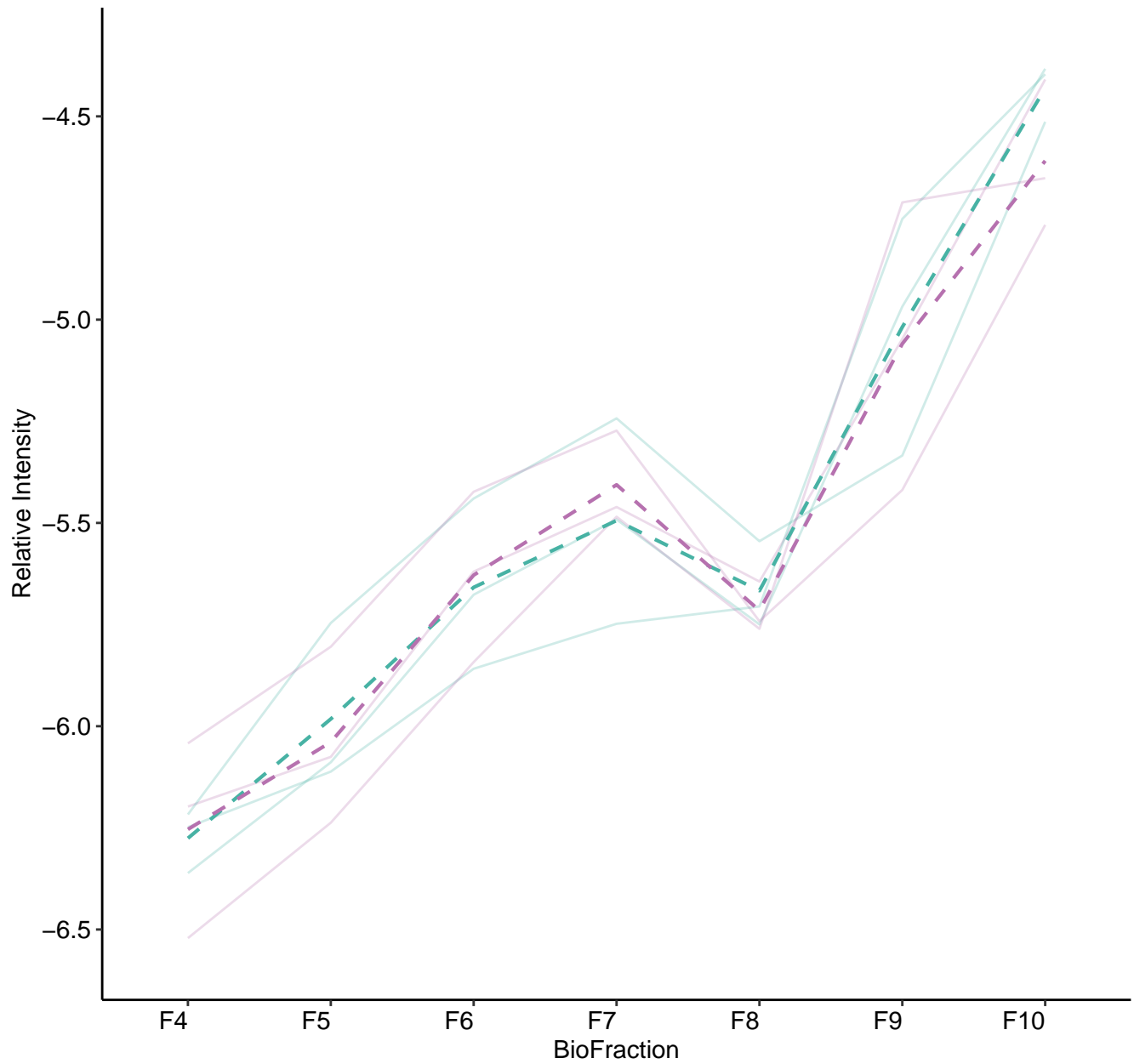
GATOR complex (n = 3)
(R2.Fixef = 0.964)



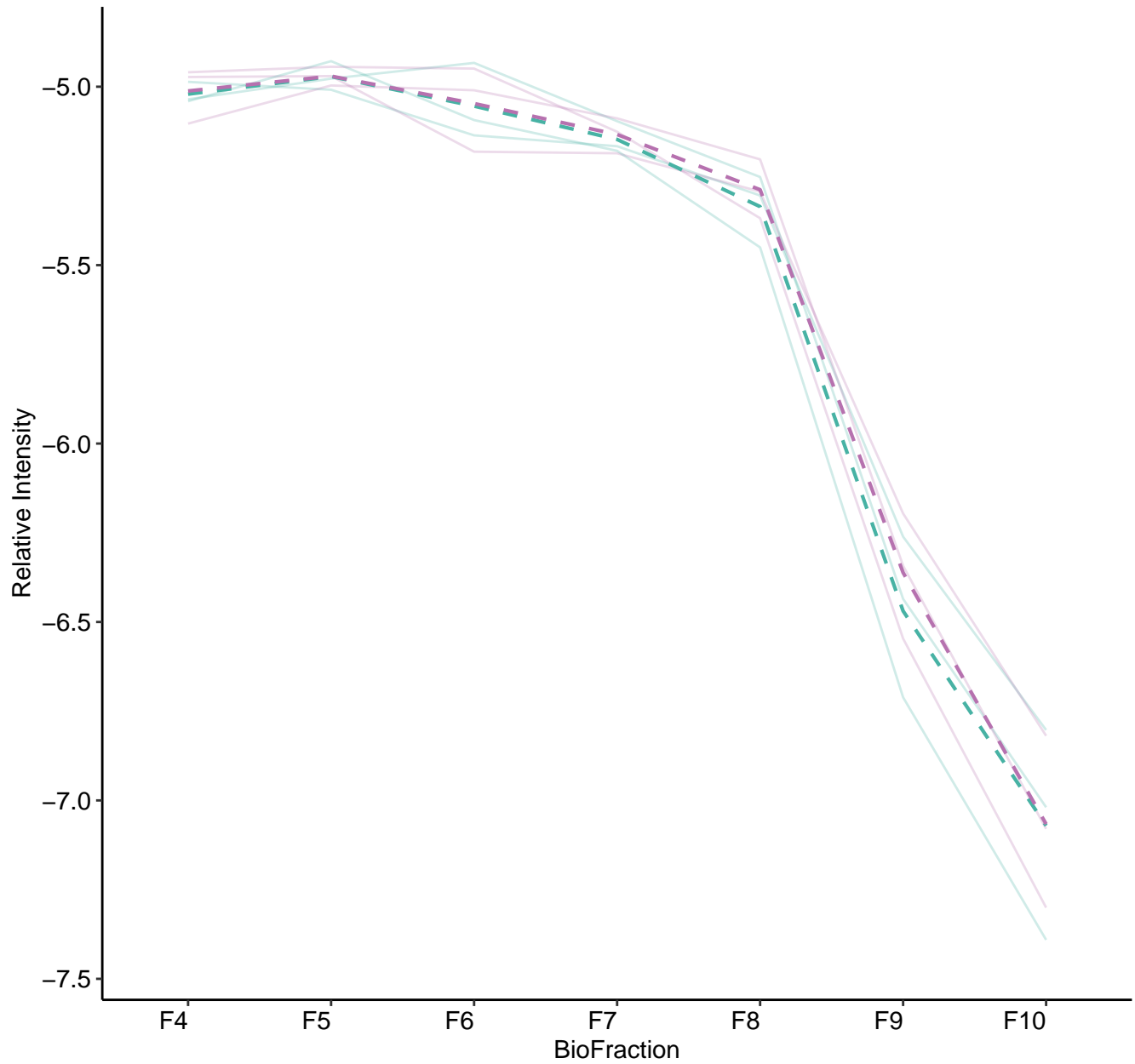
GATOR2 complex (n = 3)
(R2.Fixef = 0.964)



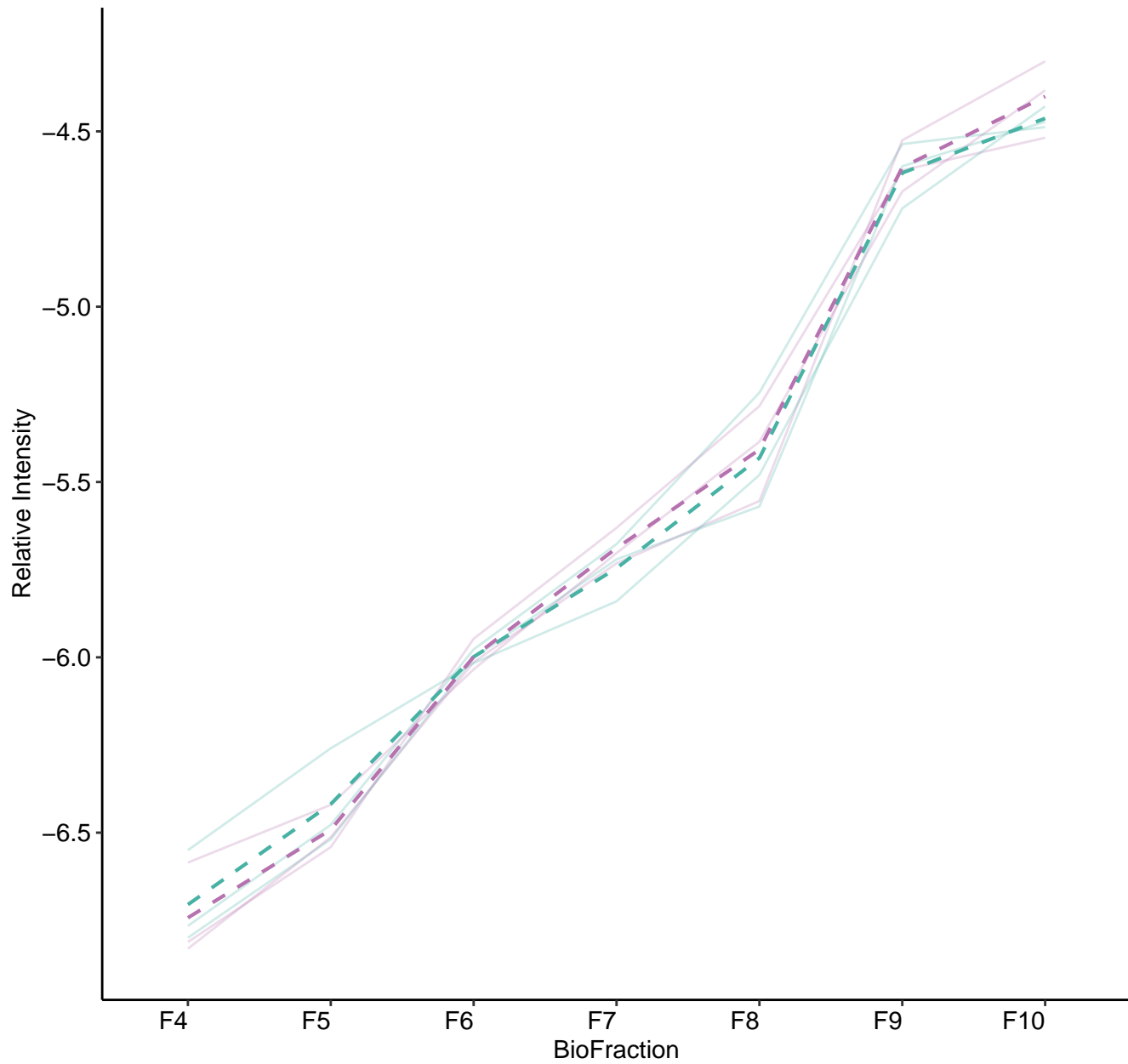
GCN5–TRRAP histone acetyltransferase complex (n = 3)
(R2.Fixef = 0.879)



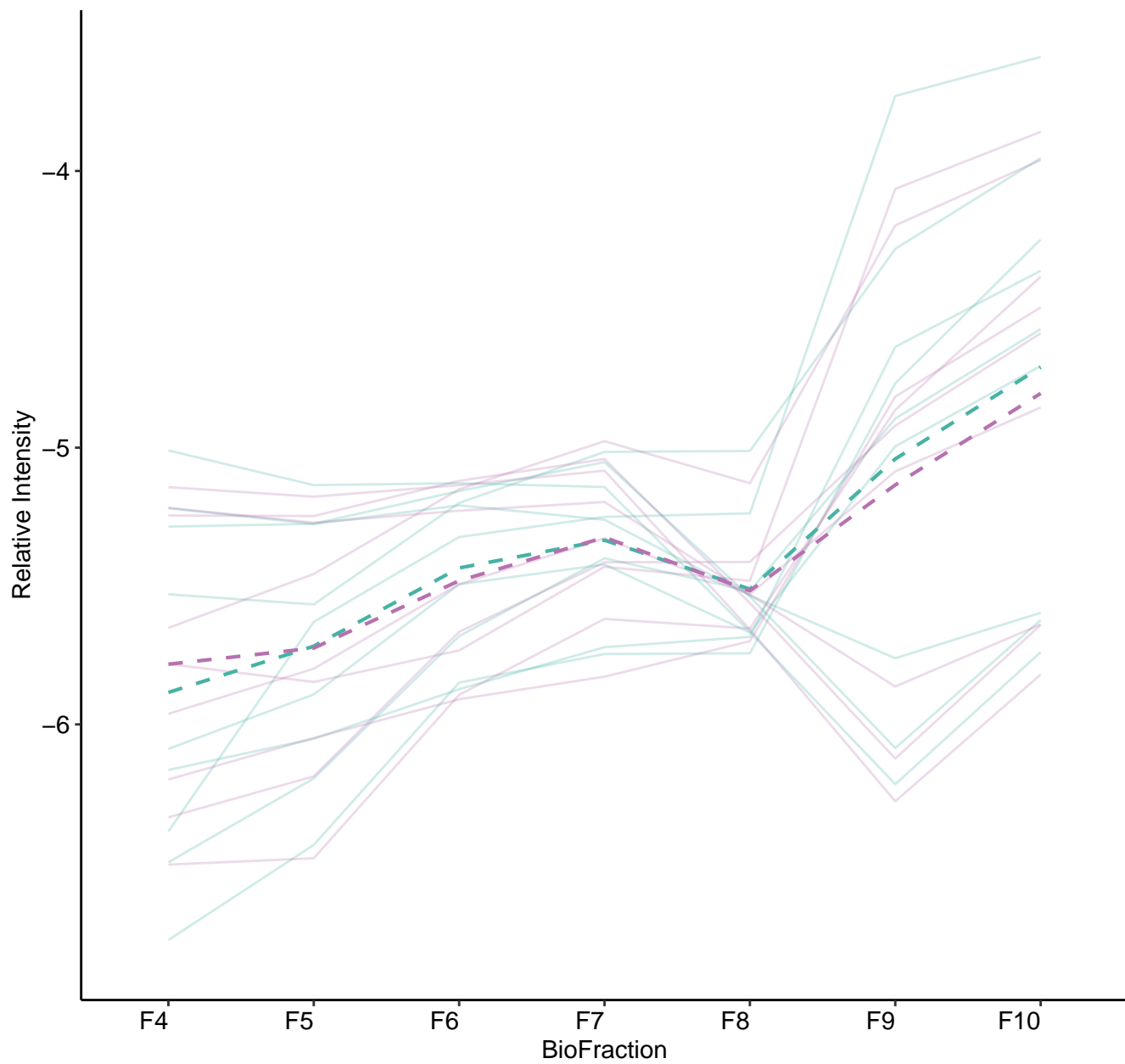
GNAS-L-GNB2-GNG12 complex (n = 3)
(R2.Fixef = 0.967)



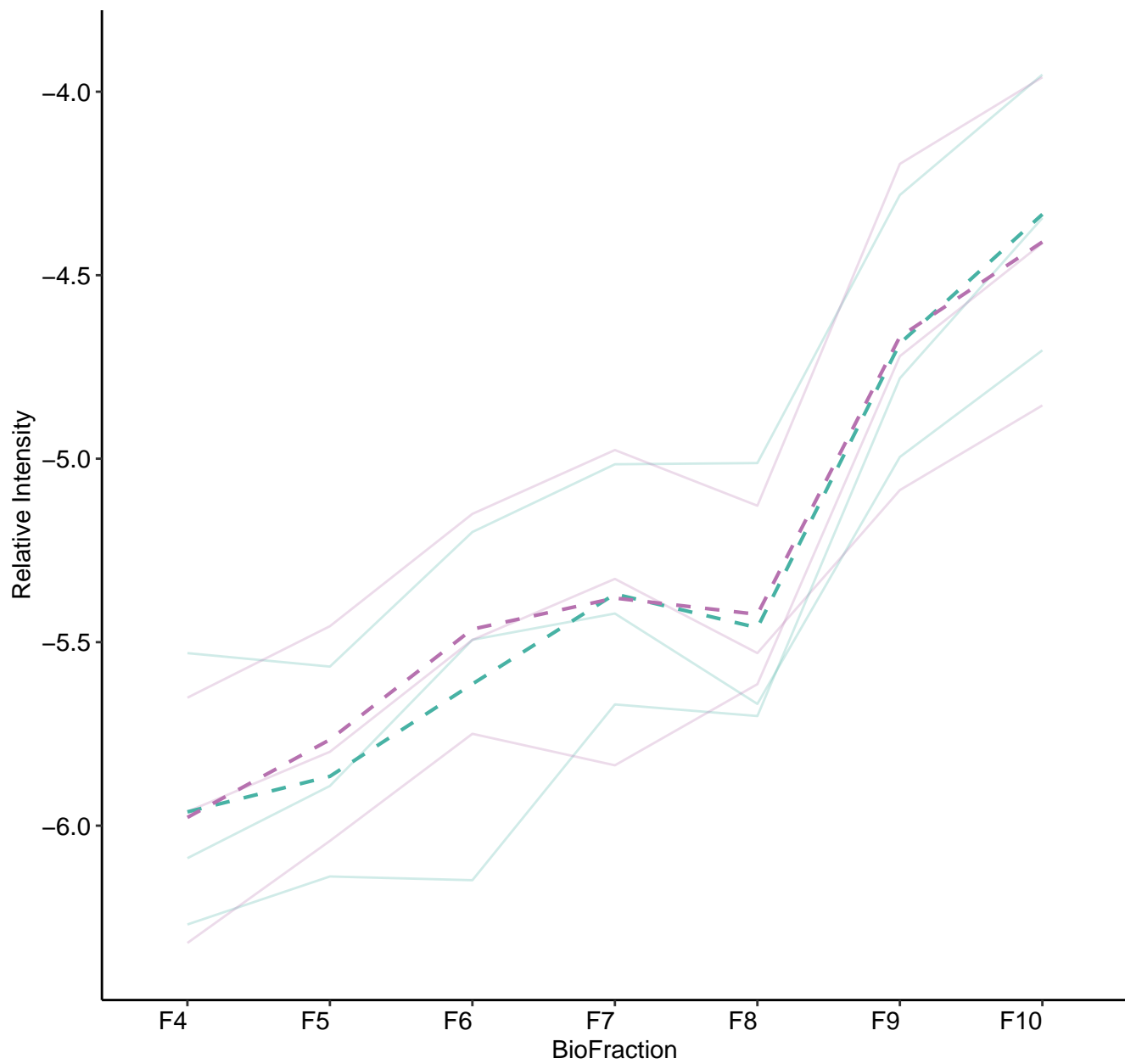
H2AX complex, isolated from cells without IR exposure (n = 3)
(R2.Fixef = 0.984)



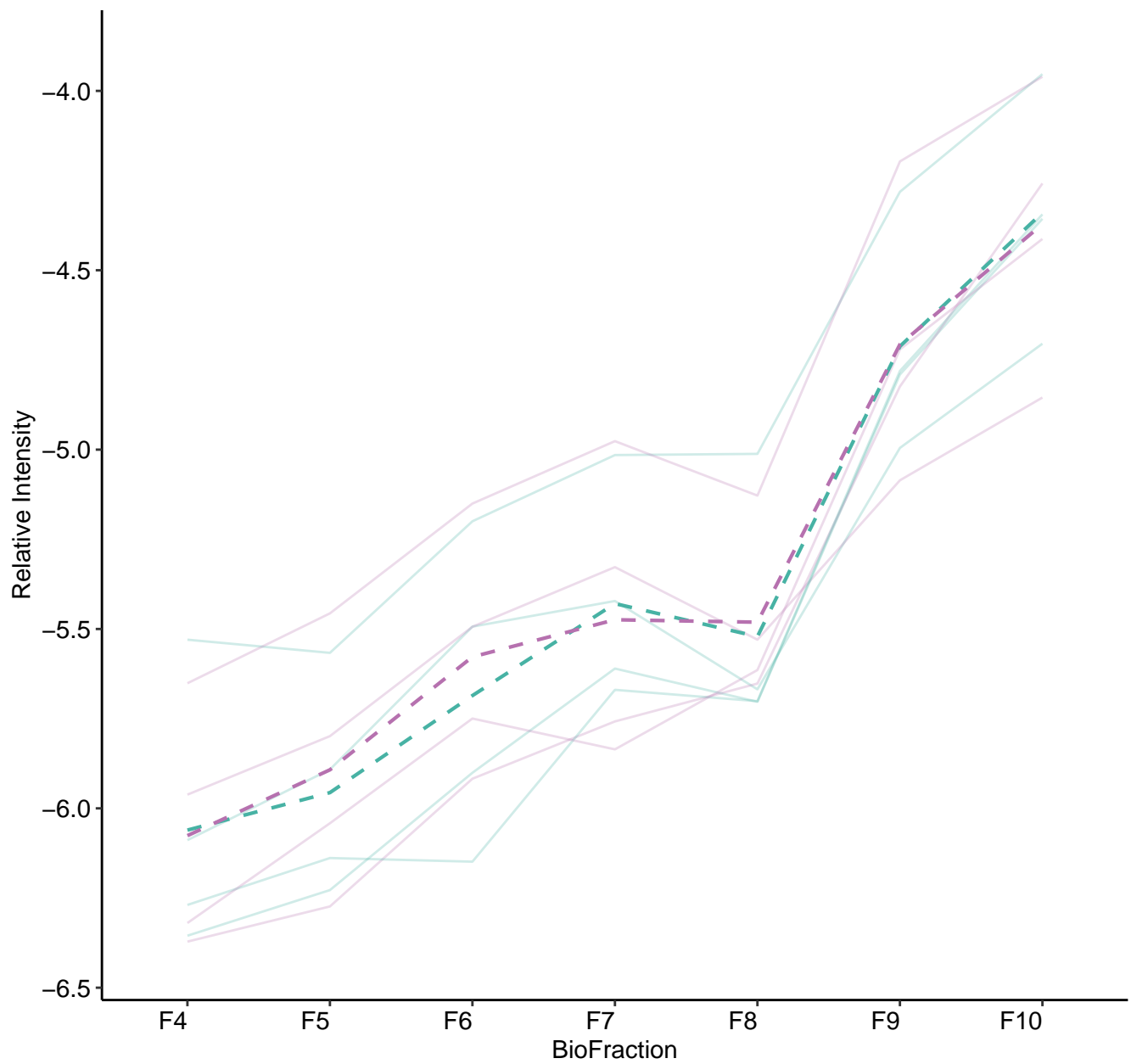
HCF-1 complex (n = 9)
(R2.Fixef = 0.295)



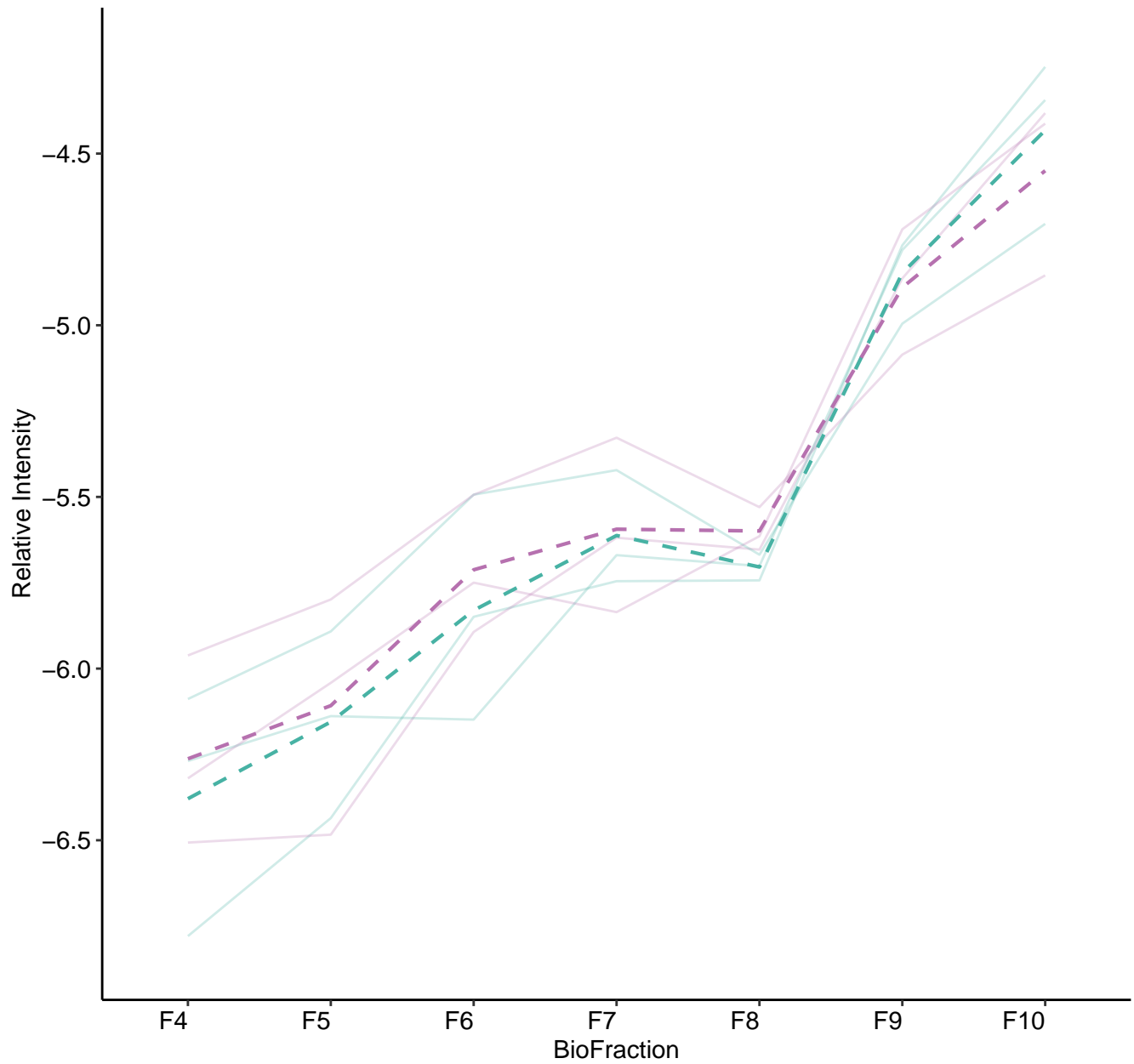
HDAC1-associated core complex cll (n = 3)
(R2.Fixef = 0.684)



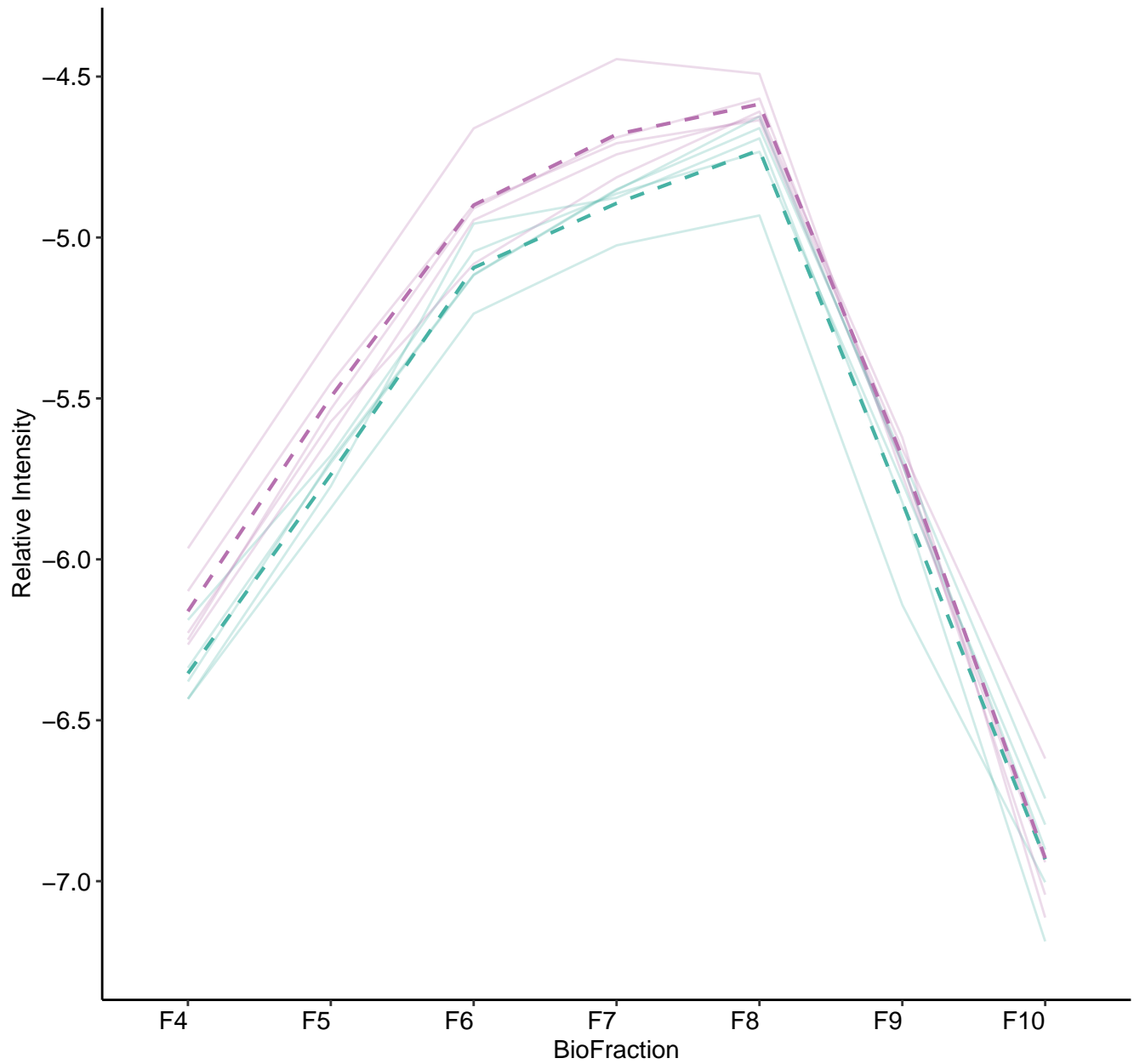
HDAC1-associated protein complex (n = 4)
(R2.Fixef = 0.746)



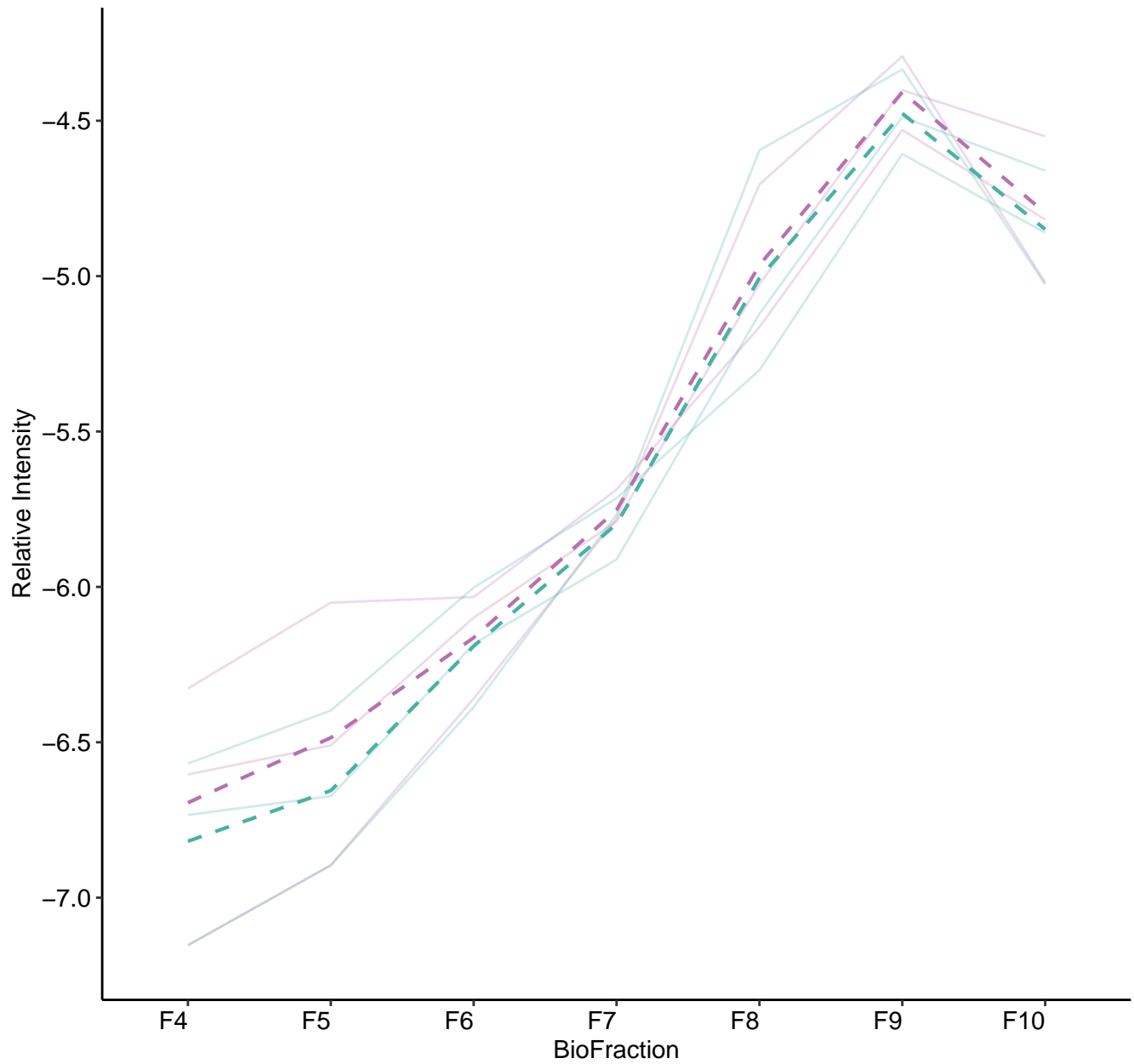
HDAC2-associated core complex (n = 3)
(R2.Fixef = 0.866)



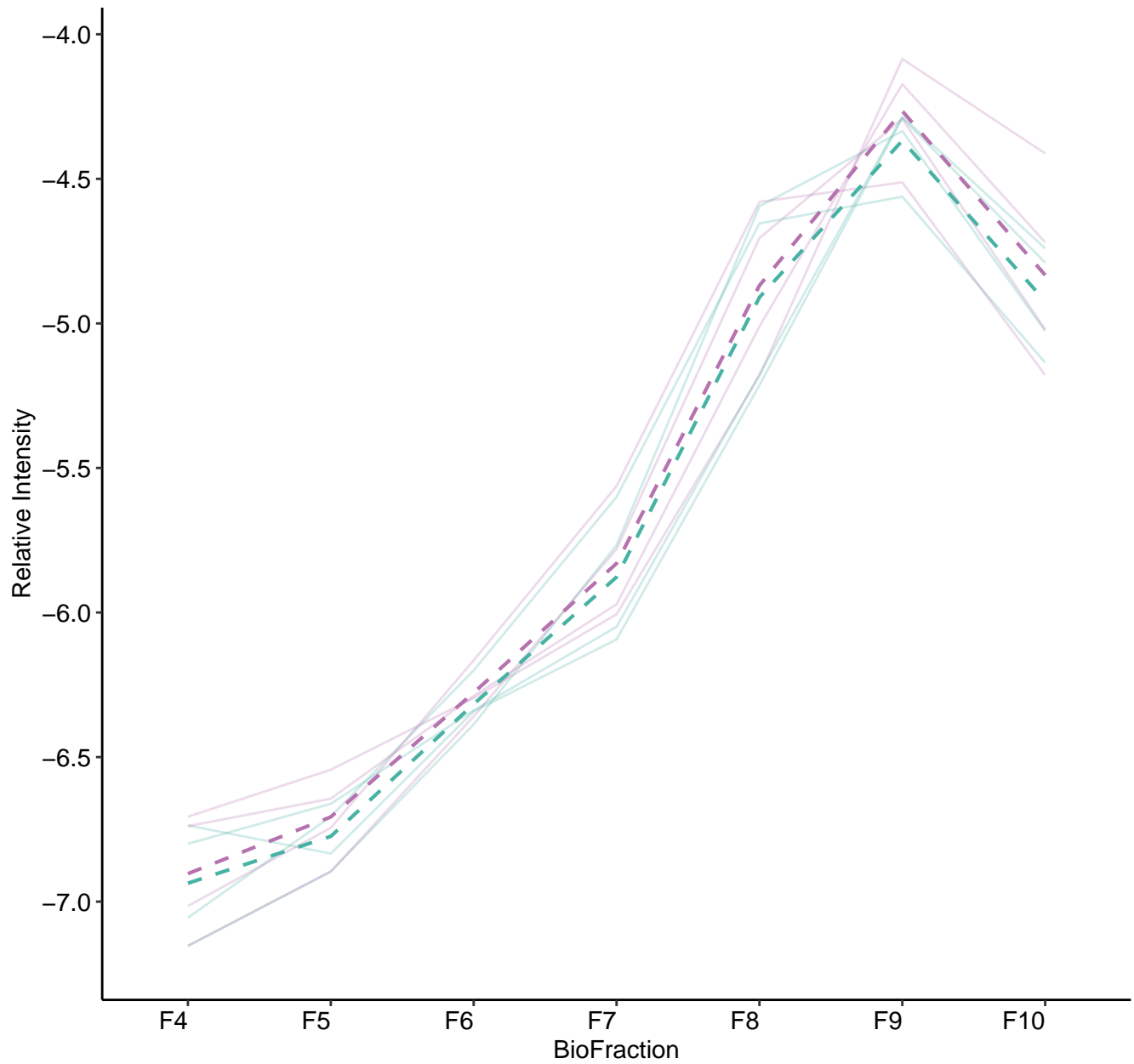
HRD1 complex (n = 5)
(R2.Fixef = 0.974)



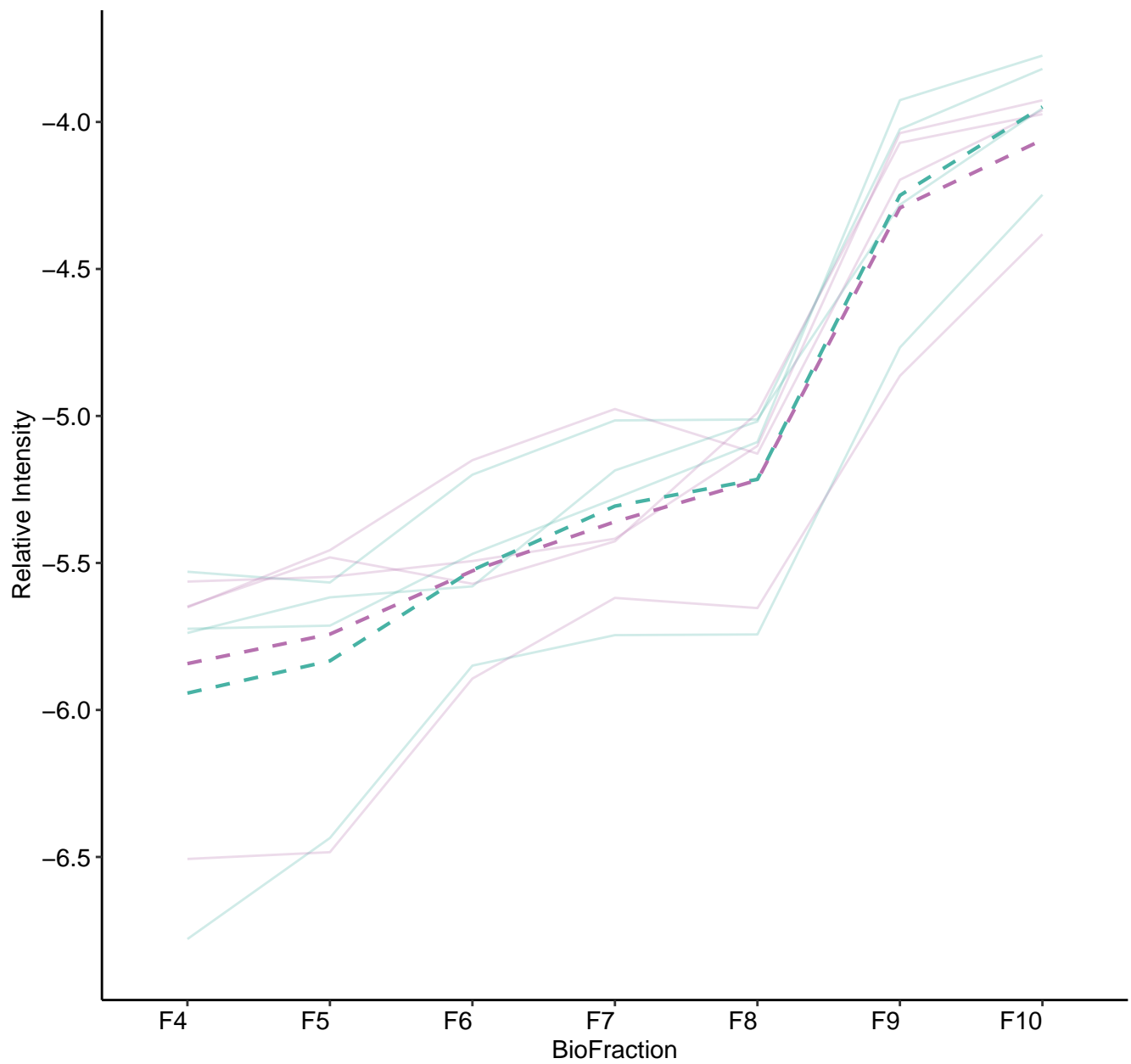
HuR-Tia1-Tiar-Hrnpu complex (n = 3)
(R2.Fixef = 0.919)



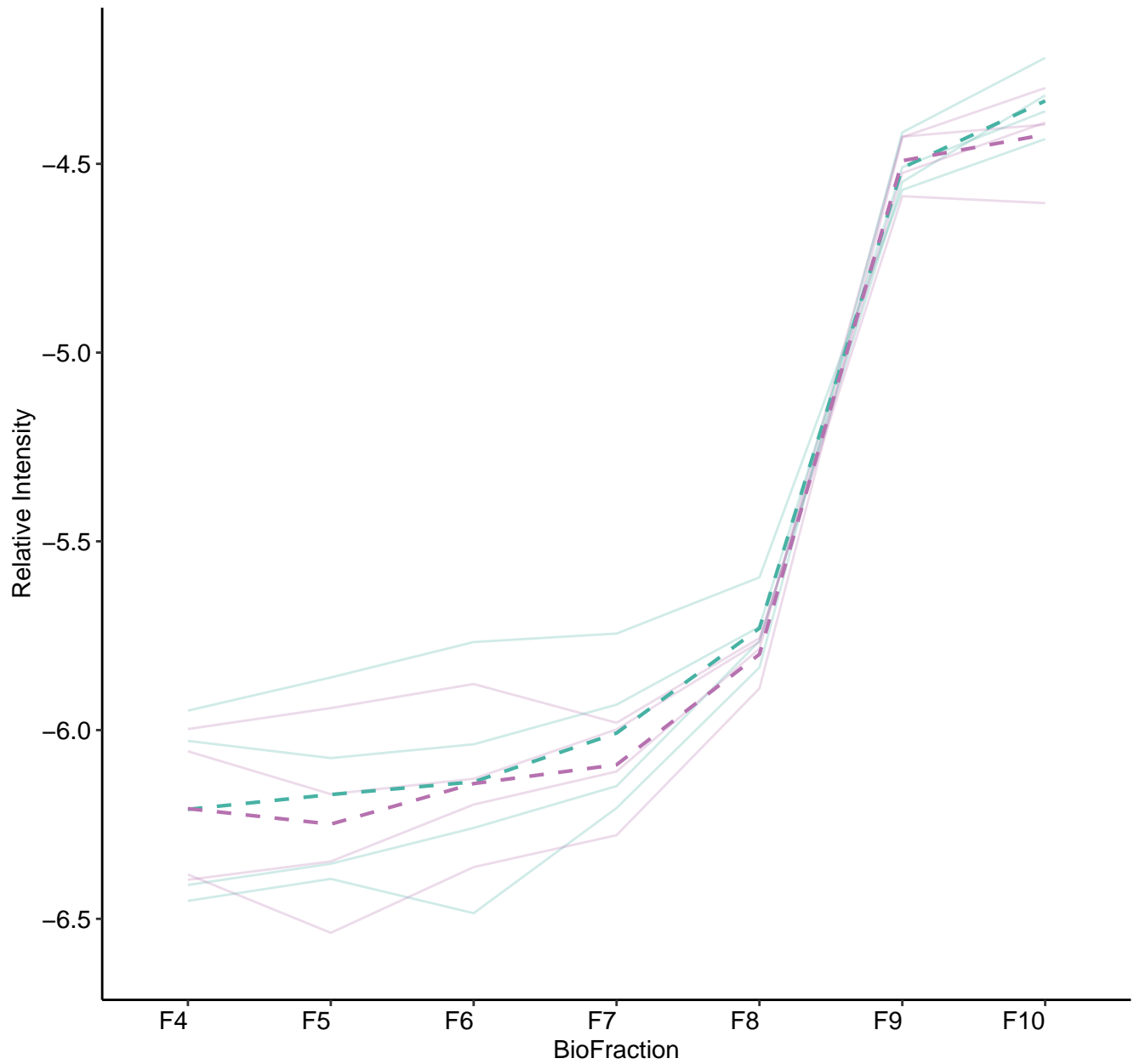
IGF2BP1 complex (n = 4)
(R2.Fixef = 0.953)



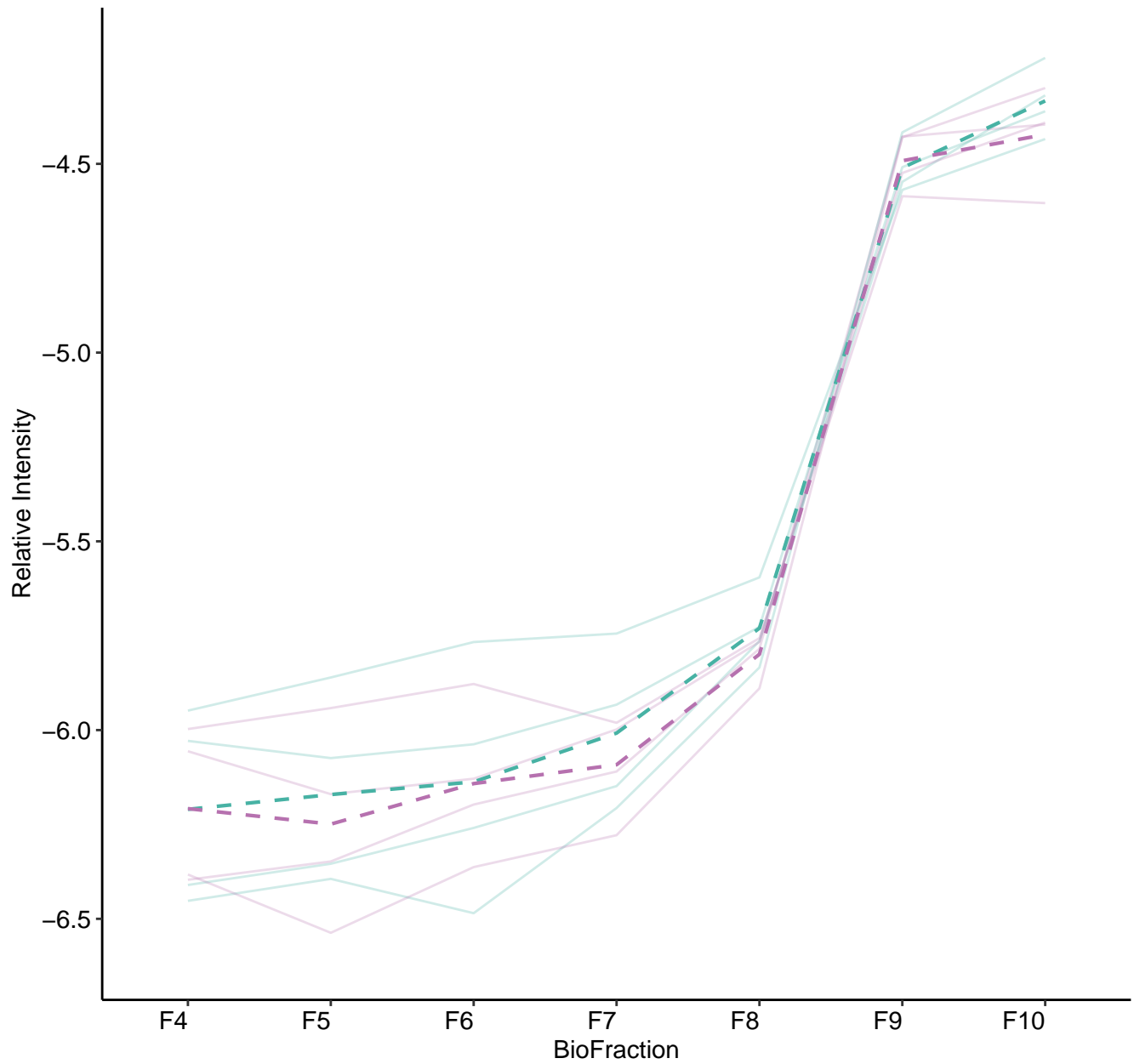
Ikaros complex (n = 4)
(R2.Fixef = 0.779)



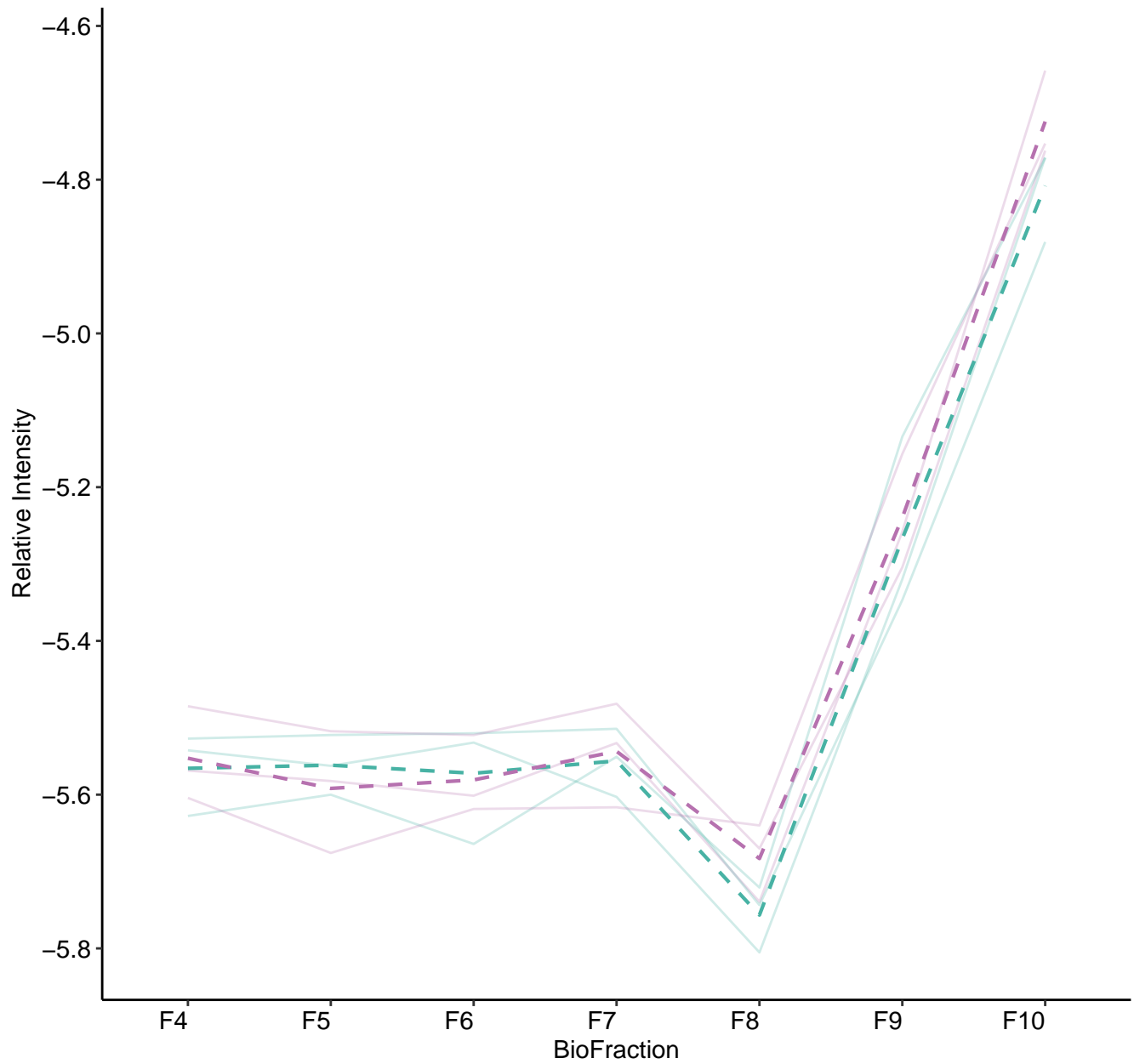
Integrator complex (n = 4)
(R2.Fixef = 0.943)



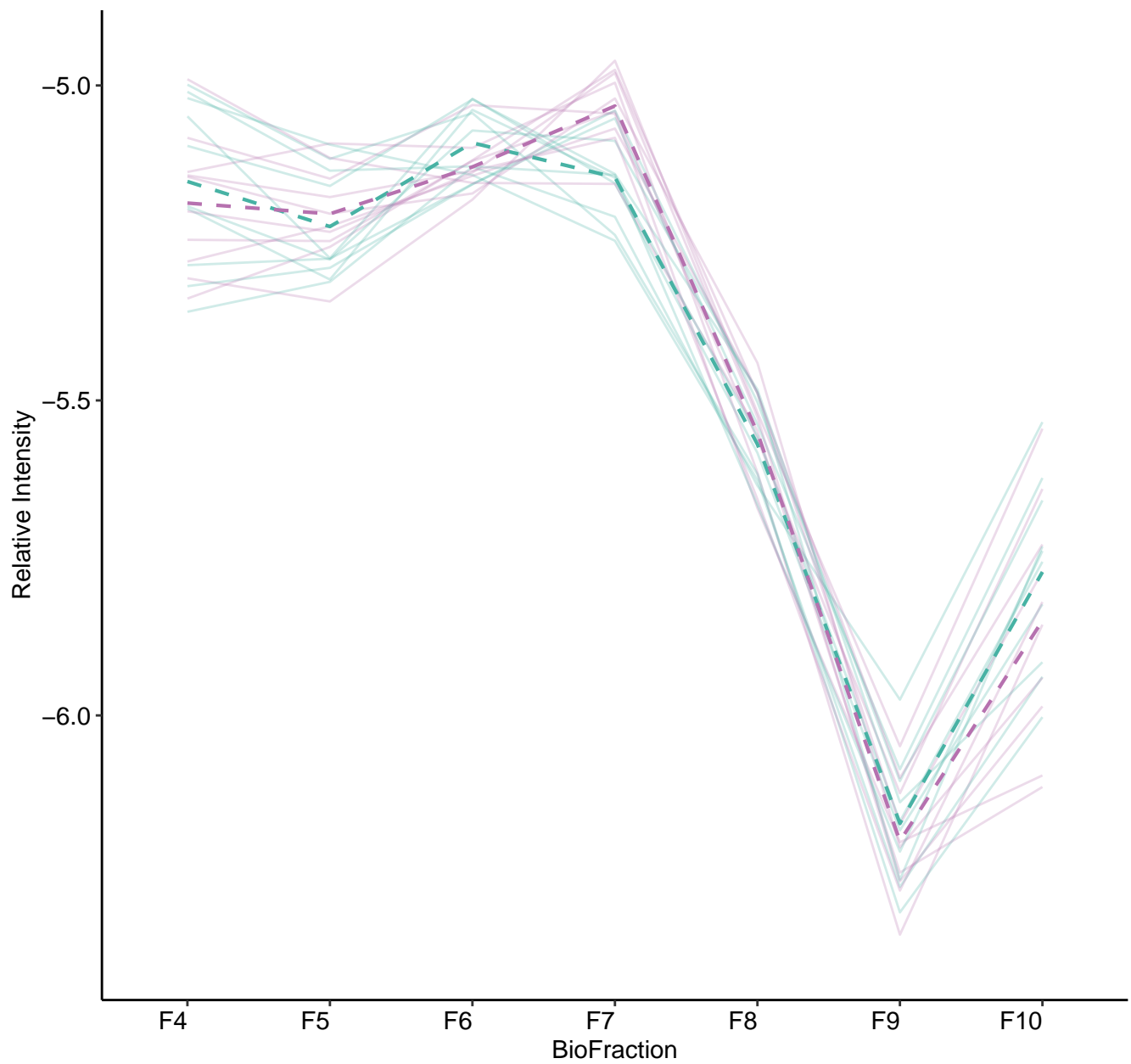
Integrator–RNAPII complex (n = 4)
(R2.Fixef = 0.943)



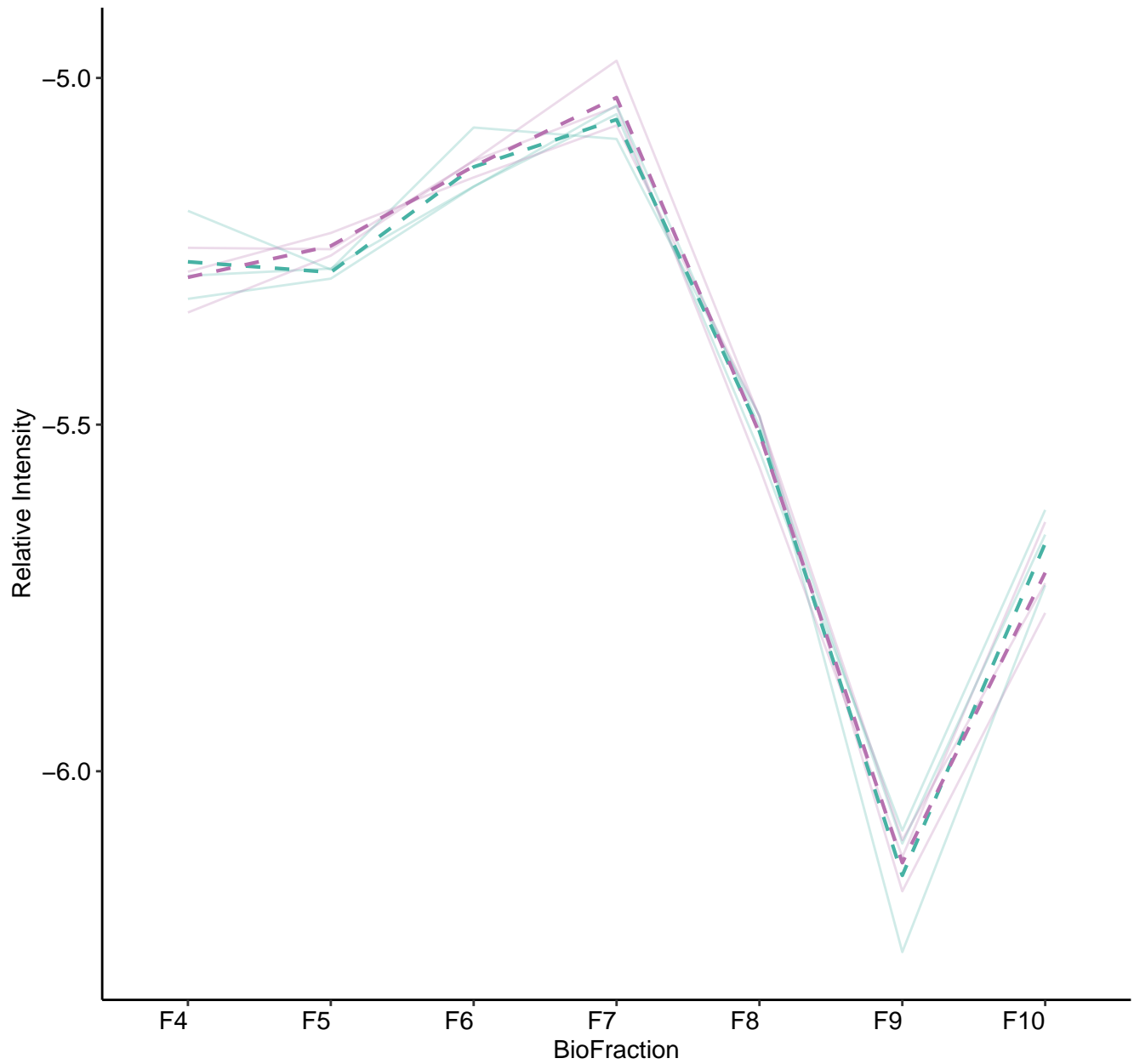
KIF3A/B-PAR-3-aPKC-PAR-6 complex (n = 3)
(R2.Fixef = 0.955)



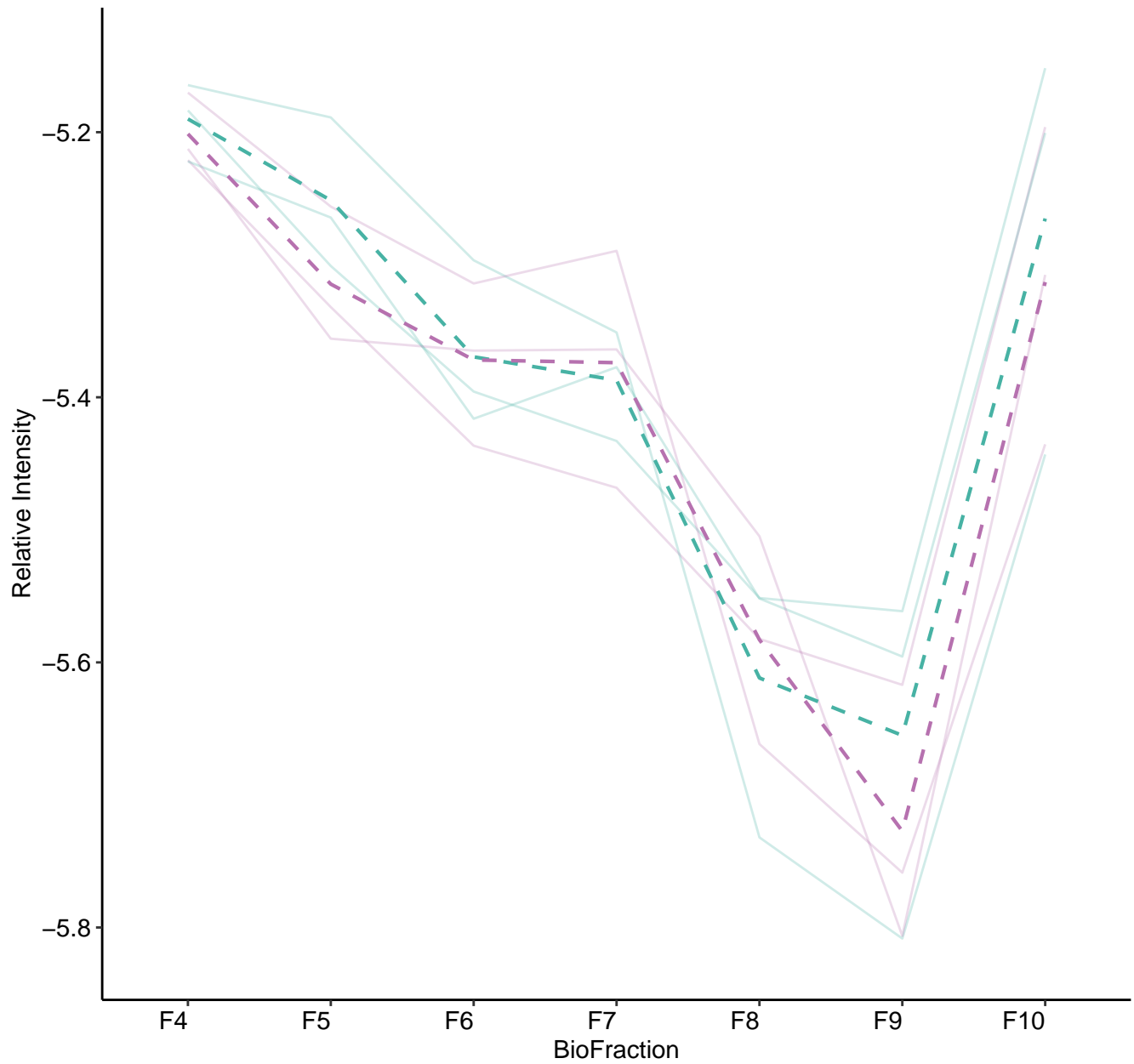
Kinase maturation complex 1 (n = 10)
(R2.Fixef = 0.938)



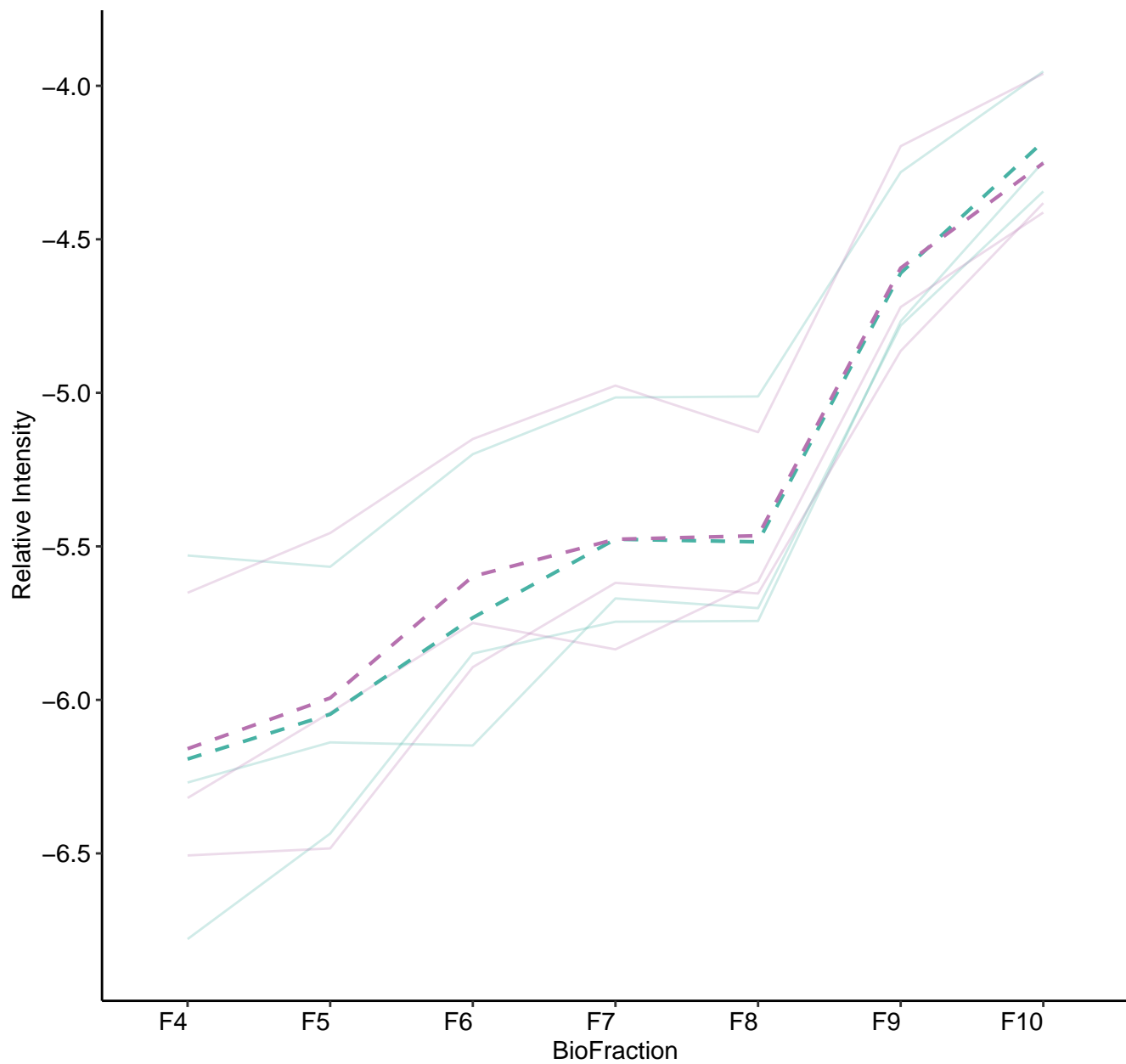
Kinase maturation complex 2 (n = 3)
(R2.Fixef = 0.982)



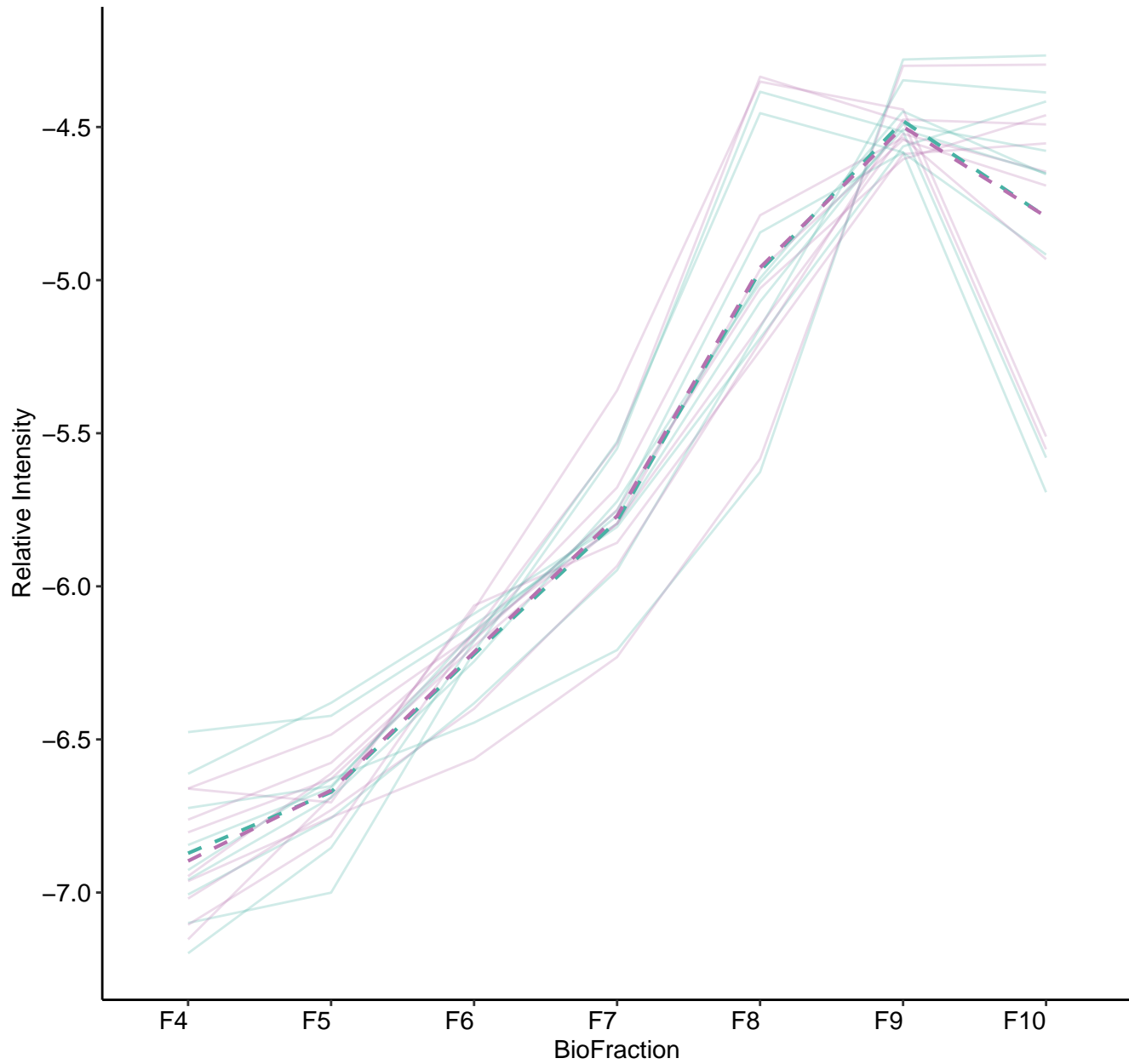
Ksr1-CK2-MEK-14-3-3 complex, PDGF treated (n = 3)
(R2.Fixef = 0.787)



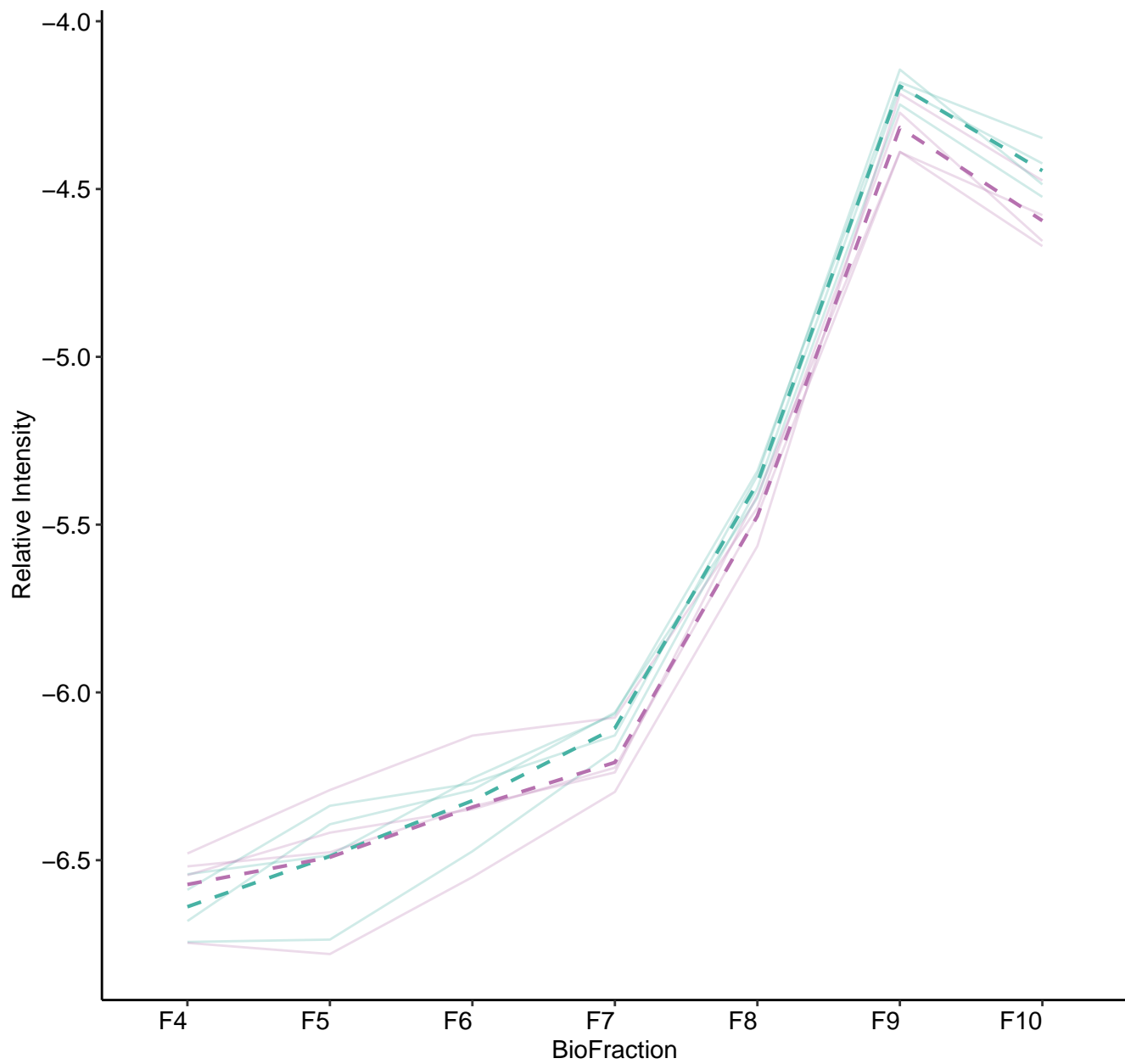
LARC complex (LCR-associated remodeling complex) (n = 3)
(R2.Fixef = 0.729)



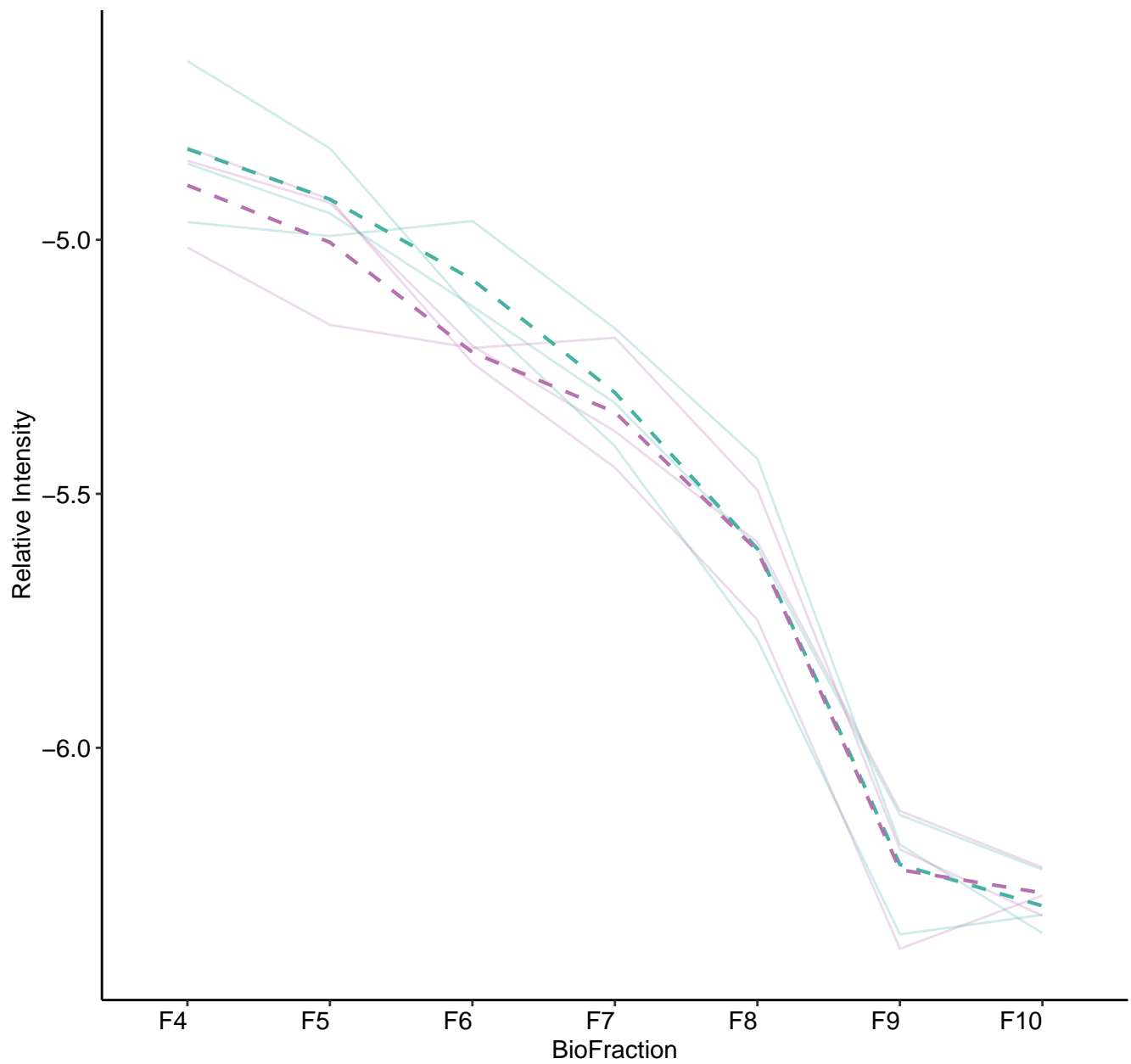
Large Drosha complex (n = 9)
(R2.Fixef = 0.91)



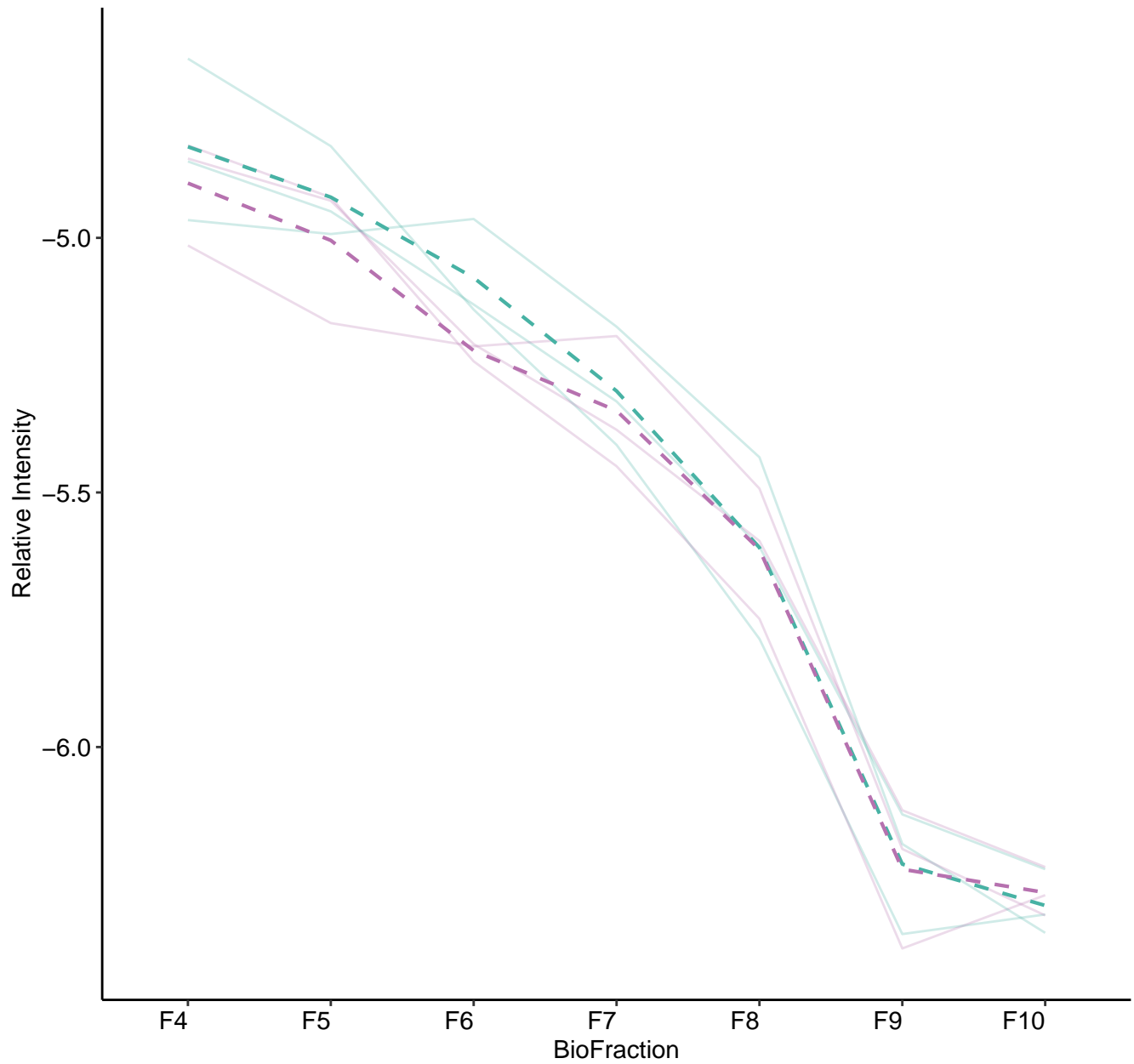
LAS1L-PELP1-TEX10-WDR18-NOL9-SEN3 complex (n = 4)
(R2.Fixef = 0.985)



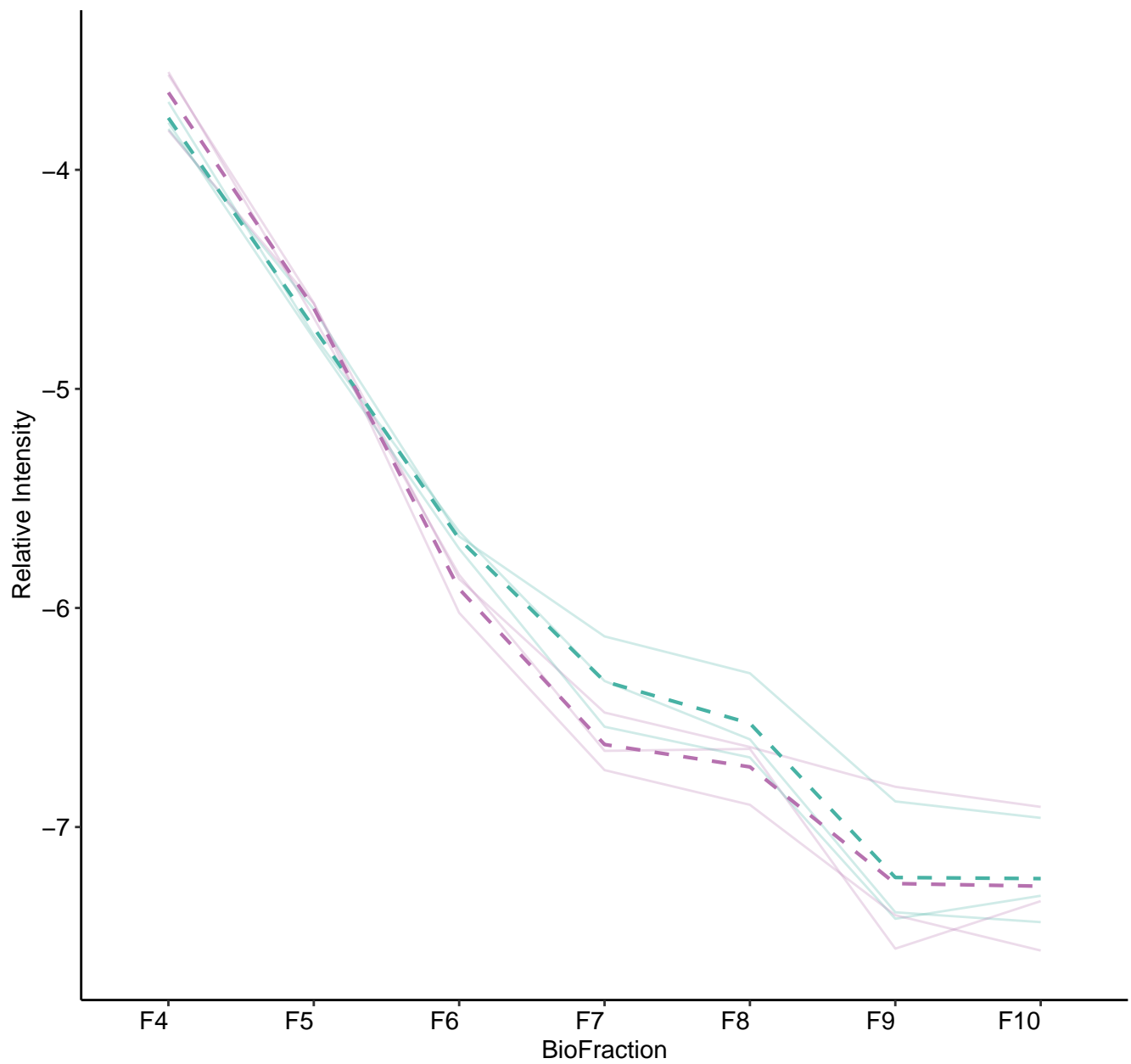
LIN2-LIN7-SAP97 complex (n = 3)
(R2.Fixef = 0.956)



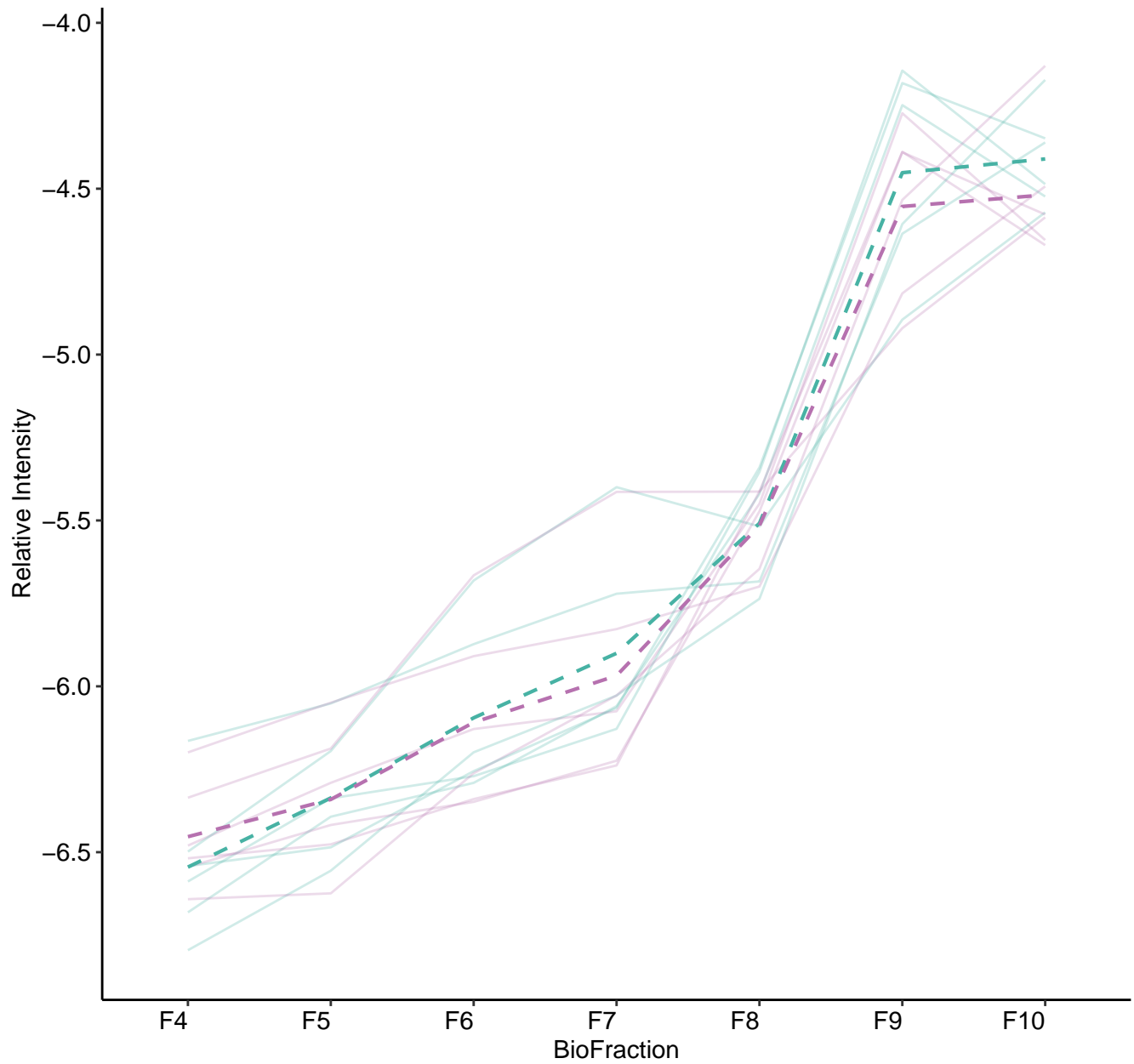
LIN2-LIN7-SAP97-MINT1 complex (n = 3)
(R2.Fixef = 0.956)



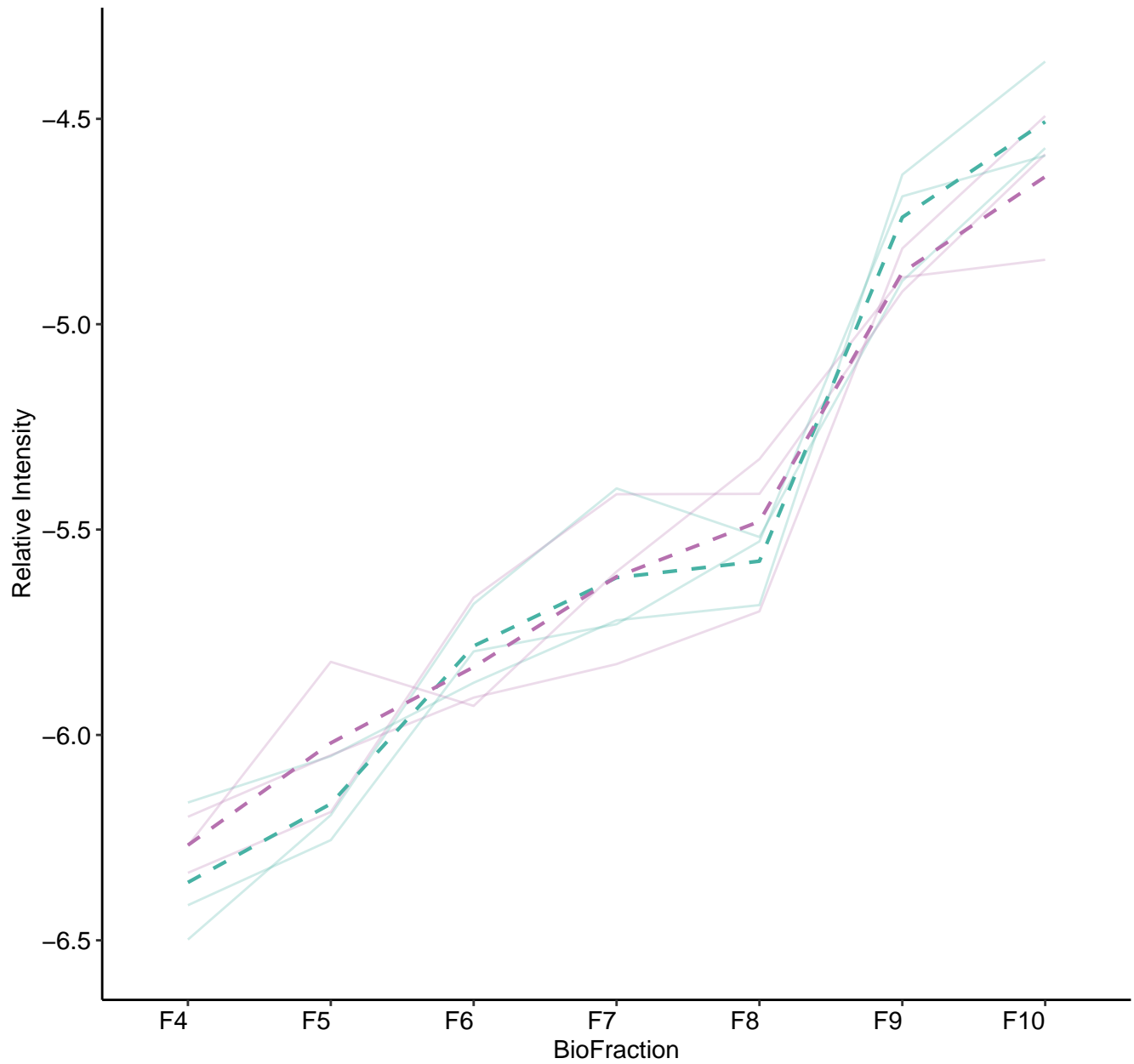
mitochondrial calcium uniporter complex (n = 3)
(R2.Fixef = 0.974)



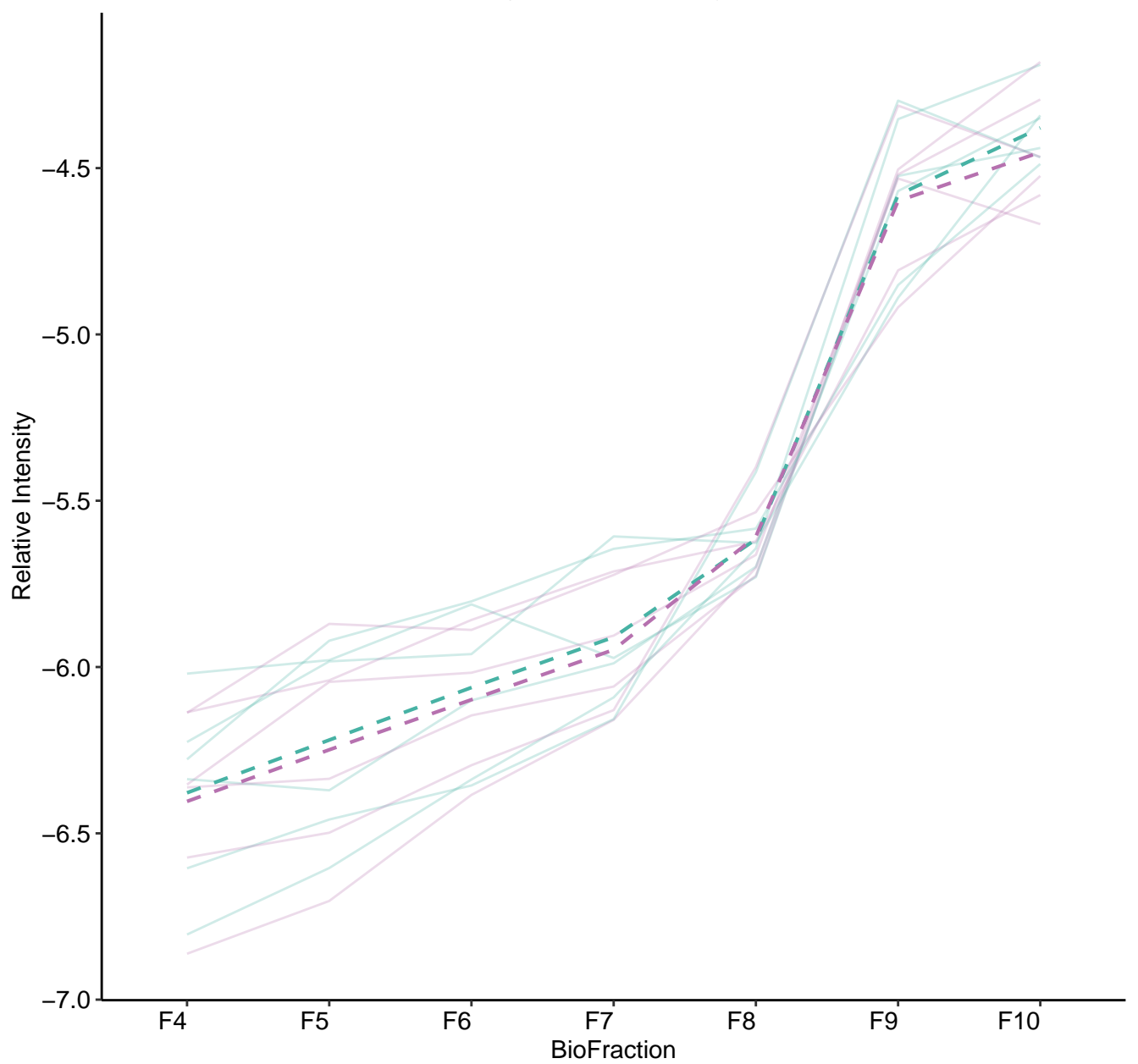
MLL1-WDR5 complex (n = 6)
(R2.Fixef = 0.921)



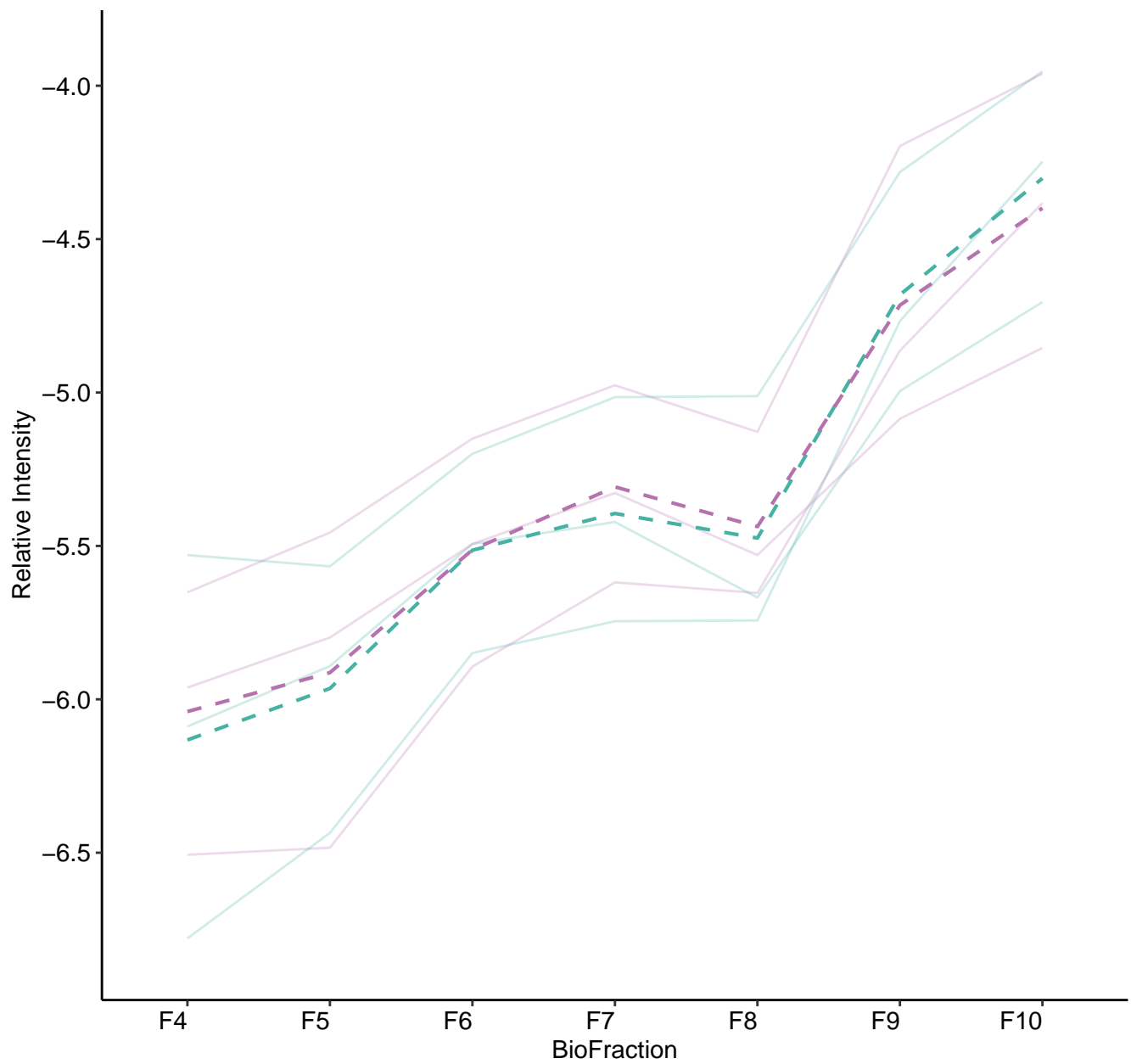
MLL2 complex (n = 3)
(R2.Fixef = 0.943)



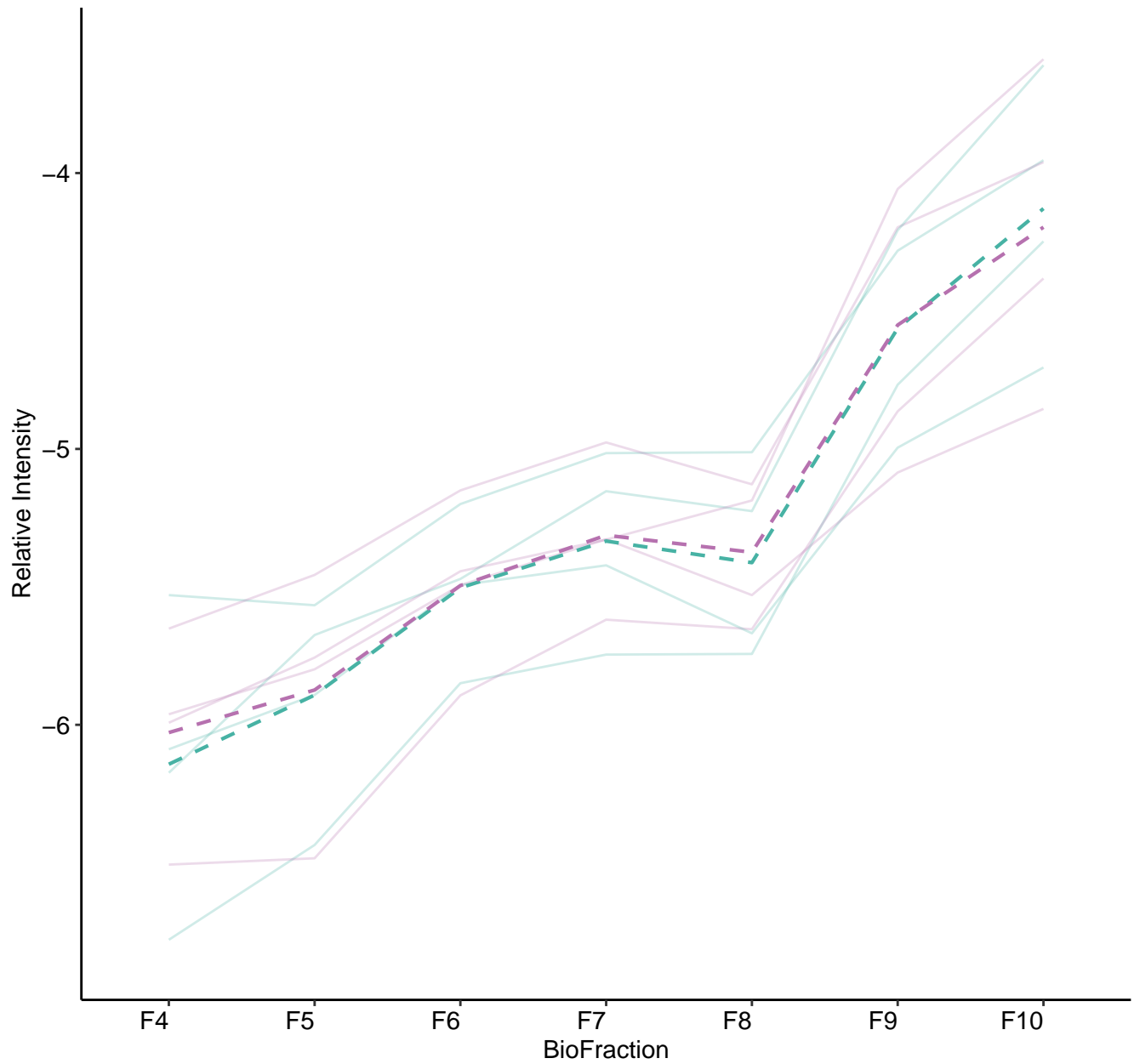
RNA decay complex (UPF1, UPF2, UPF3B, DCP2, XRN1, XRN2, EXOSC2, EXOSC4, EXOSC10, PARP1)
(R2.Fixef = 0.913)



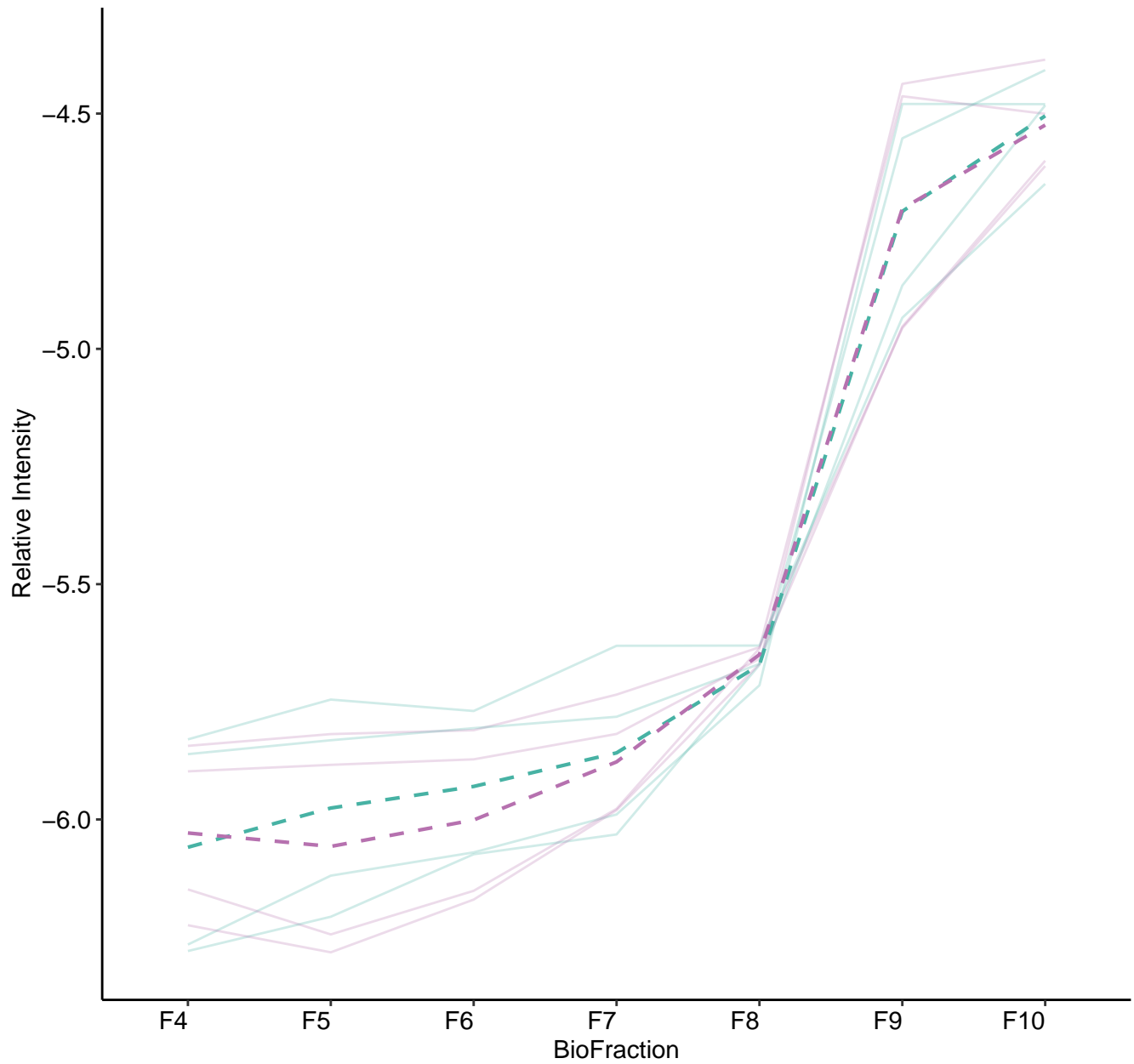
mSin3A complex (n = 3)
(R2.Fixef = 0.663)



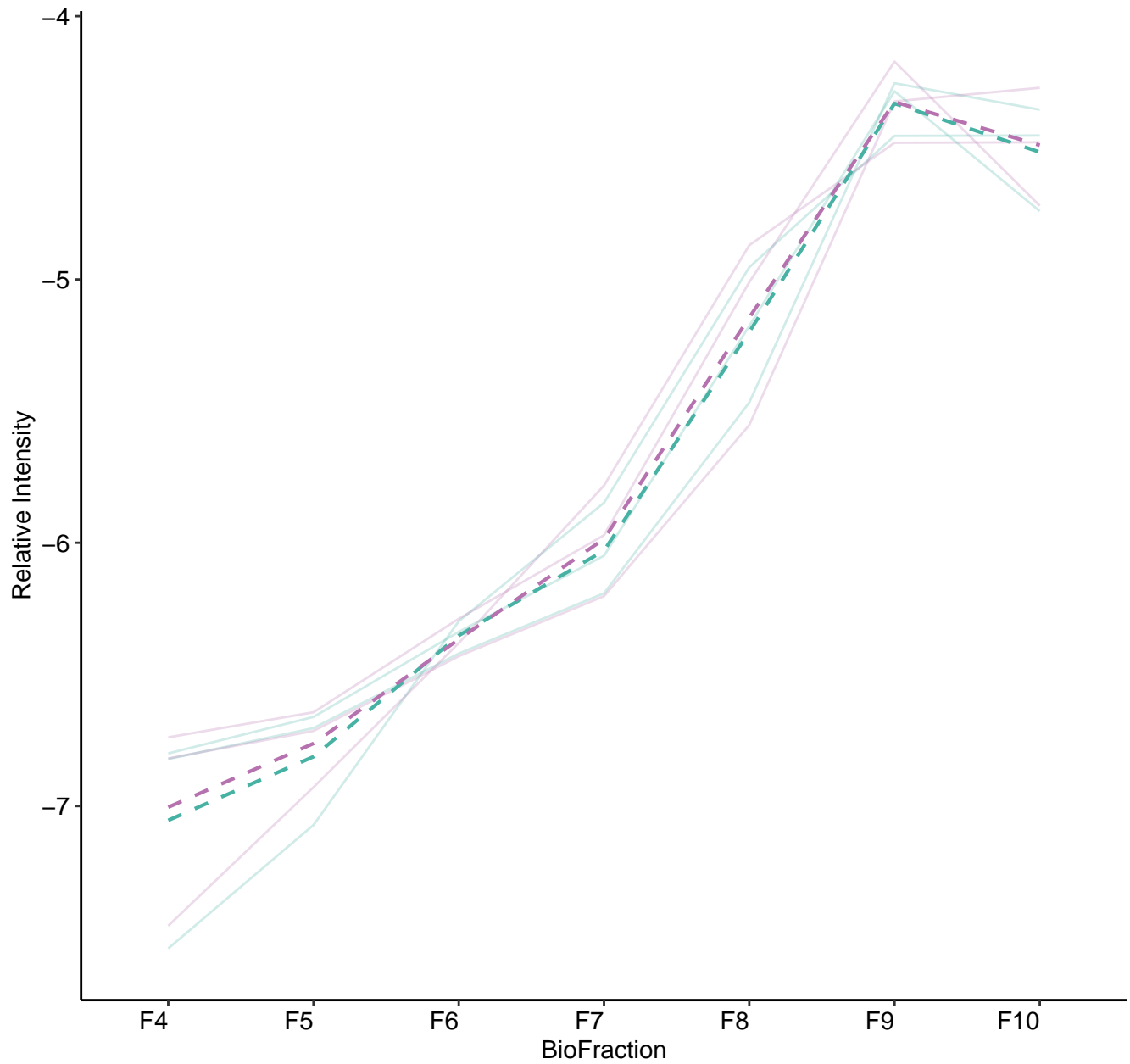
MTA1 complex (n = 4)
(R2.Fixef = 0.729)



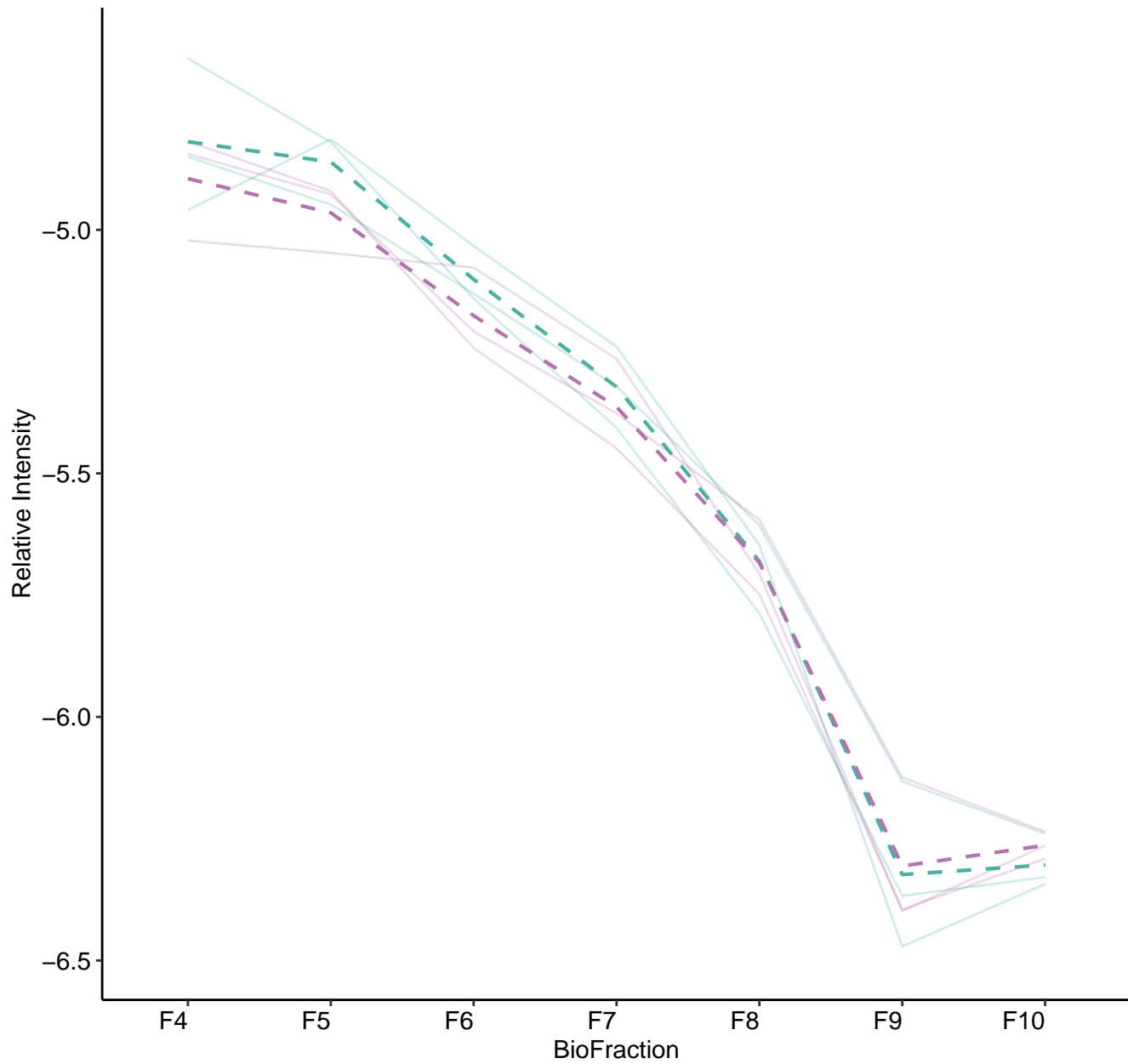
mTORC2 complex (n = 4)
(R2.Fixef = 0.916)



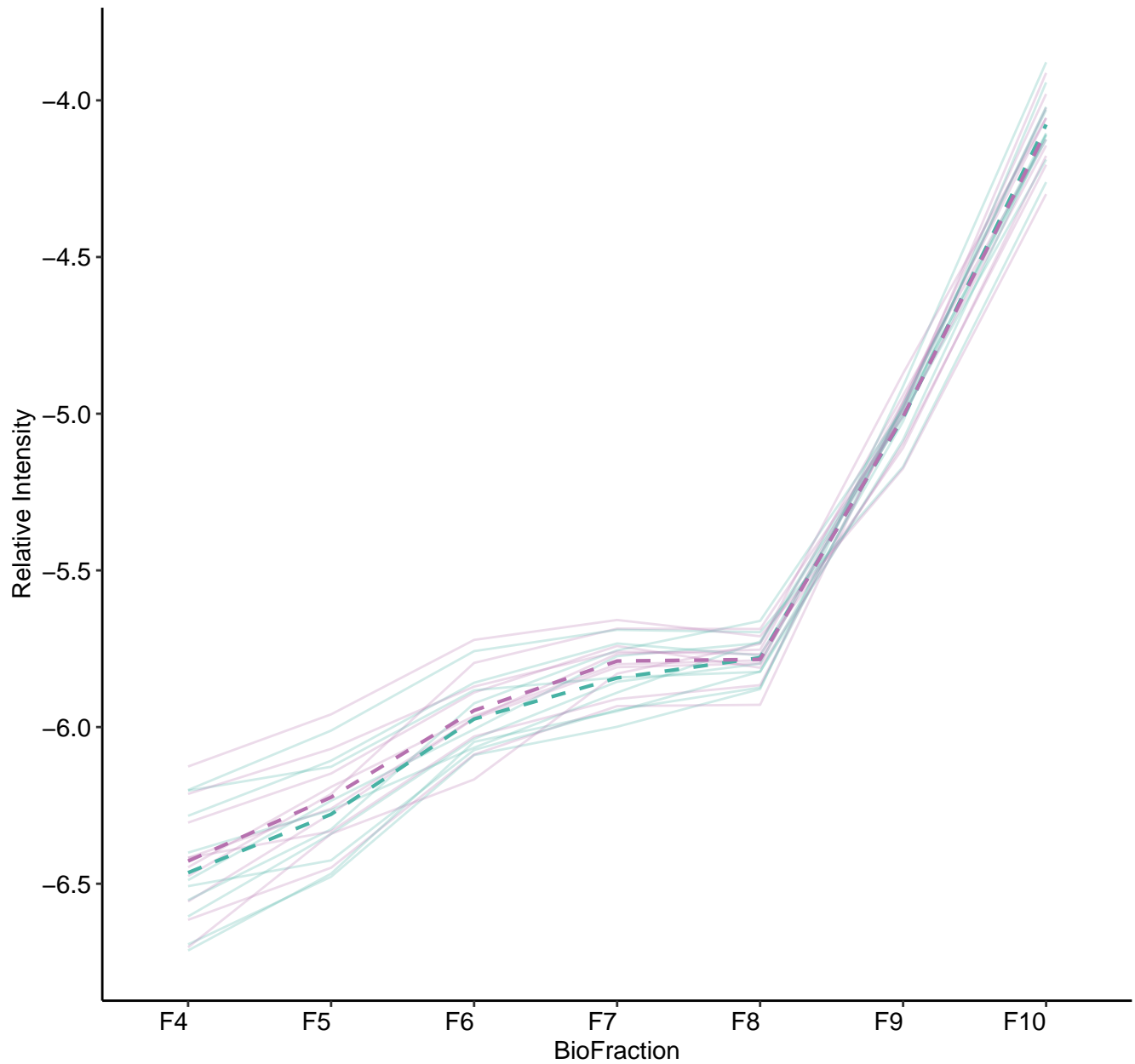
Multiprotein complex (mRNA turnover) (n = 3)
(R2.Fixef = 0.947)



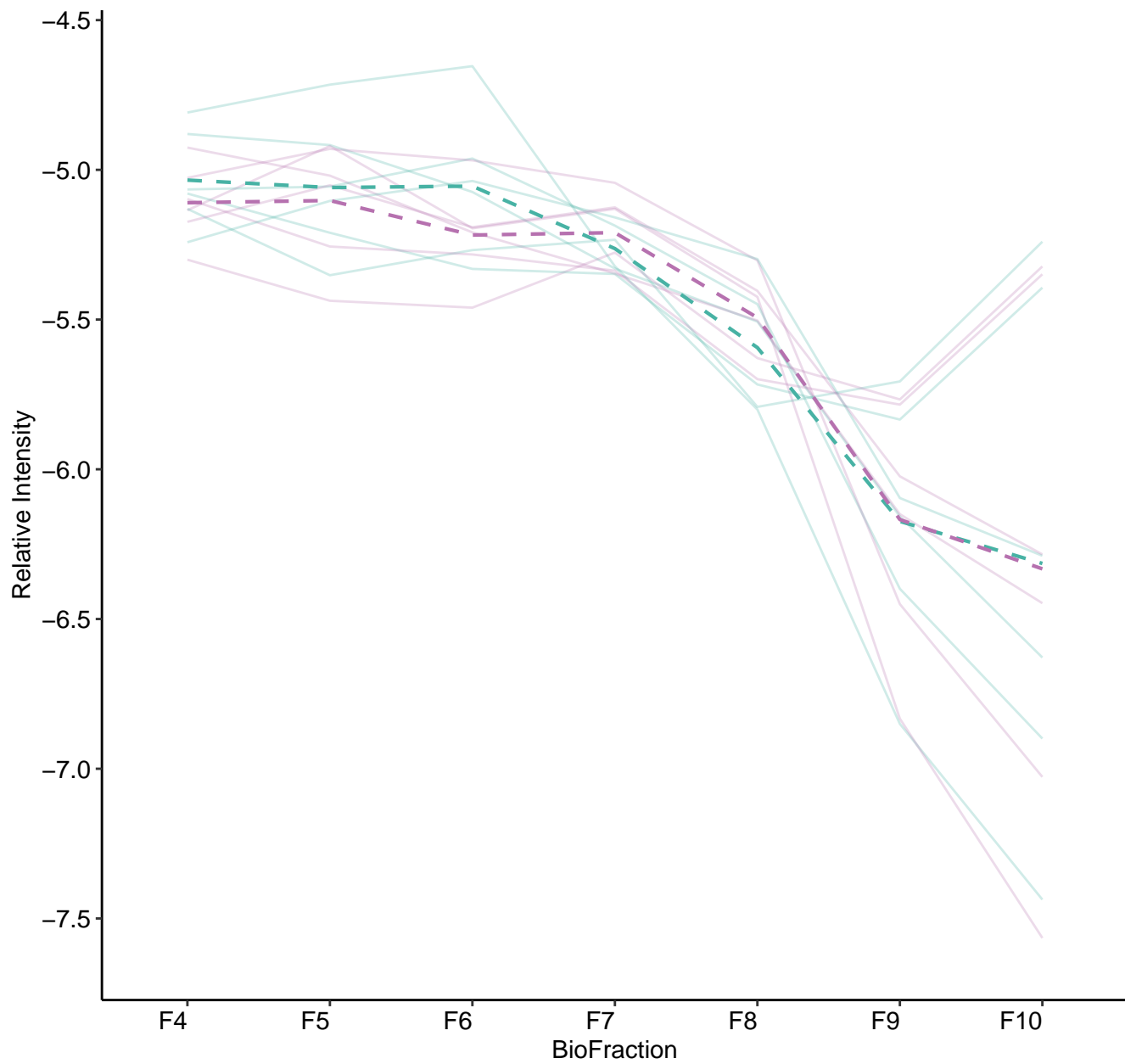
Multiprotein trafficking complex (n = 3)
(R2.Fixef = 0.968)



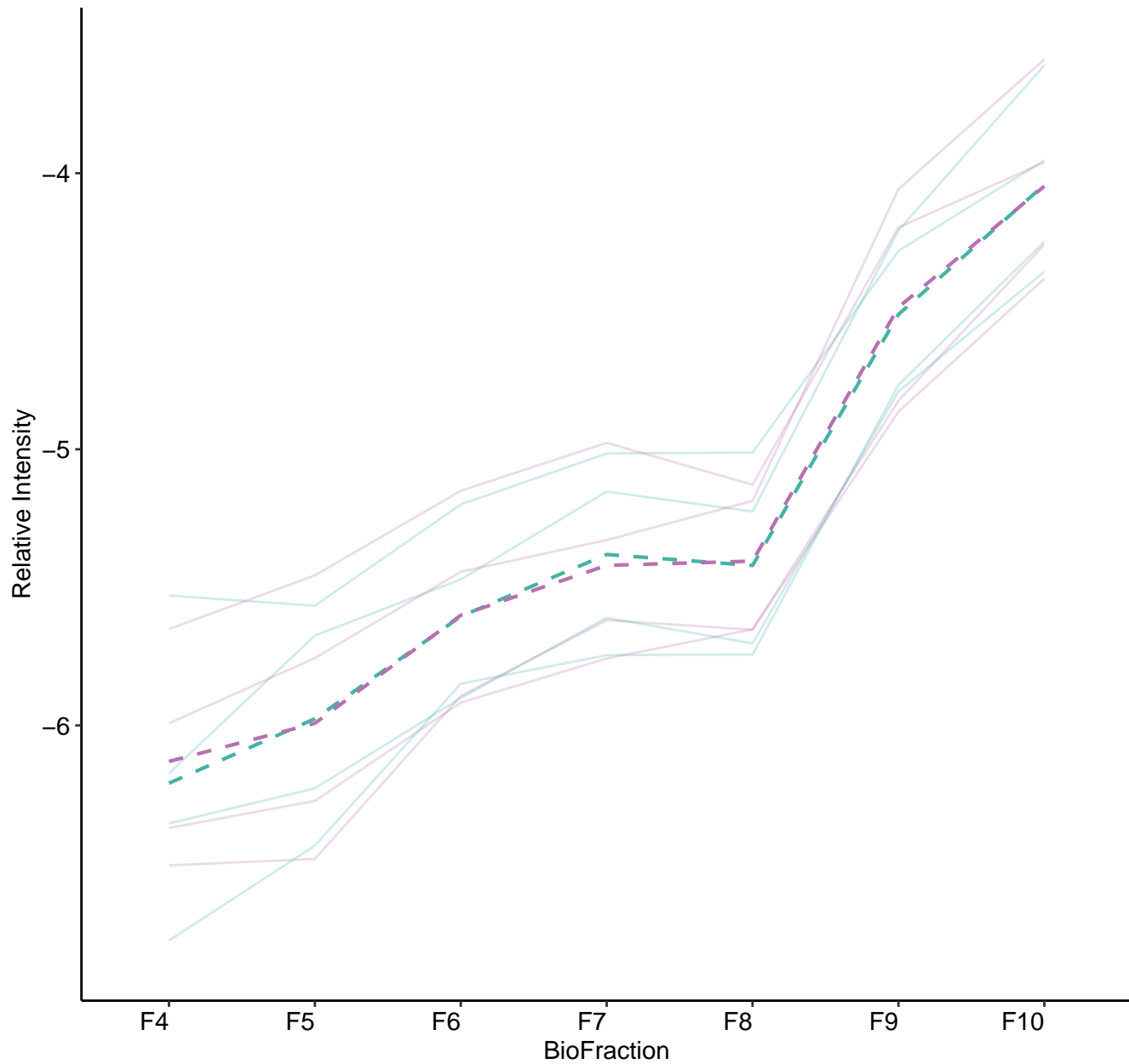
Multisynthetase complex (n = 10)
(R2.Fixef = 0.974)



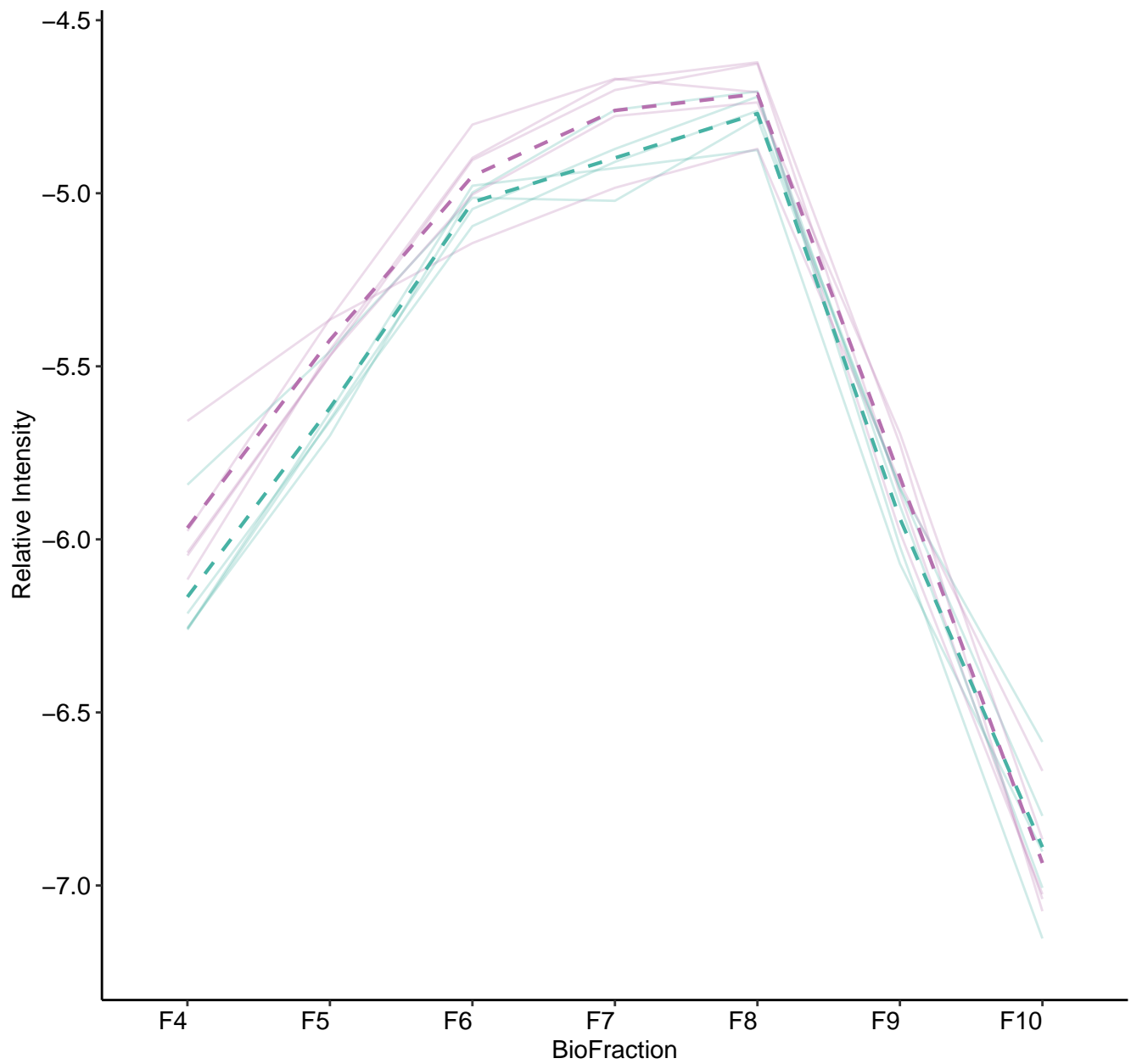
Ncam-Fgfr4 signaling complex (n = 6)
(R2.Fixef = 0.612)



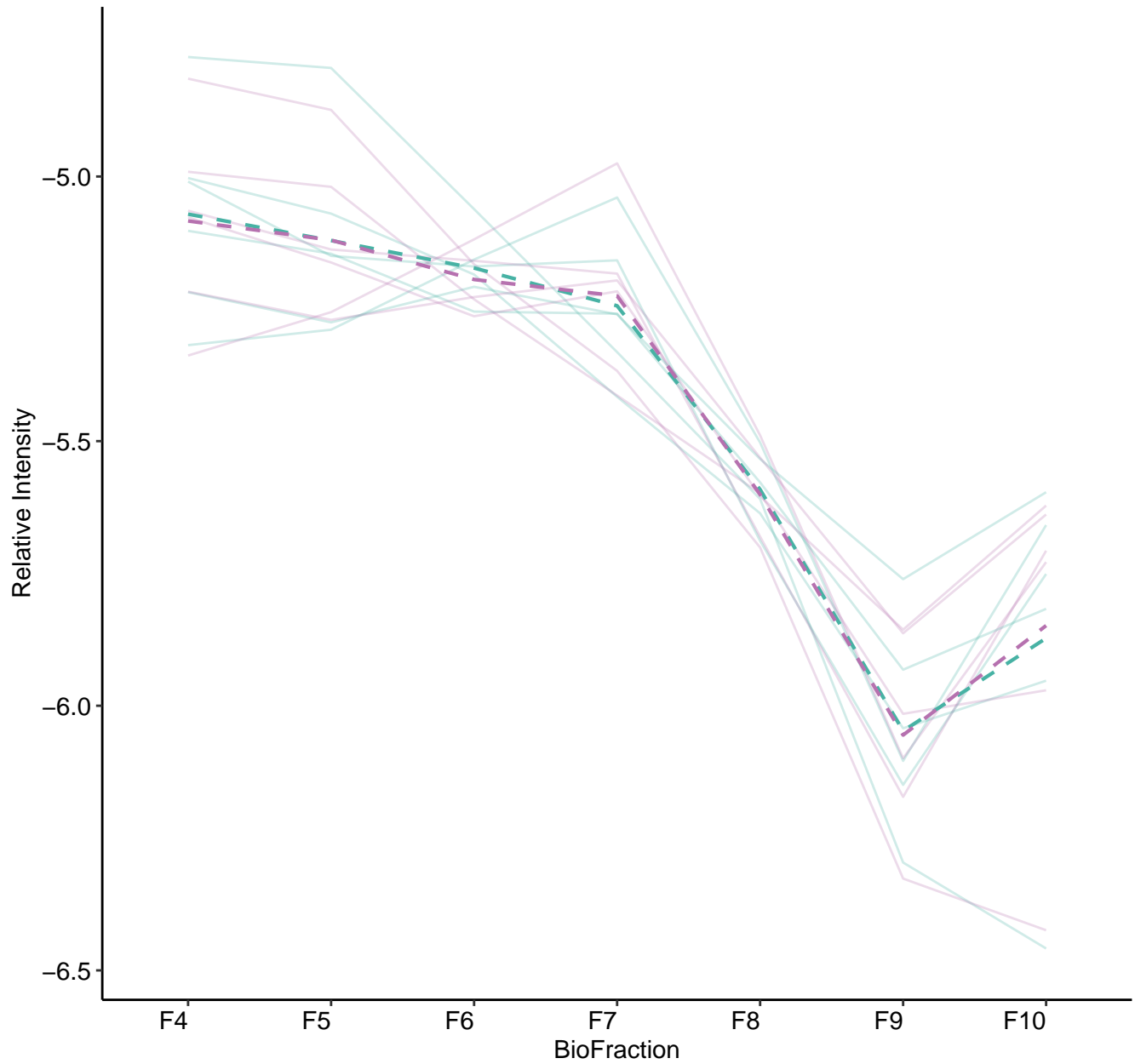
NRD complex (Nucleosome remodeling and deacetylation complex) (n = 4)
(R2.Fixef = 0.781)



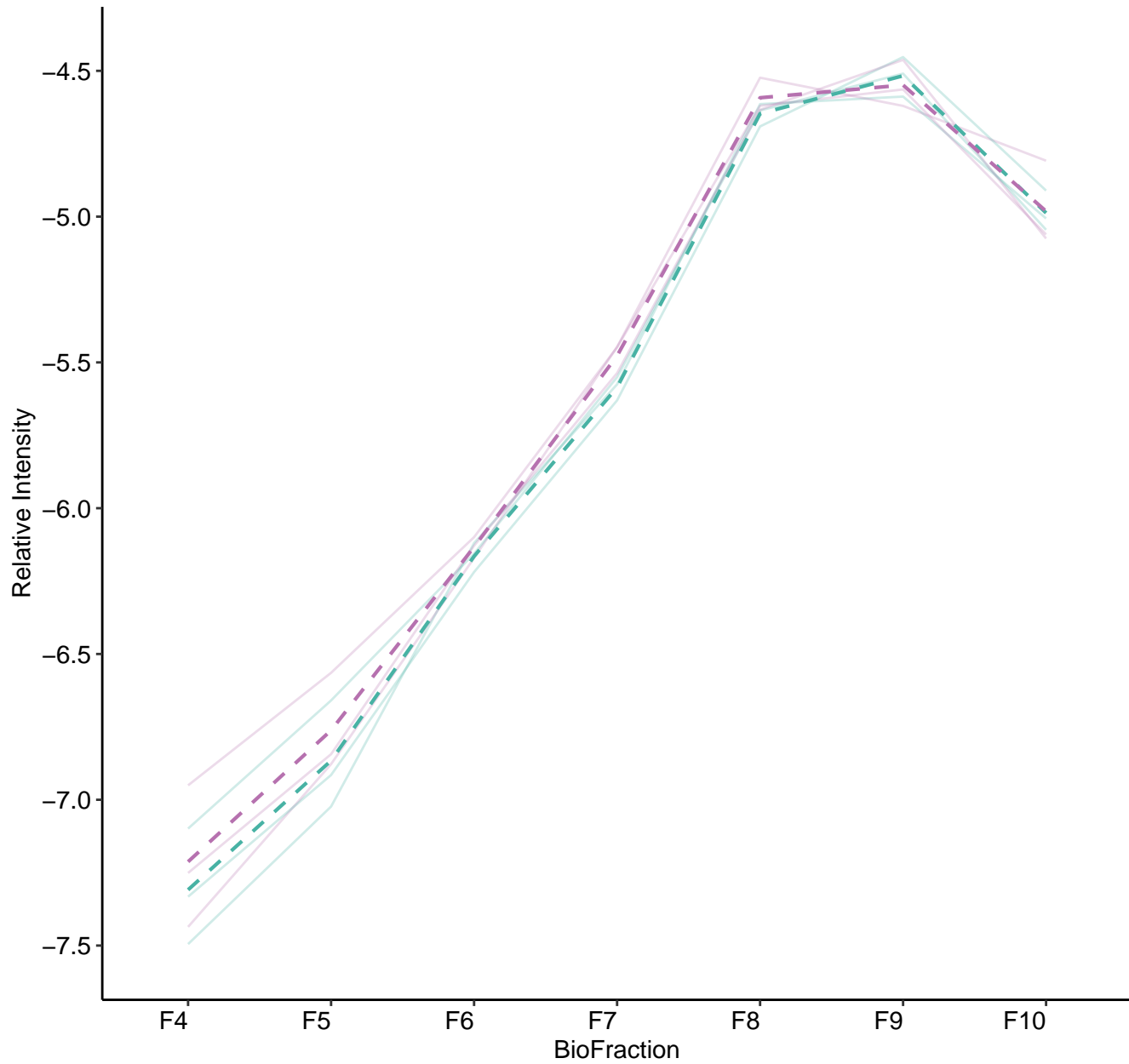
Oligosaccharyltransferase complex (Stt3A variant) (n = 5)
(R2.Fixef = 0.97)



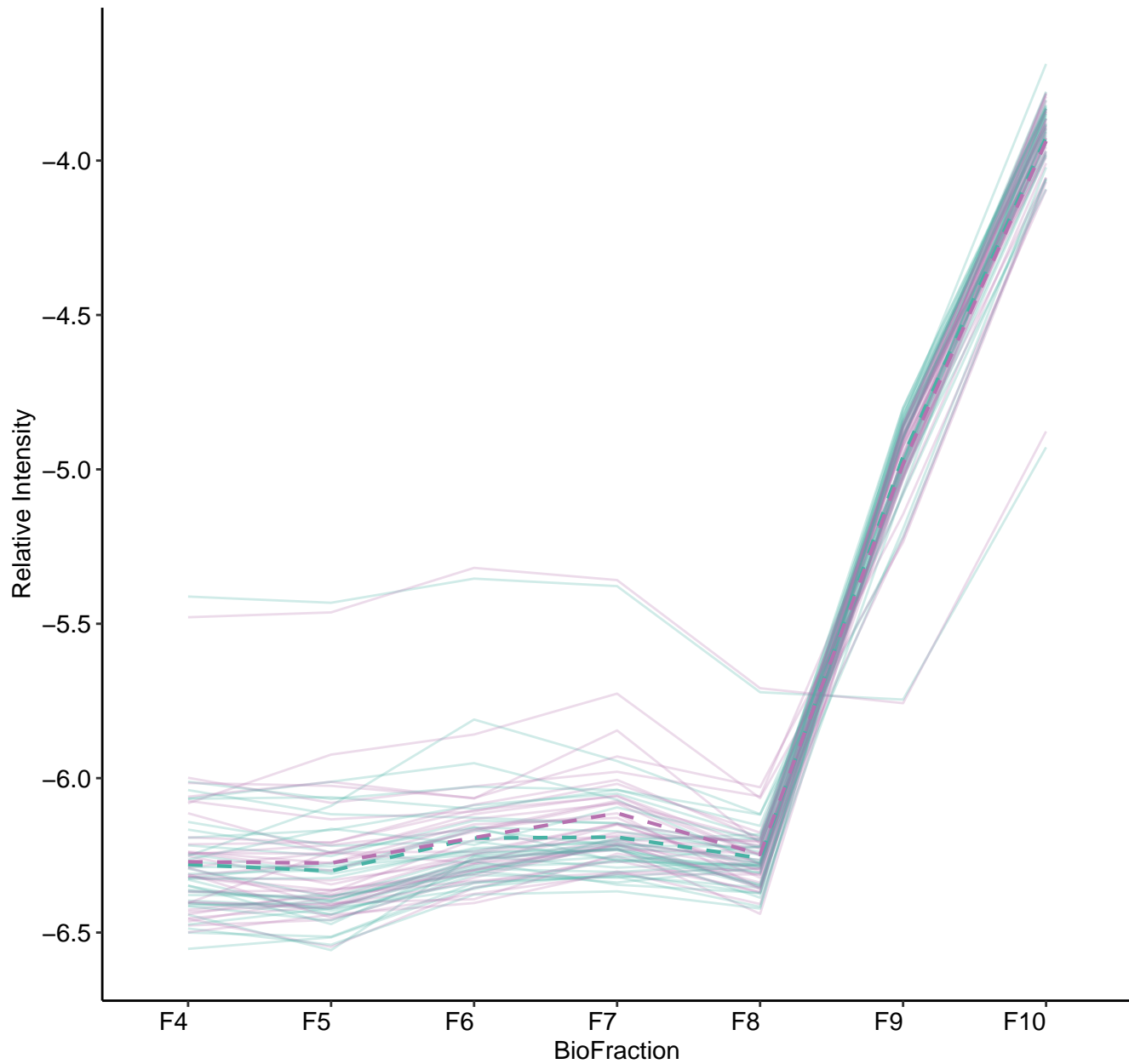
P2X7 receptor signalling complex (n = 6)
(R2.Fixef = 0.806)



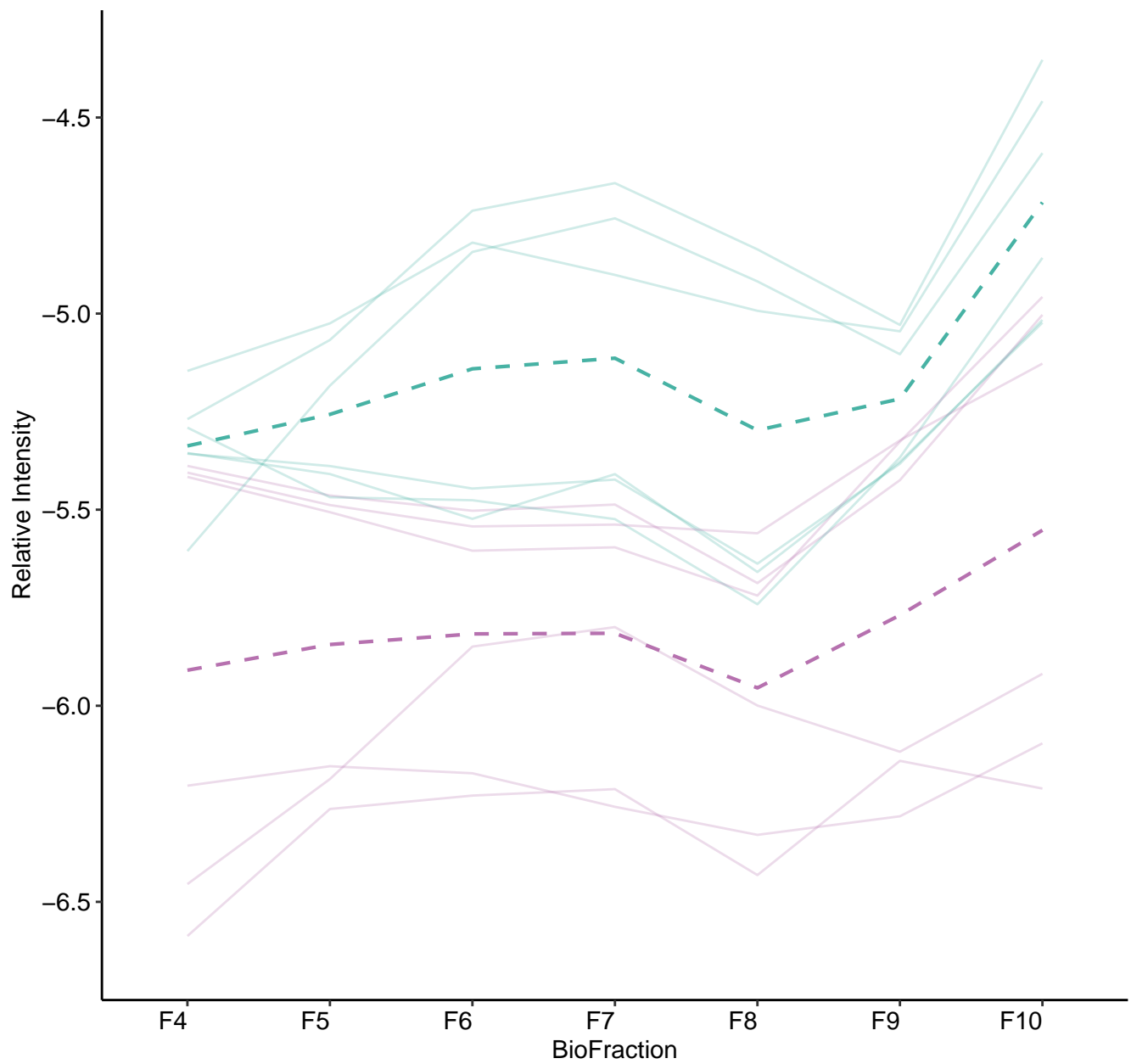
p54(nrb)-PSF-matrin3 complex (n = 3)
(R2.Fixef = 0.985)



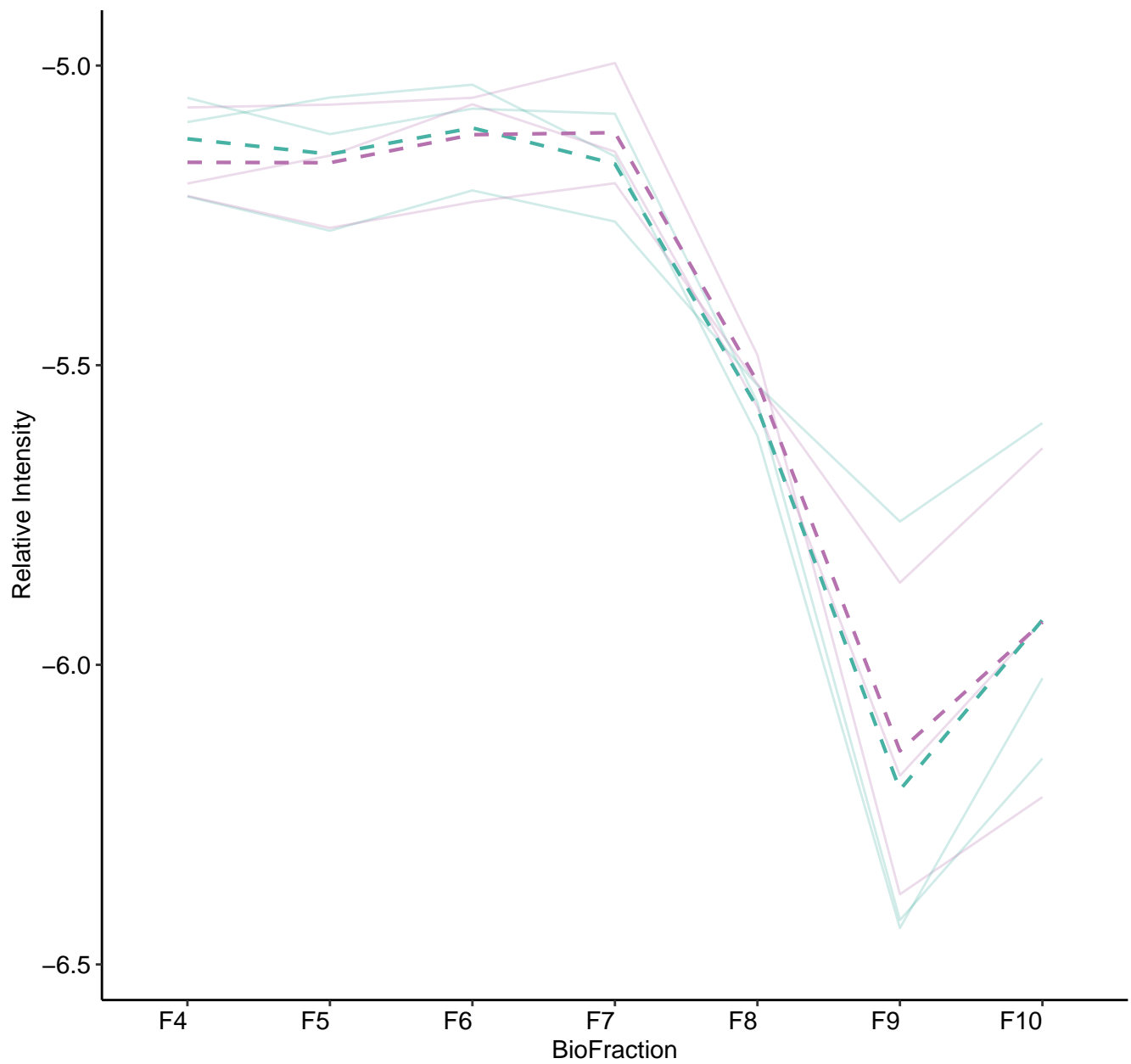
PA700-20S-PA28 complex (n = 34)
(R2.Fixef = 0.956)



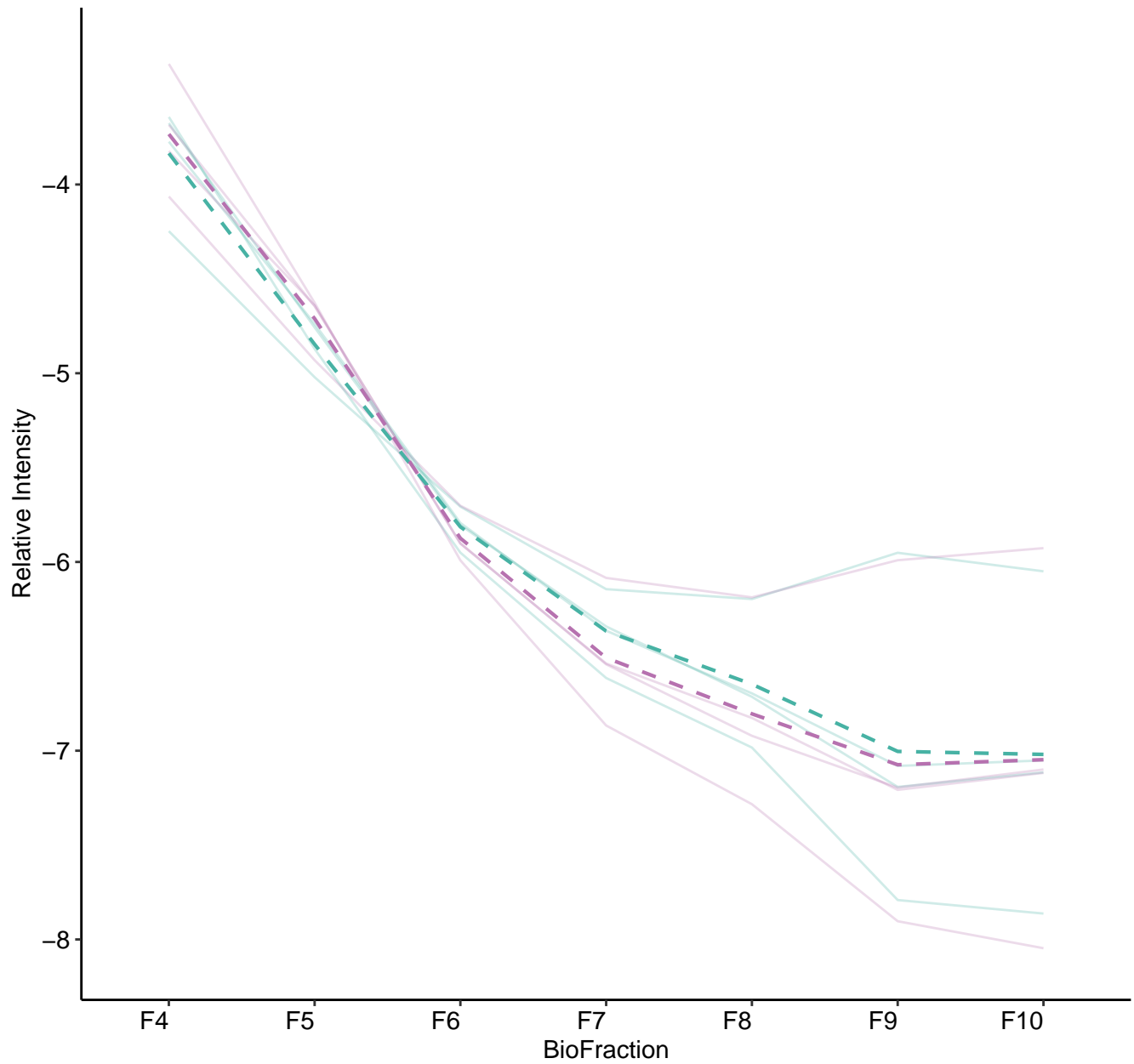
PI4K2A-WASH complex (n = 6)
(R2.Fixef = 0.483)



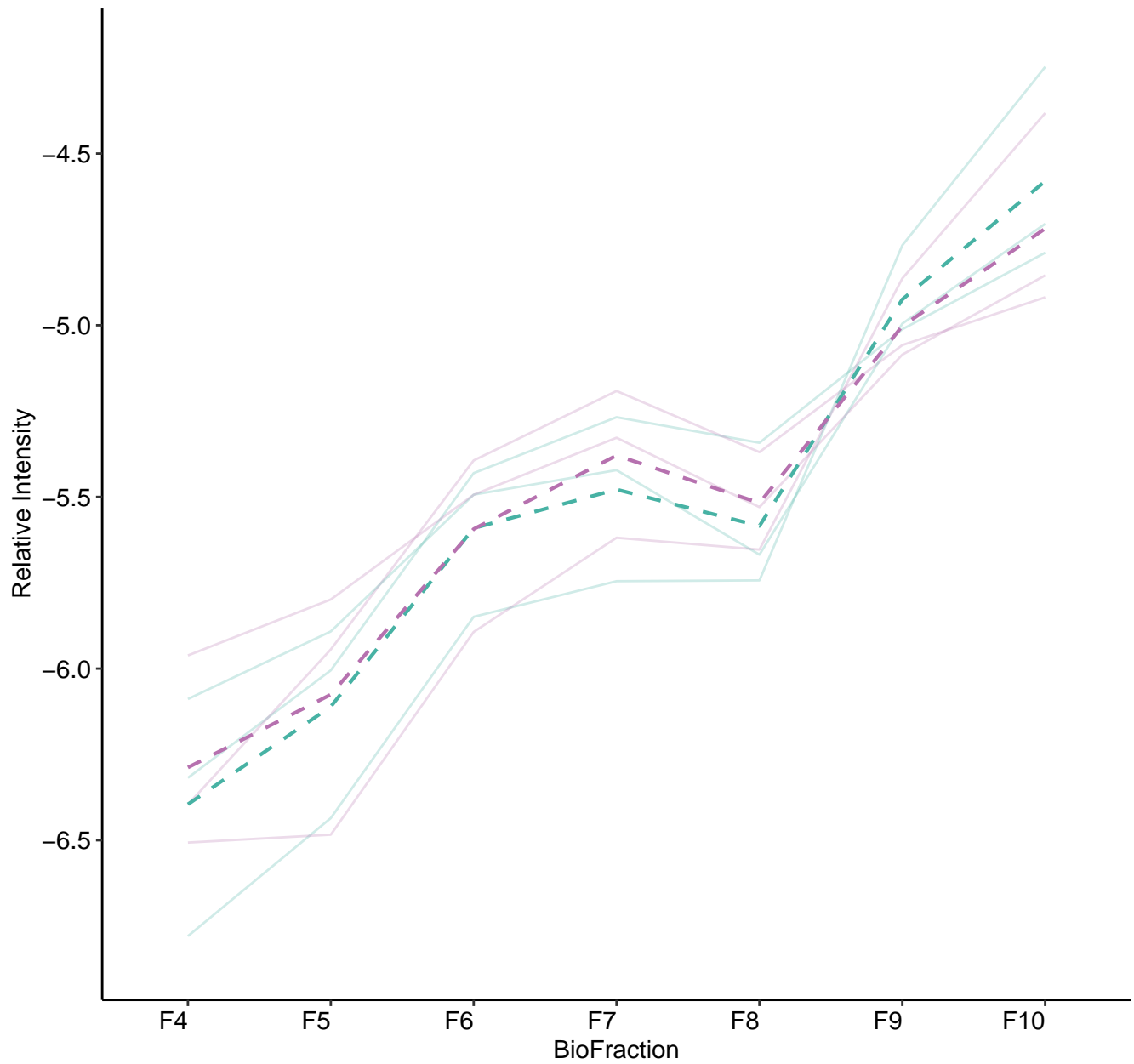
Profilin 1 complex (n = 3)
(R2.Fixef = 0.836)



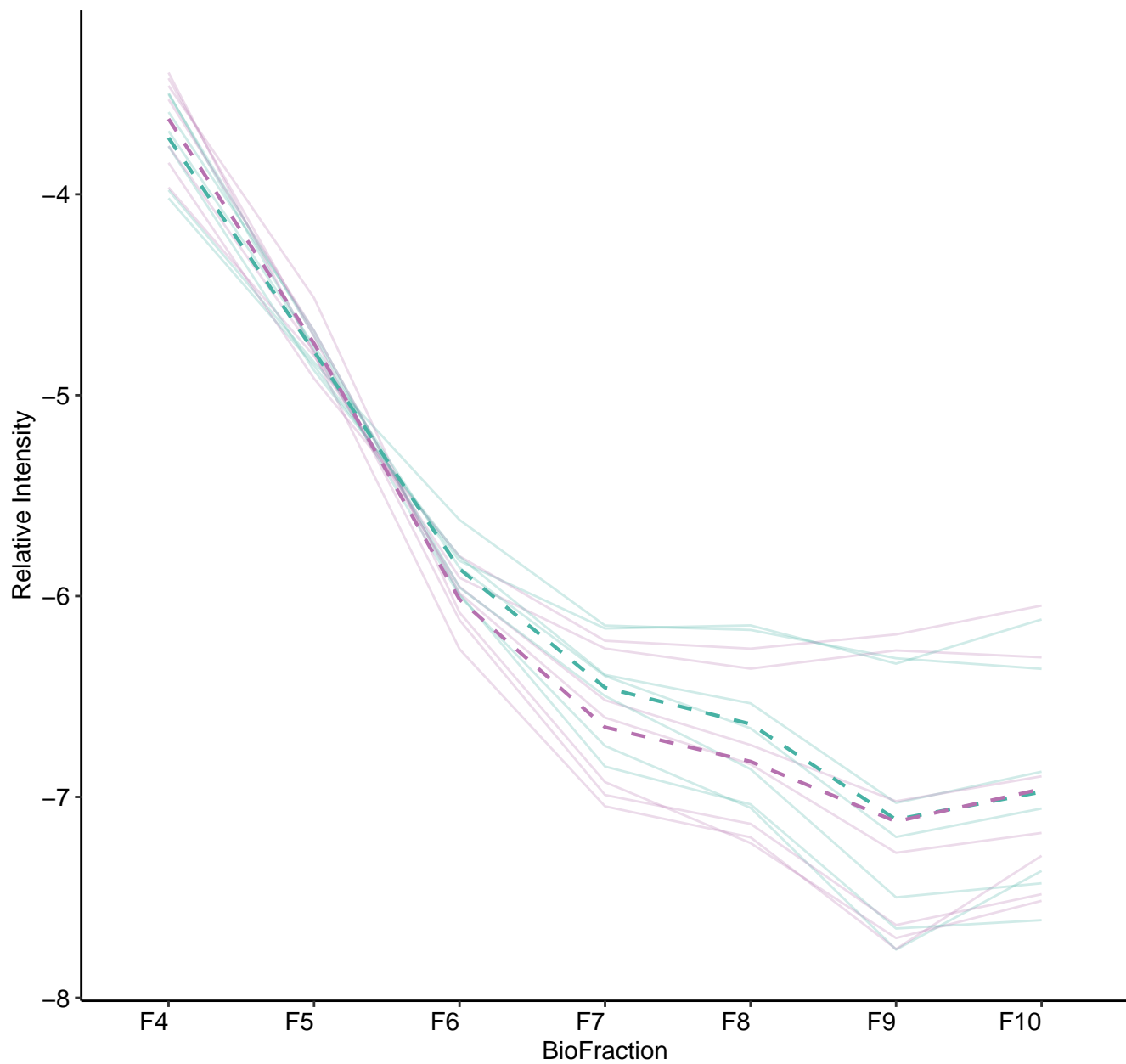
prohibitin 2 complex, mitochondrial (n = 4)
(R2.Fixef = 0.855)



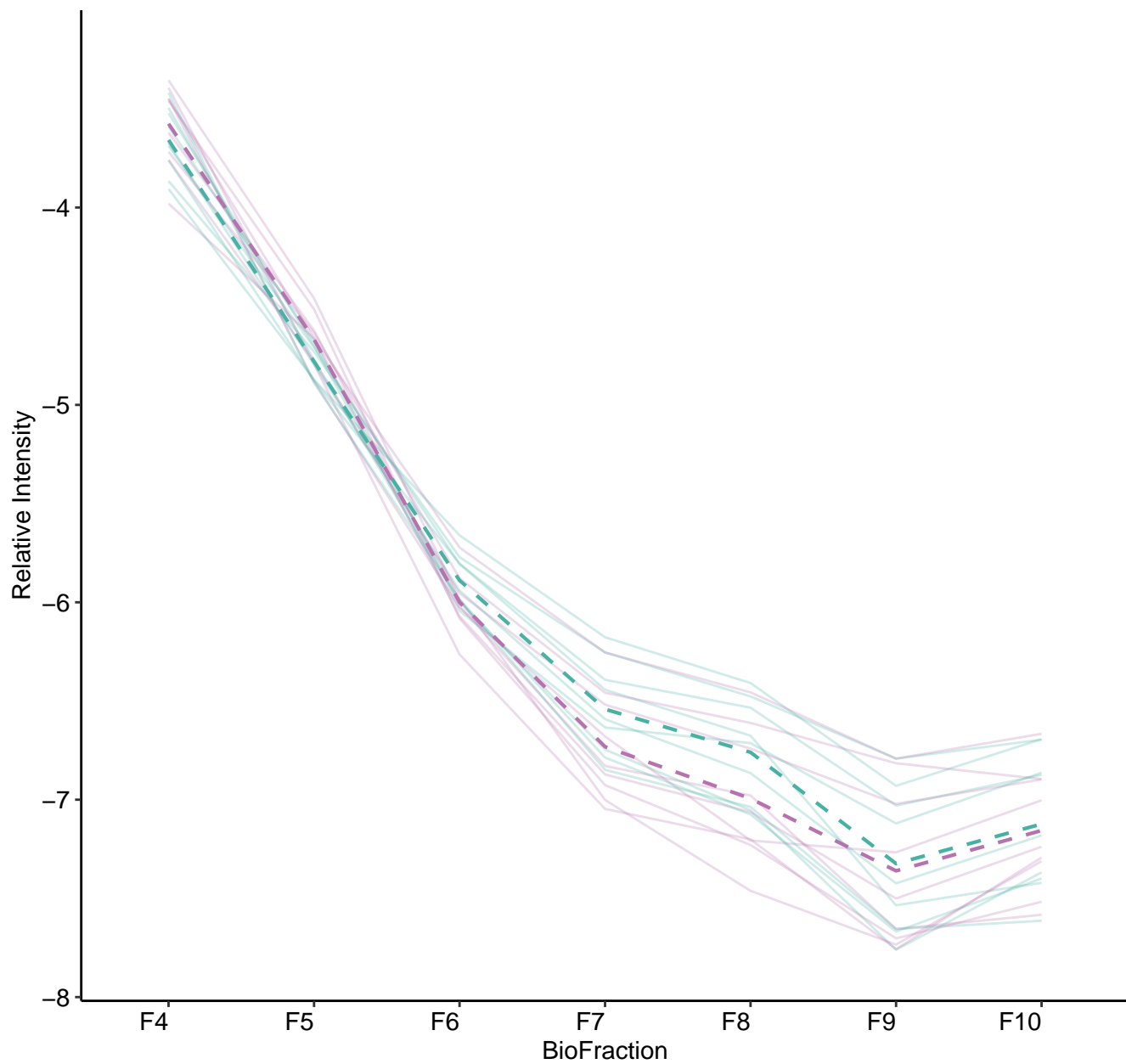
PYR complex (n = 3)
(R2.Fixef = 0.824)



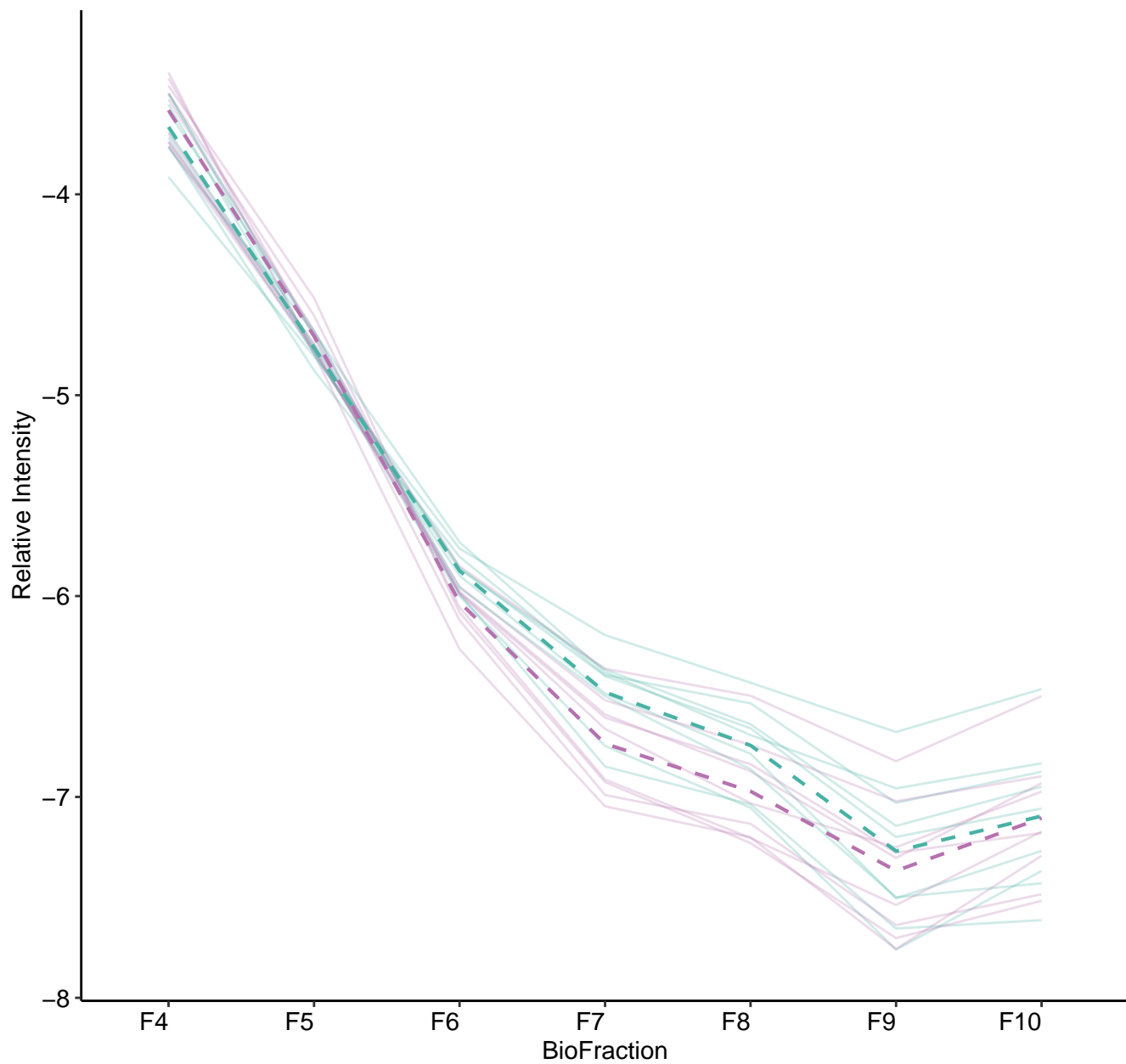
Respiratory chain complex I (early intermediate NDUFAF1 assembly), mitochondrial (n = 7)
(R2.Fixef = 0.906)



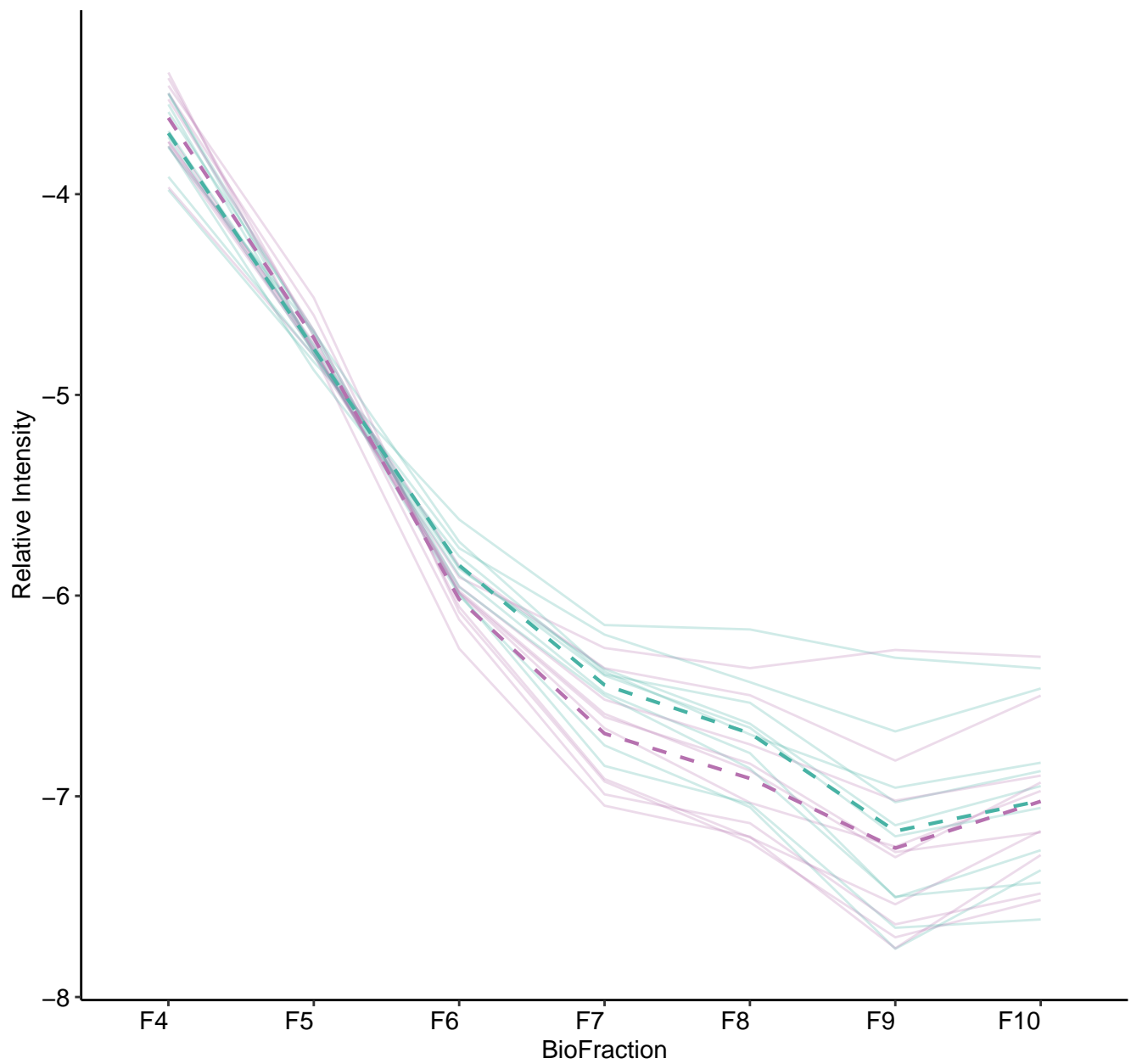
Respiratory chain complex I (gamma subunit) mitochondrial (n = 9)
(R2.Fixef = 0.961)



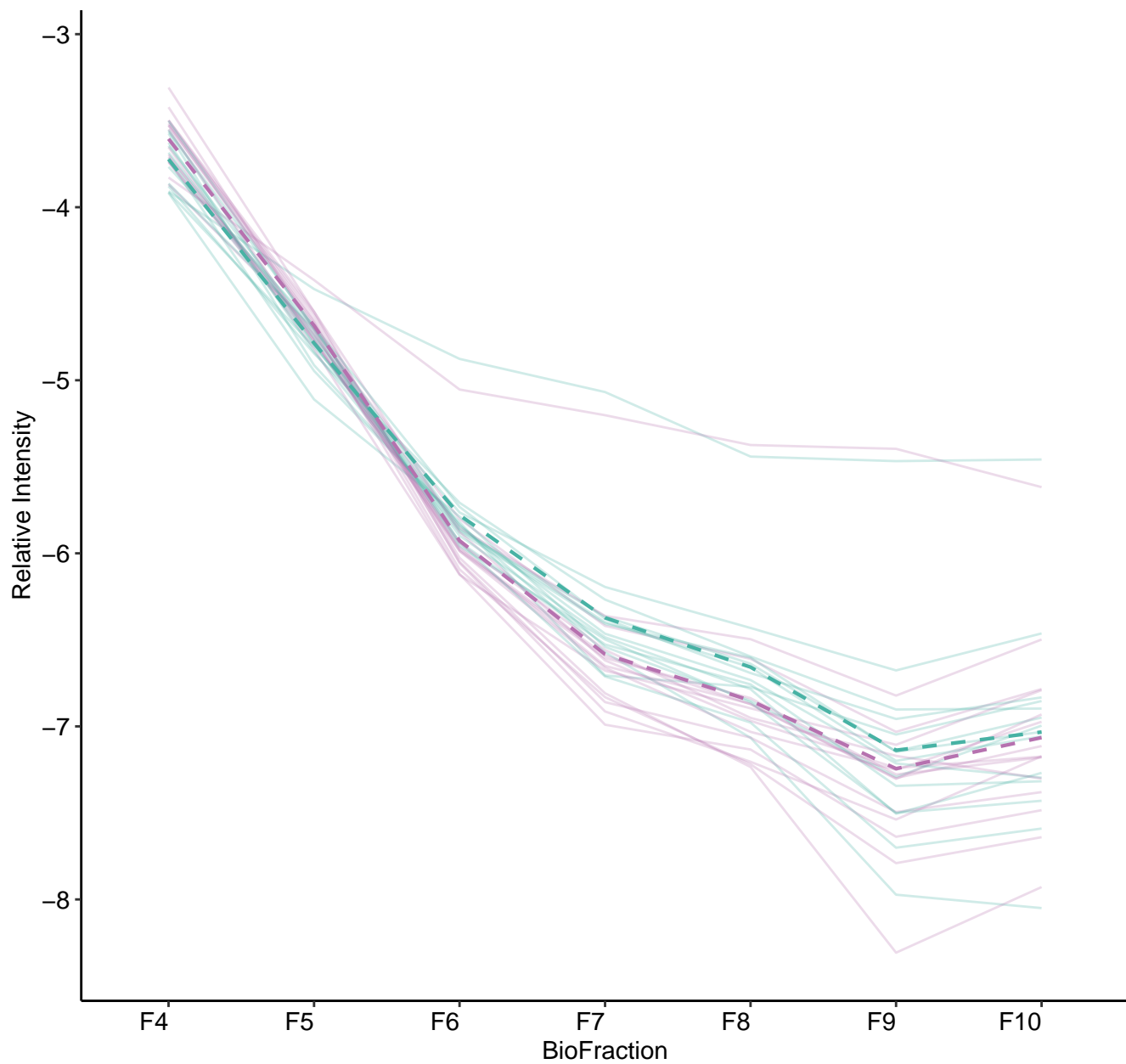
Respiratory chain complex I (intermediate VII/650kD), mitochondrial (n = 9)
(R2.Fixef = 0.968)



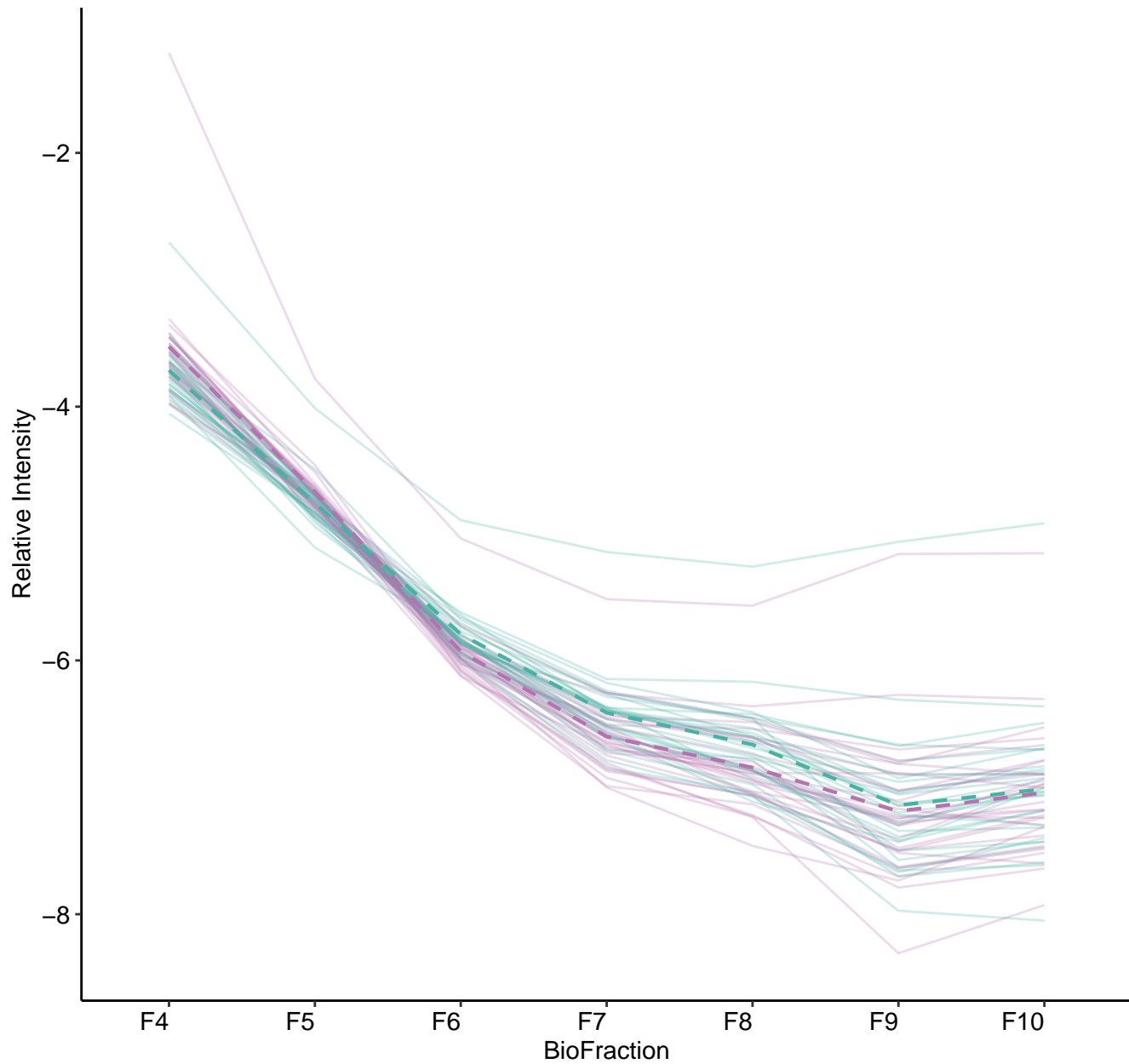
Respiratory chain complex I (intermediate), mitochondrial (n = 10)
(R2.Fixef = 0.95)



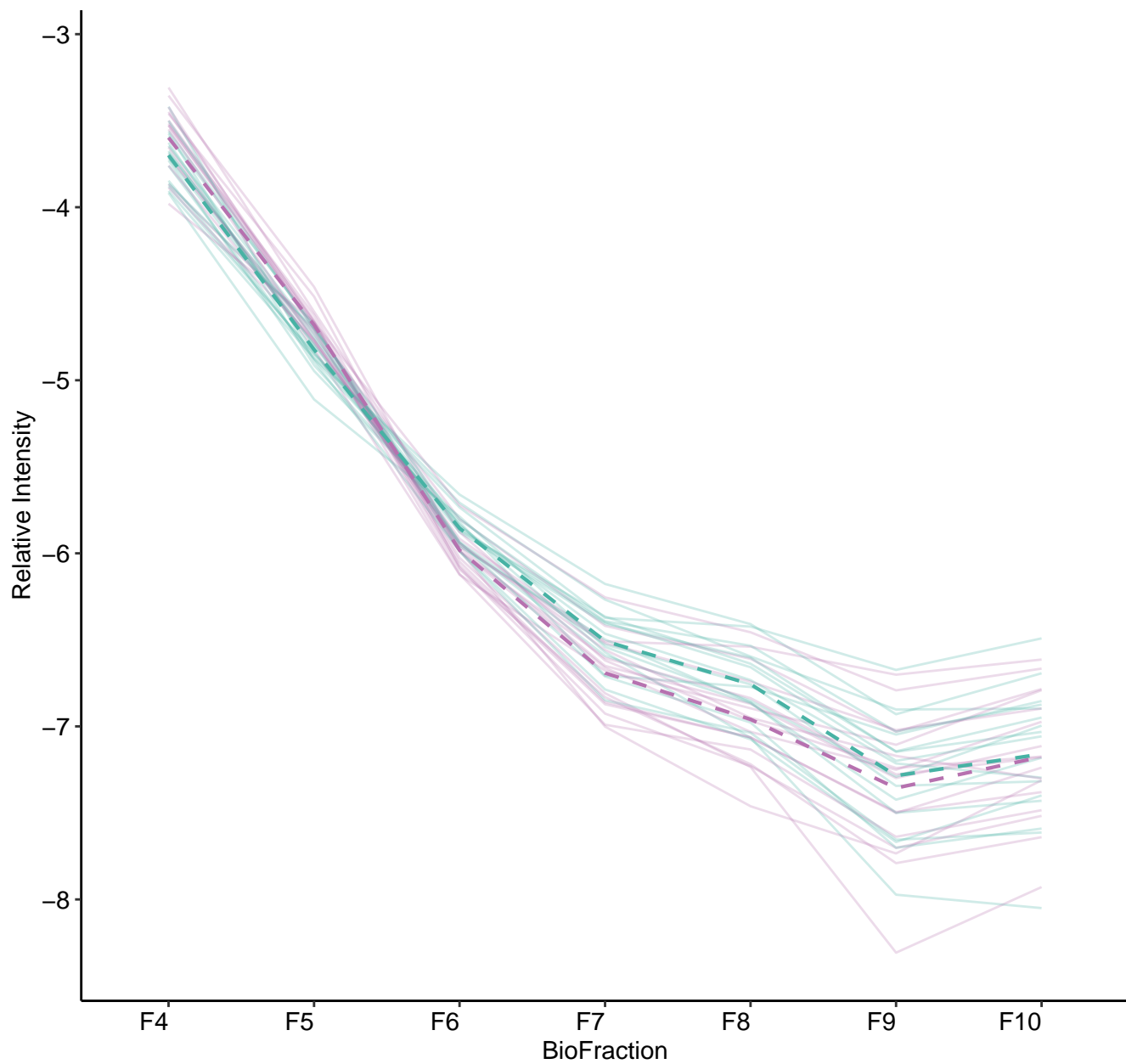
Respiratory chain complex I (lambda subunit) mitochondrial (n = 15)
(R2.Fixef = 0.905)



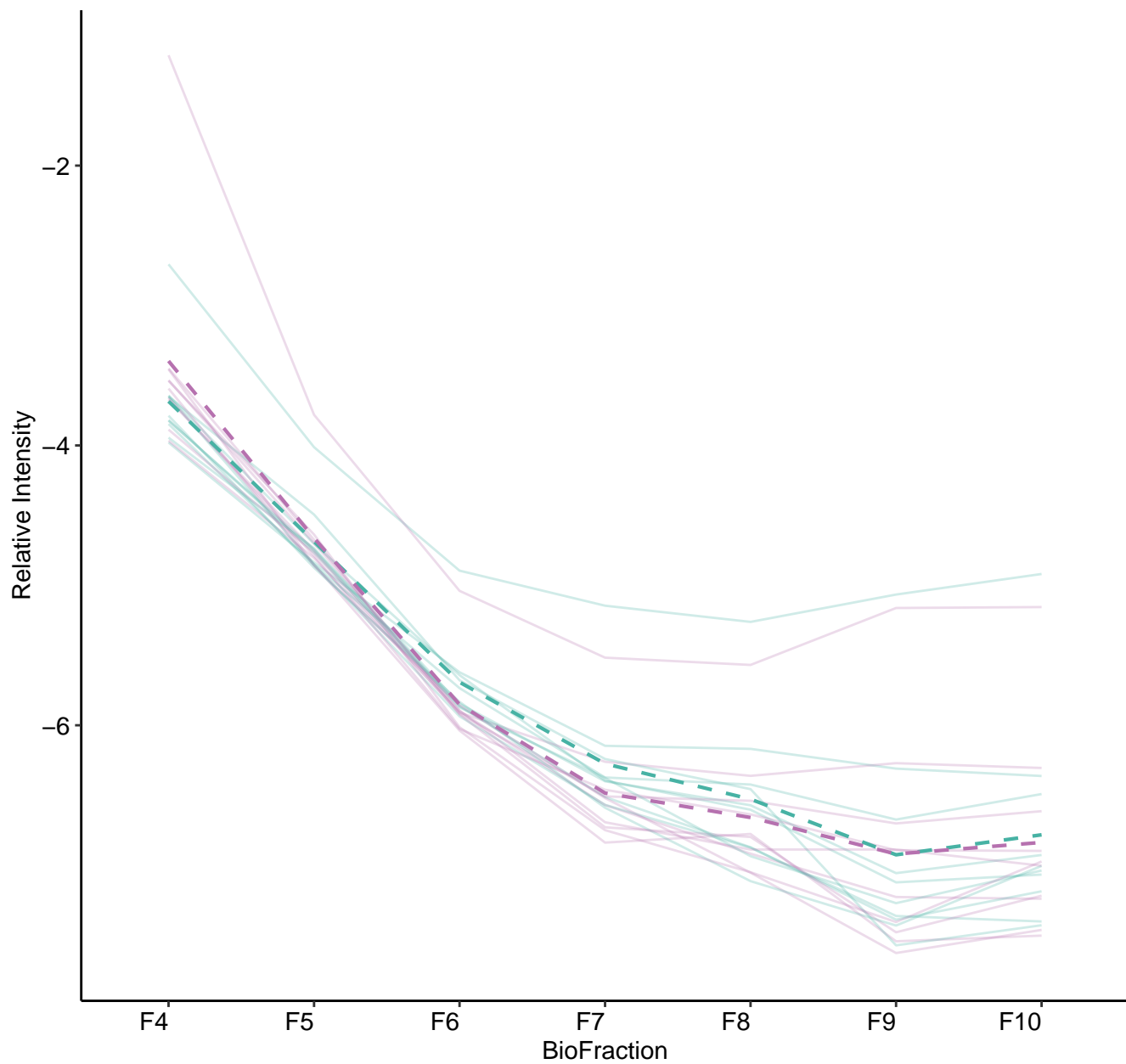
Respiratory chain complex I (nuclear encoded subunits), mitochondrial (n = 29)
(R2.Fixef = 0.913)



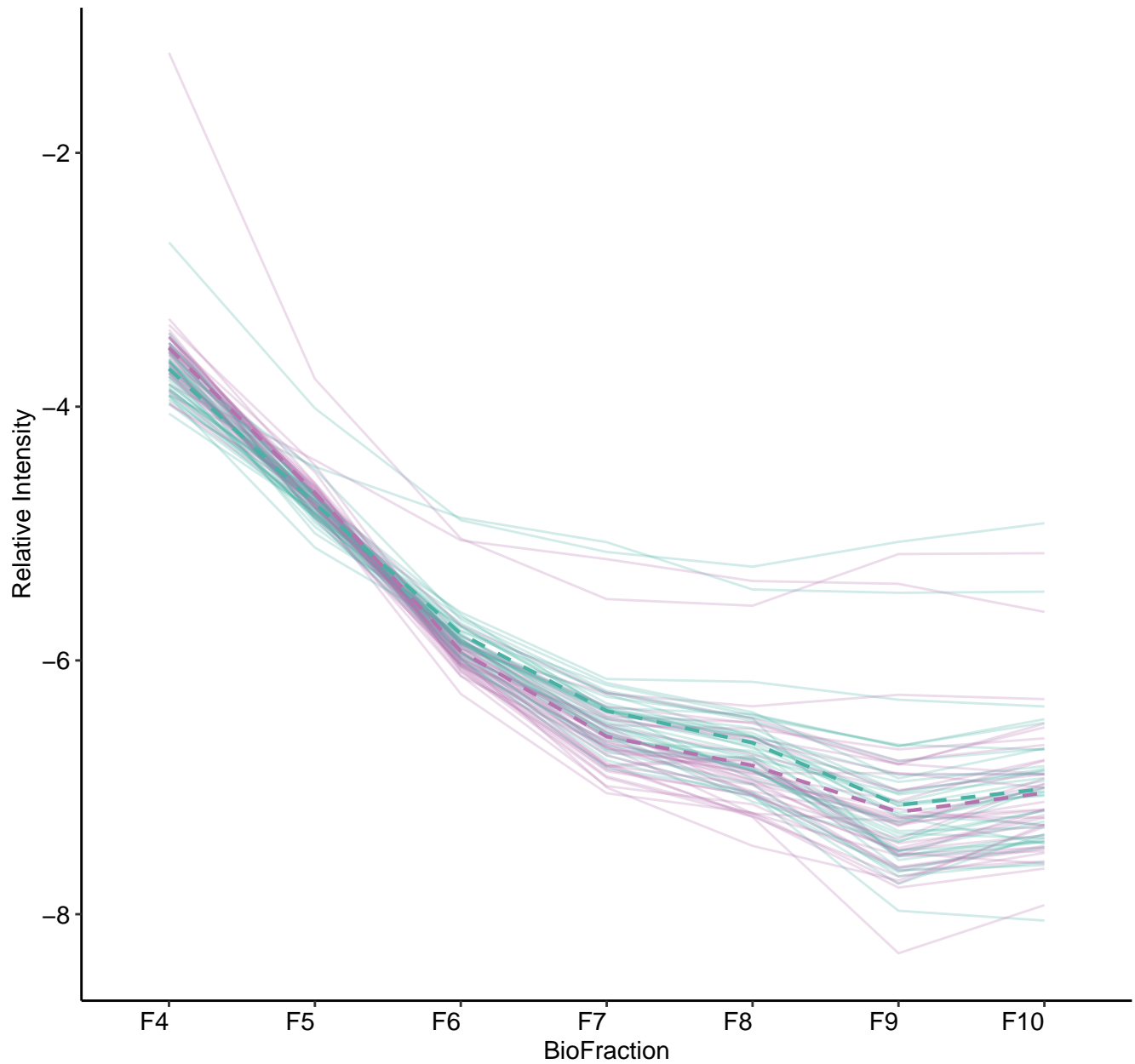
Respiratory chain complex I (subcomplex I alpha), mitochondrial (n = 17)
(R2.Fixef = 0.964)



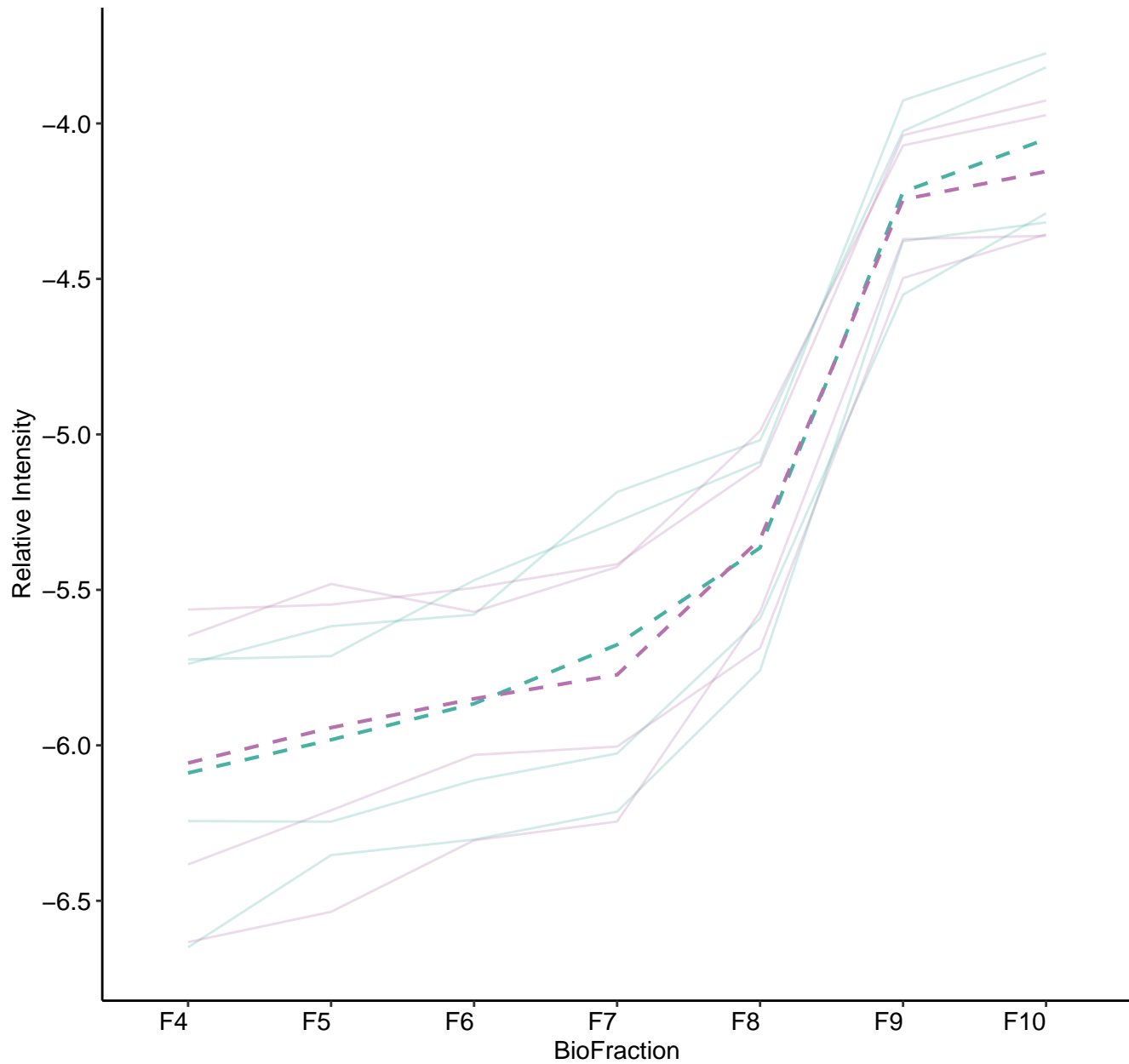
Respiratory chain complex I (subcomplex I beta), mitochondrial (n = 10)
(R2.Fixef = 0.827)



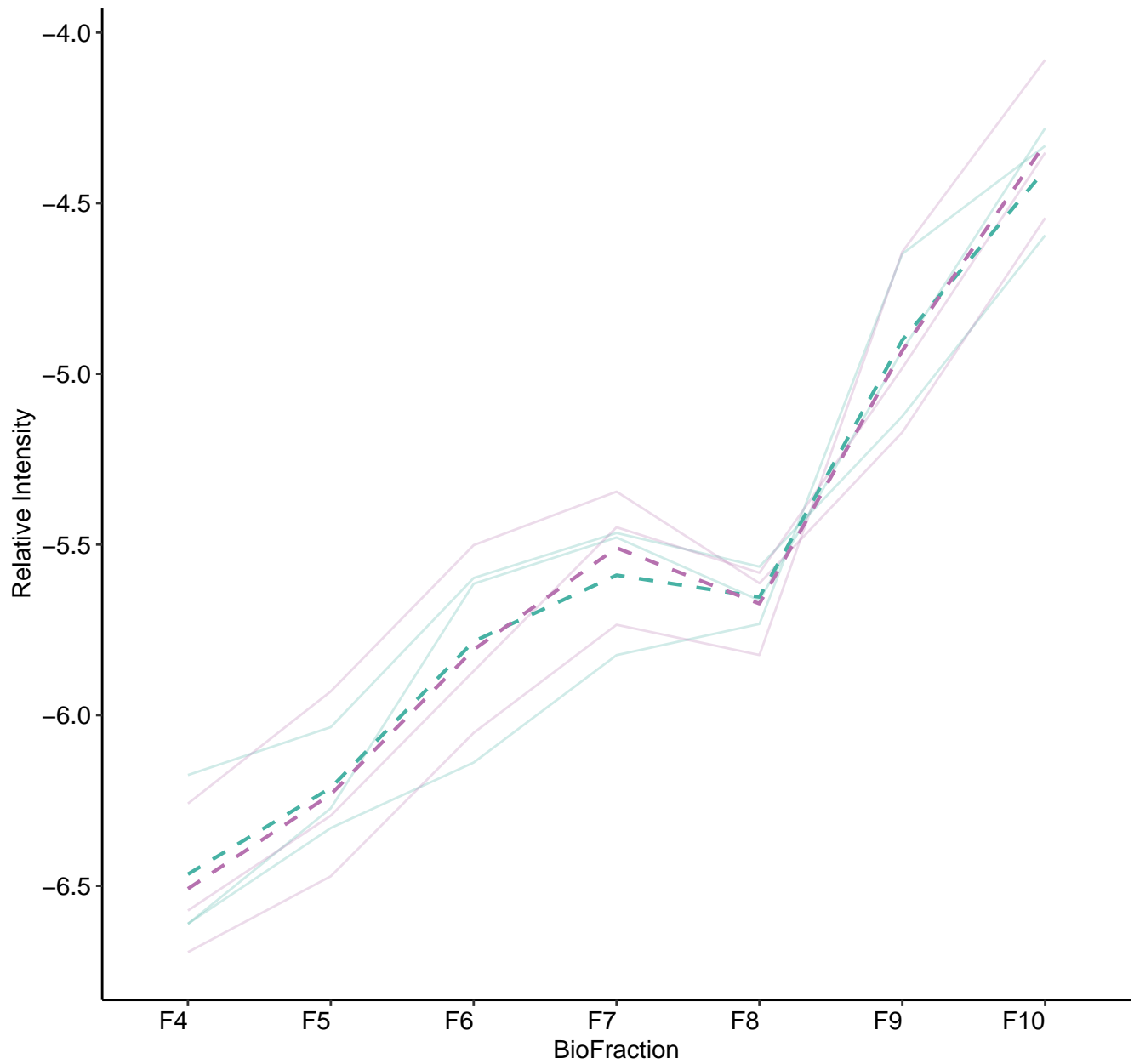
Respiratory chain complex I, mitochondrial (n = 38)
(R2.Fixef = 0.904)



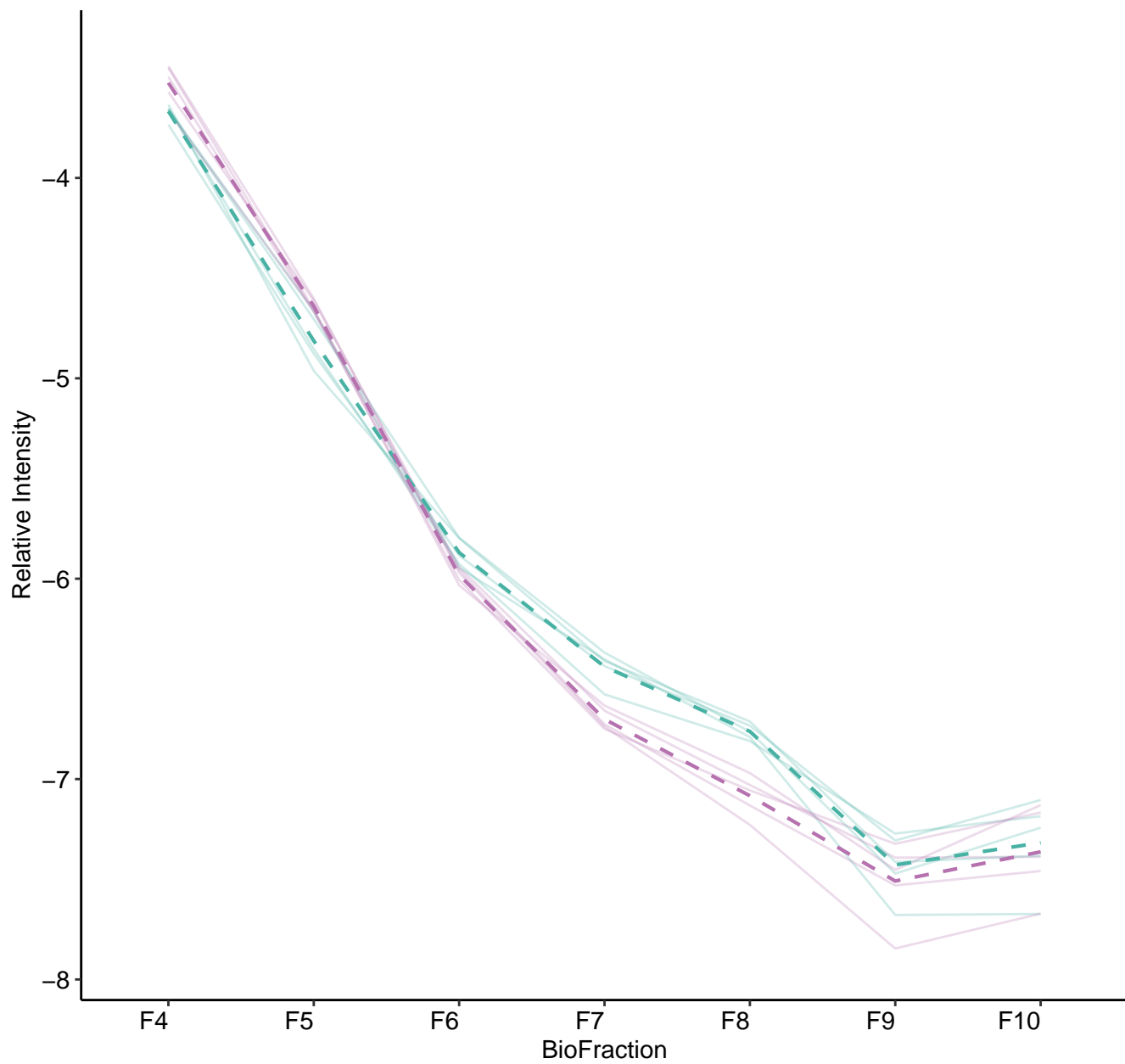
RNA polymerase II complex, chromatin structure modifying (n = 4)
(R2.Fixef = 0.794)



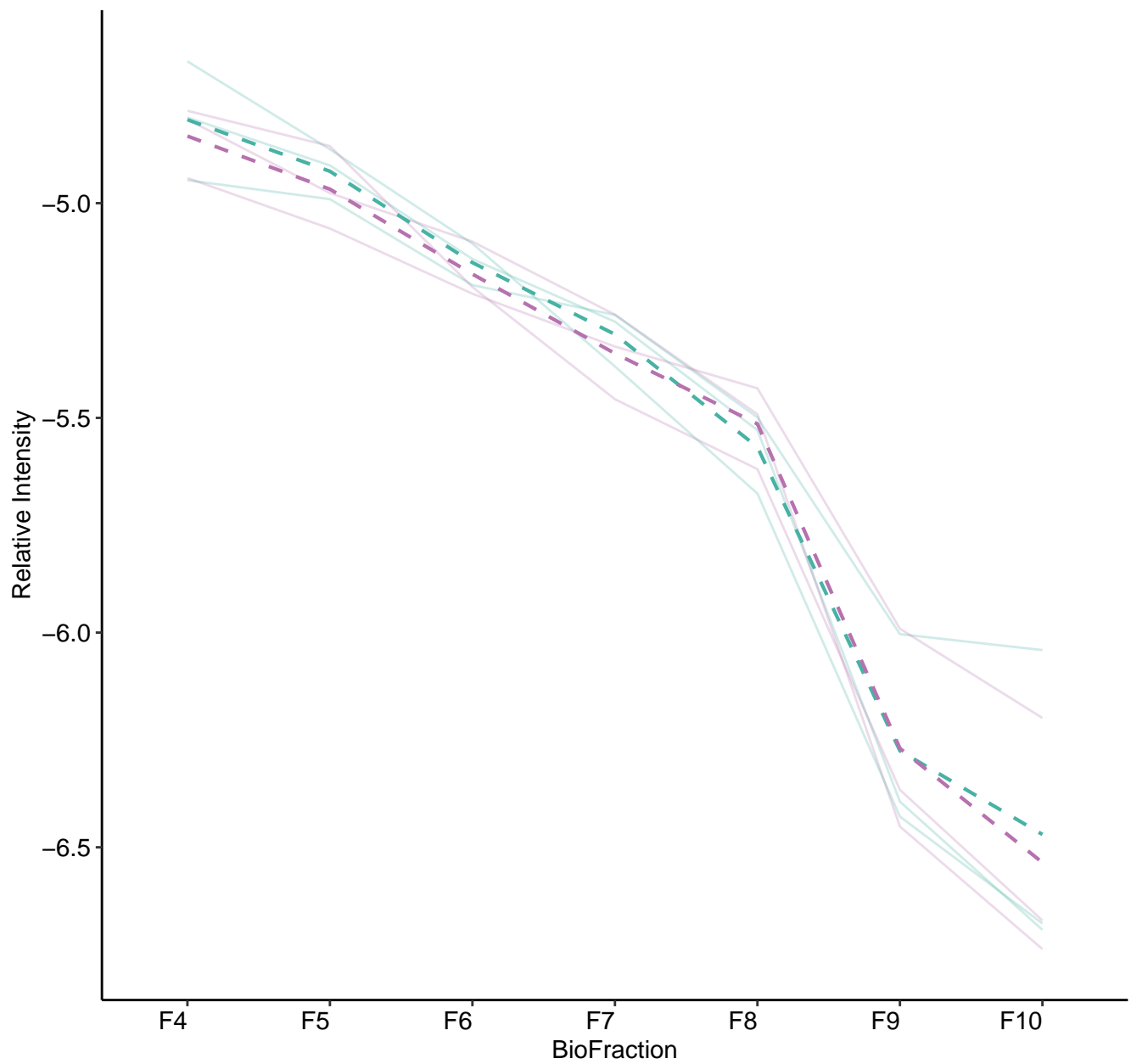
RNA polymerase II holoenzyme complex (n = 3)
(R2.Fixef = 0.903)



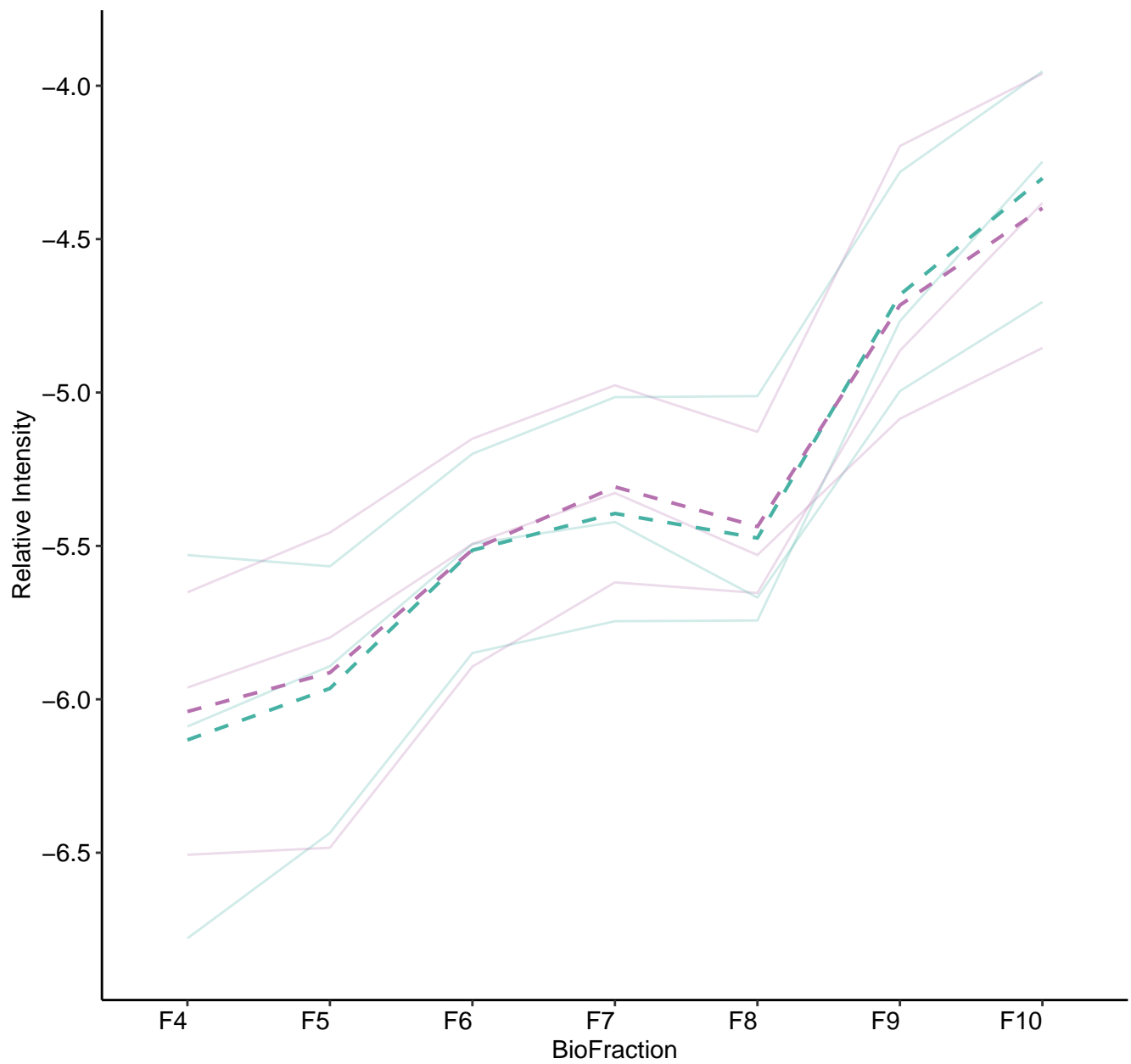
SDH-mABC1-PIC-ANT-ATPase complex (n = 5)
(R2.Fixef = 0.991)



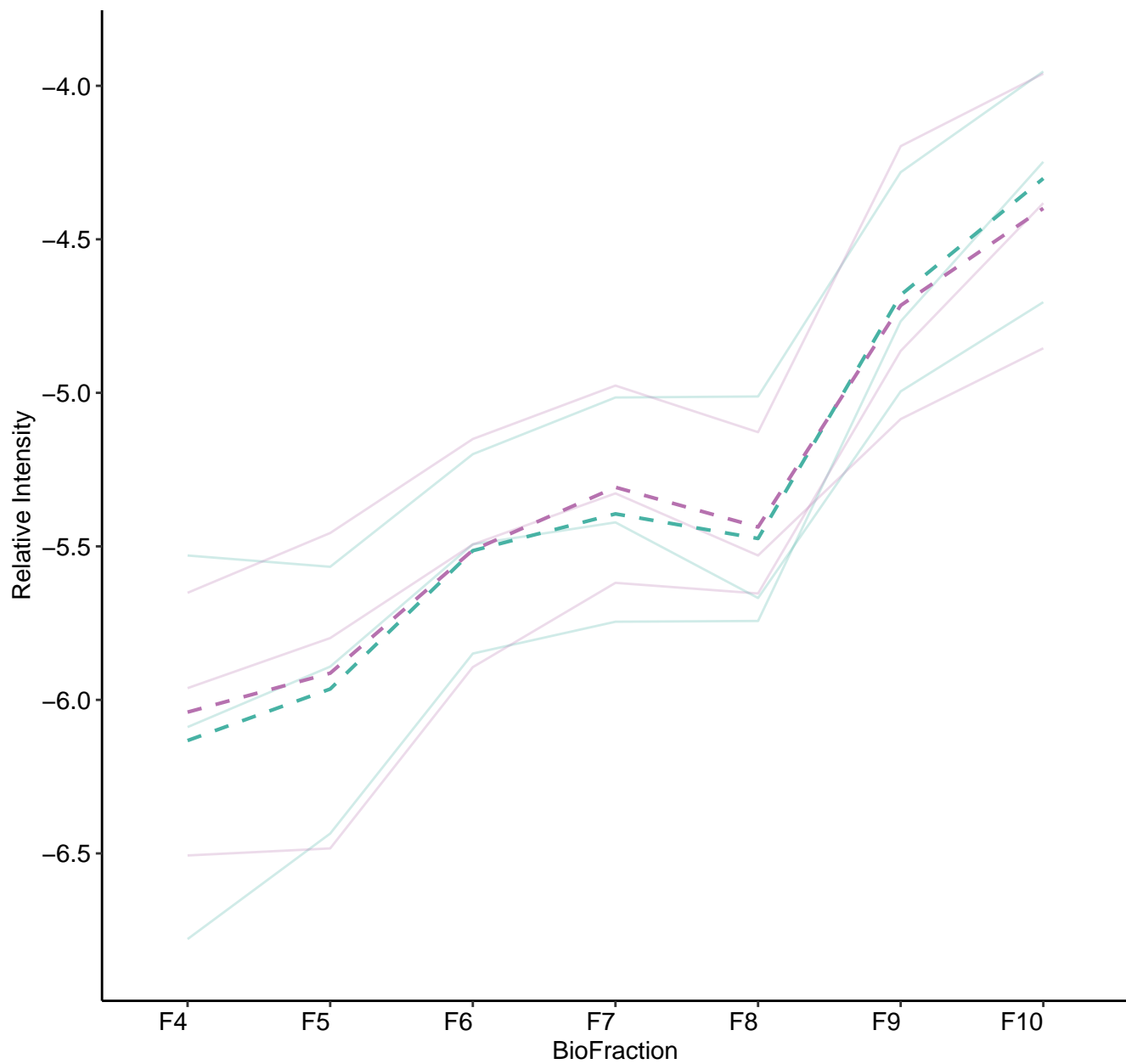
Septin complex (n = 3)
(R2.Fixef = 0.925)



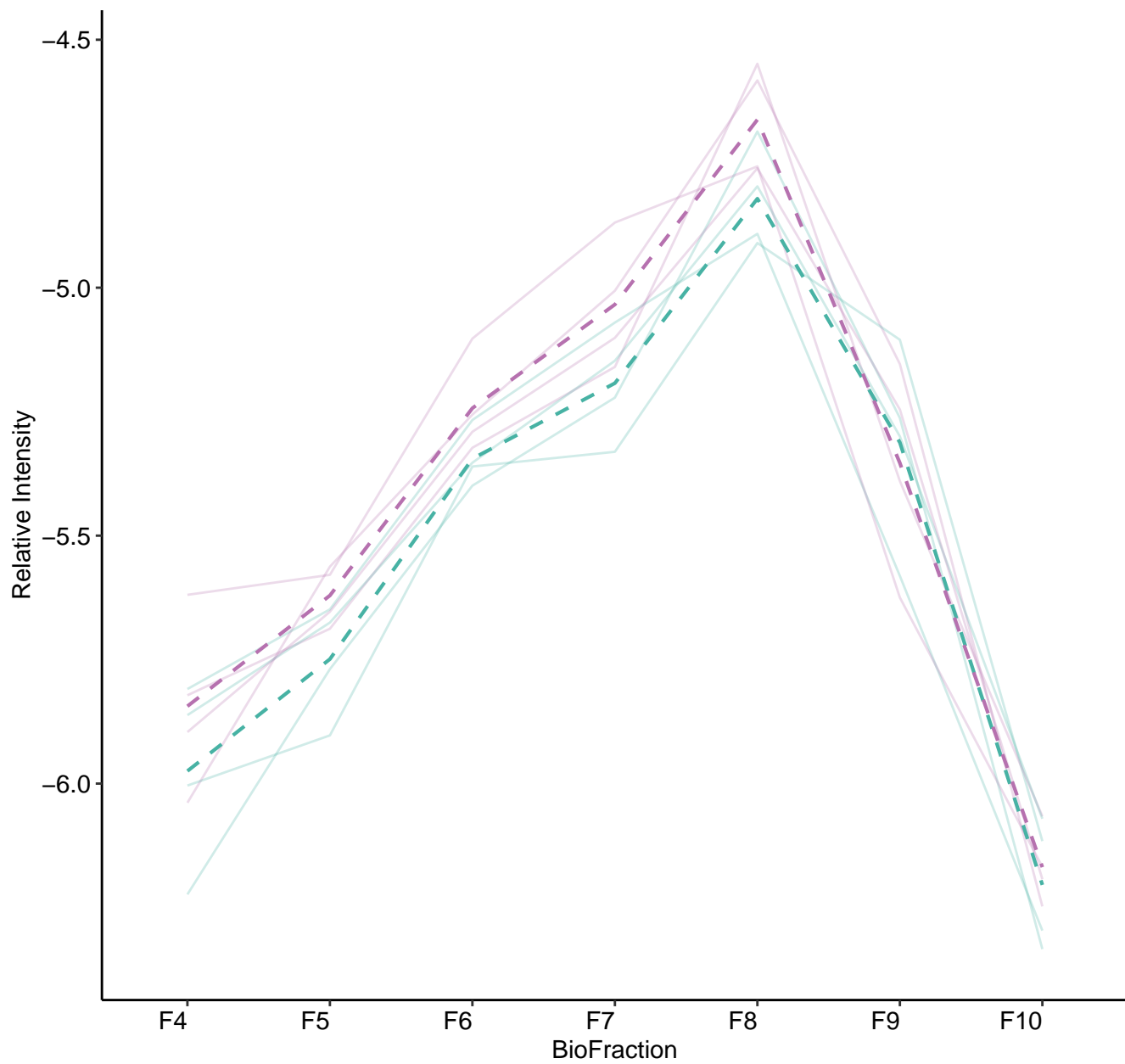
SIN3 complex (n = 3)
(R2.Fixef = 0.663)



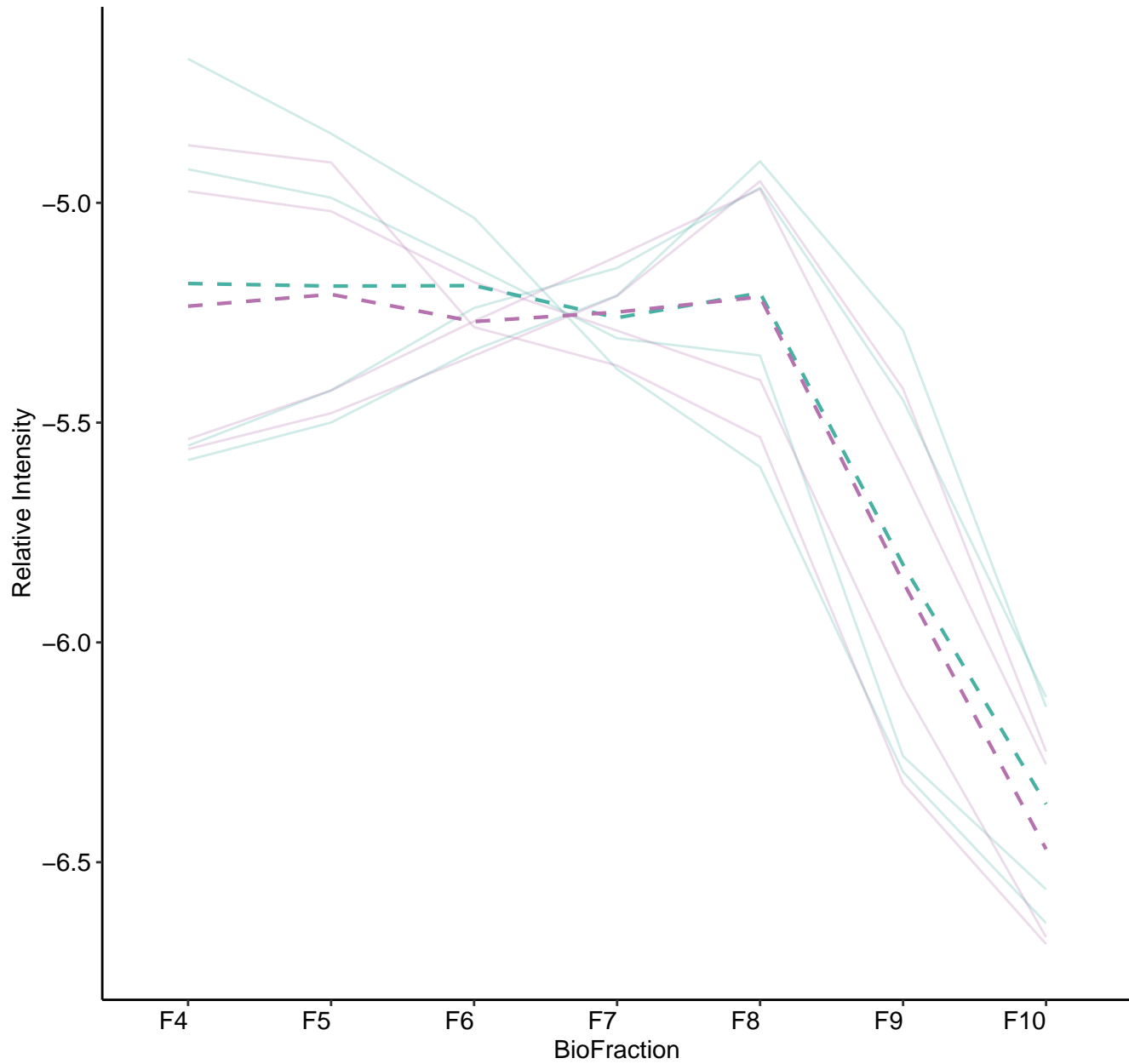
SIN3-HDAC-SAP30-ARID4 complex (n = 3)
(R2.Fixef = 0.663)



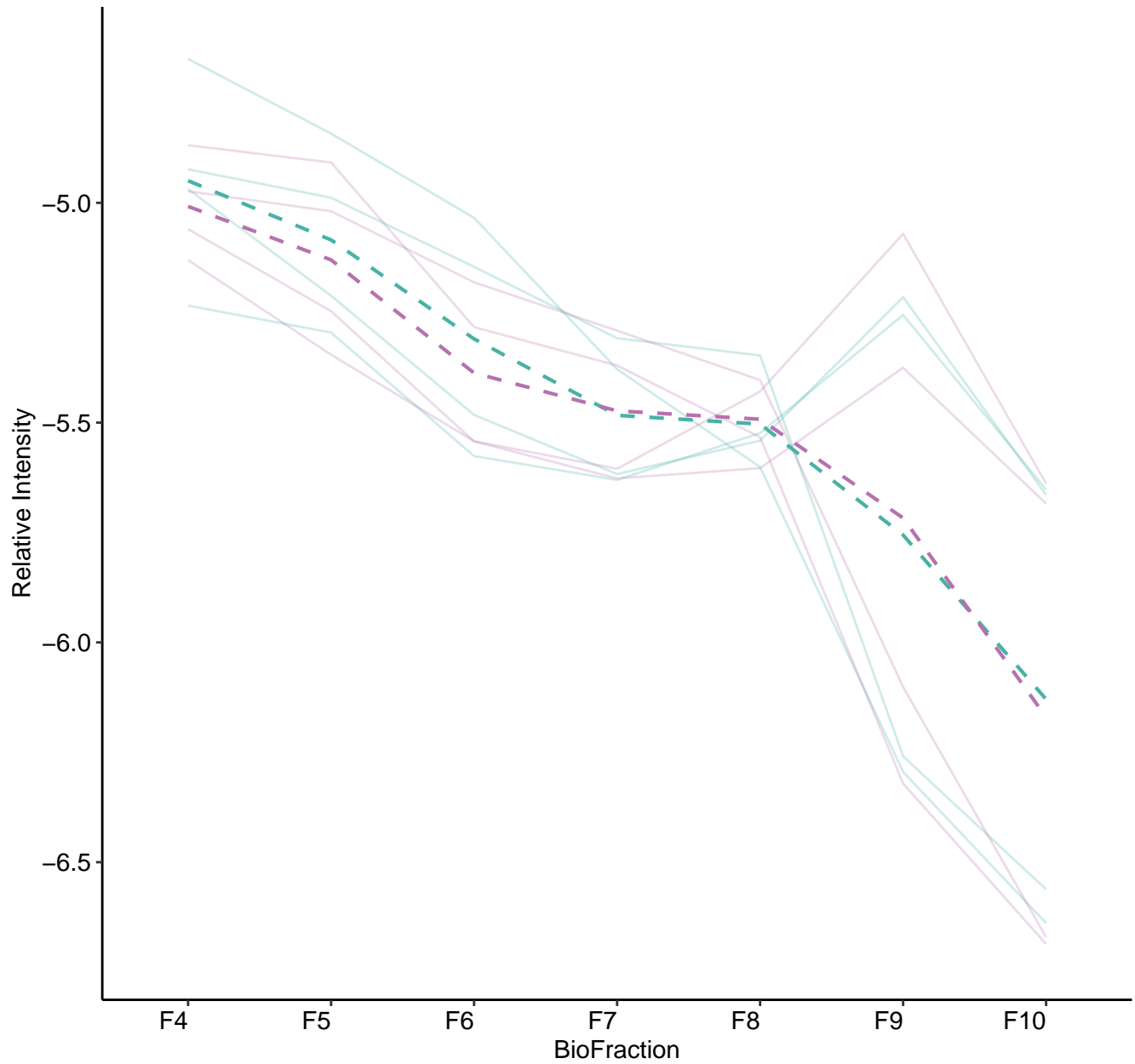
SNARE complex (Bet1, Gosr1, Gosr2, Scfd1, Stx5, Sec22b) (n = 4)
(R2.Fixef = 0.924)



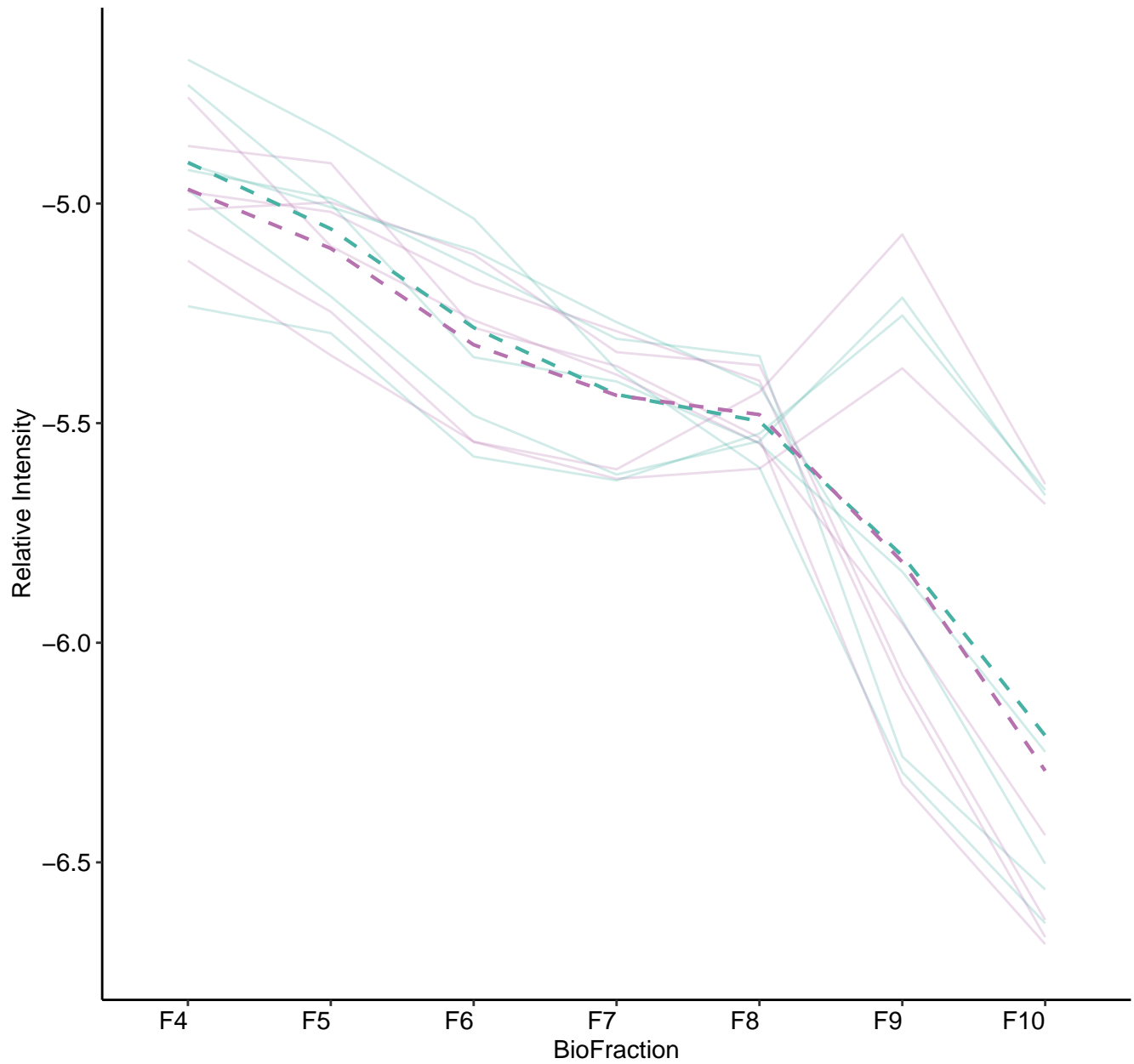
SNARE complex (Stx1a, Napa, Snap25, Vamp2, Nsf) (n = 4)
(R2.Fixef = 0.671)



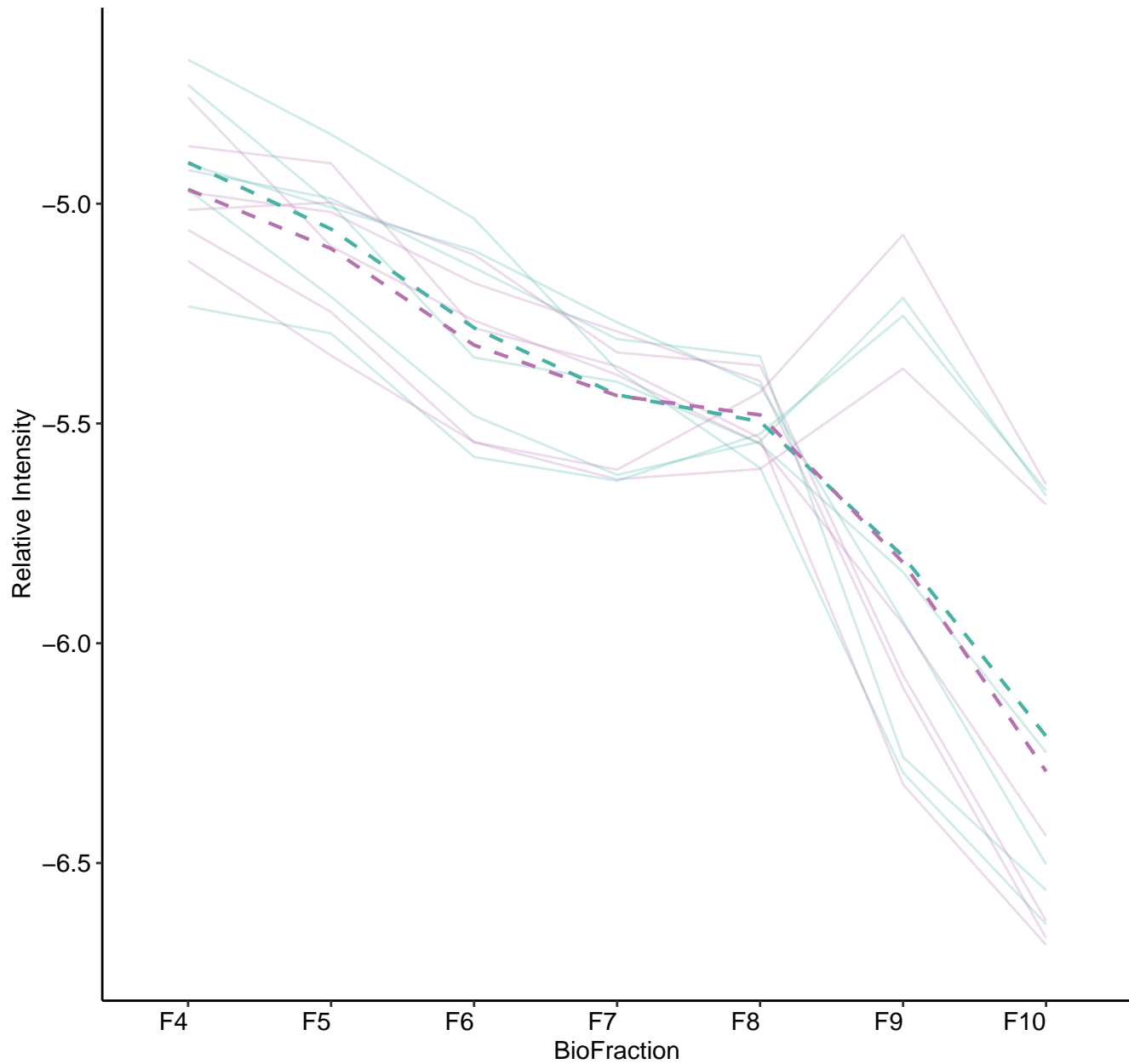
SNARE complex (Stx1a, Snap25, Vamp2, Syt1) (n = 4)
(R2.Fixef = 0.529)



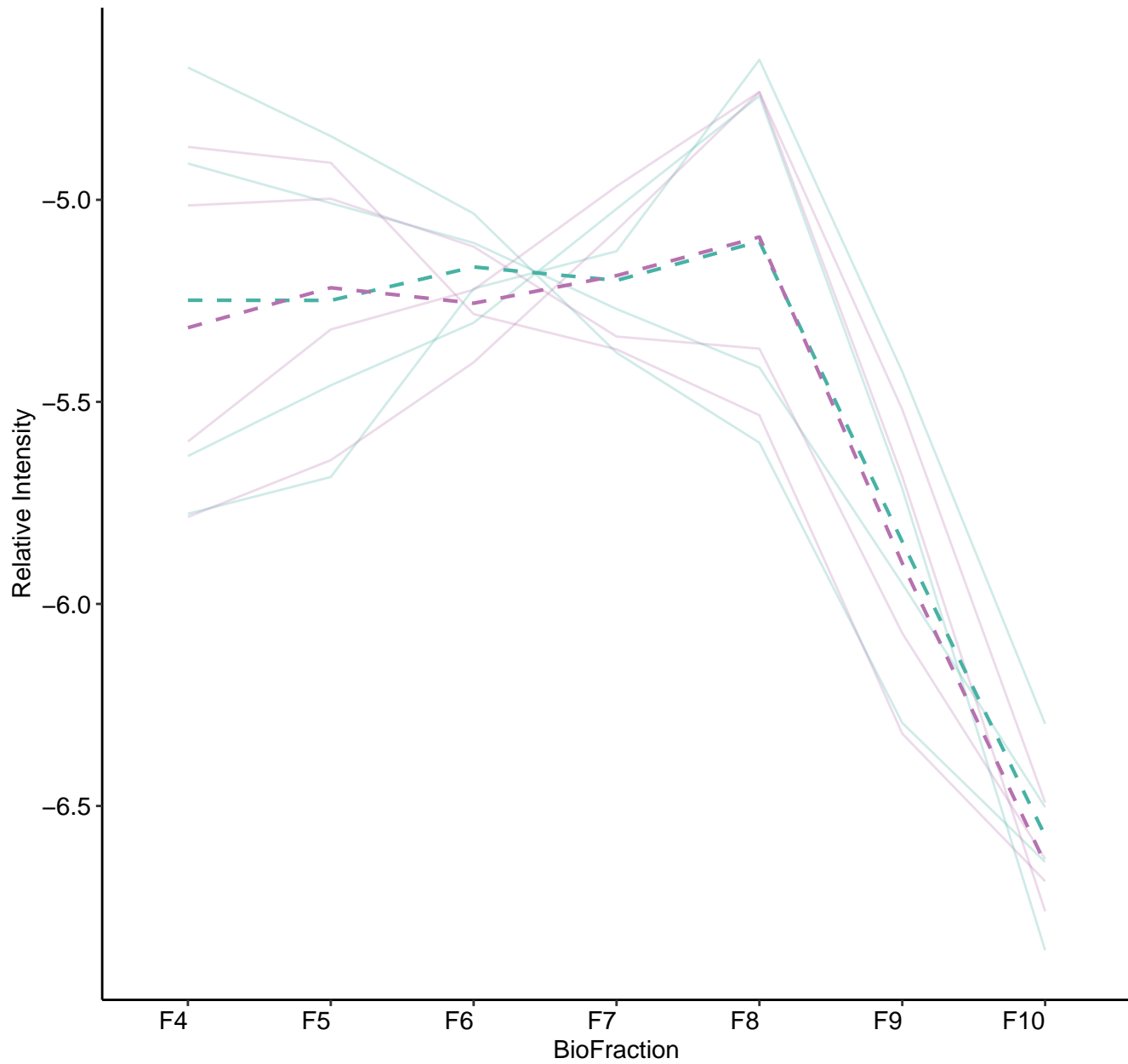
SNARE complex (STX1a, STX1b, SNAP25, RAB3a, SYT1, VAMP2, CPLX2) (n = 6)
(R2.Fixef = 0.676)



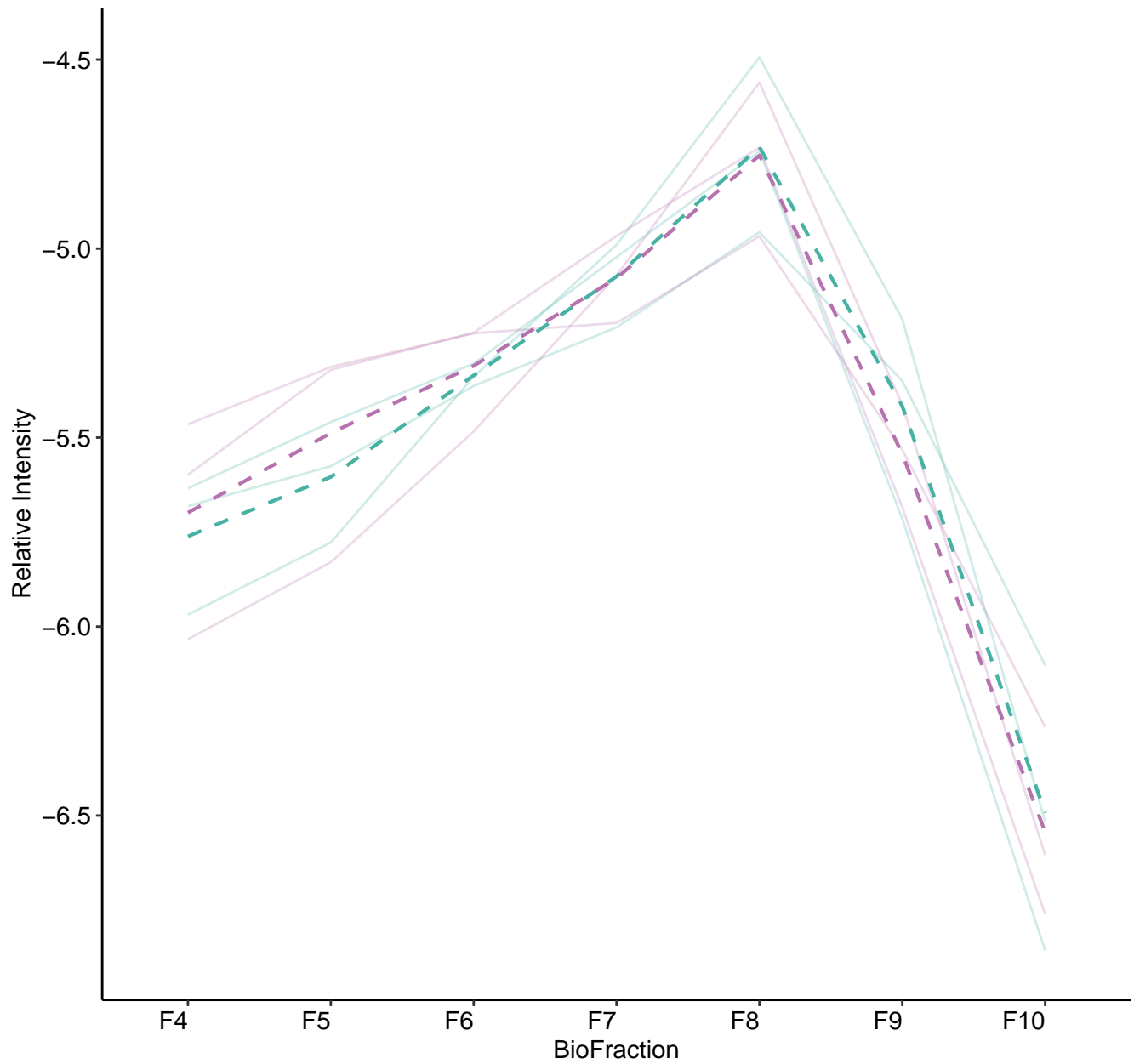
SNARE complex (STX1A, STX1B, SNAP25, RAB3A, SYT1, VAMP2) (n = 6)
(R2.Fixef = 0.676)



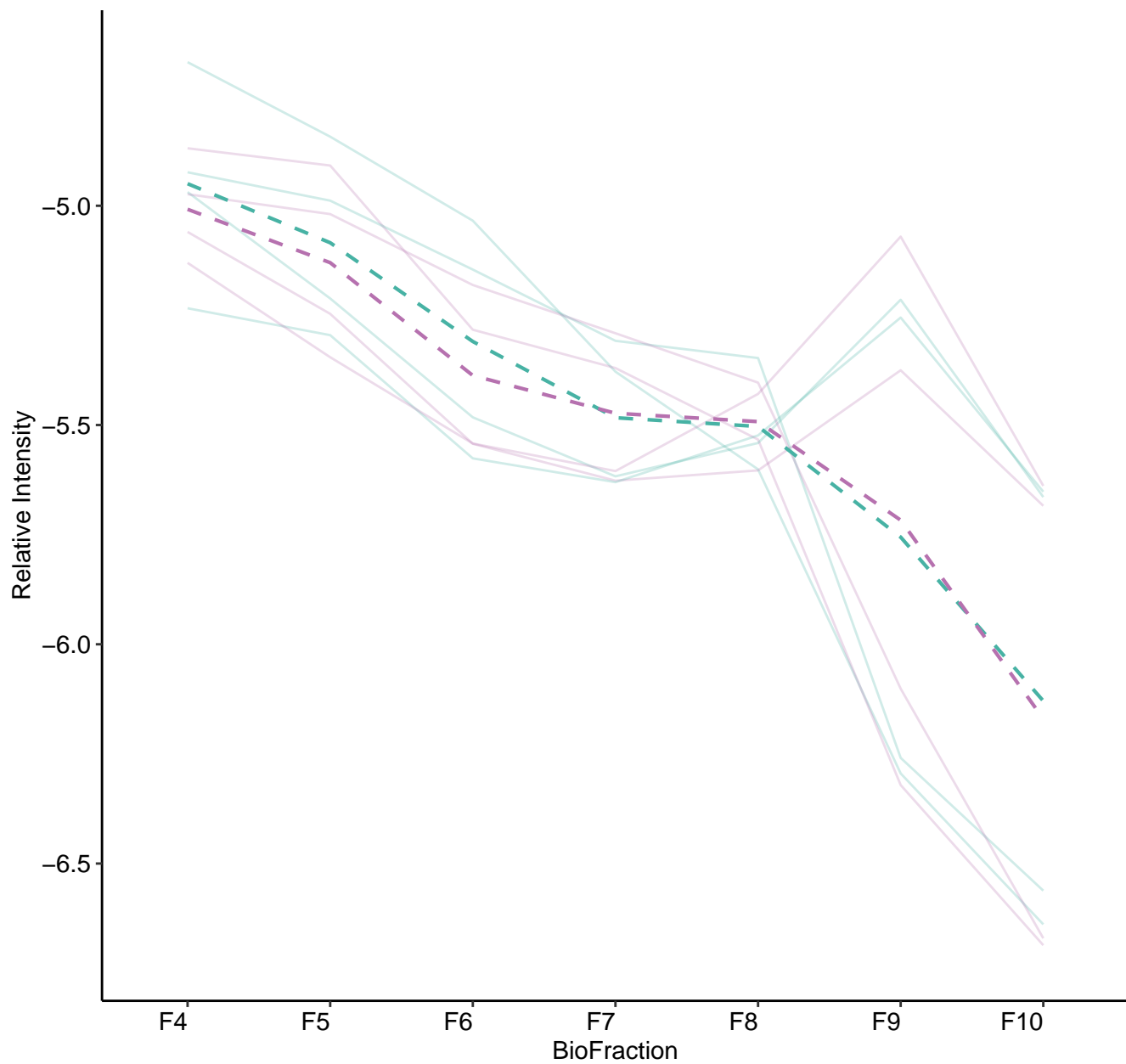
SNARE complex (Stx1a,1b,7,13, Rtn1, Vamp2) (n = 4)
(R2.Fixef = 0.702)



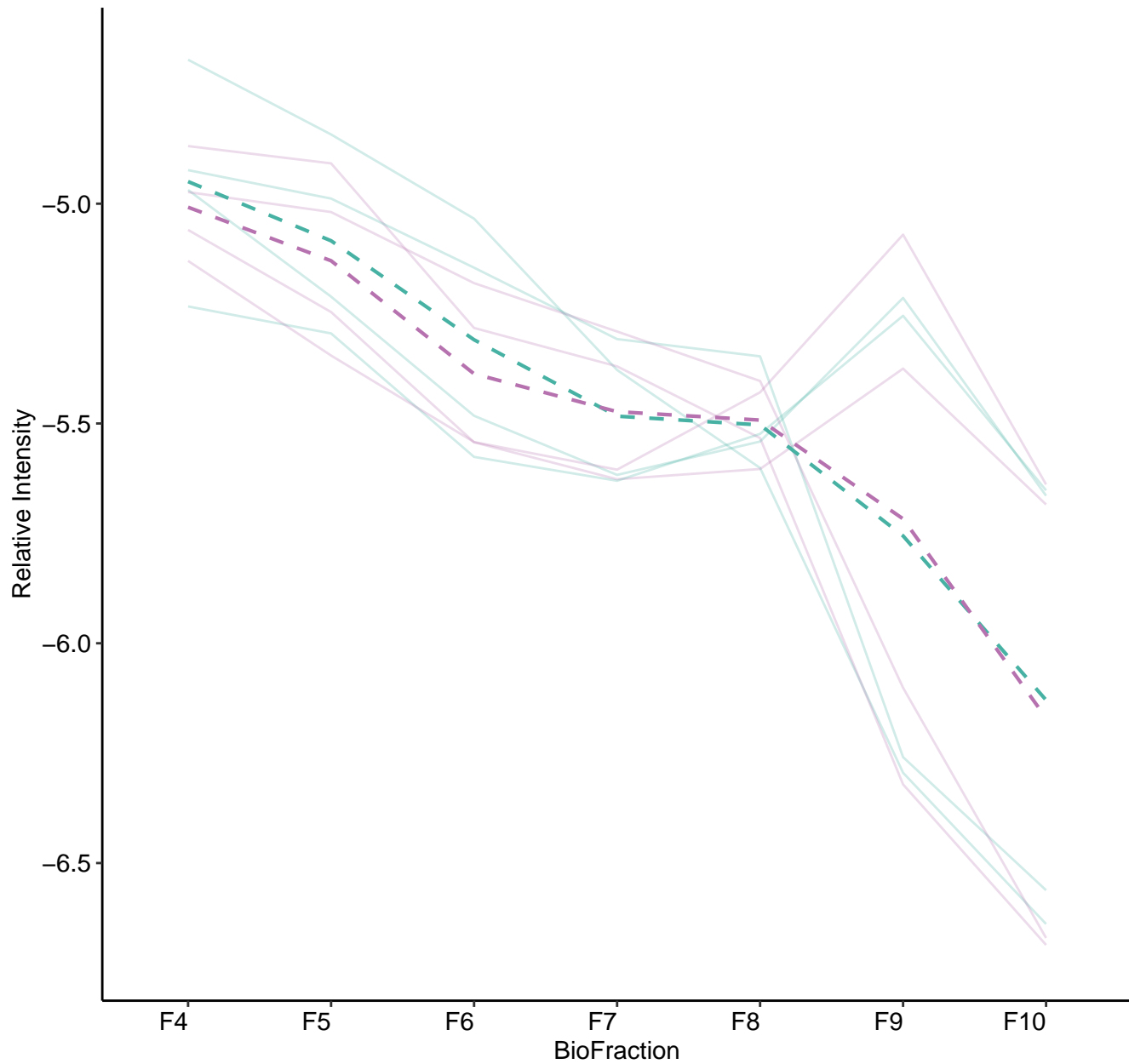
SNARE complex (Stx4, Stx6, Stx7, Vamp3, Vamp7, Vamp8, Vti1b) (n = 3)
(R2.Fixef = 0.85)



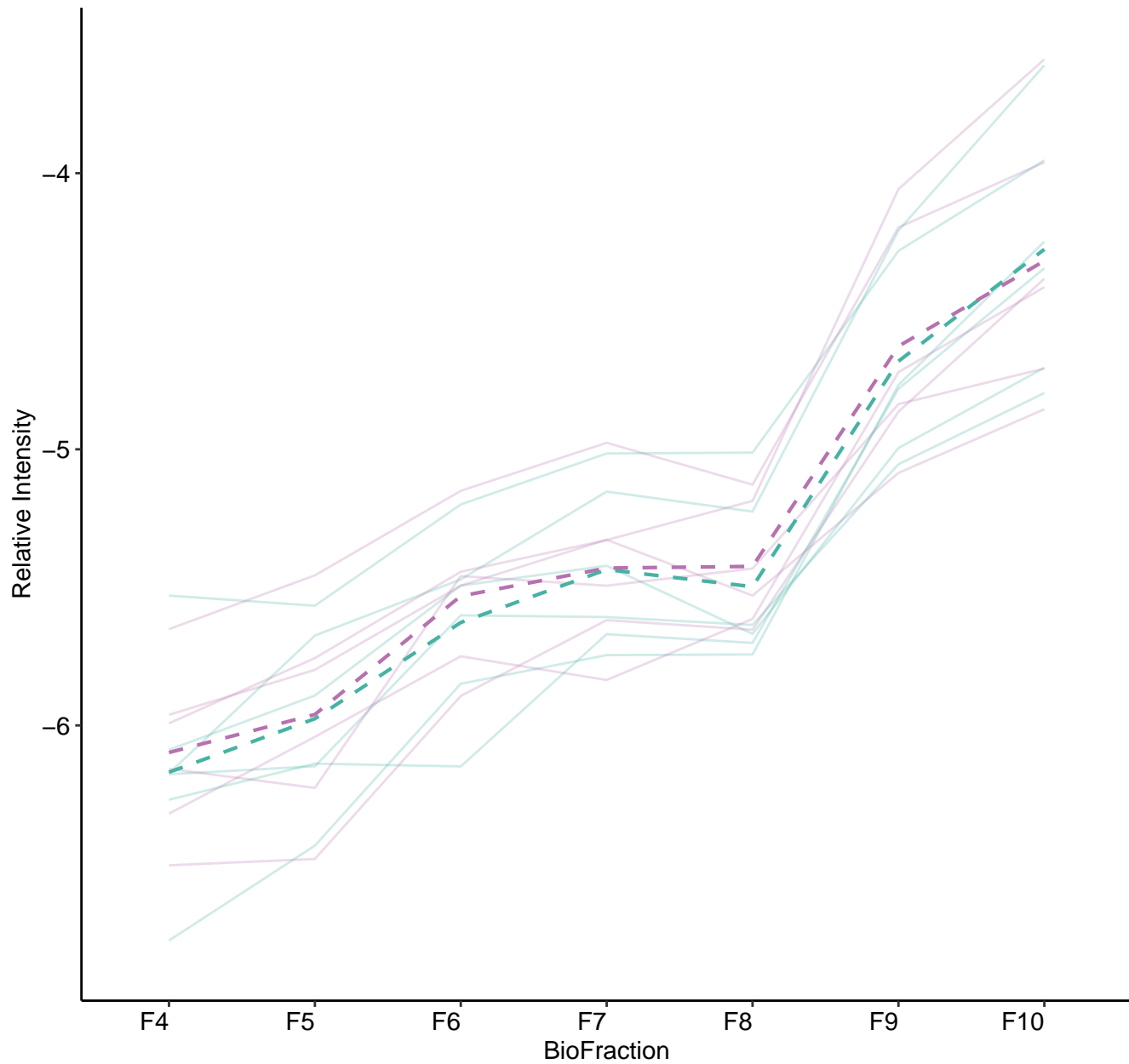
SNARE complex (Vamp2, Snap25, Stx1a, Syt1, Cplx1) (n = 4)
(R2.Fixef = 0.529)



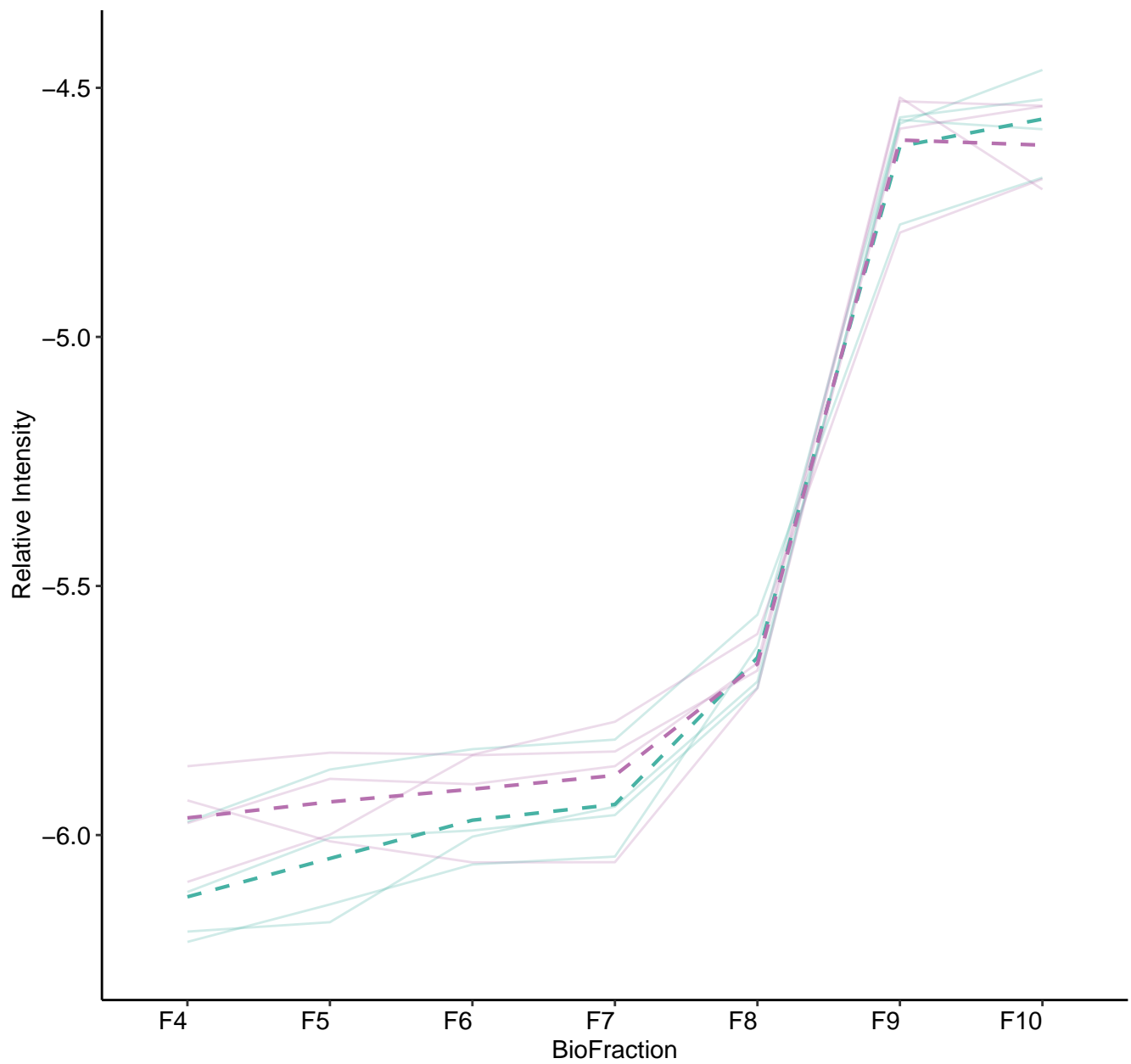
SNARE complex (Vamp2, Snap25, Stx1a, Syt1, Cplx2) (n = 4)
(R2.Fixef = 0.529)



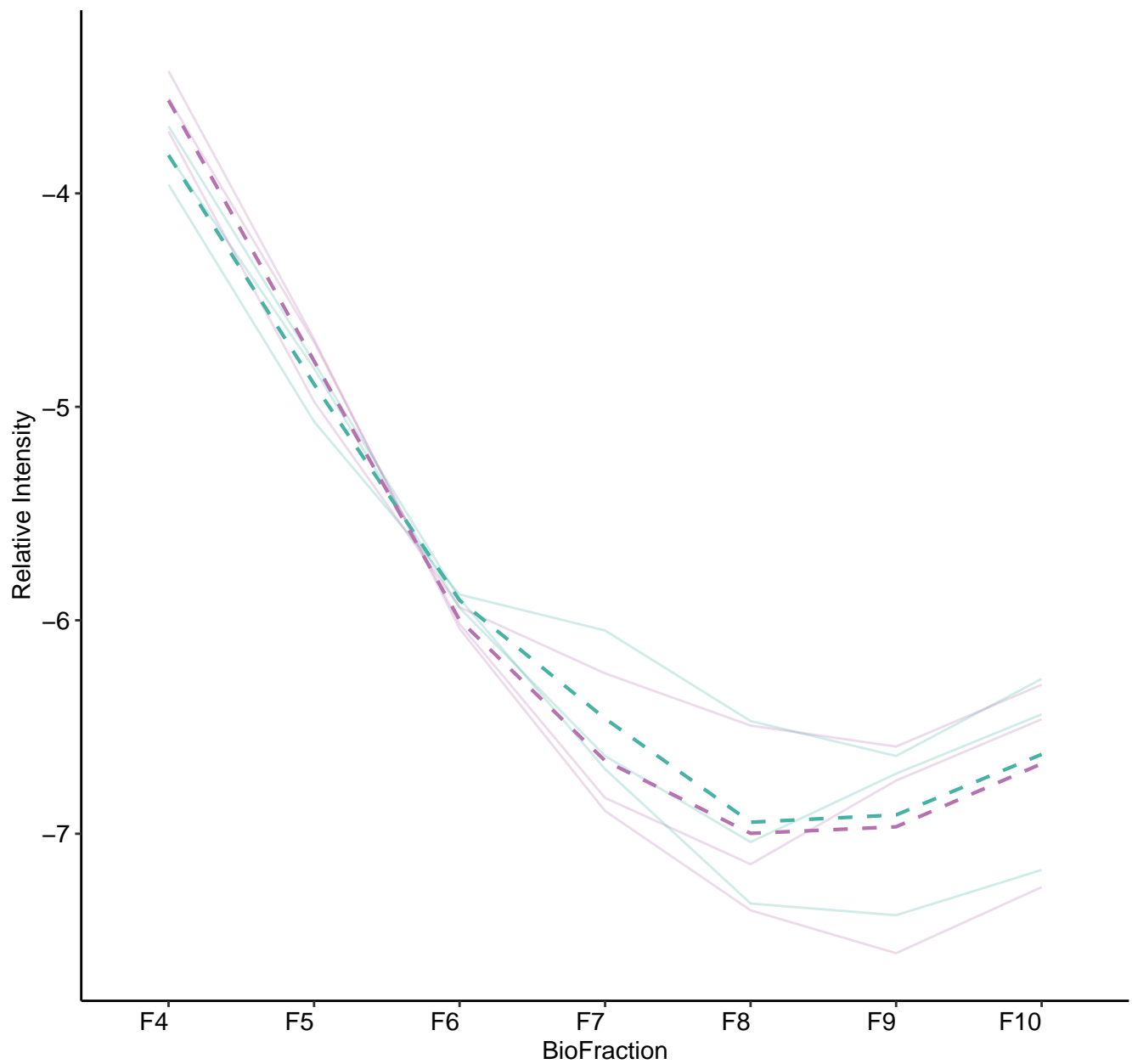
SNF2h-cohesin-NuRD complex (n = 6)
(R2.Fixef = 0.761)



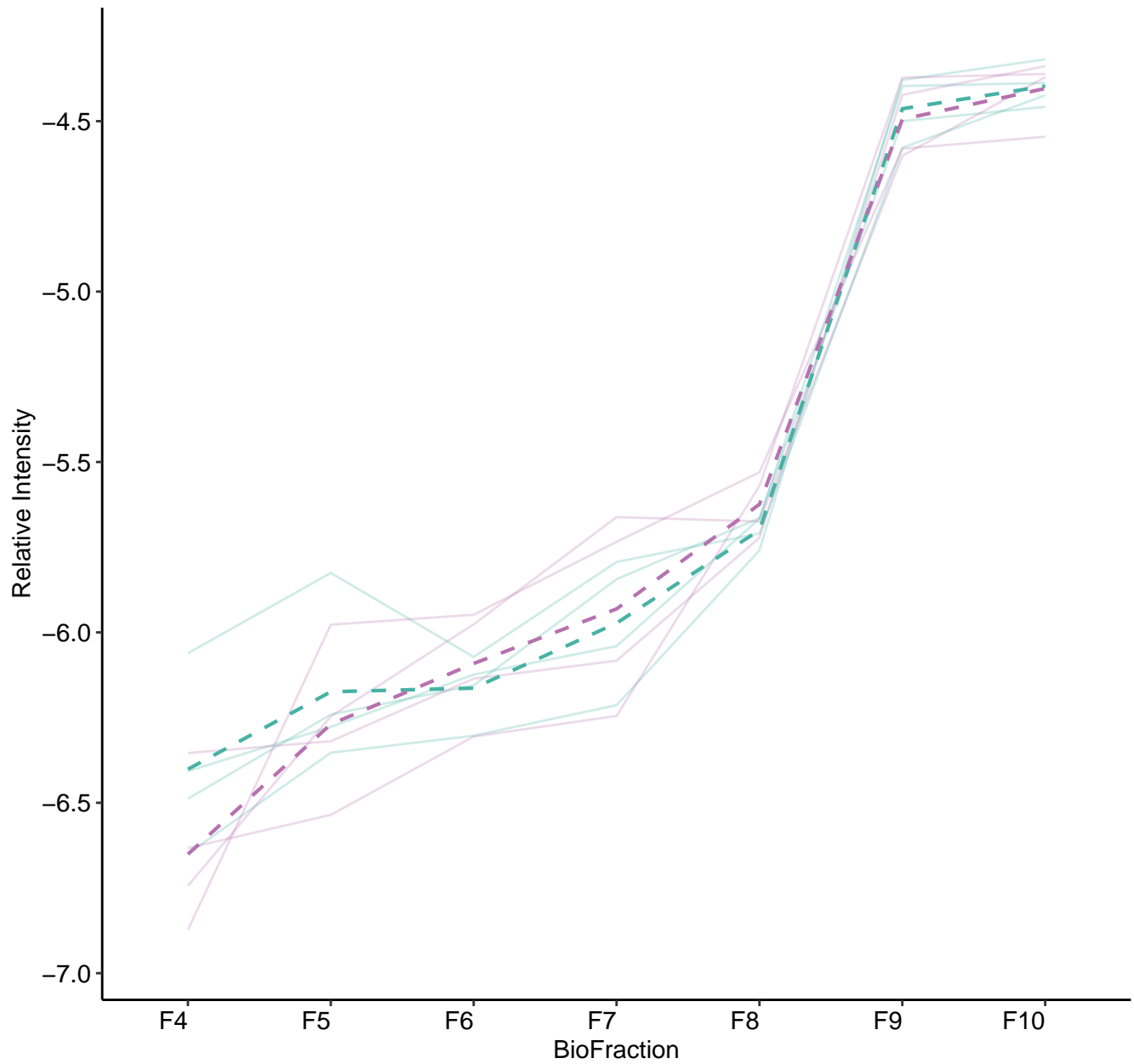
SOG complex (n = 4)
(R2.Fixef = 0.973)



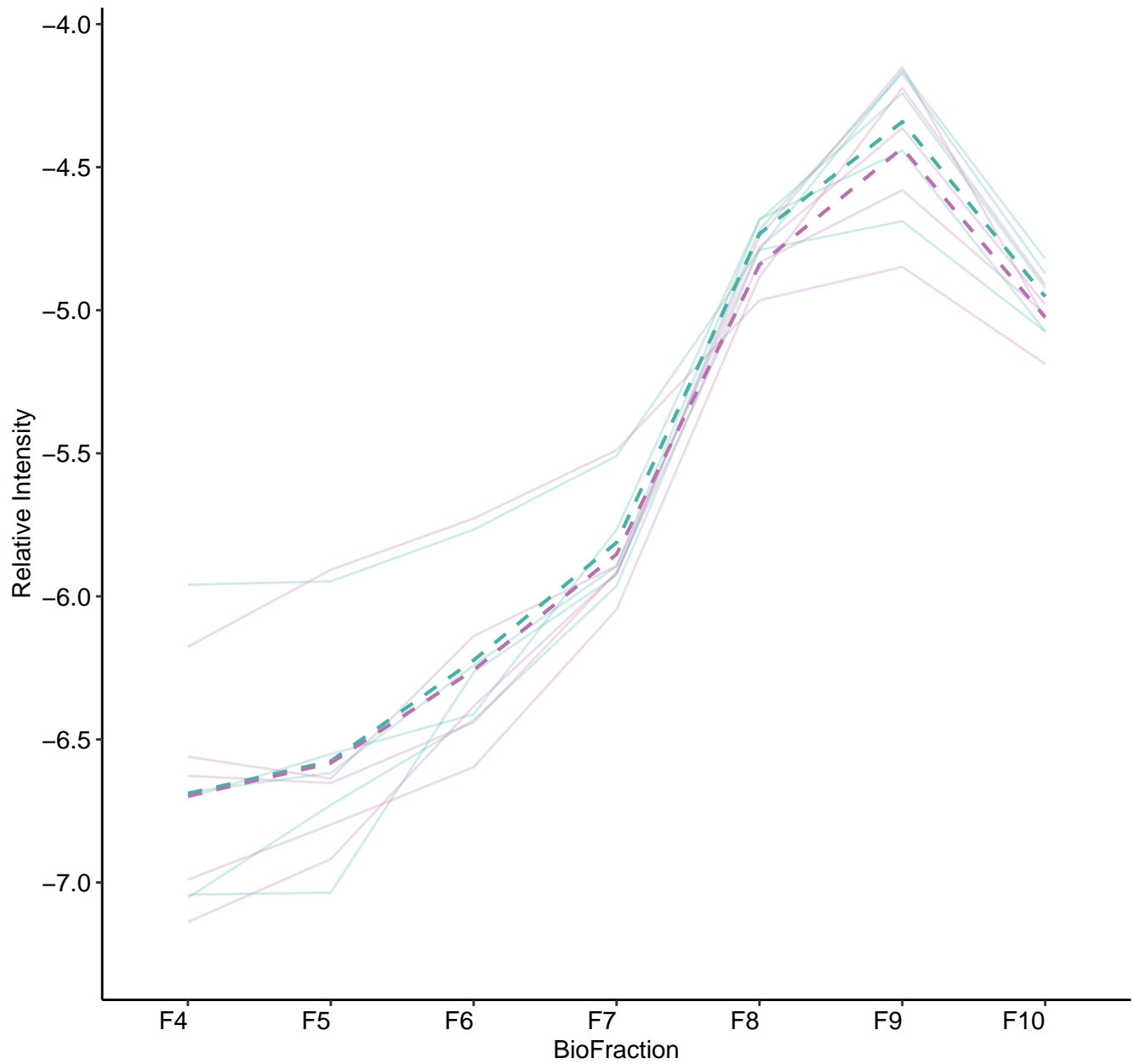
Succinyl-CoA synthetase, ADP-forming (n = 3)
(R2.Fixef = 0.919)



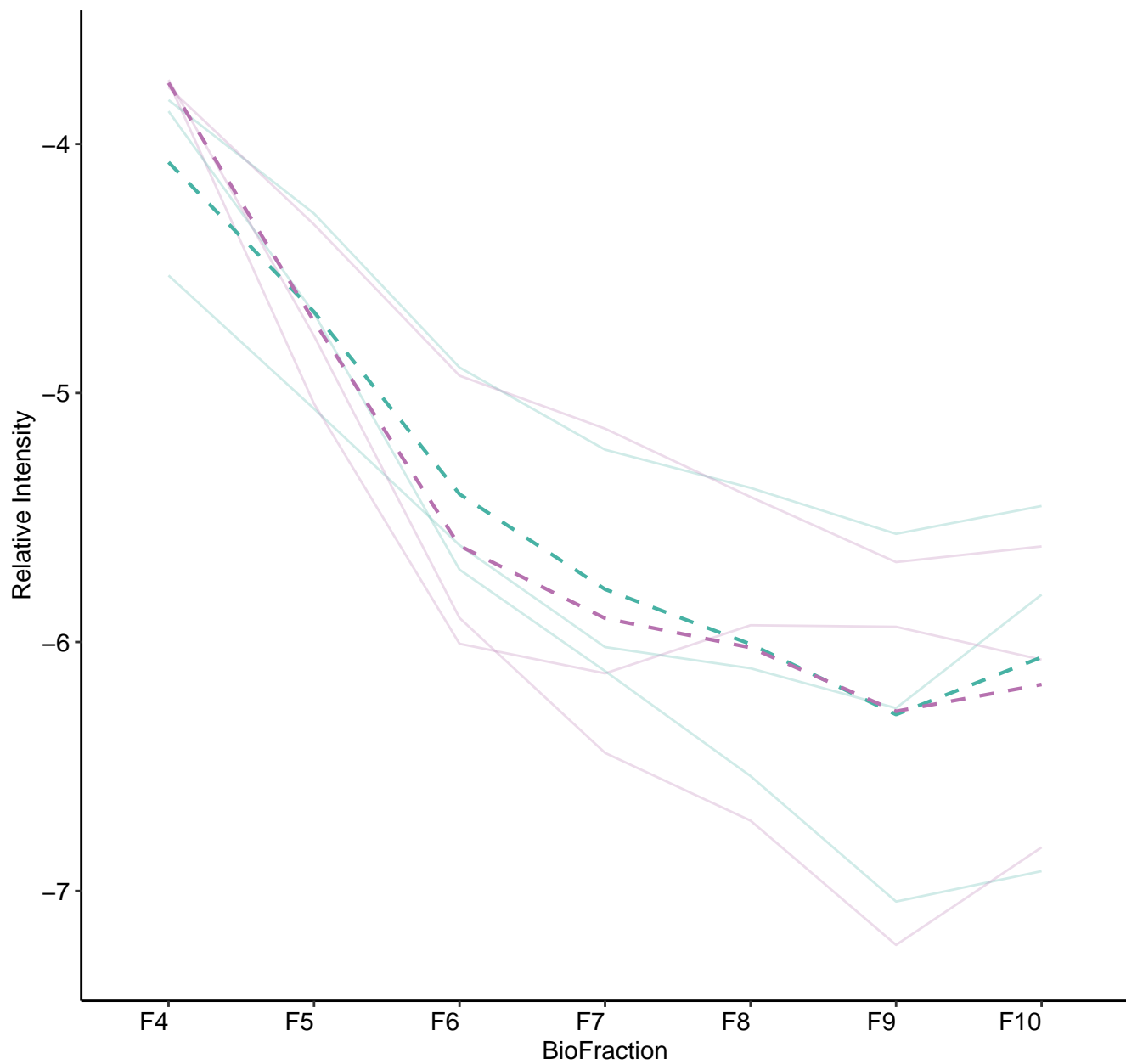
TFIIH transcription factor complex (n = 4)
(R2.Fixef = 0.955)



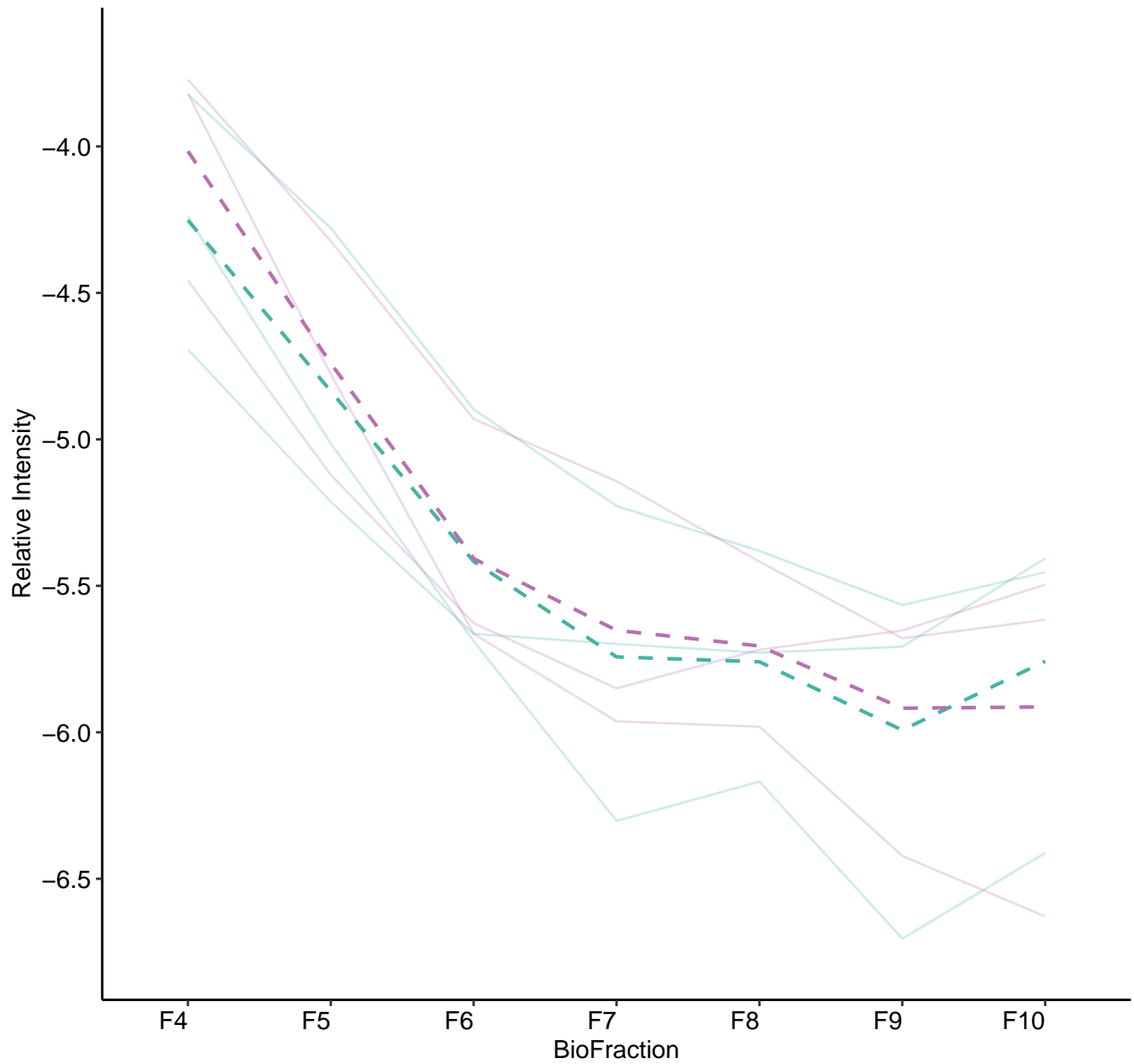
THO complex (n = 5)
(R2.Fixef = 0.905)



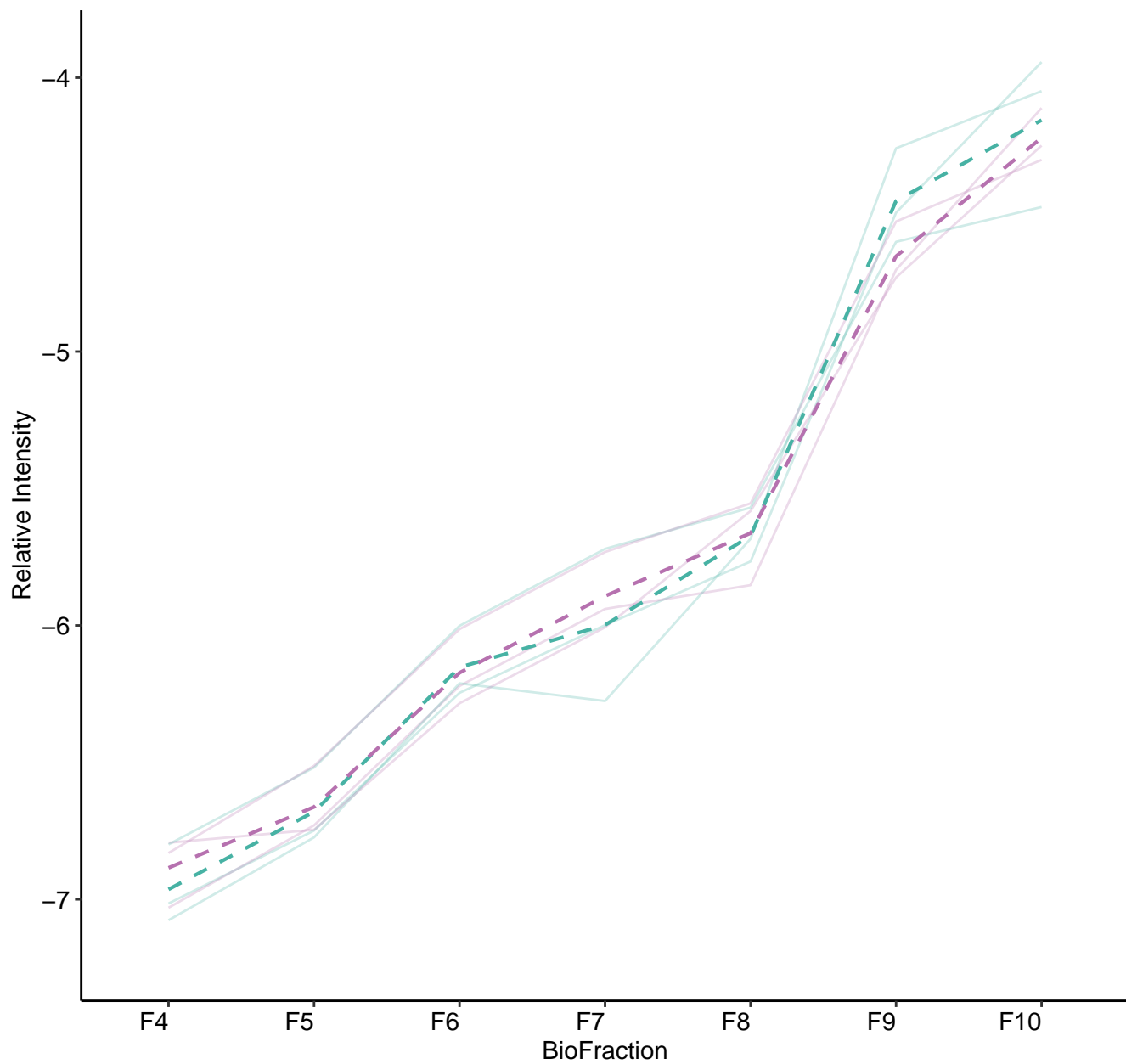
TIM23-TOM22-TIM50-HSD3B2 complex (n = 3)
(R2.Fixef = 0.669)



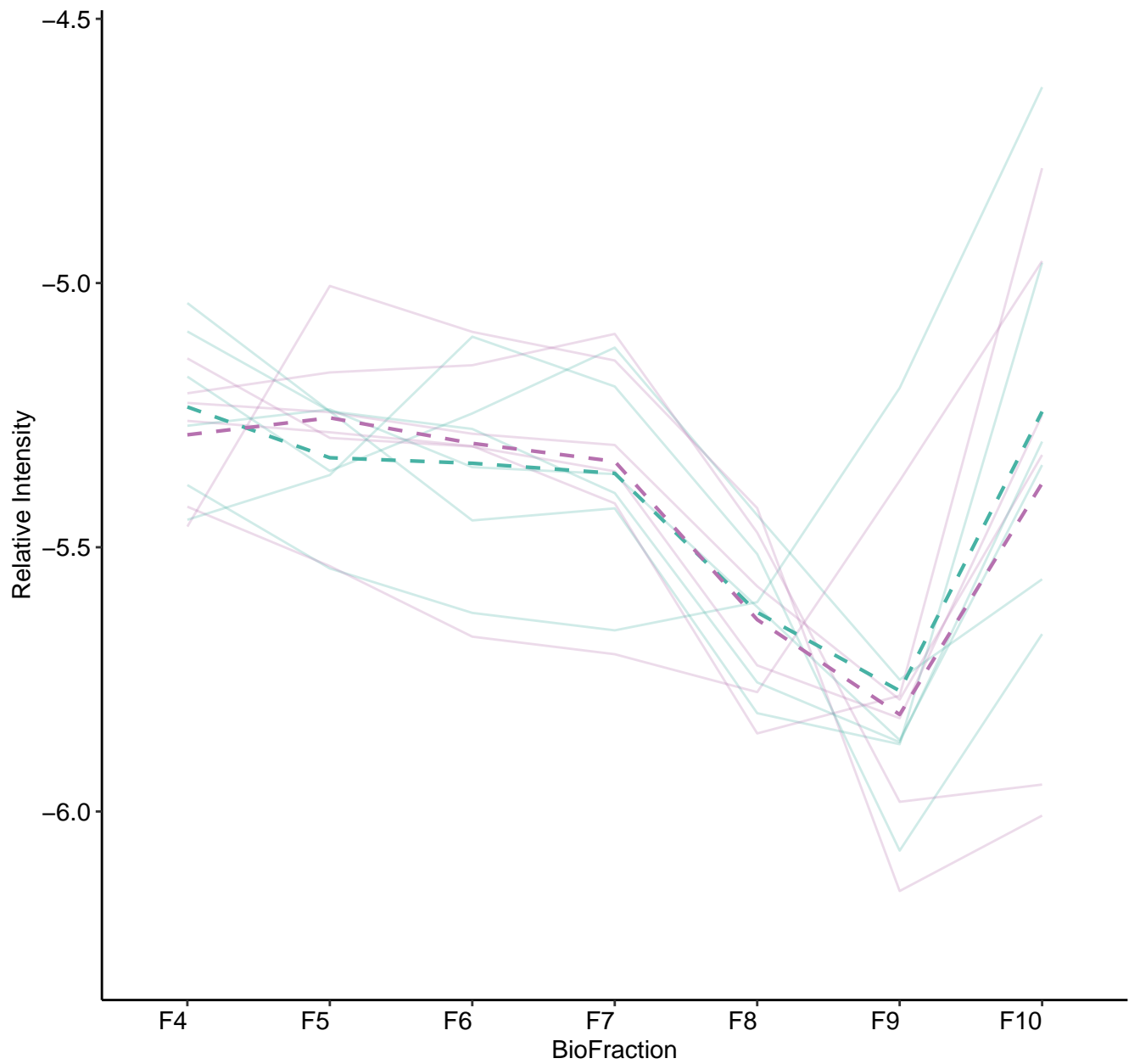
TIMM8A–TIMM13–TIMM23 complex (n = 3)
(R2.Fixef = 0.639)



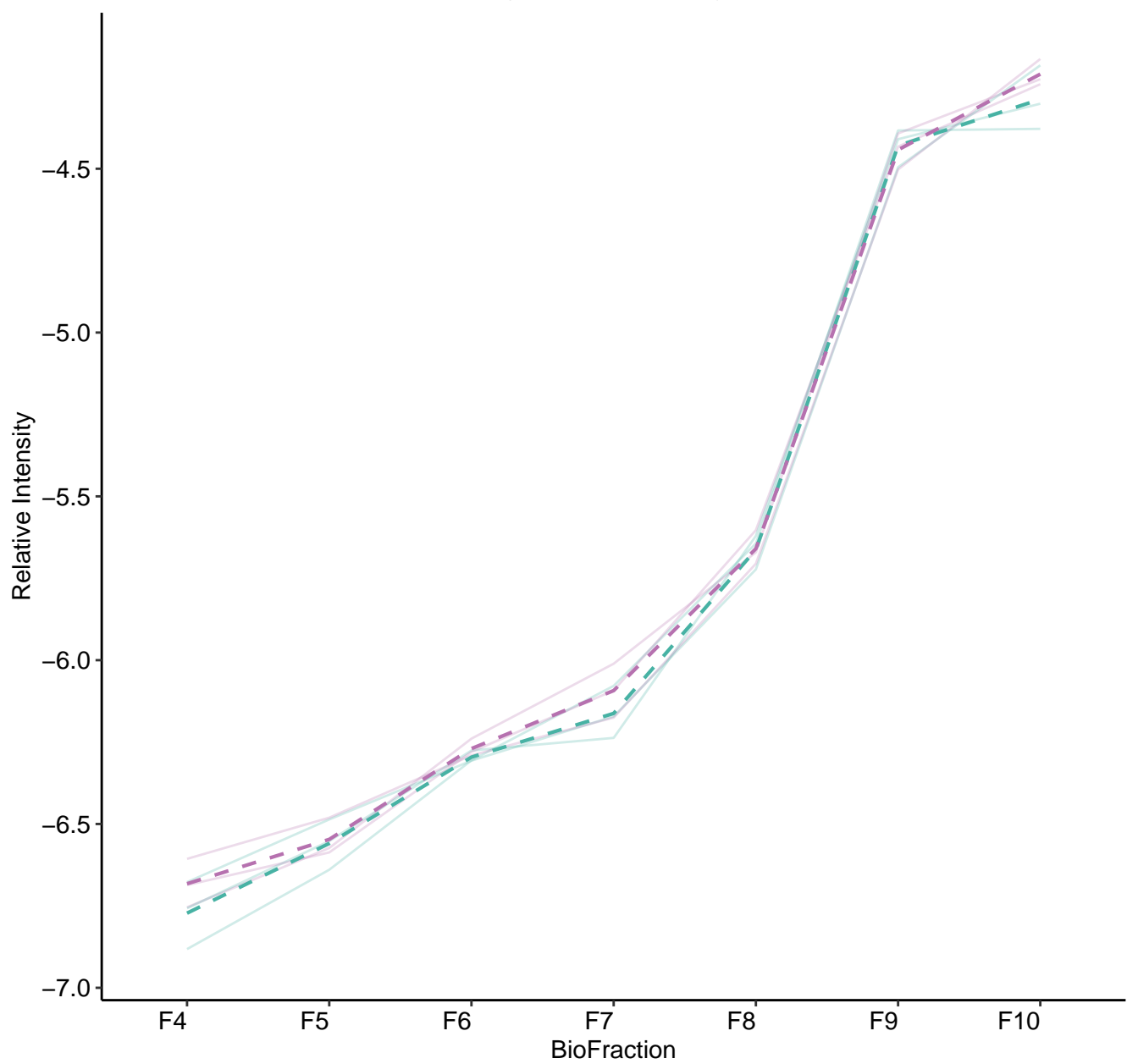
TLE1 corepressor complex (MASH1 promoter-corepressor complex) (n = 3)
(R2.Fixef = 0.972)



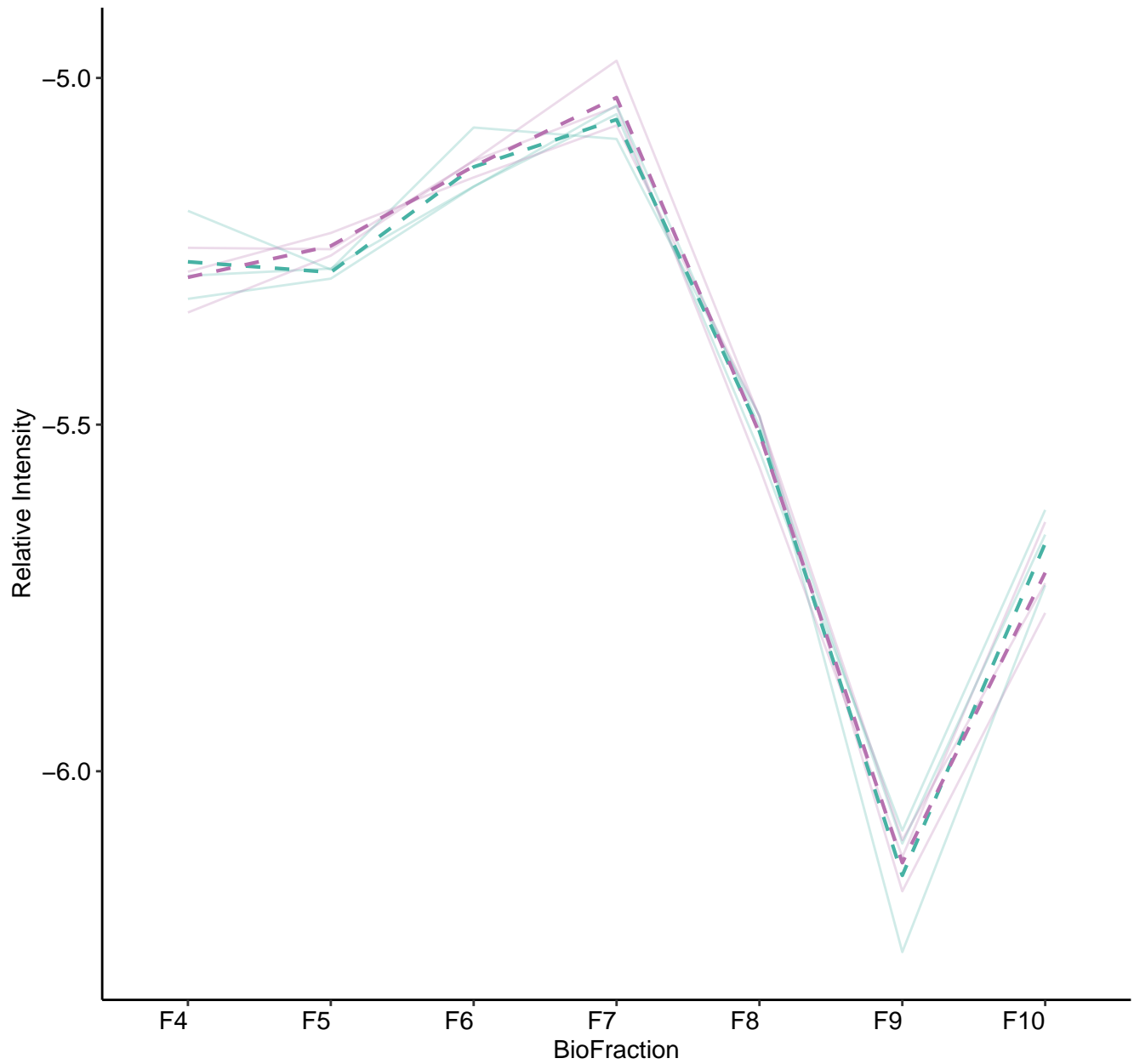
ppa B signaling complex (CHUK, BTRC, NFKB2, PPP6C, REL, CUL1, IKBKE, SAPS2, SAPS1, ANKRD
(R2.Fixef = 0.382)



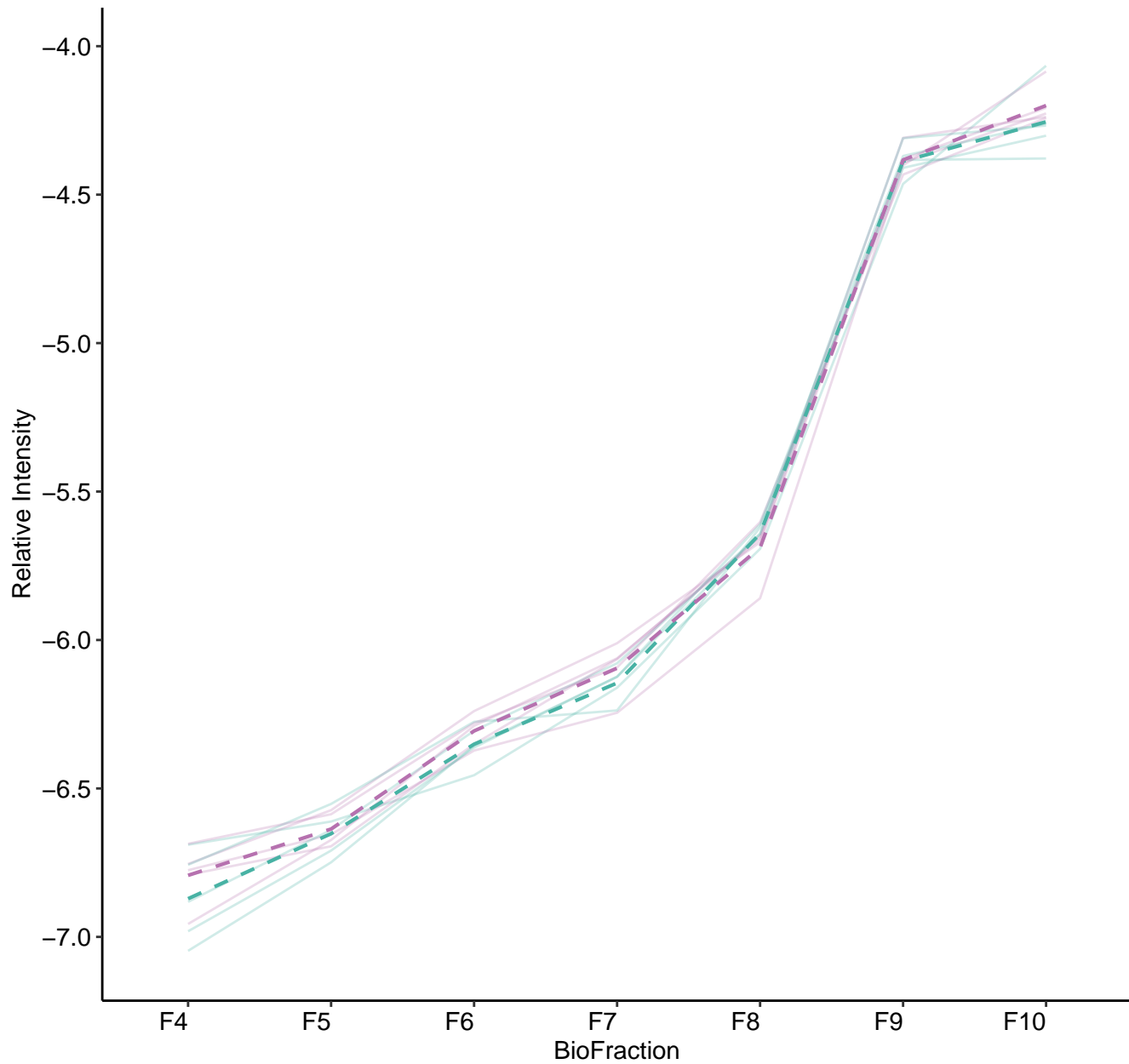
plex (RPL6, RPL30, RPS13, CHUK, DDX3X, NFKB2, NFKBIB, REL, IKBKG, NFKB1, MAP3K8, RELB, (R2.Fixef = 0.995)



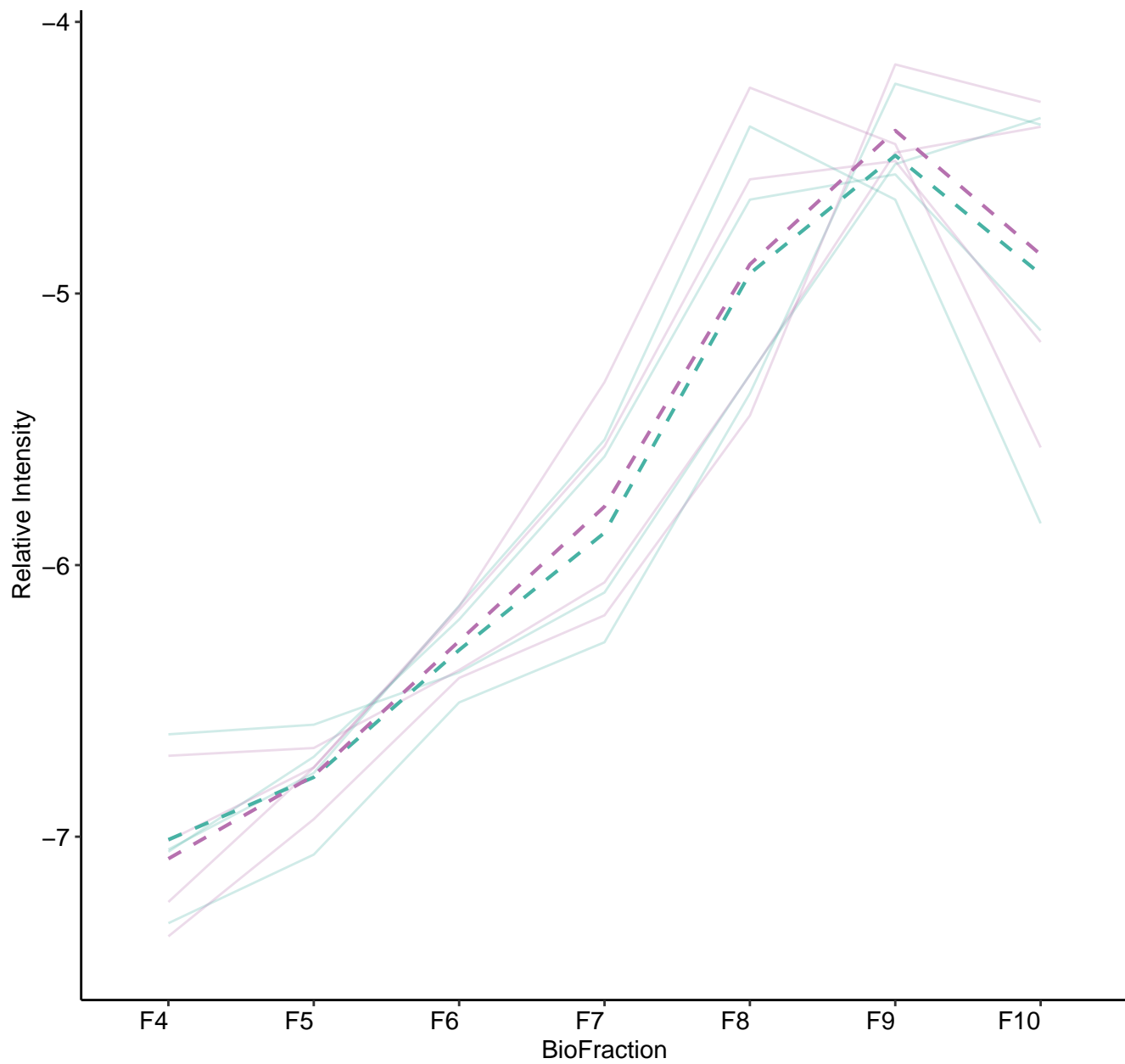
TNF- α /NF- κ B signaling complex 10 (n = 3)
(R2.Fixef = 0.982)



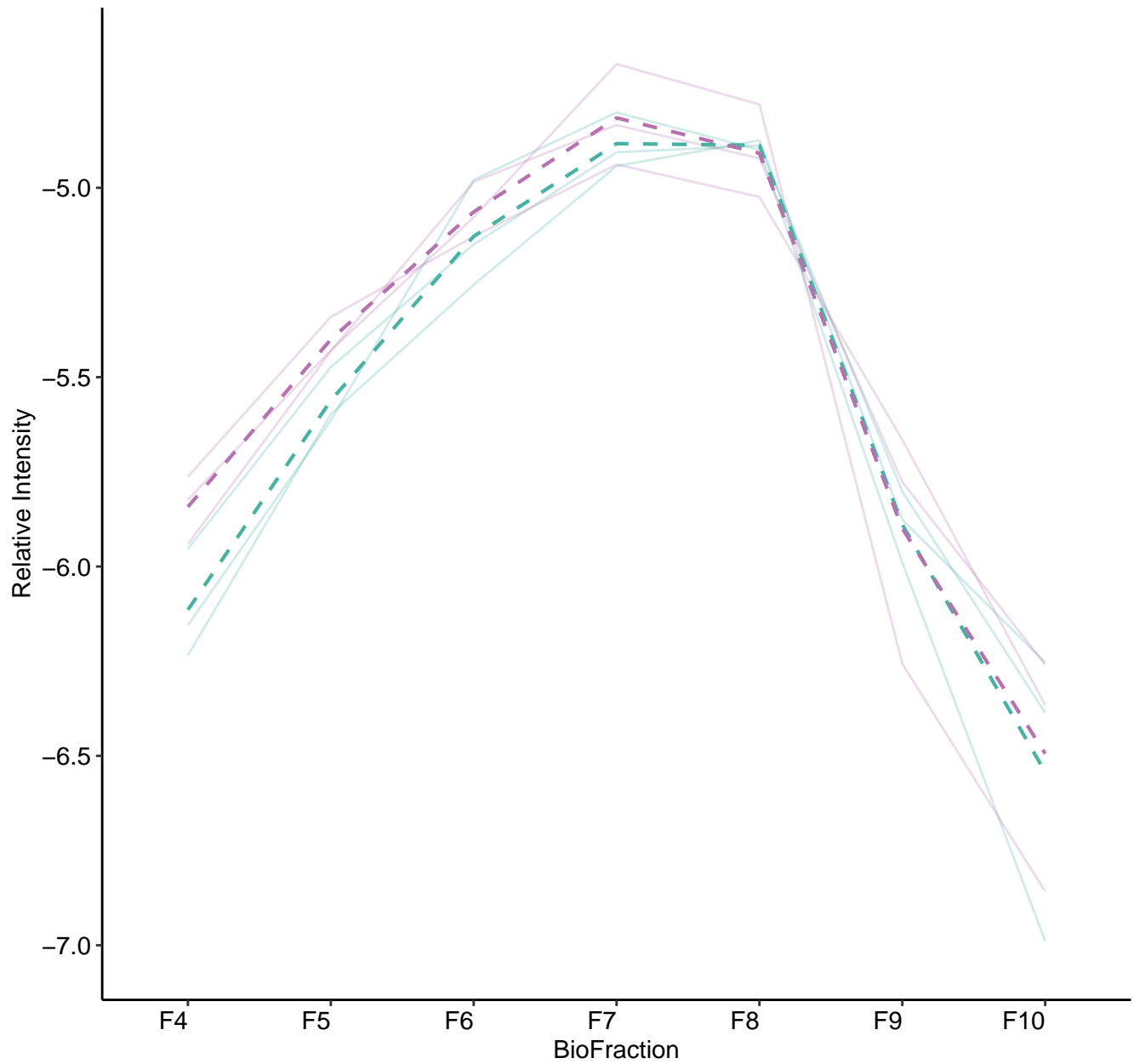
TNF- α /NF- κ B signaling complex 6 (n = 5)
(R2.Fixef = 0.993)



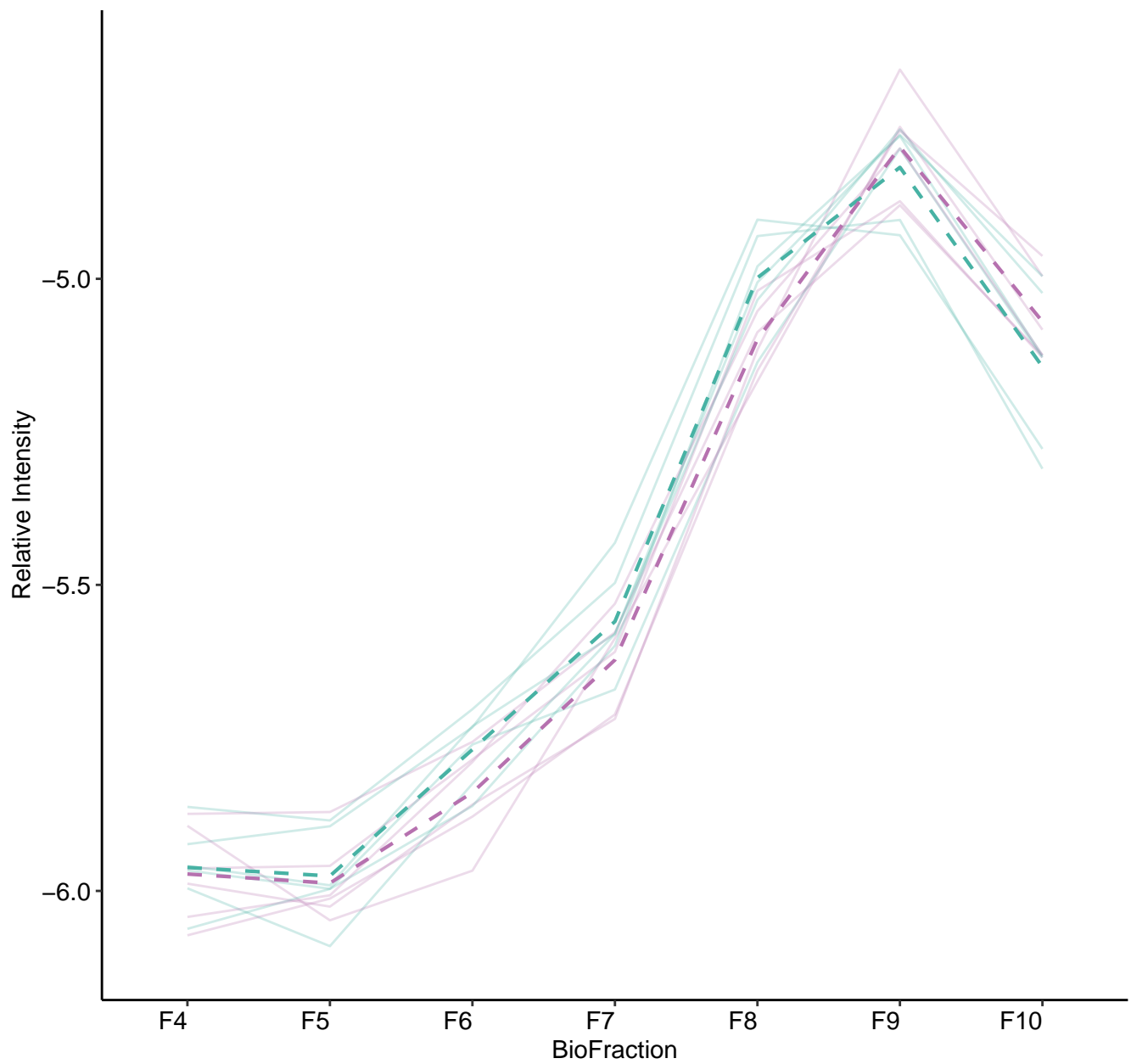
Toposome (n = 4)
(R2.Fixef = 0.858)



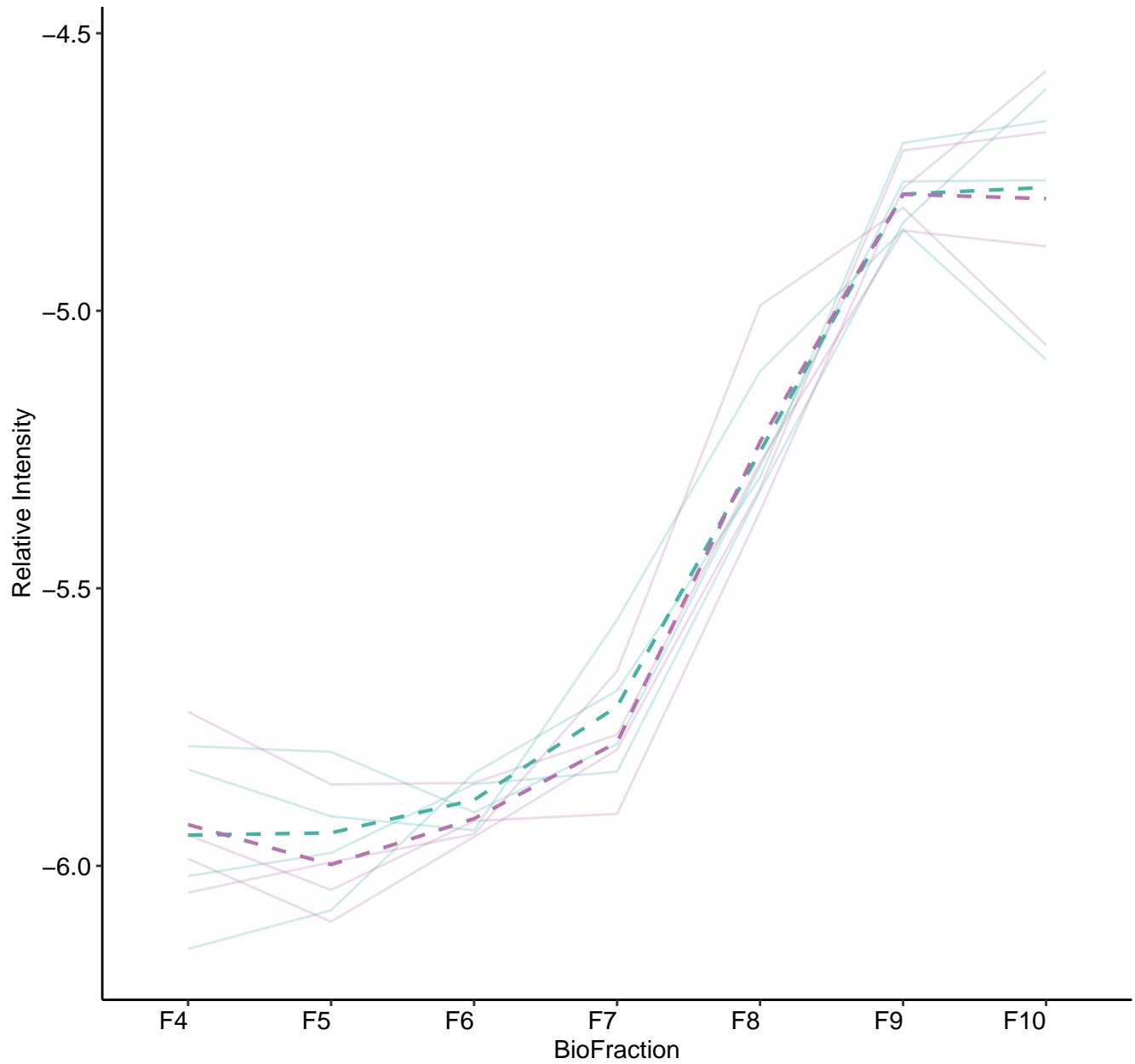
Translocon-associated protein (TRAP) complex (n = 3)
(R2.Fixef = 0.913)



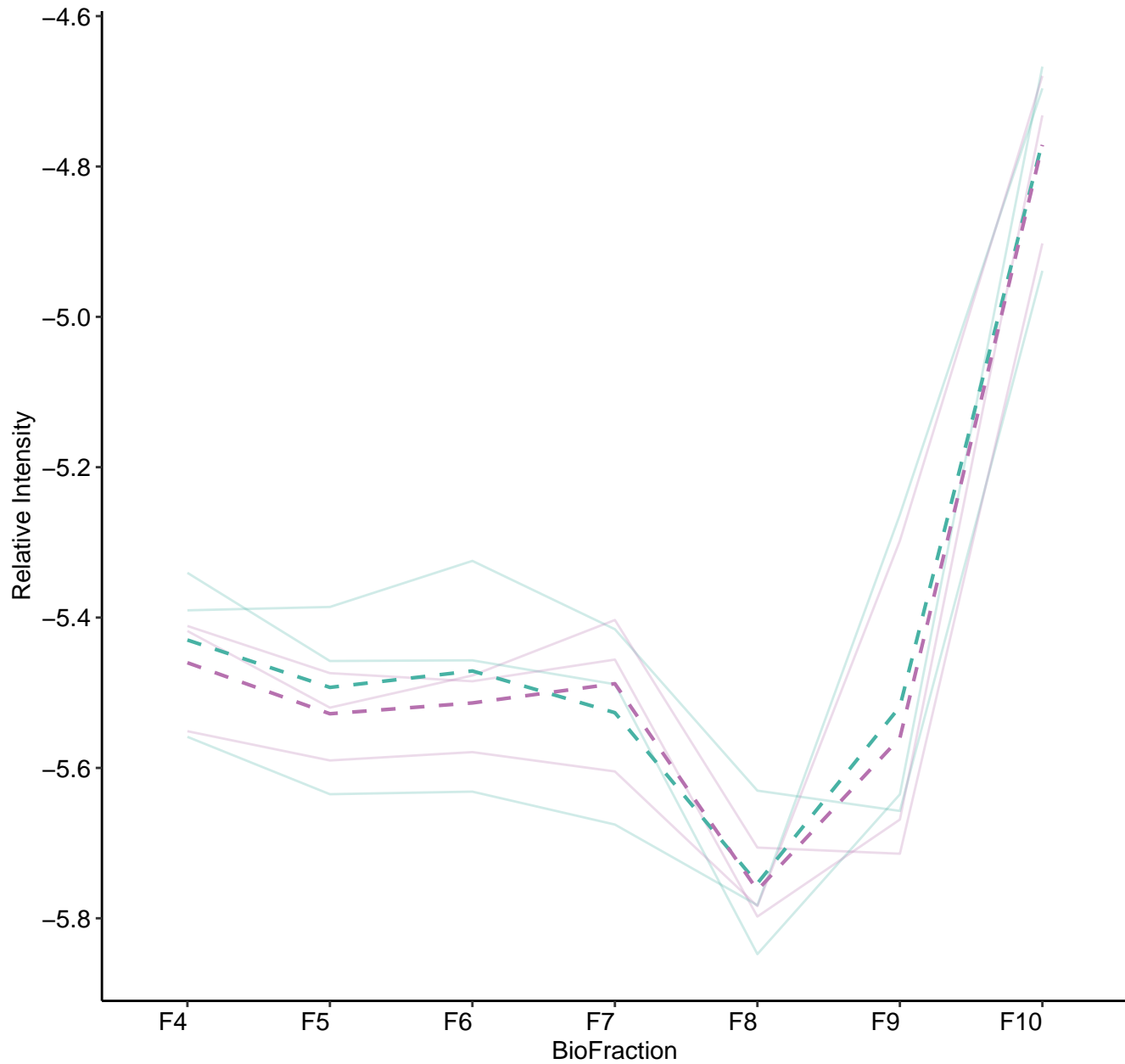
TRAPP complex (n = 6)
(R2.Fixef = 0.97)



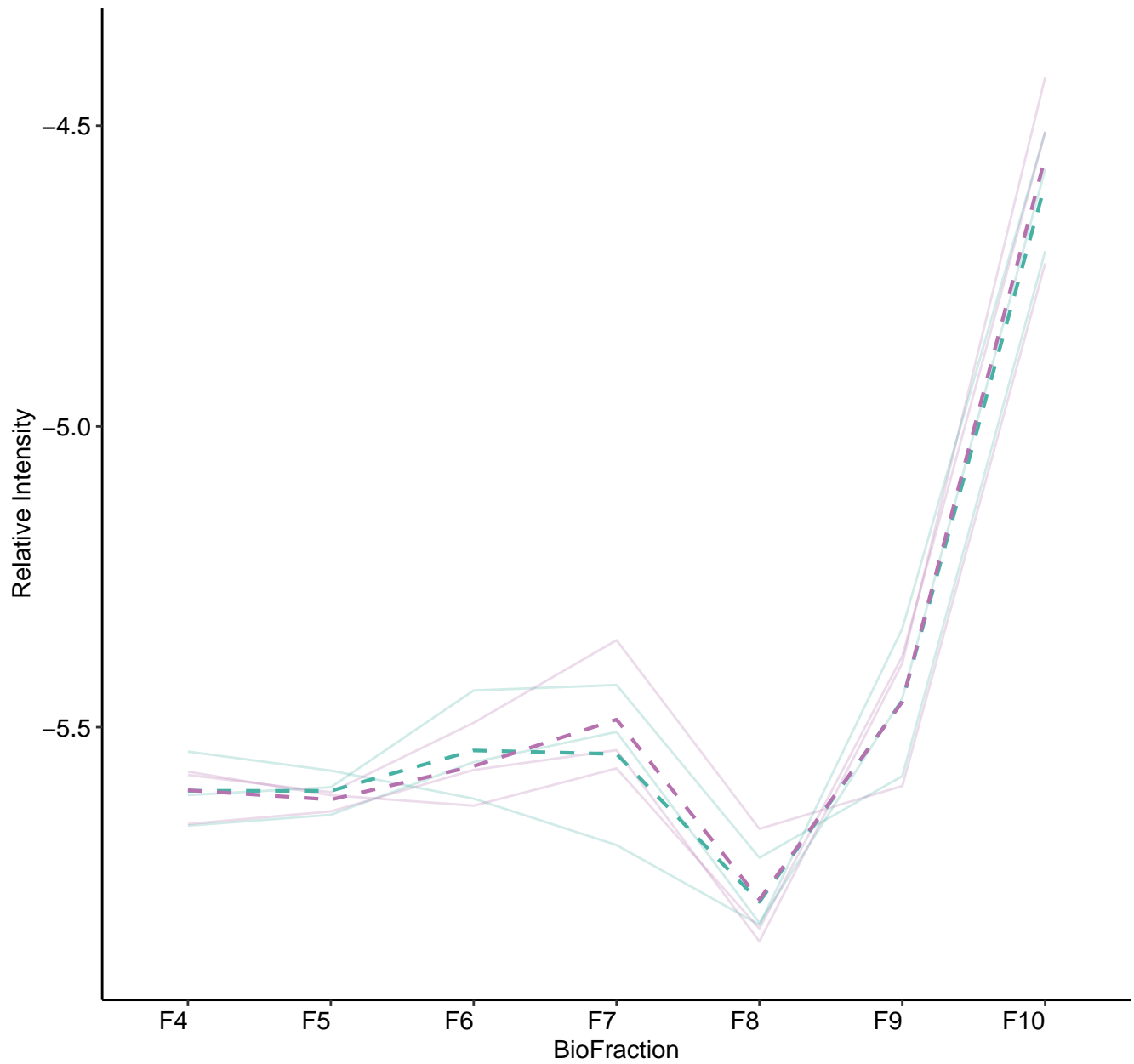
Tubulin polyglutamylase complex (n = 4)
(R2.Fixef = 0.934)



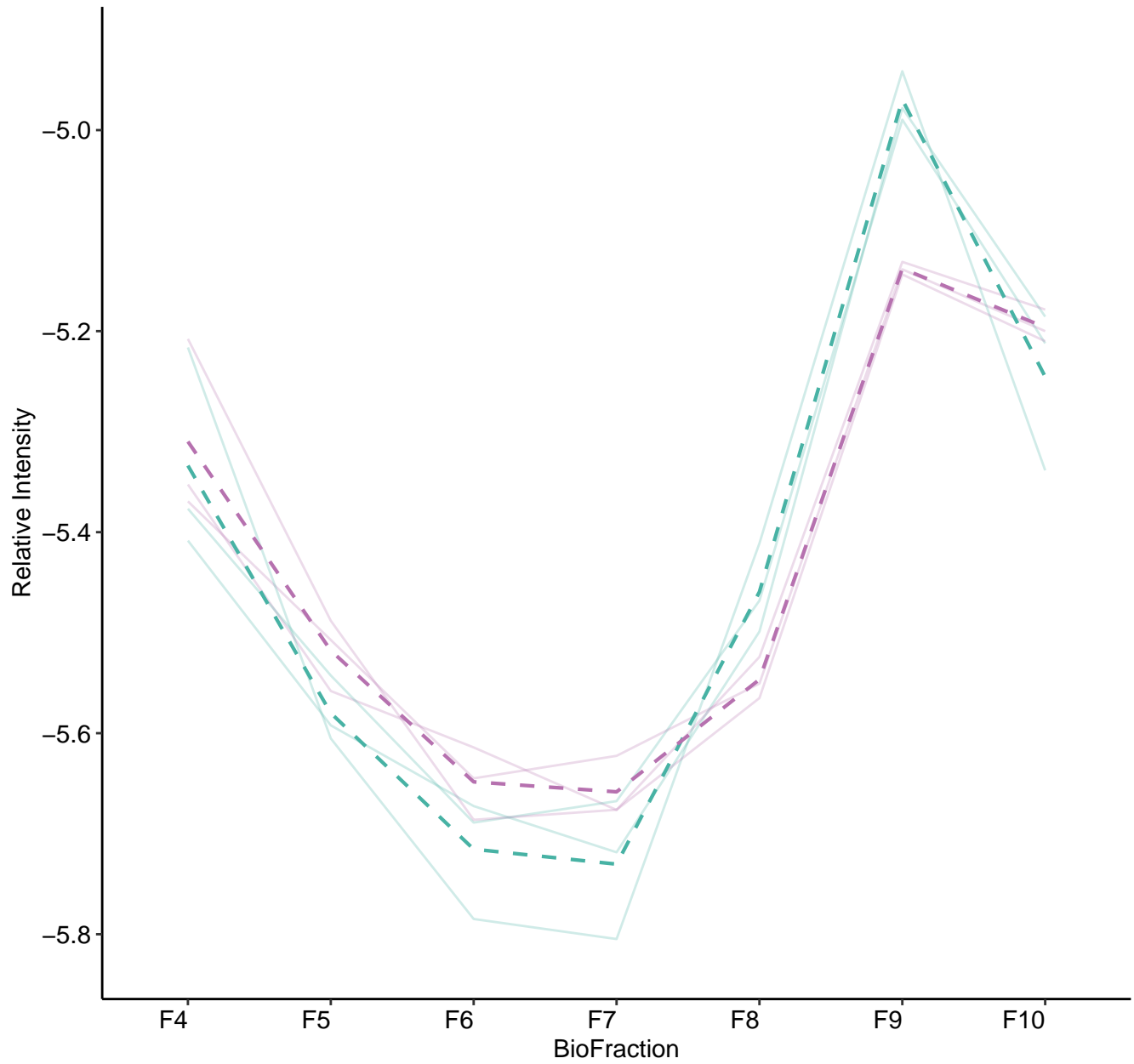
Ubiquitin E3 ligase (CSN1, CSN8, HRT1, SKP1, SKP2, CUL1, CUL2, CUL3) (n = 3)
(R2.Fixef = 0.827)



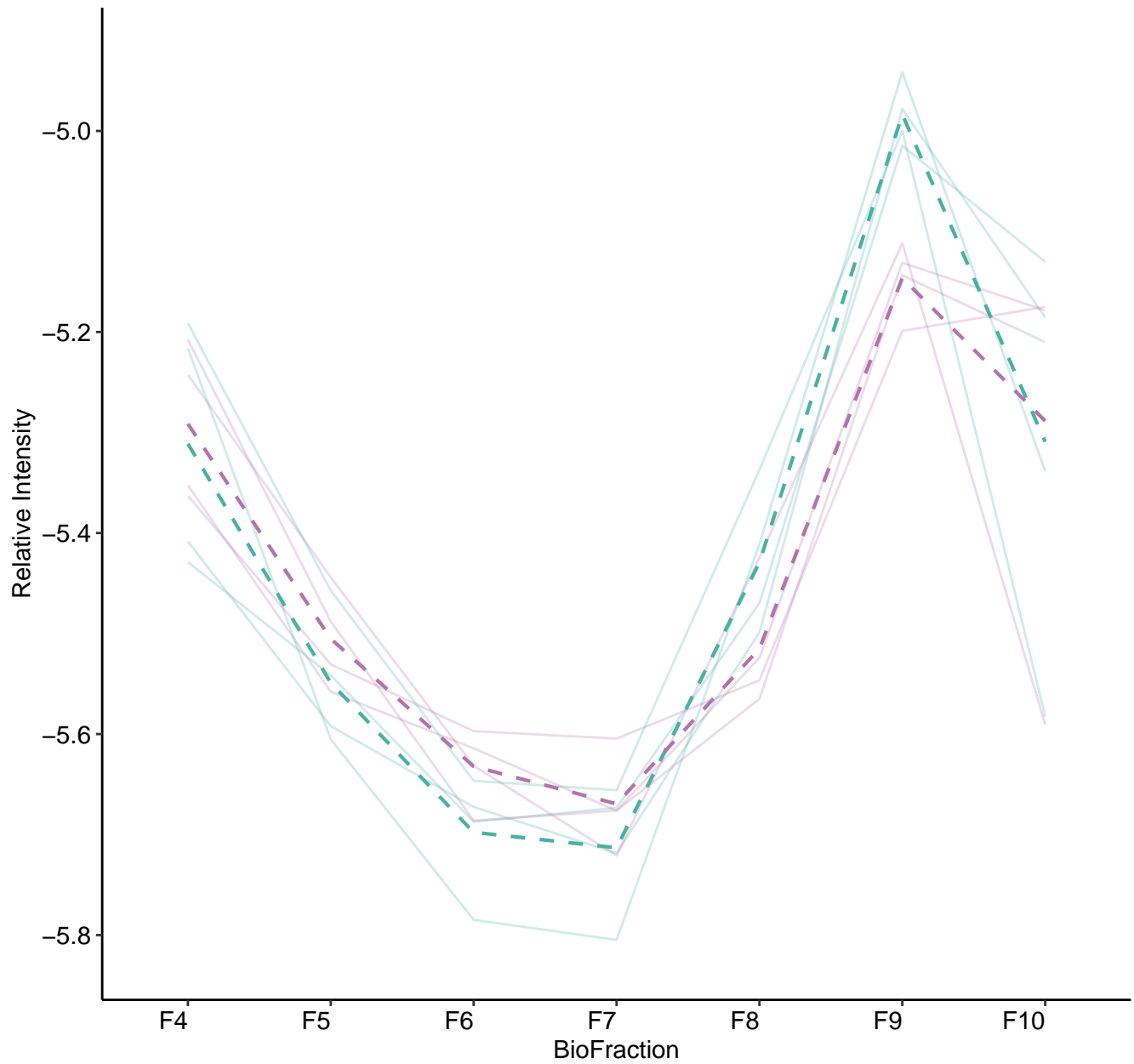
Ubiquitin E3 ligase (DDB1, DDB2, CUL4A, CUL4B, RBX1) (n = 3)
(R2.Fixef = 0.936)



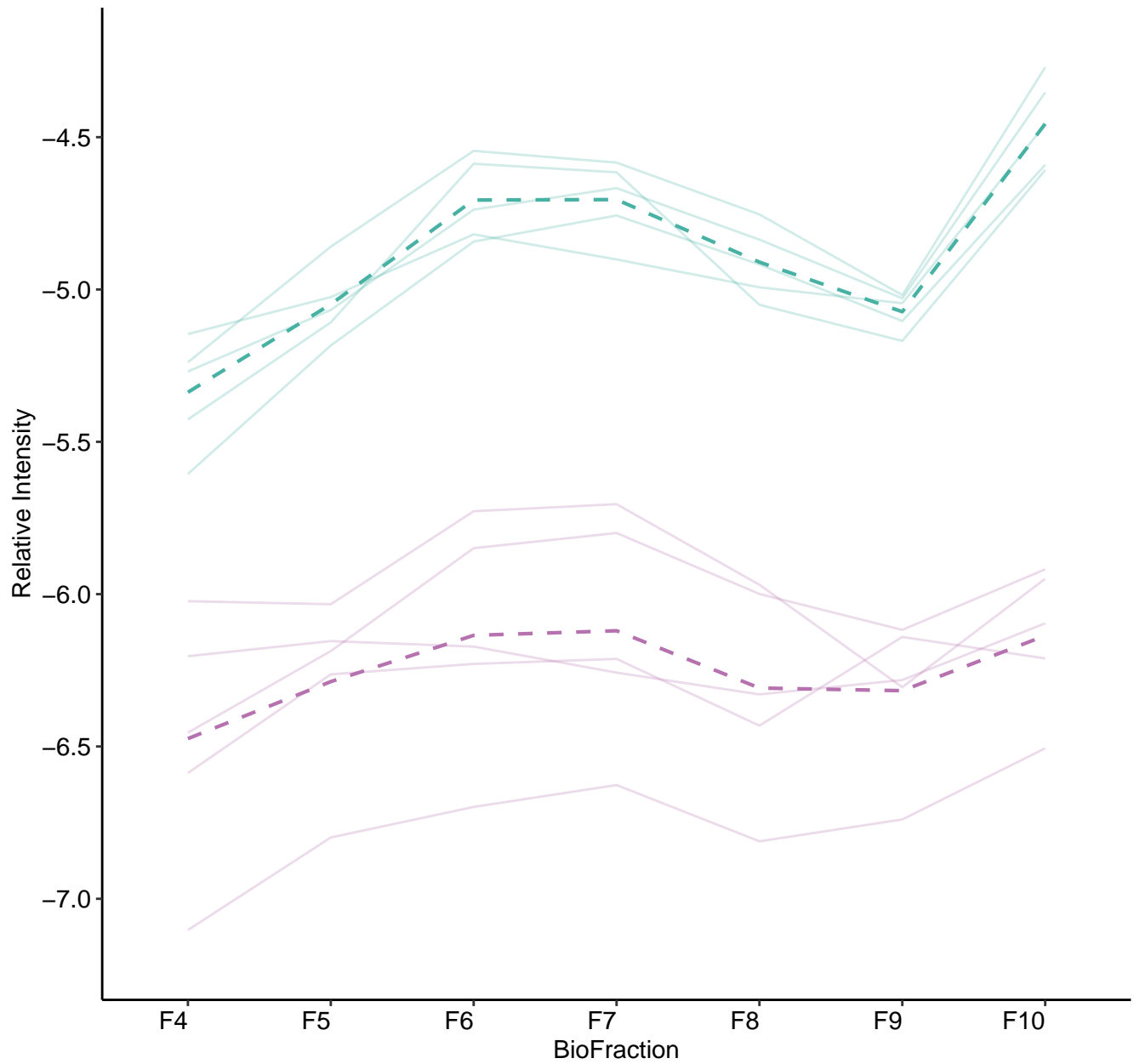
v-ATPase-Ragulator-AXIN/LKB1-AMPK complex (n = 3)
(R2.Fixef = 0.947)



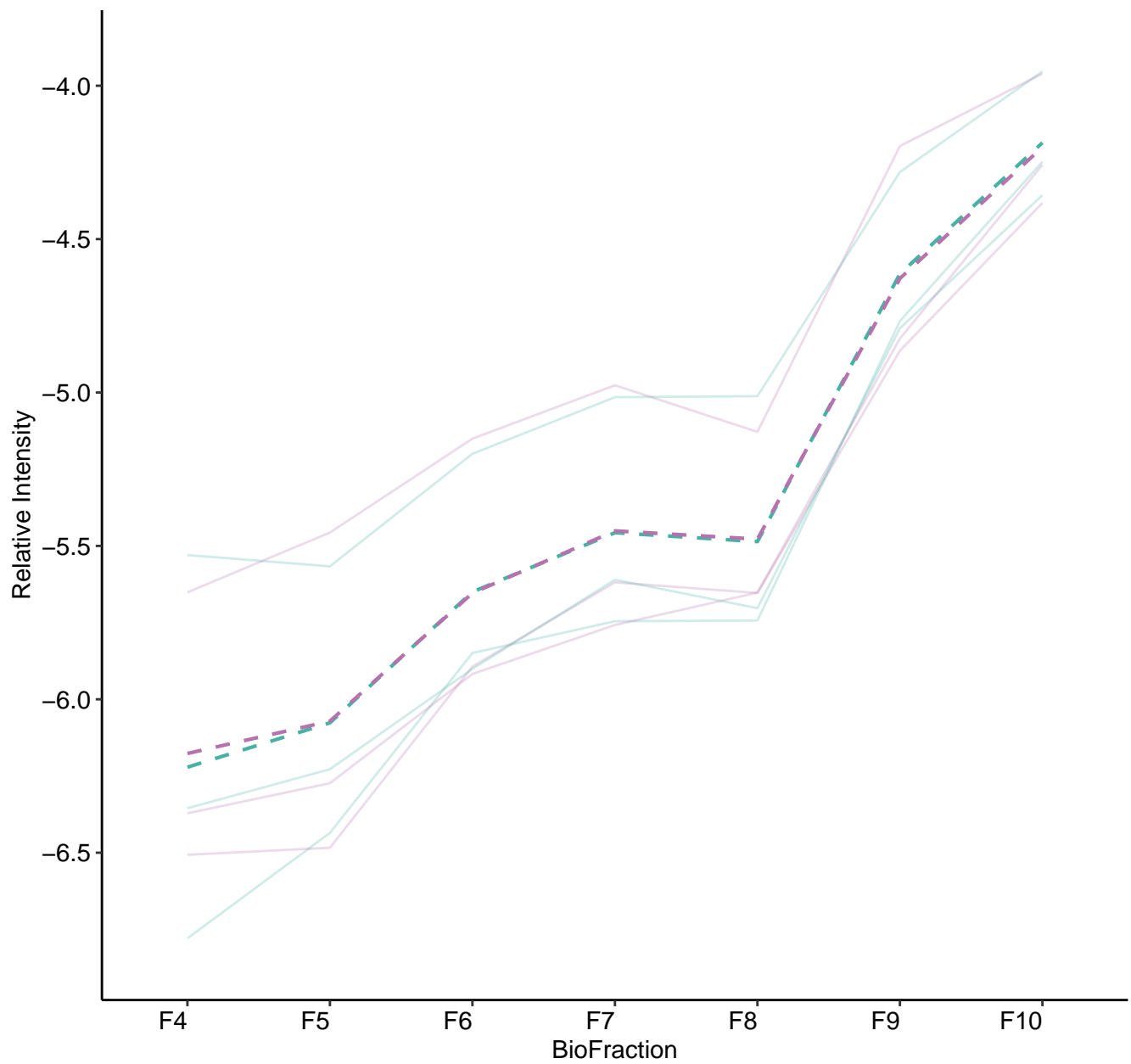
Vacuolar ATPase (n = 4)
(R2.Fixef = 0.826)



WASH-CAPZalpa/beta complex (n = 5)
(R2.Fixef = 0.889)



XFIM complex (n = 3)
(R2.Fixef = 0.737)



RAVE (n = 3)
(R2.Fixef = 0.991)

