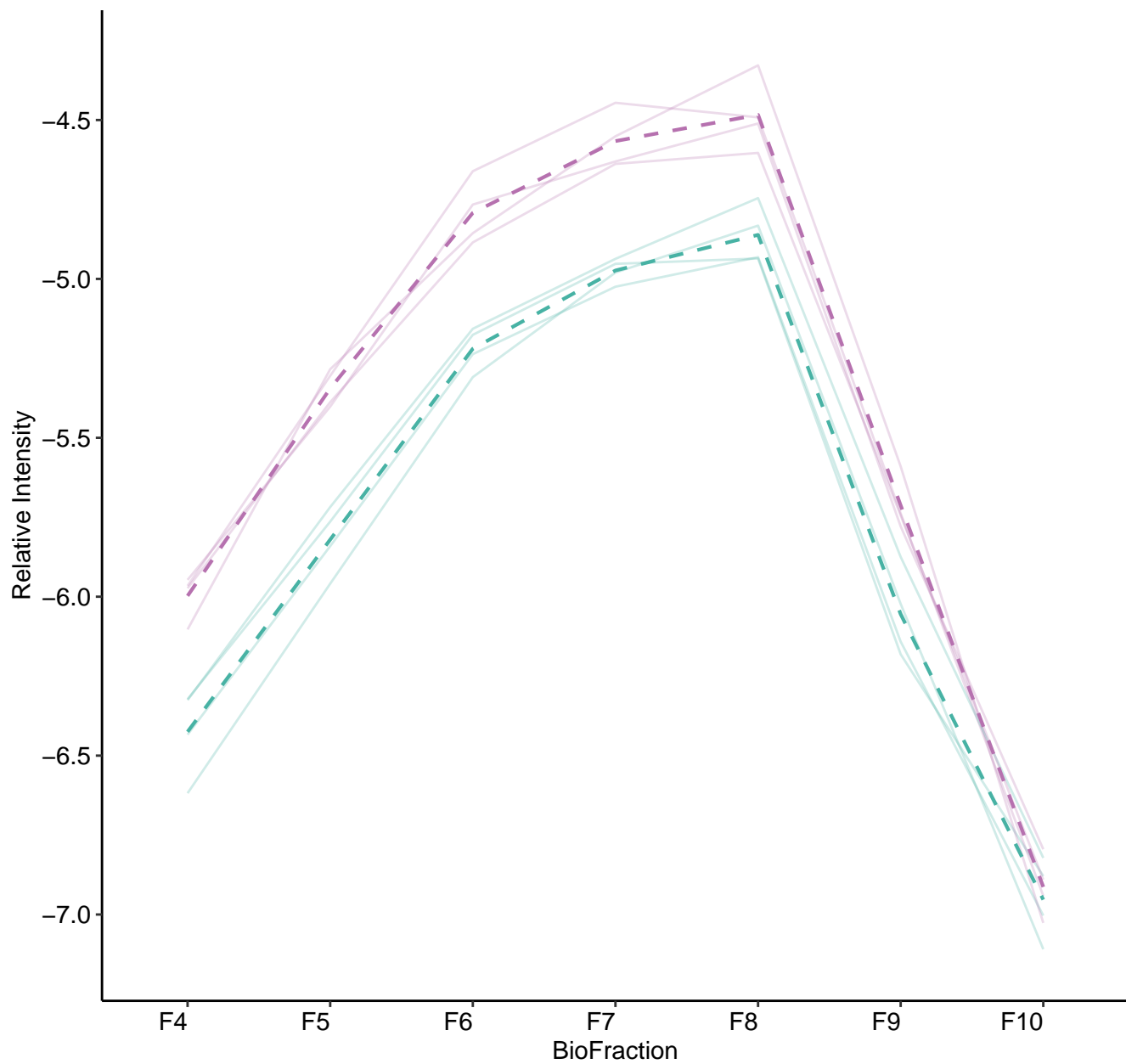
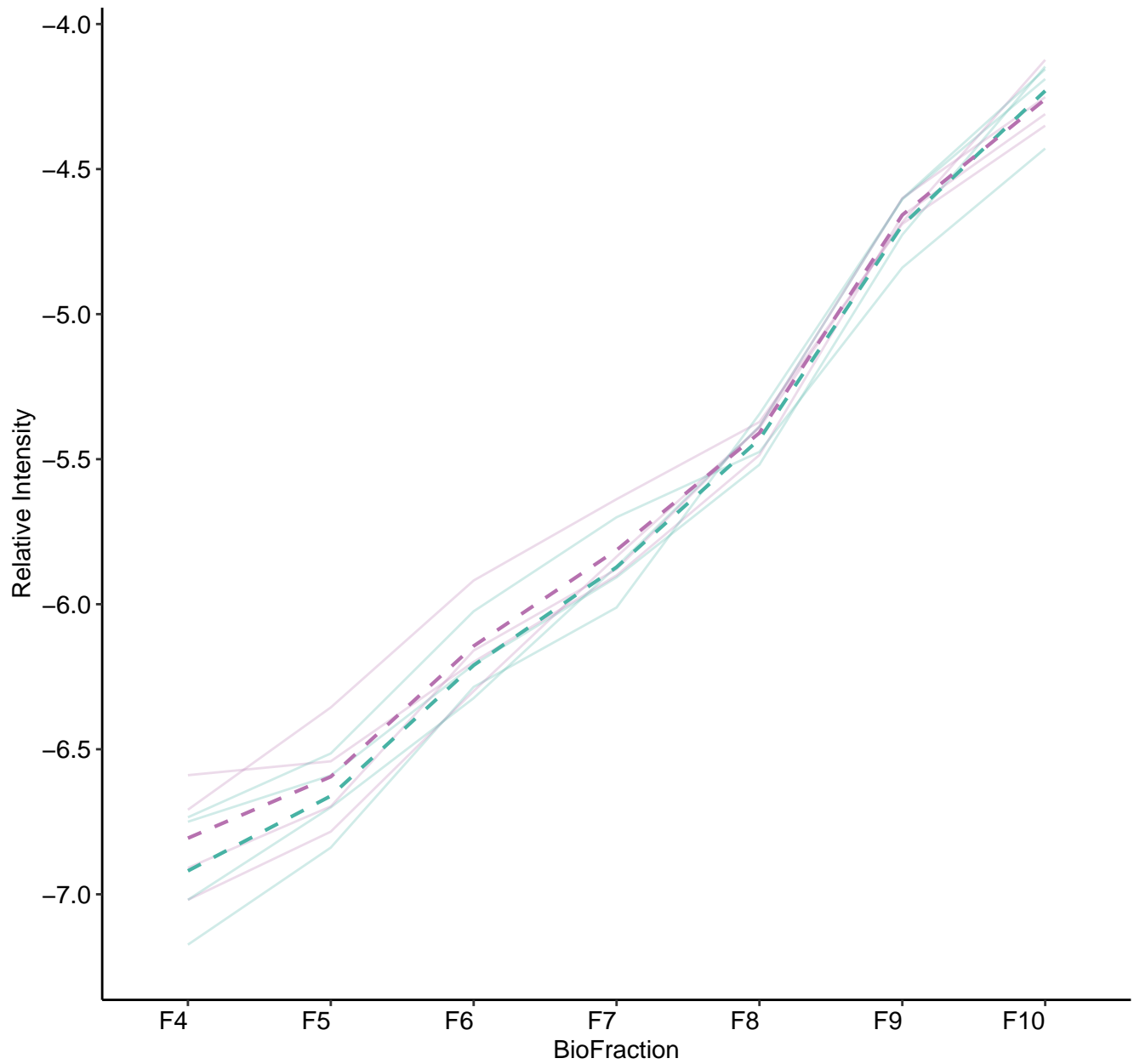


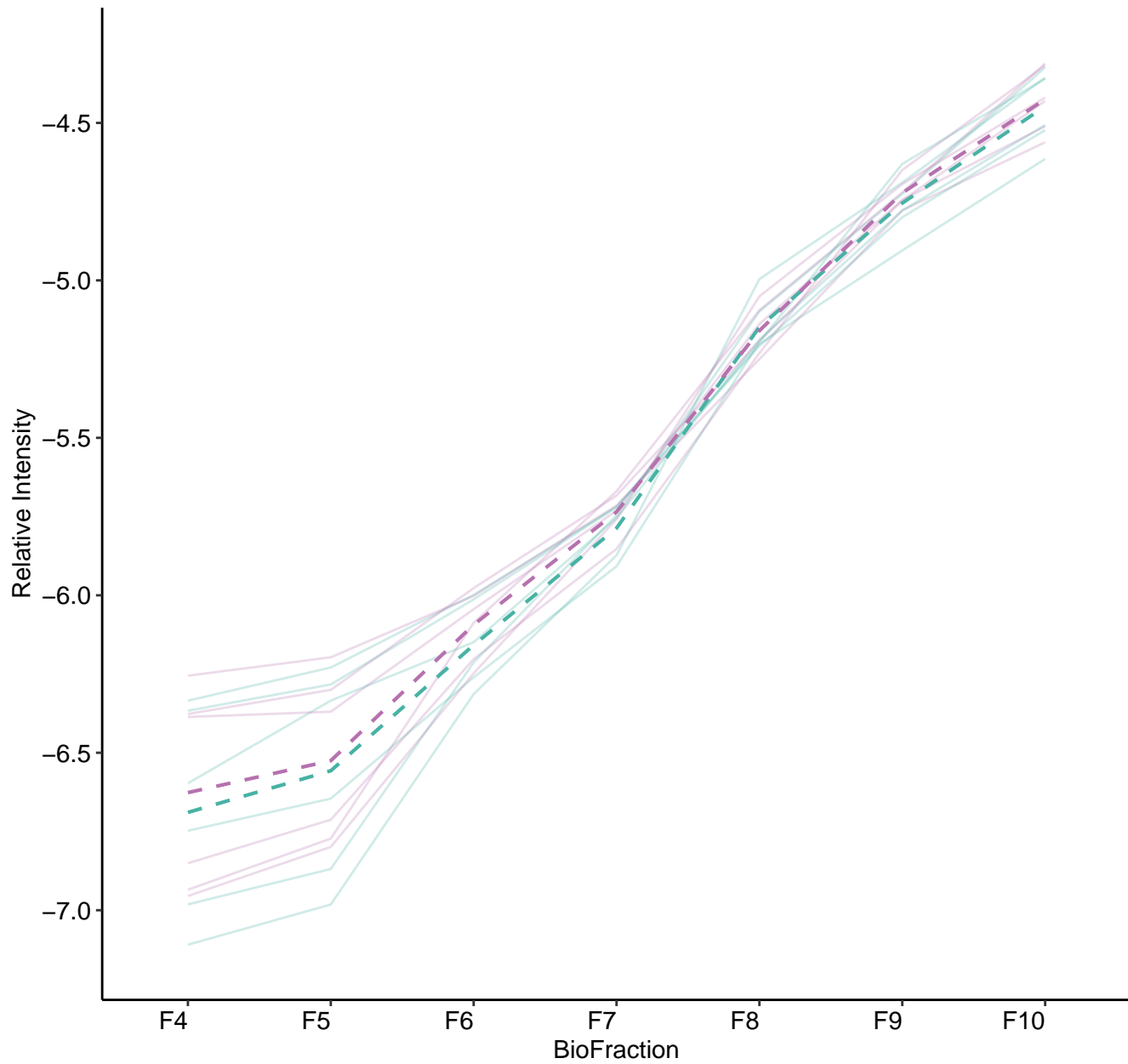
(ER)-localized multiprotein complex, Ig heavy chains associated (n = 4)
(R2.Fixef = 0.985)



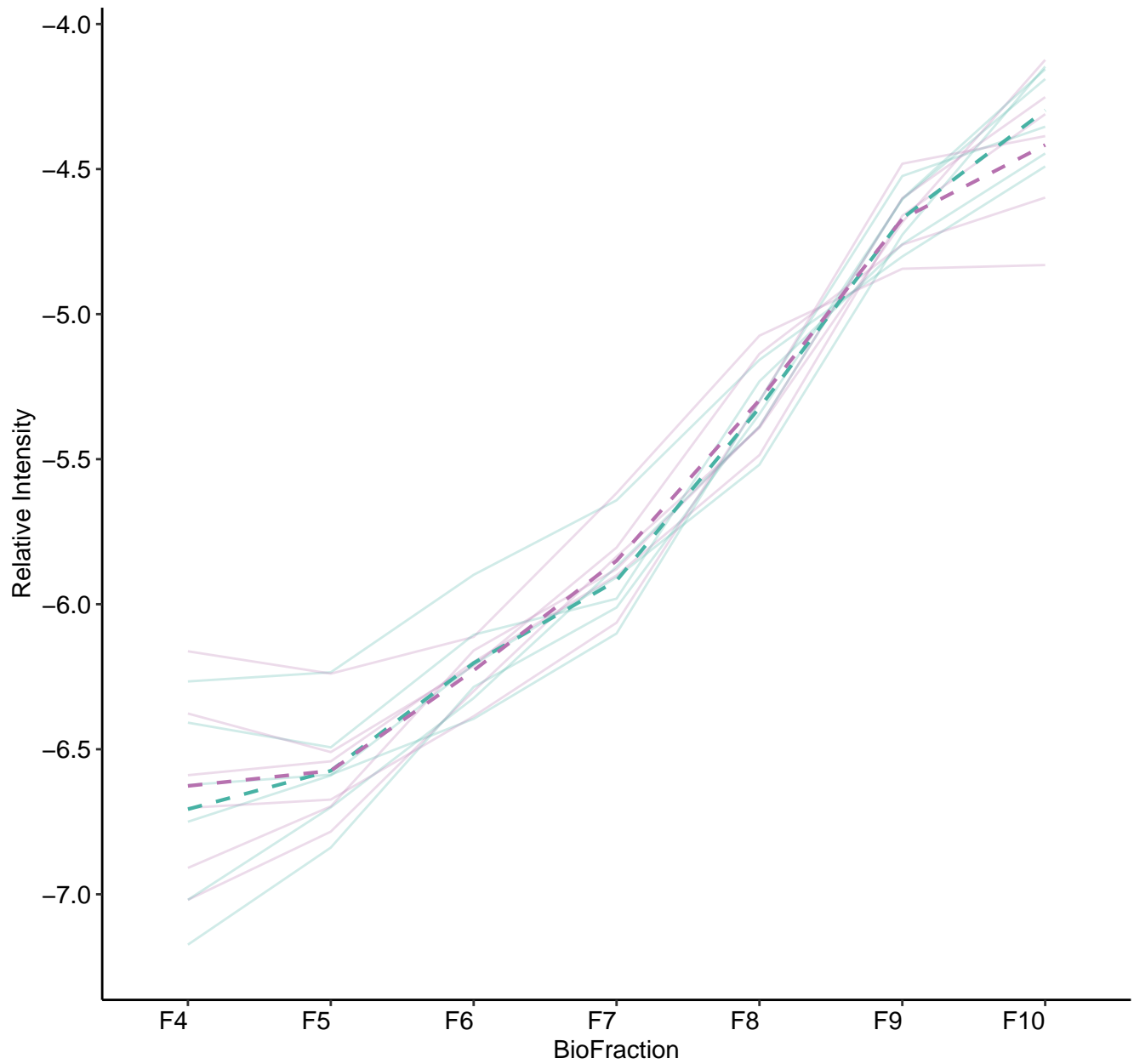
12S U11 snRNP (n = 4)
(R2.Fixef = 0.978)



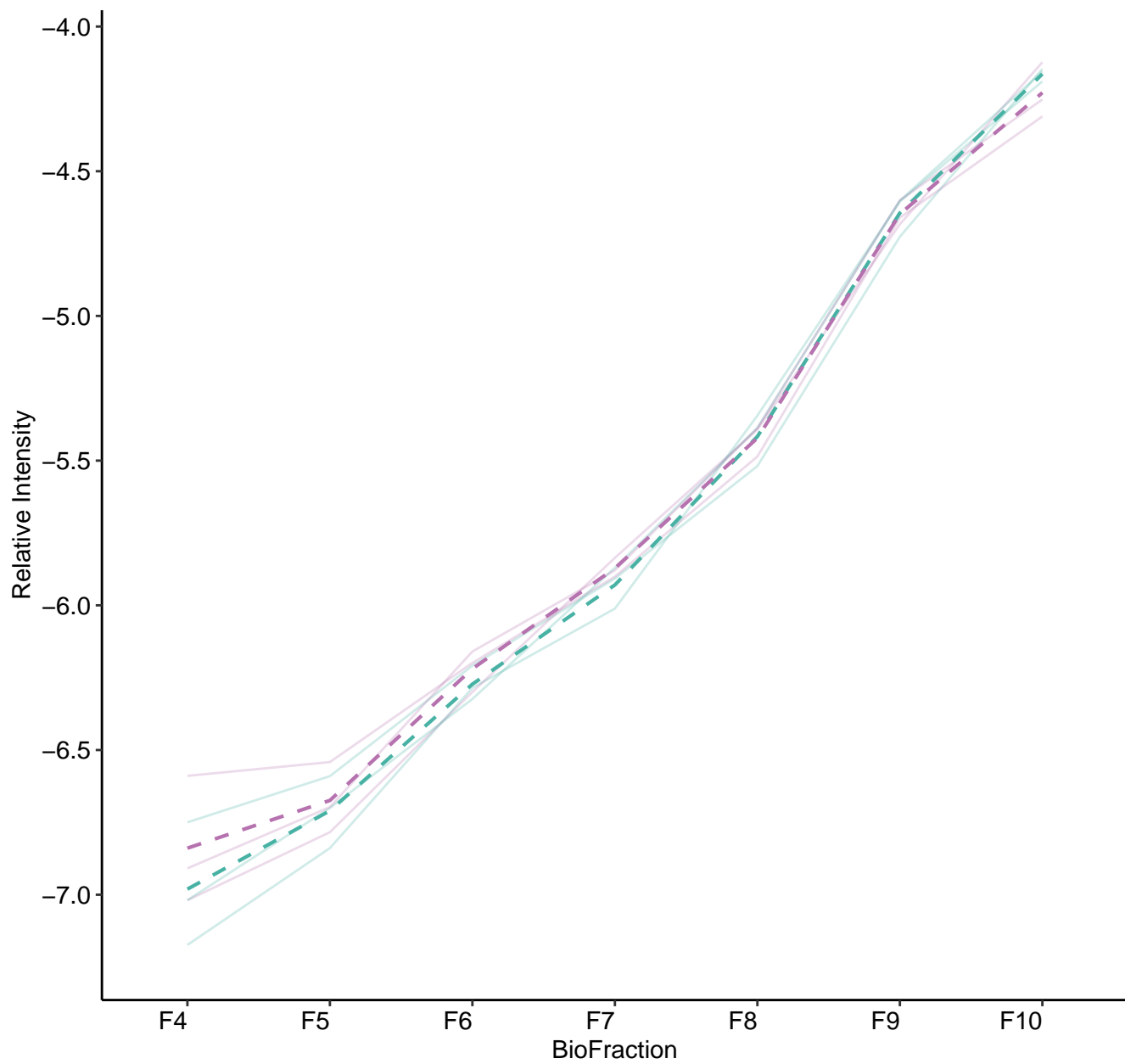
17S U2 snRNP (n = 6)
(R2.Fixef = 0.952)



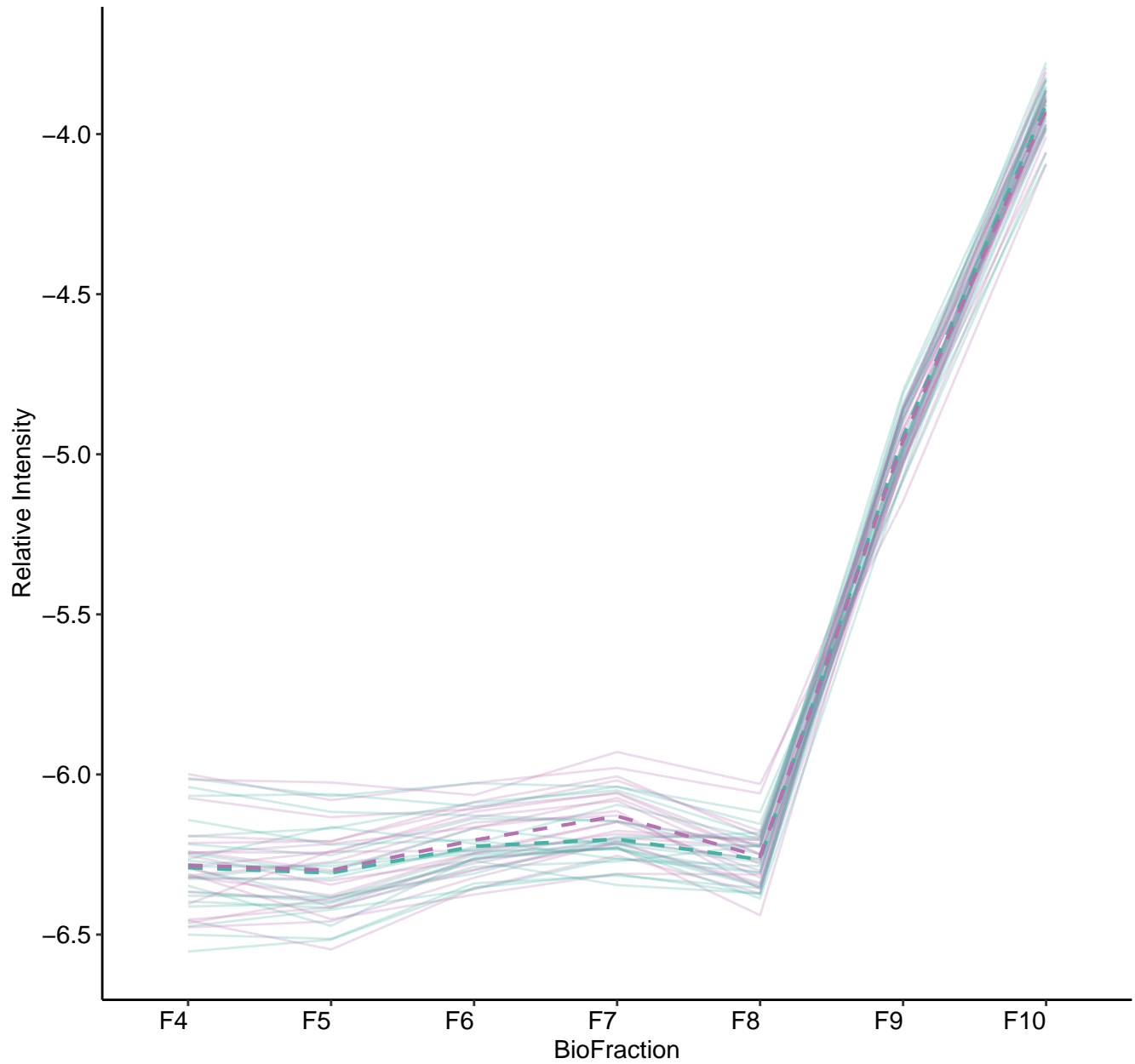
18S U11/U12 snRNP (n = 6)
(R2.Fixef = 0.949)



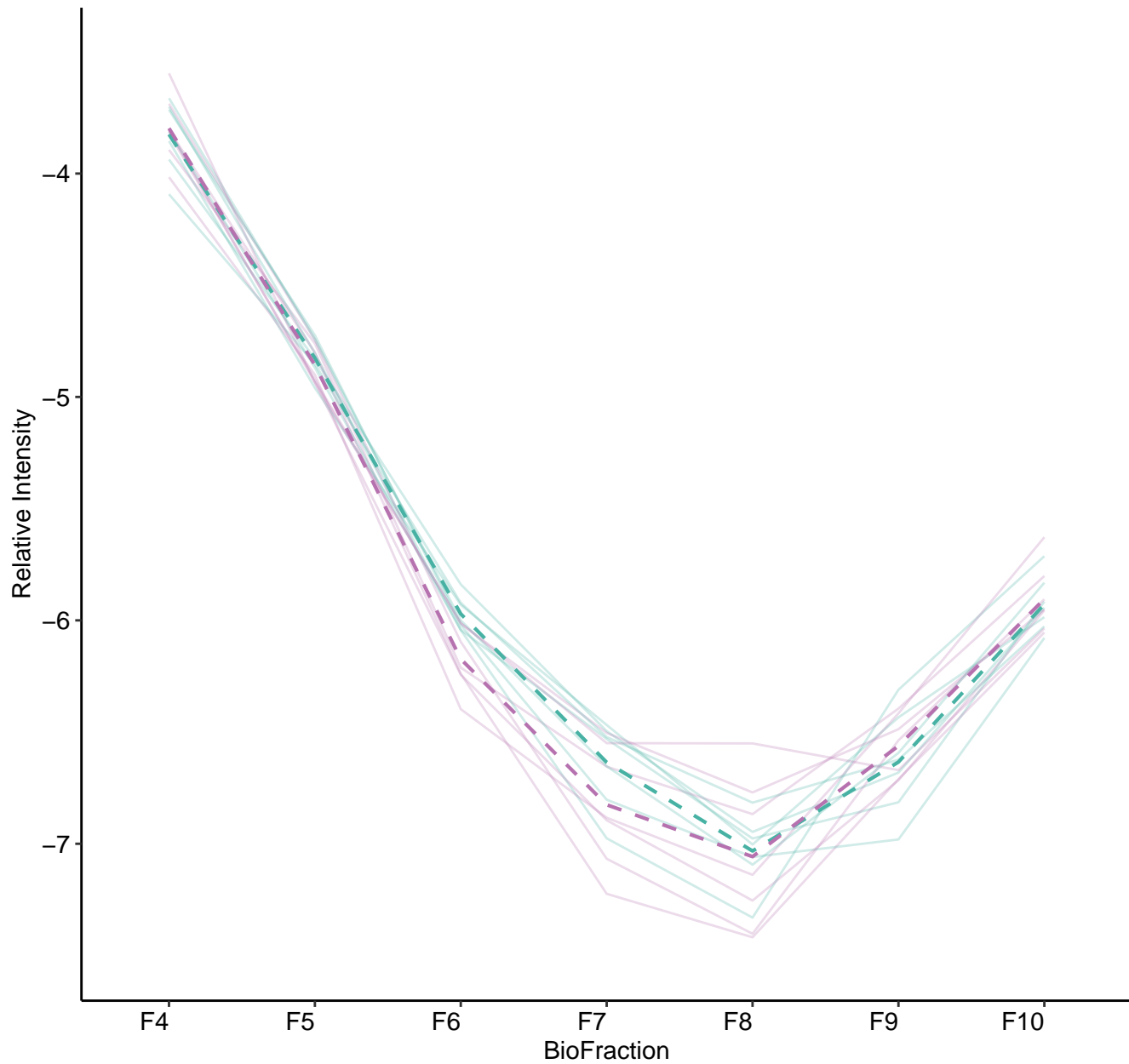
20S methylosome and RG-containing Sm protein complex (n = 3)
(R2.Fixef = 0.987)



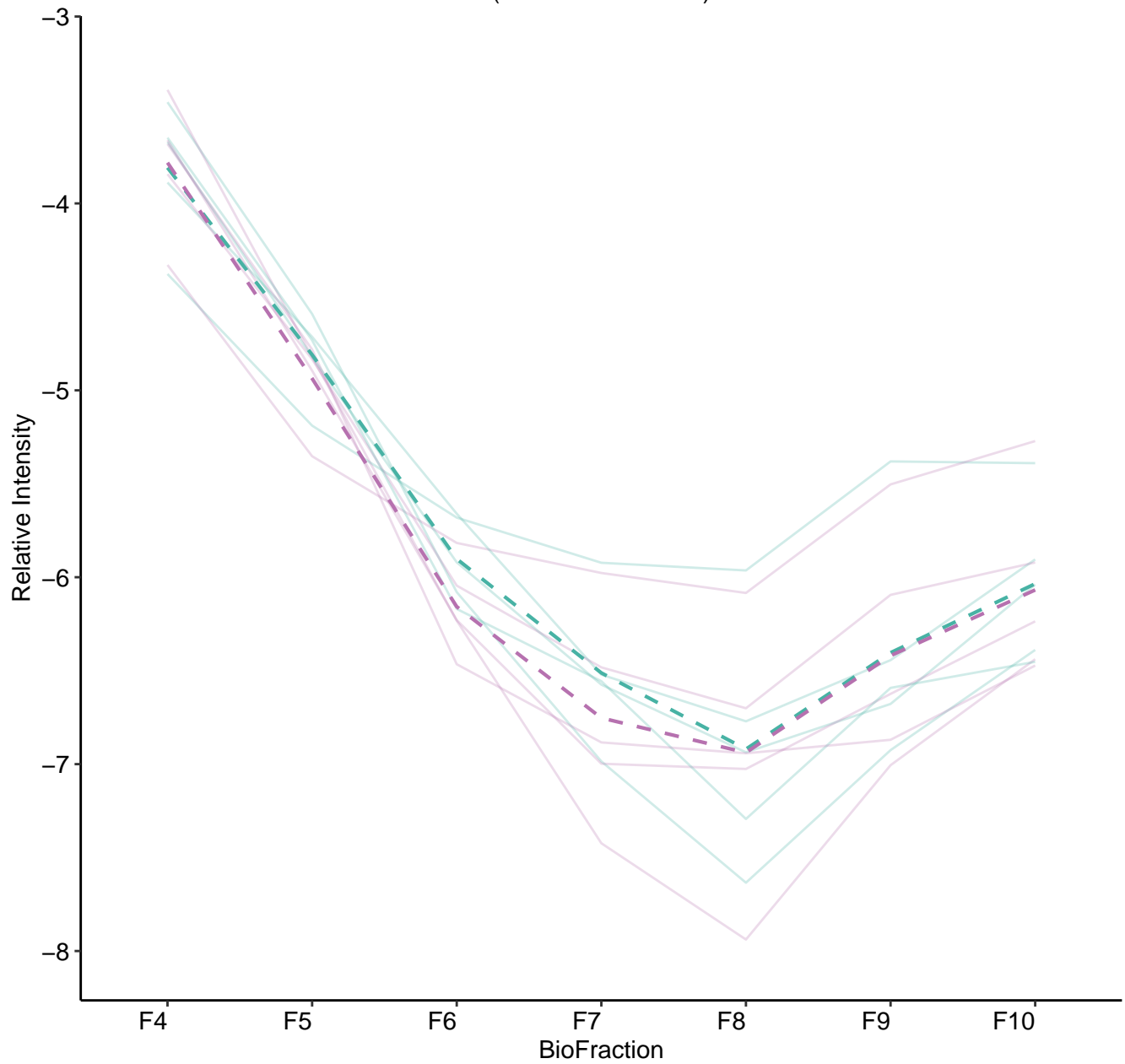
26S proteasome (n = 21)
(R2.Fixef = 0.986)



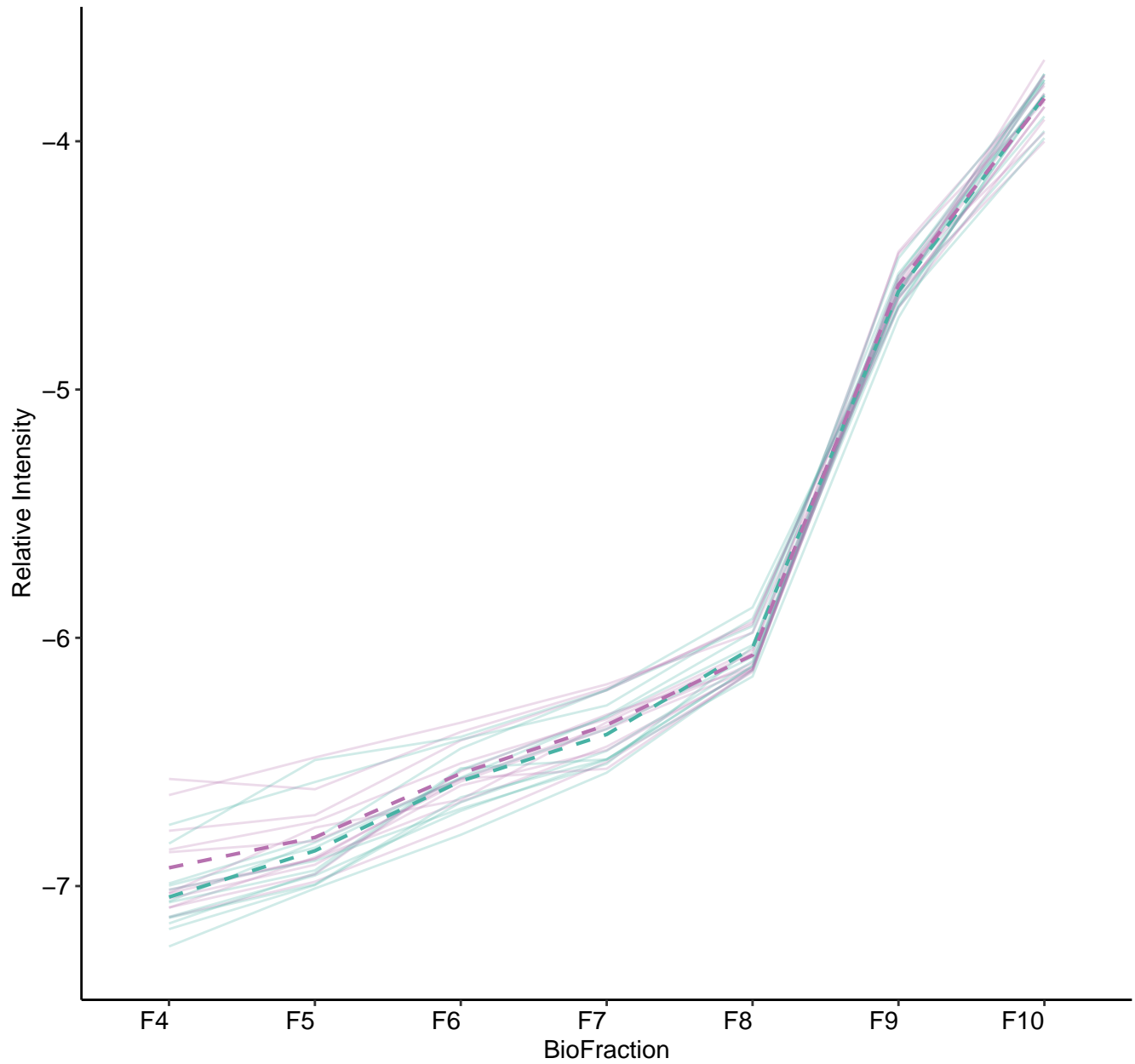
28S ribosomal subunit, mitochondrial (n = 7)
(R2.Fixef = 0.974)



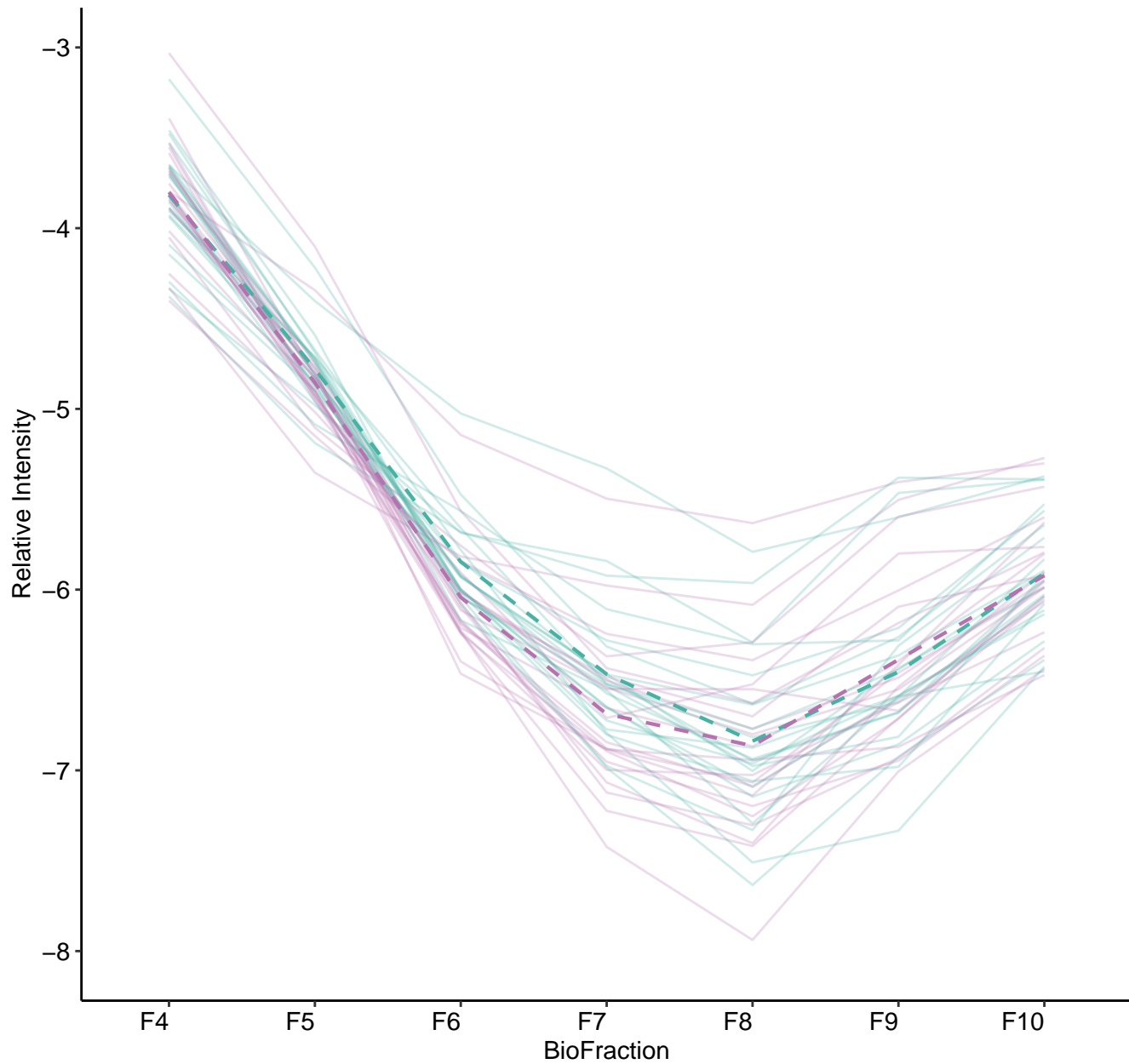
39S ribosomal subunit, mitochondrial (n = 5)
(R2.Fixef = 0.836)



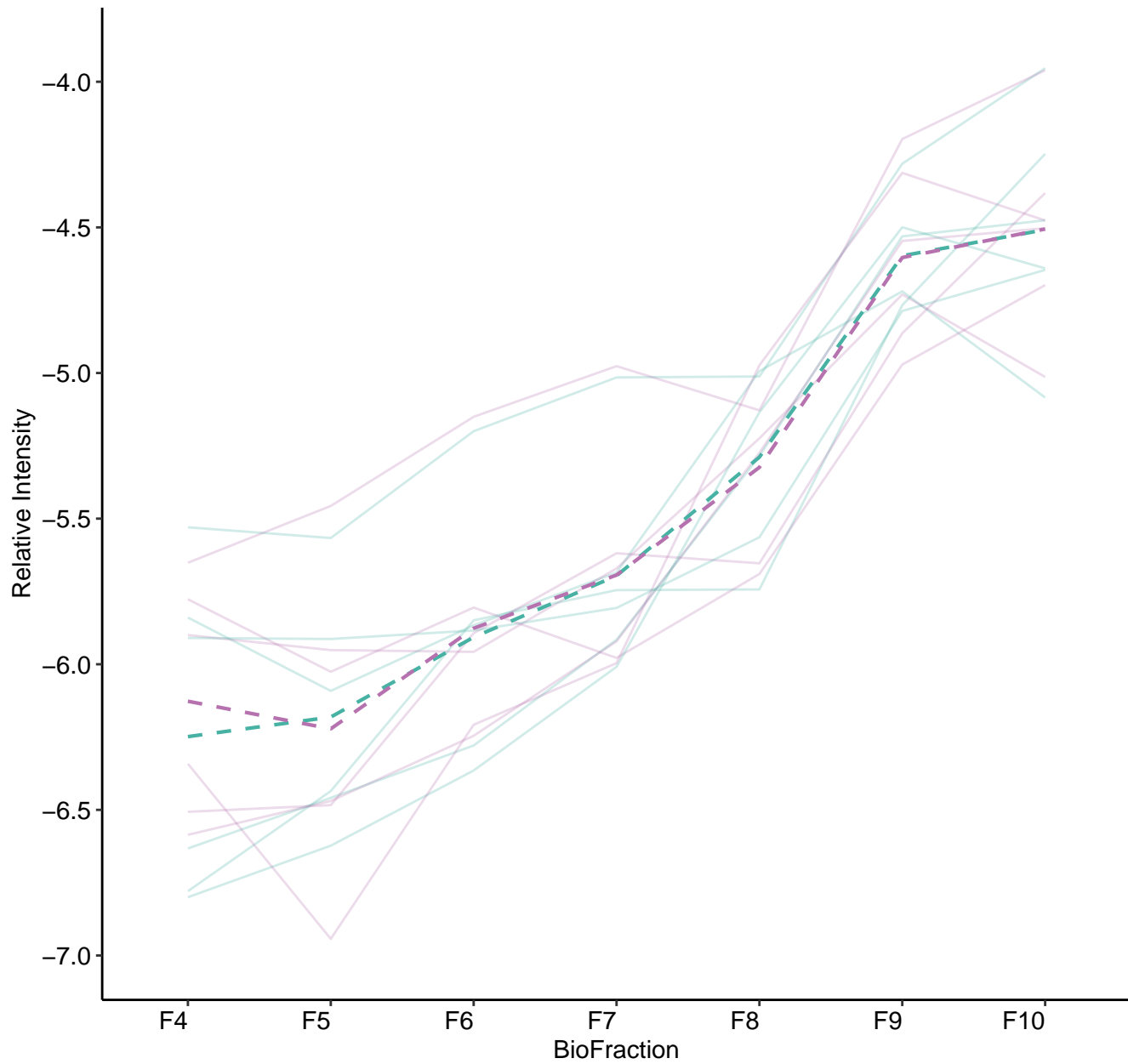
40S ribosomal subunit, cytoplasmic (n = 12)
(R2.Fixef = 0.989)



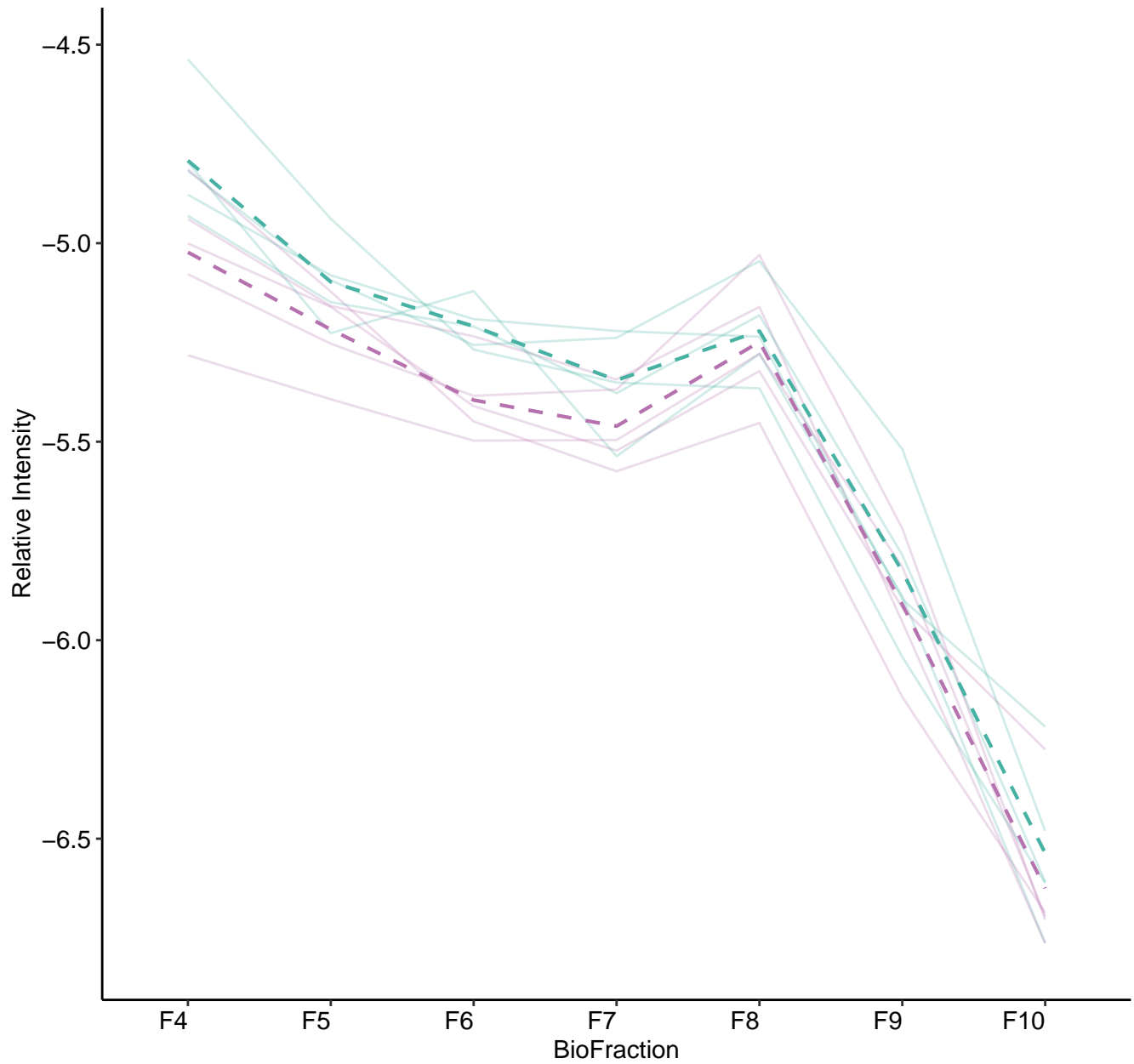
55S ribosome, mitochondrial (n = 22)
(R2.Fixef = 0.881)



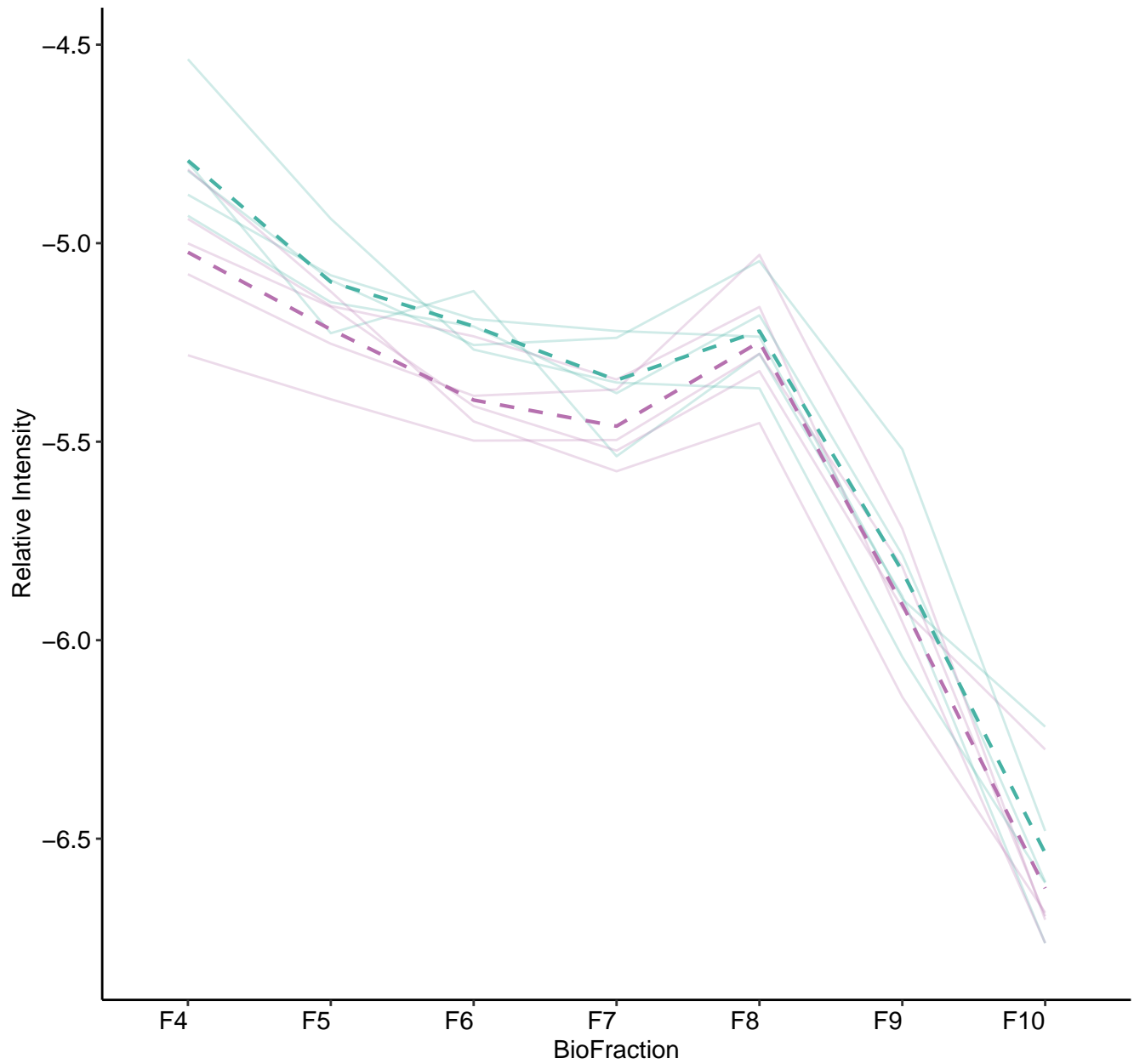
ALL-1 supercomplex (n = 6)
(R2.Fixef = 0.745)



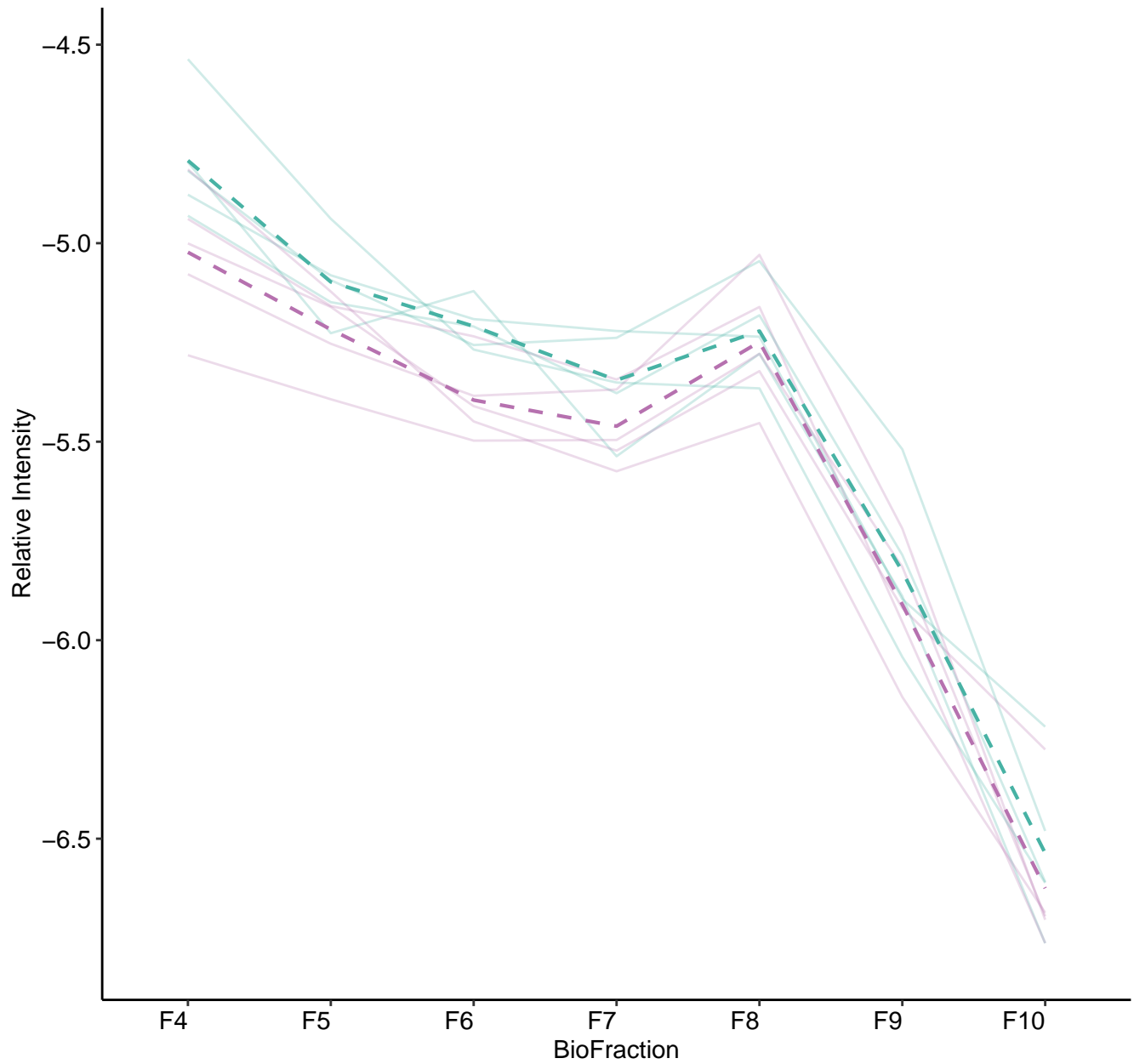
AMPA receptor complex (anti-GluA1-a) (n = 5)
(R2.Fixef = 0.928)



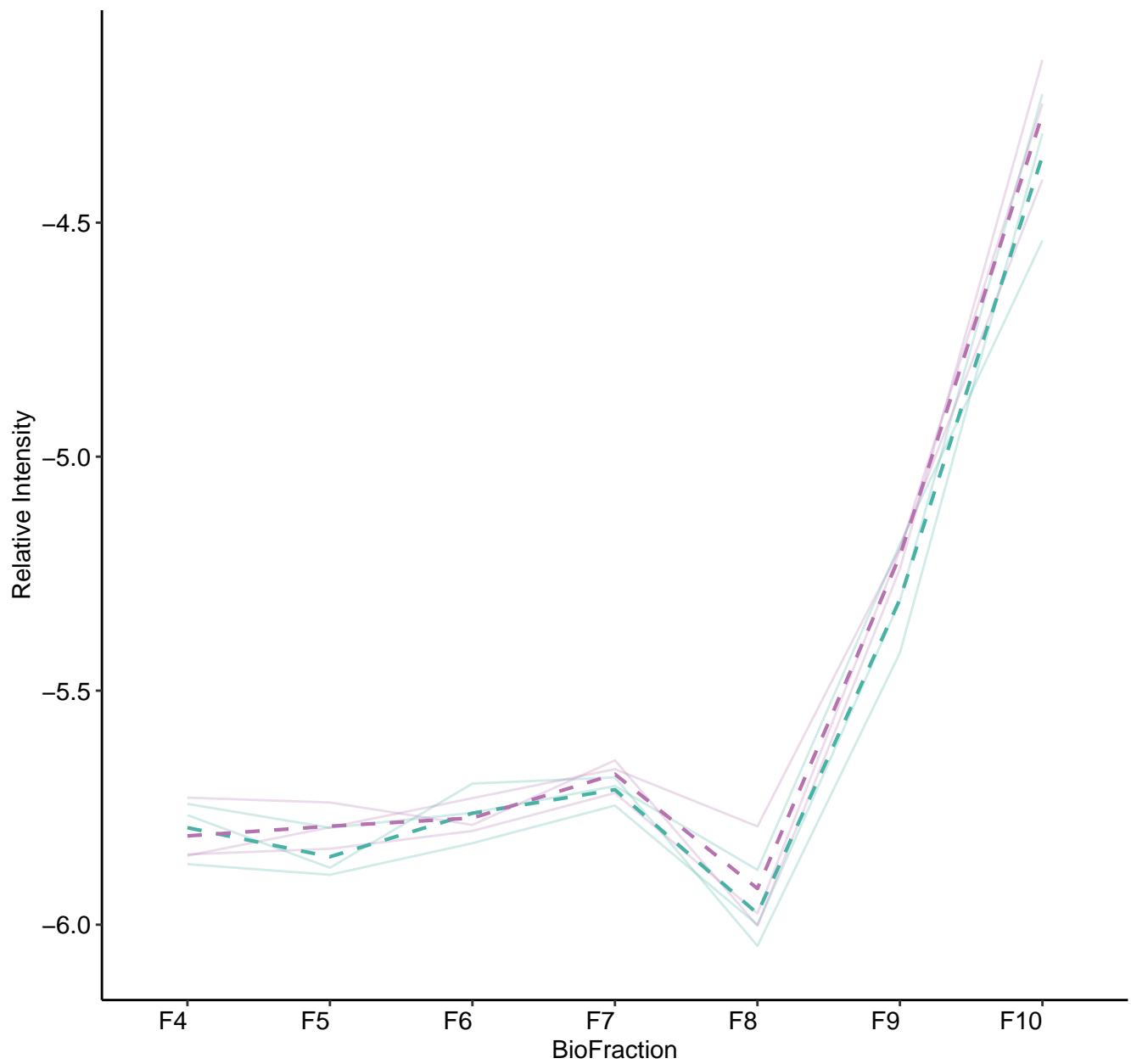
AMPA receptor complex (anti-GluA1-c) (n = 5)
(R2.Fixef = 0.928)



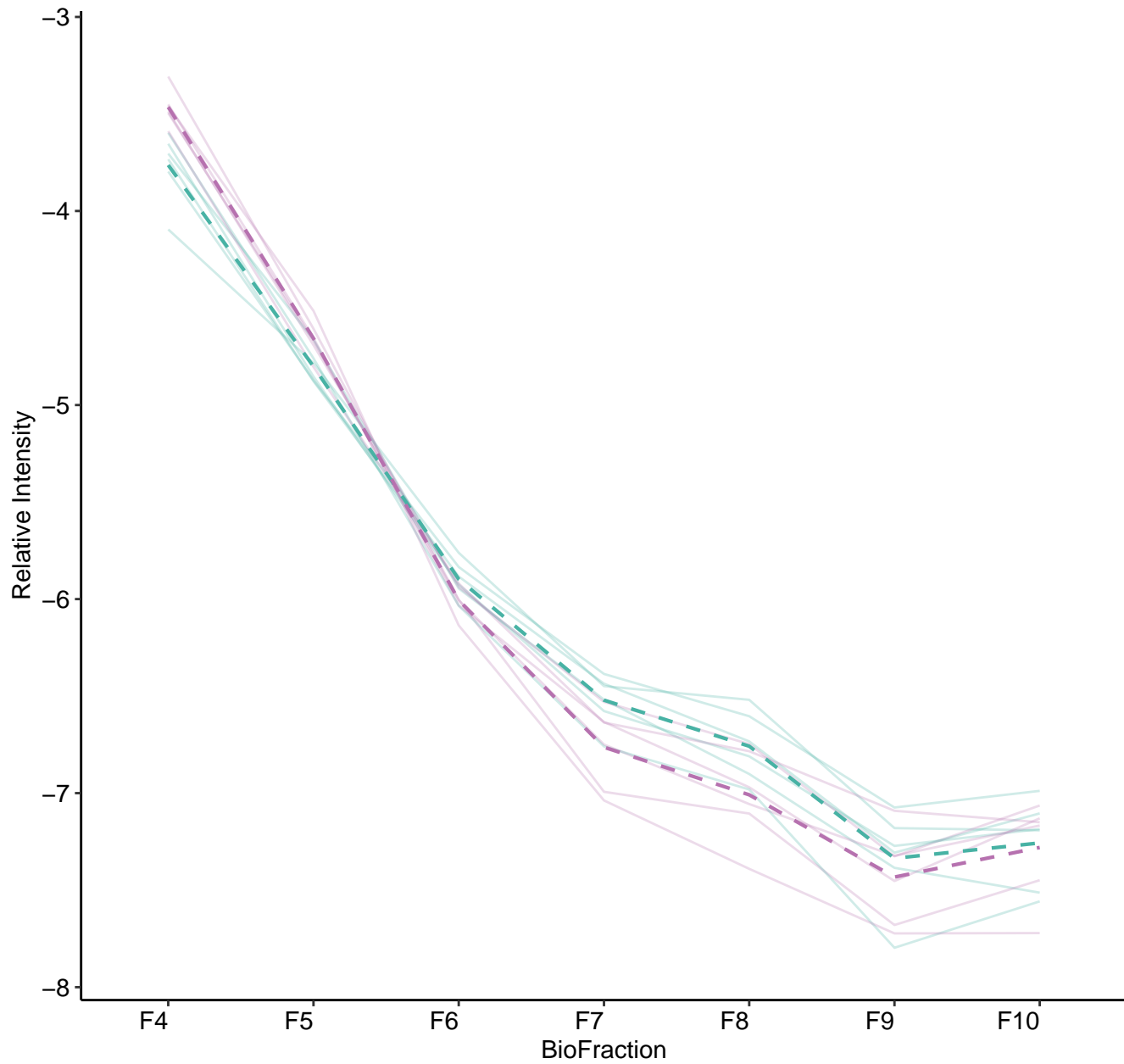
AMPA receptor complex (anti-GluA2-a) (n = 5)
(R2.Fixef = 0.928)



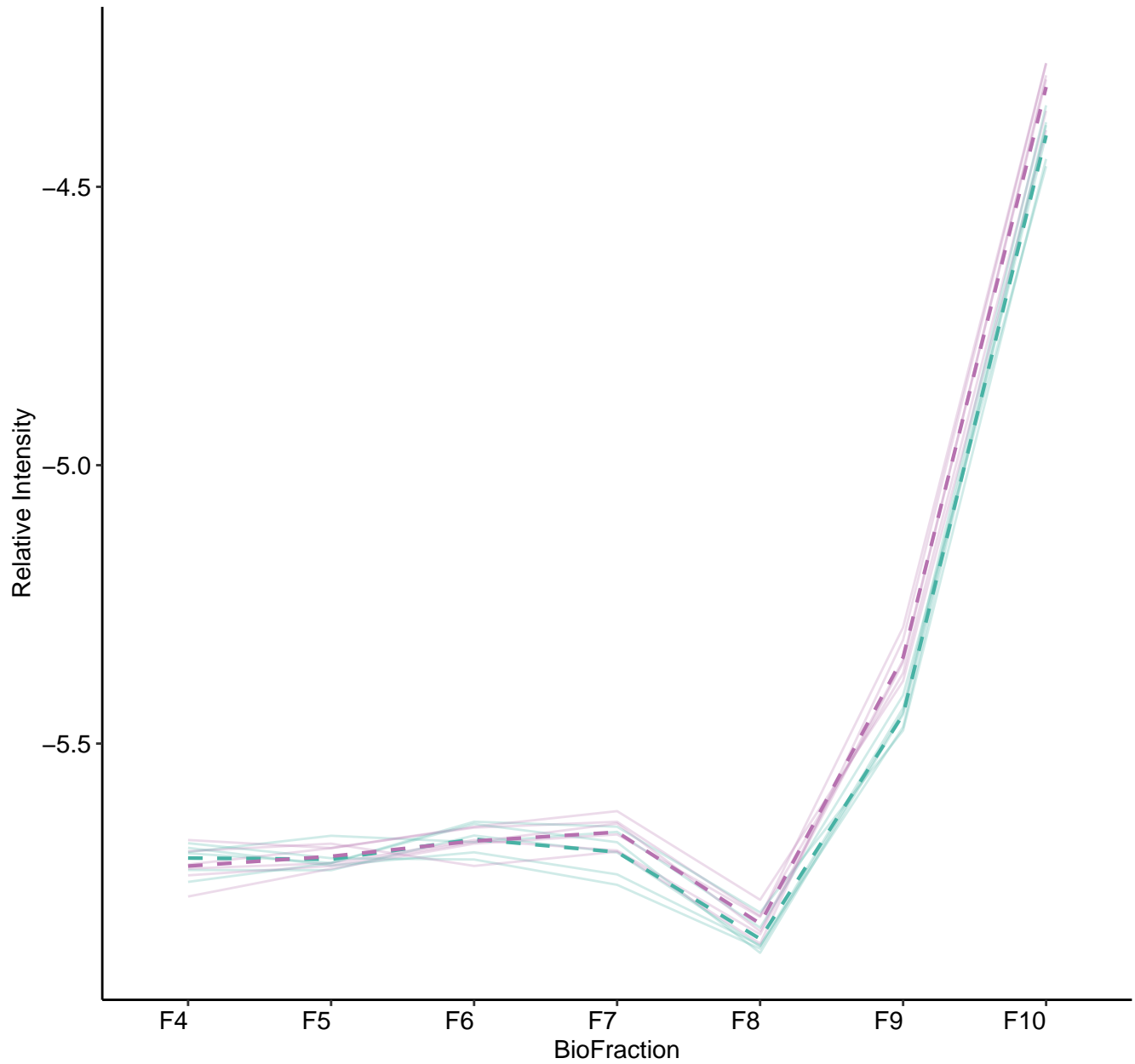
Anaphase-promoting complex (n = 3)
(R2.Fixef = 0.976)



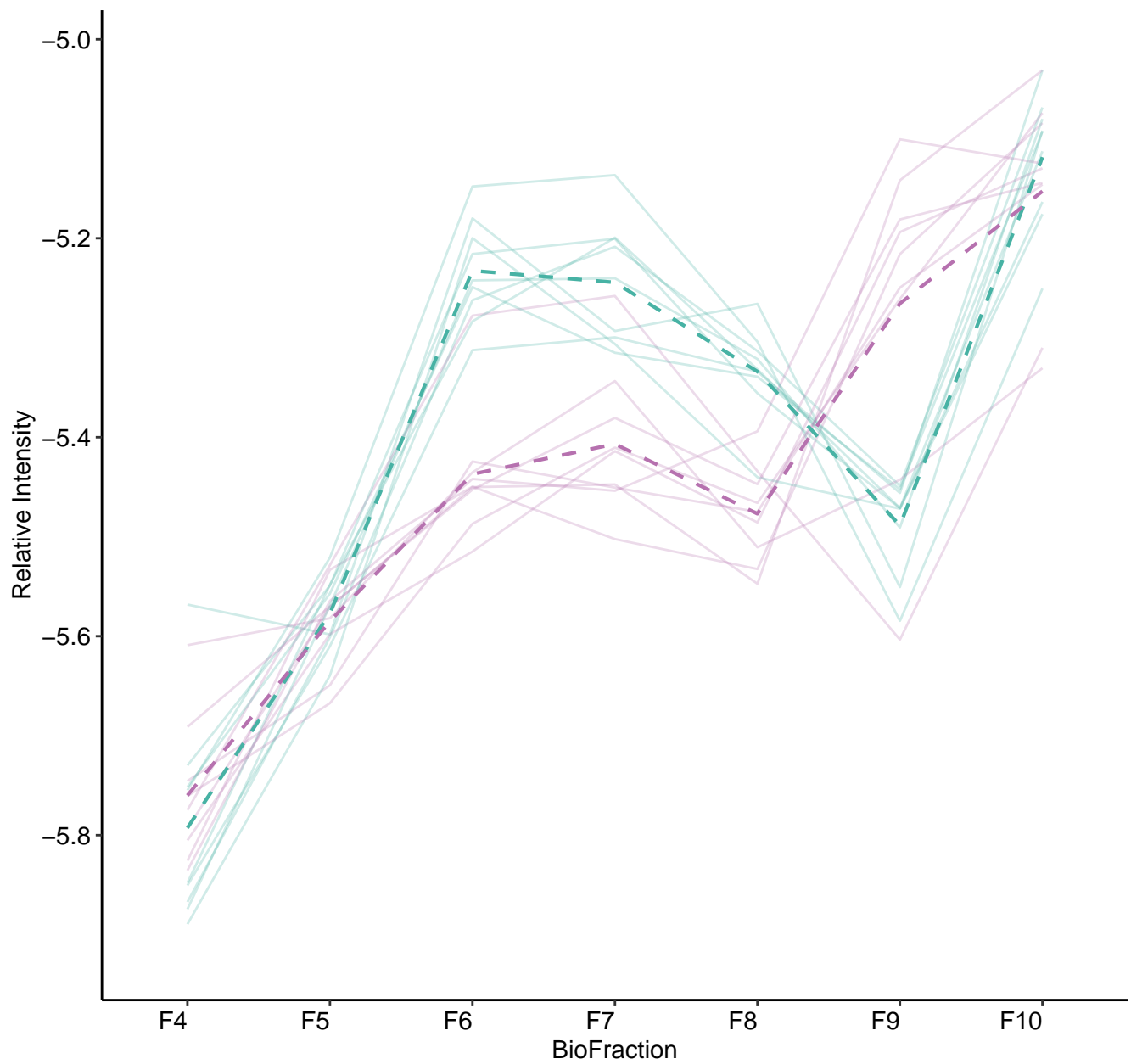
ATP synthasome (n = 6)
(R2.Fixef = 0.982)



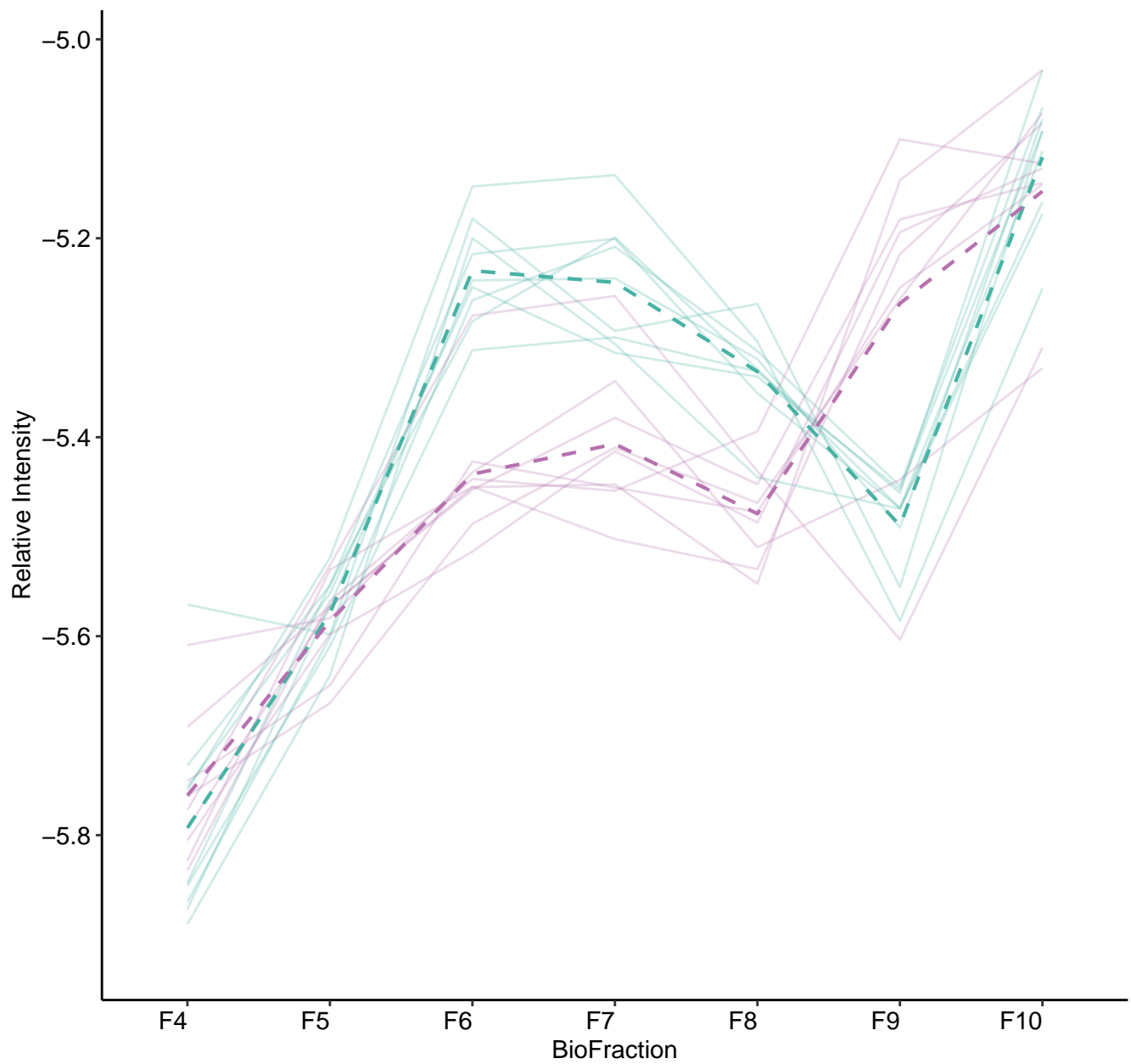
BBS-chaperonin complex (n = 6)
(R2.Fixef = 0.995)



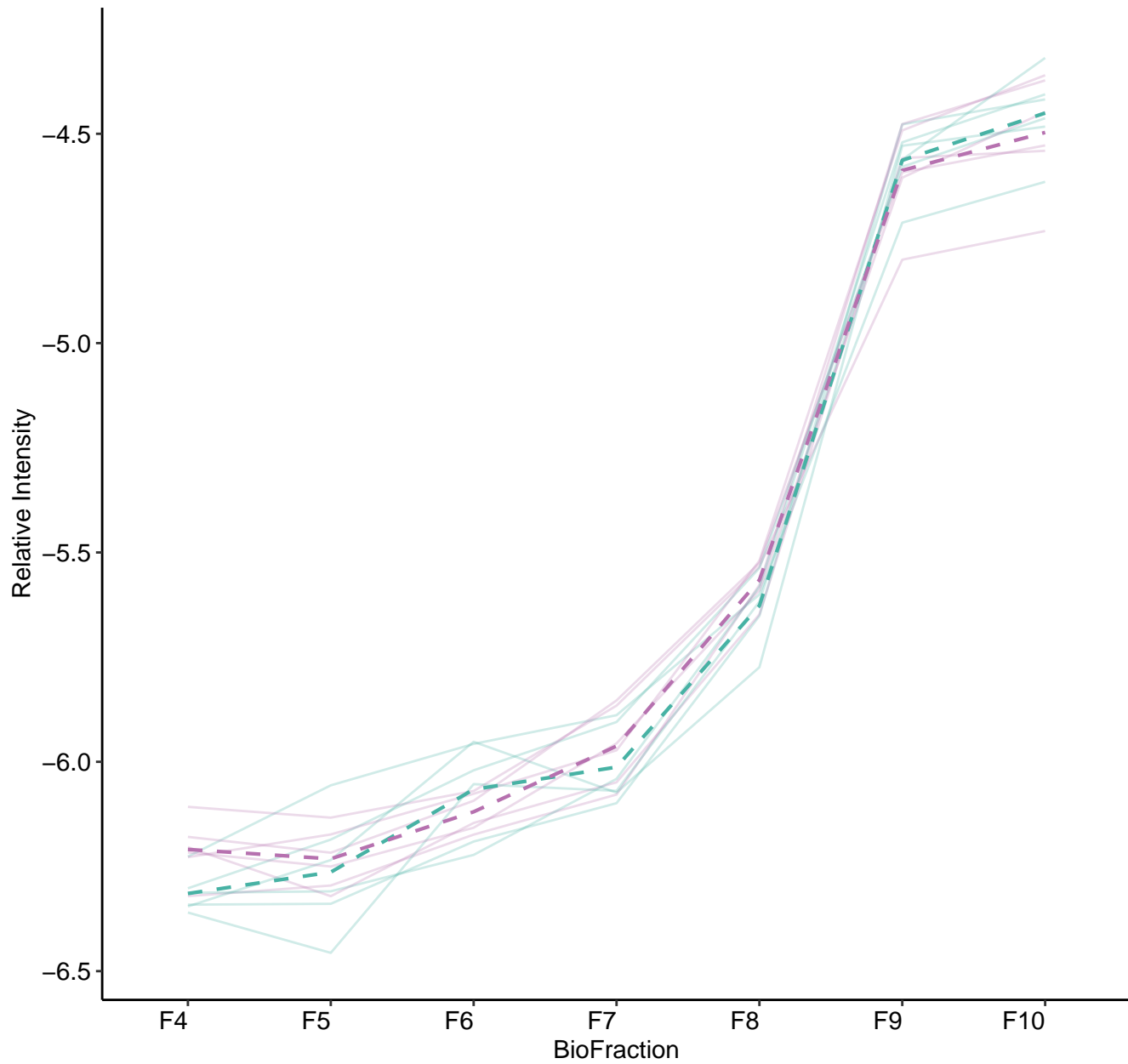
CCC complex (n = 9)
(R2.Fixef = 0.875)



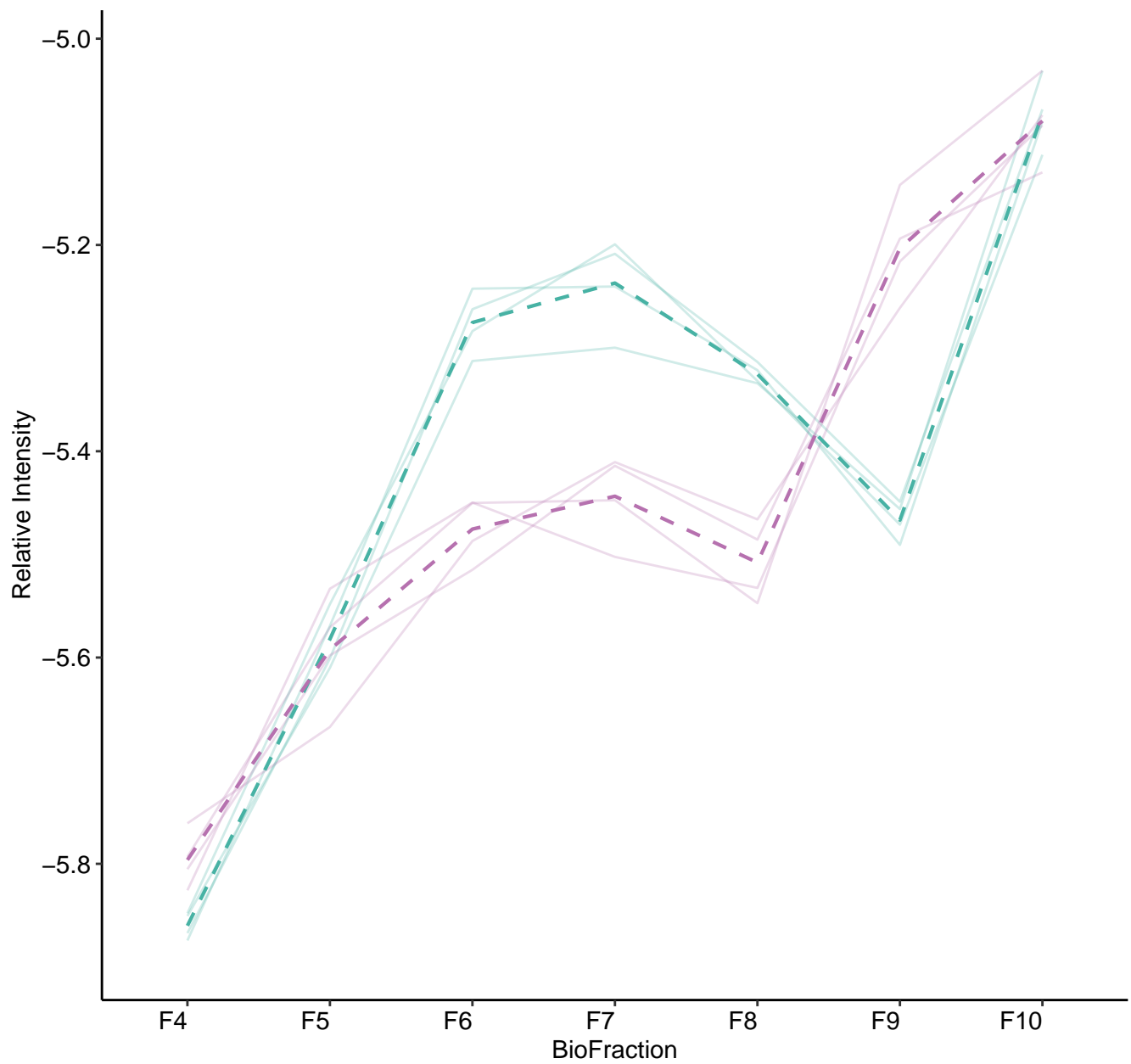
CCC-Wash (WASH1, FAM21C) complex (n = 9)
(R2.Fixef = 0.875)



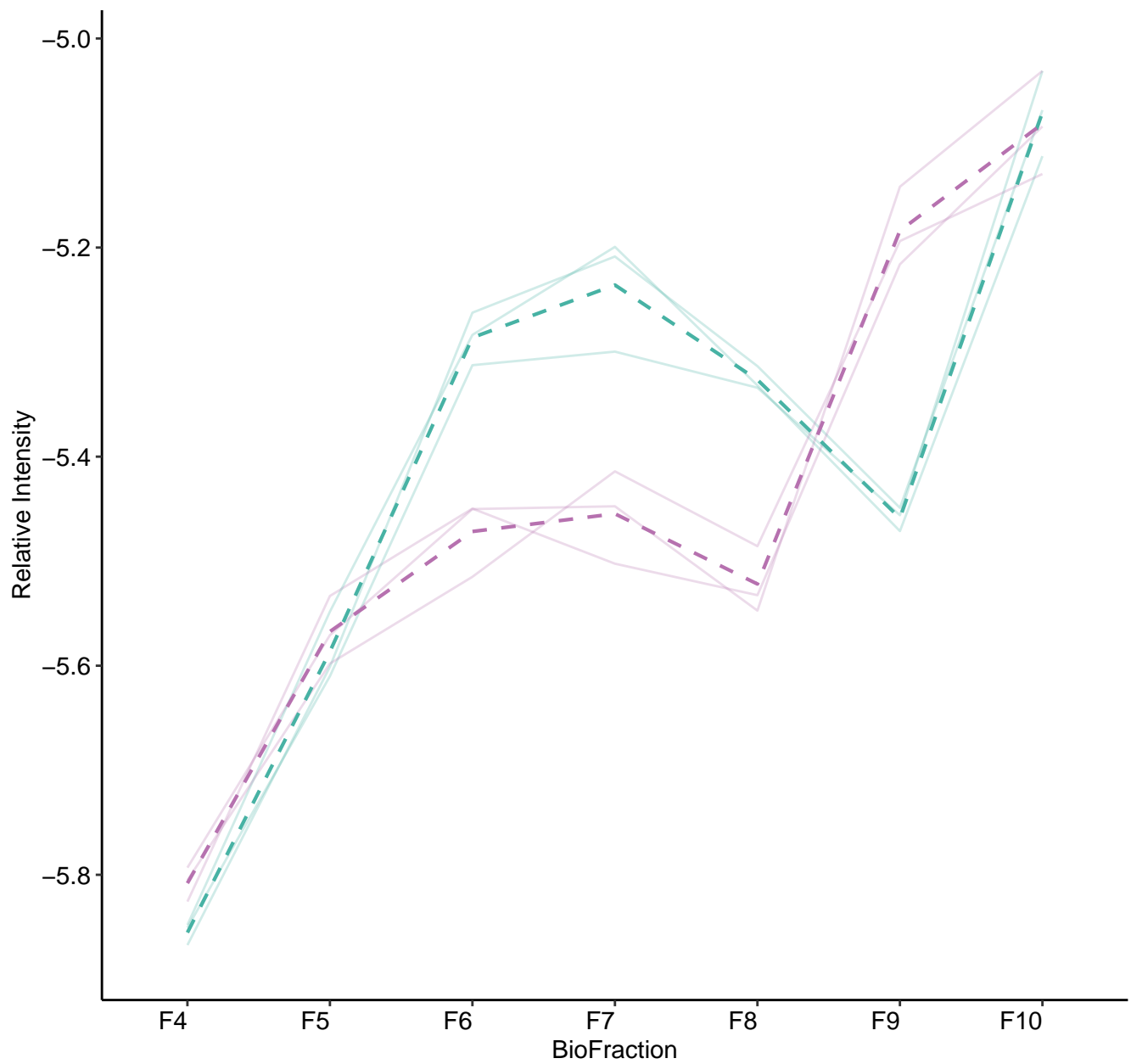
CCR4–NOT–CNOT7–CNOT6L complex (n = 6)
(R2.Fixef = 0.983)



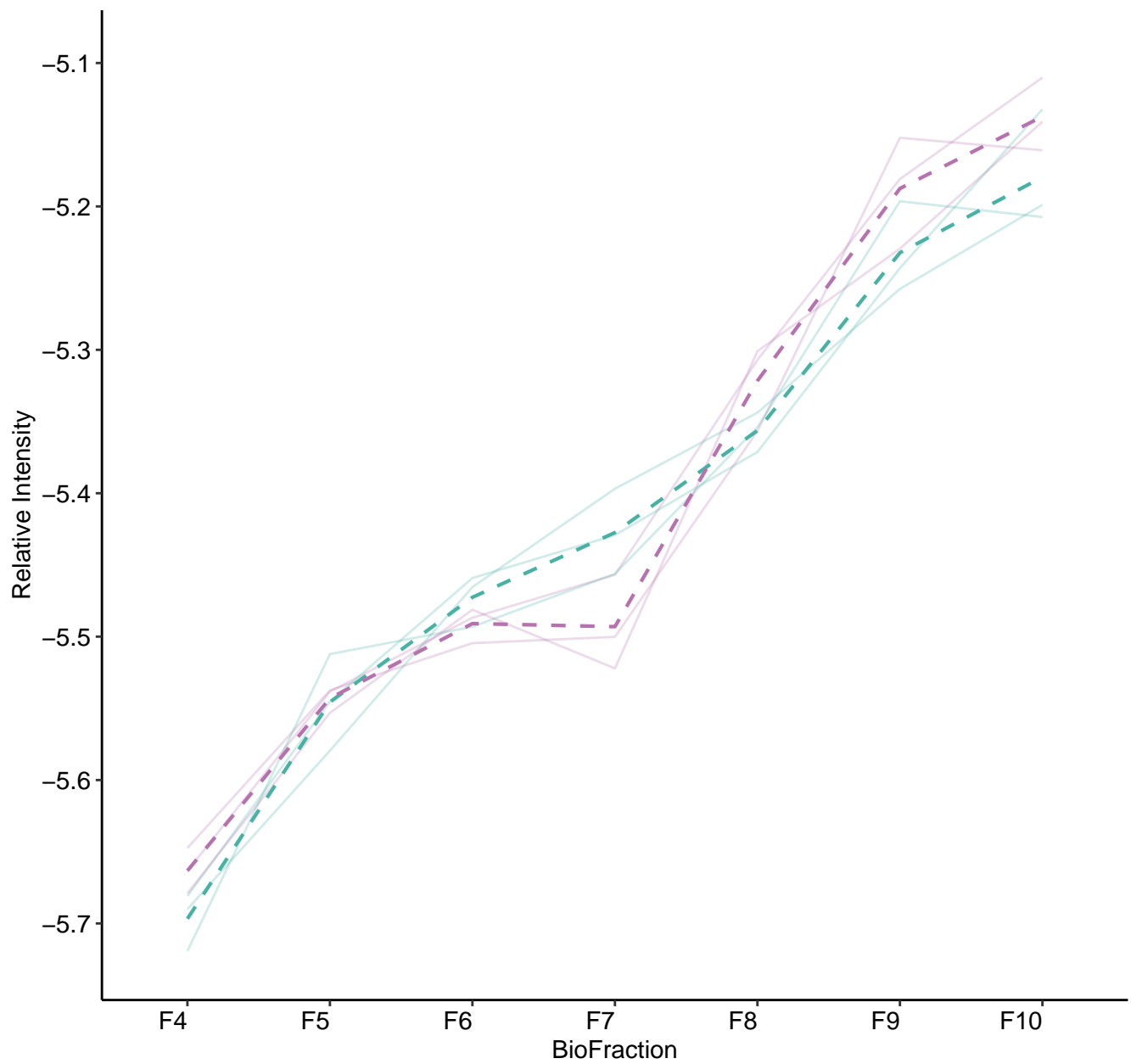
COMMD1-CCDC22-CCDC93-C16orf62 complex (n = 4)
(R2.Fixef = 0.977)



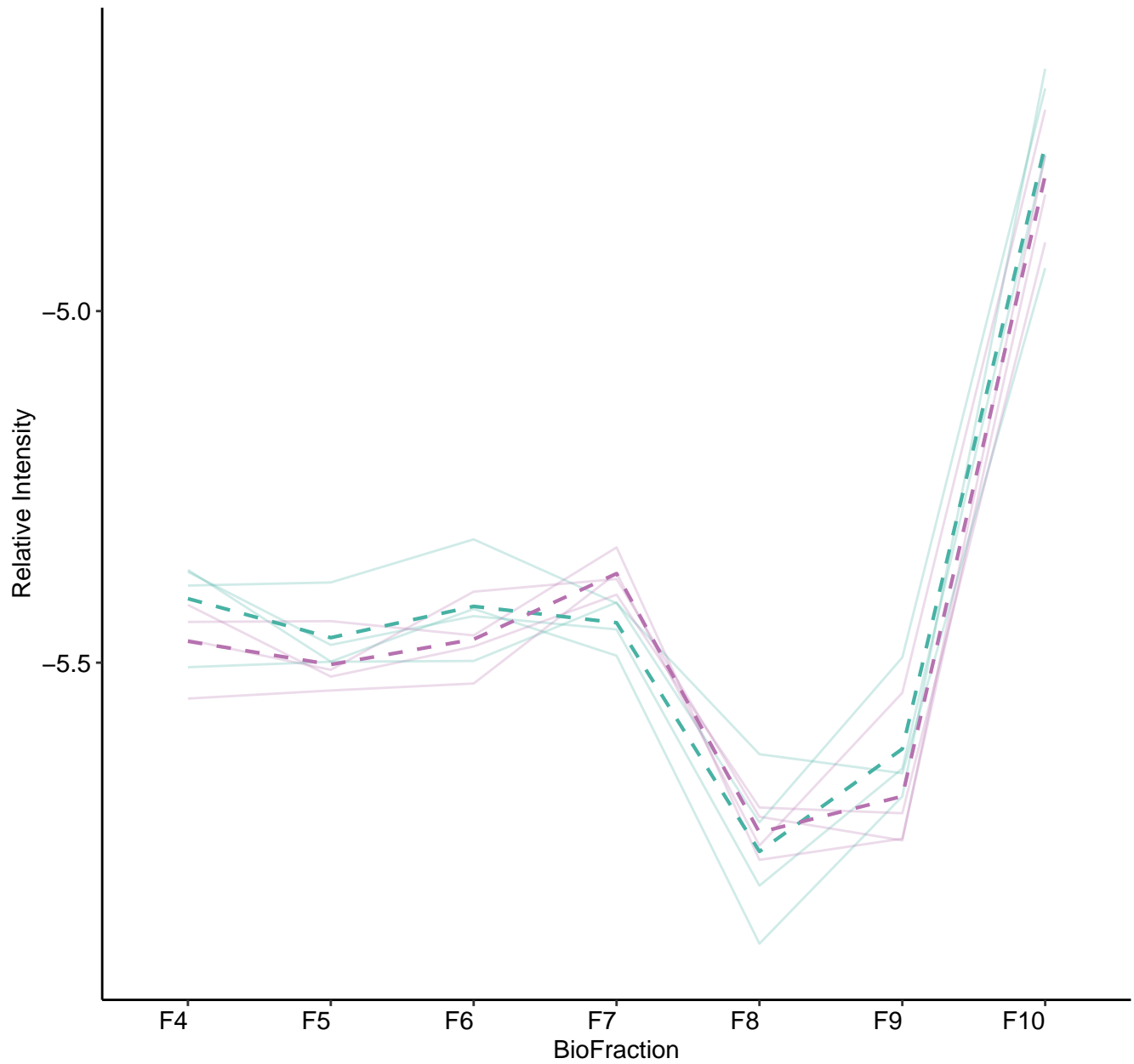
COMMD1-CCDC93-C16orf62 complex (n = 3)
(R2.Fixef = 0.979)



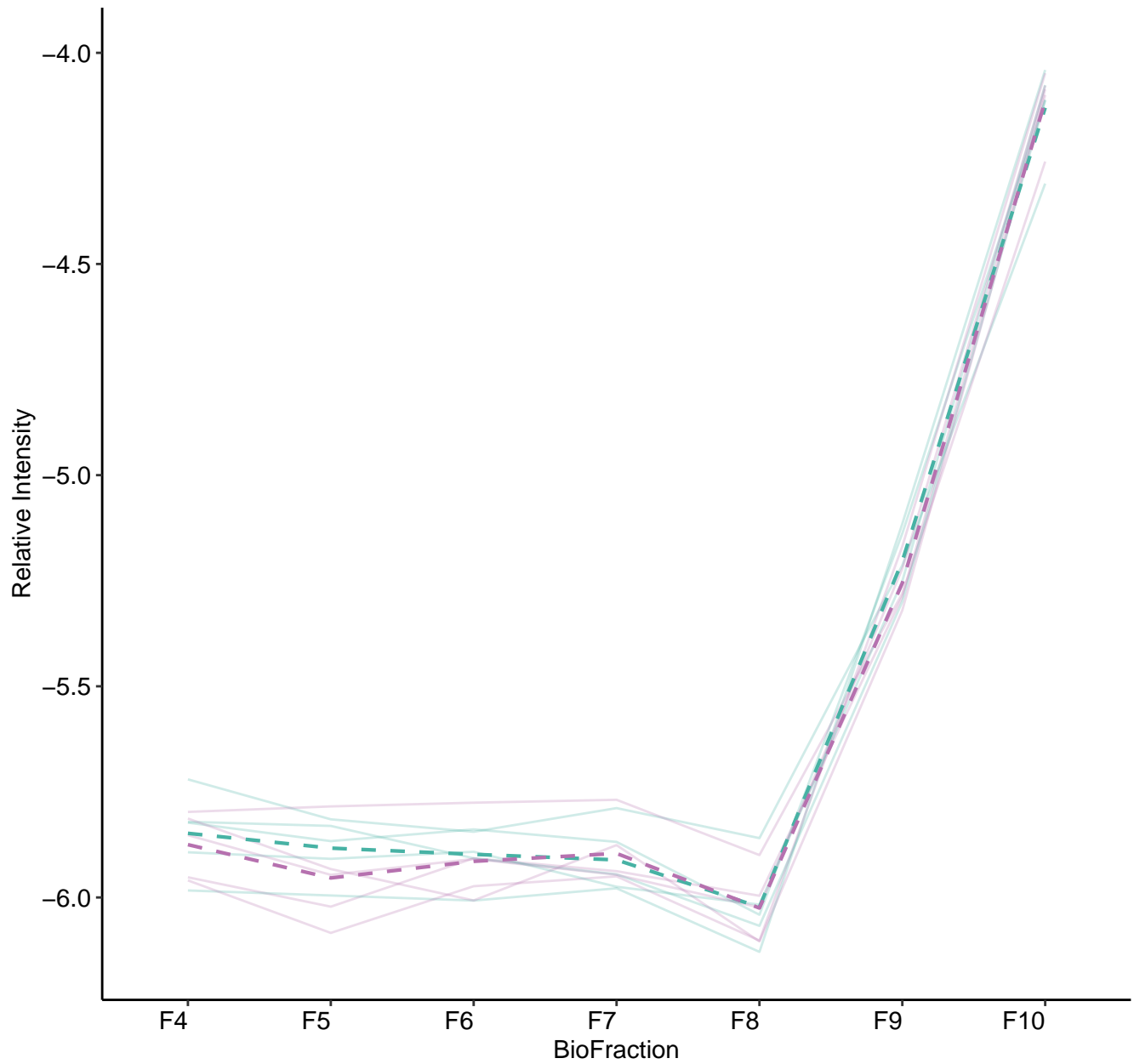
CORVET complex (n = 3)
(R2.Fixef = 0.976)



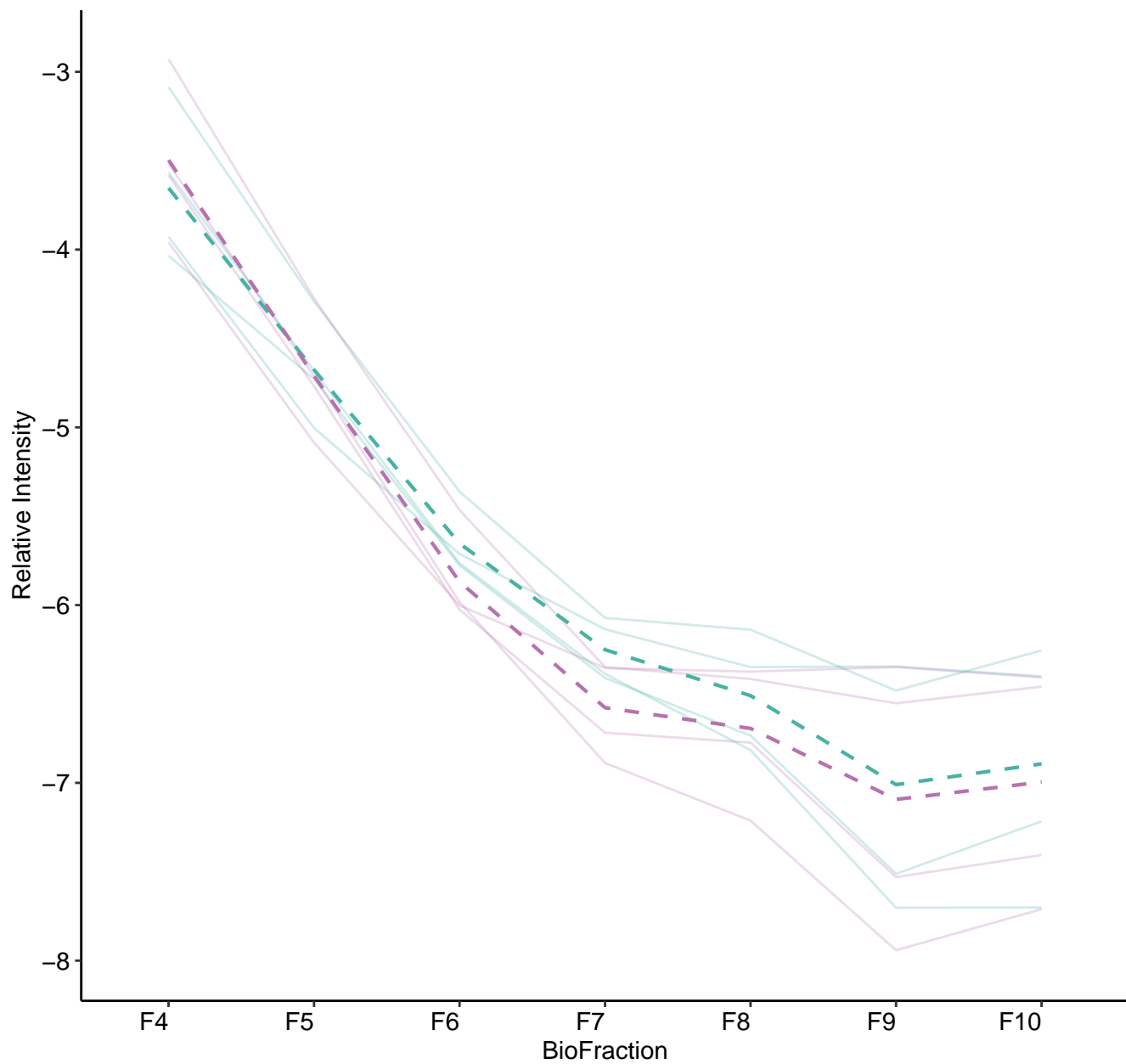
CSA-POLIIa complex (n = 4)
(R2.Fixef = 0.938)



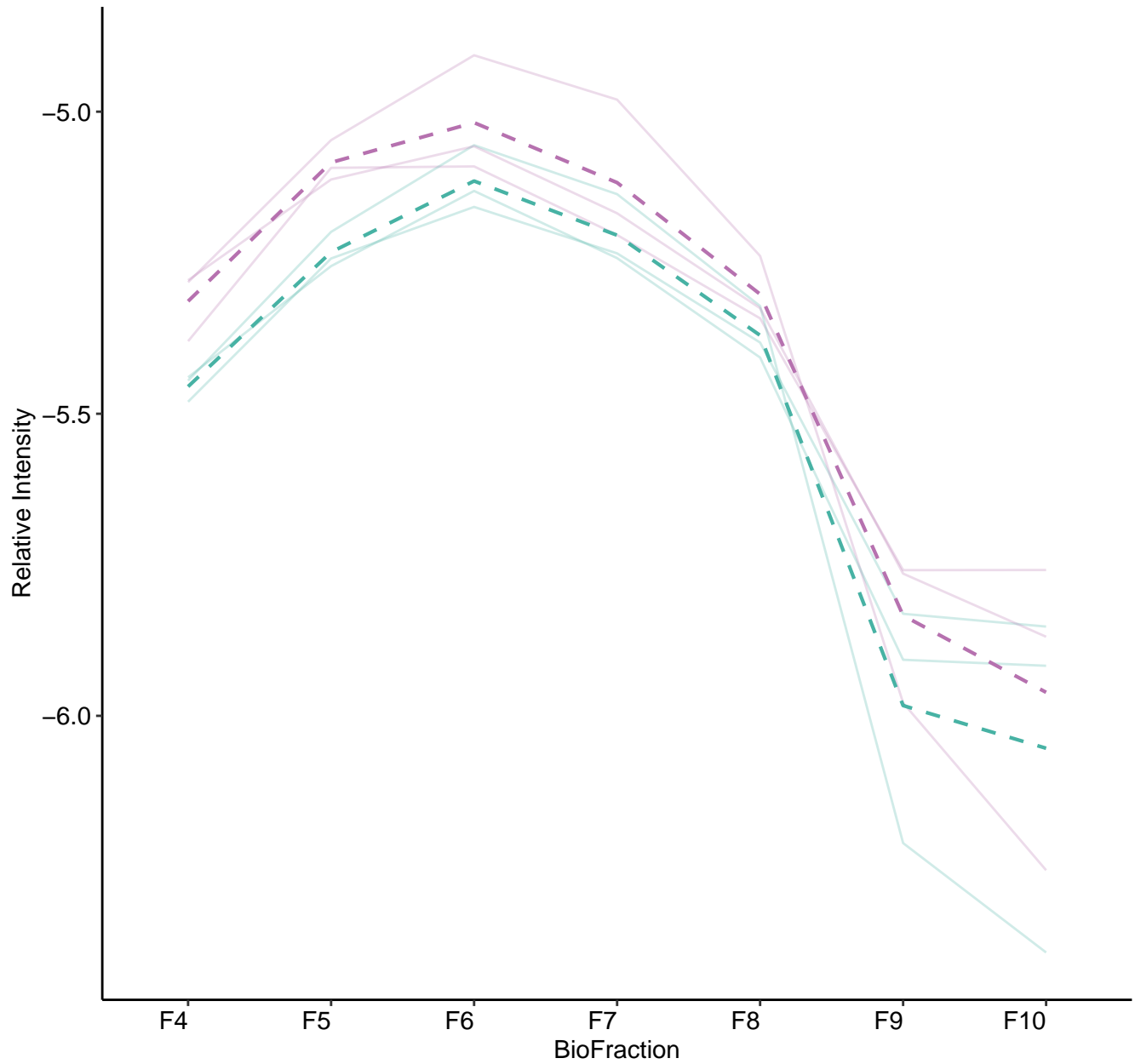
CTLH complex (n = 5)
(R2.Fixef = 0.983)



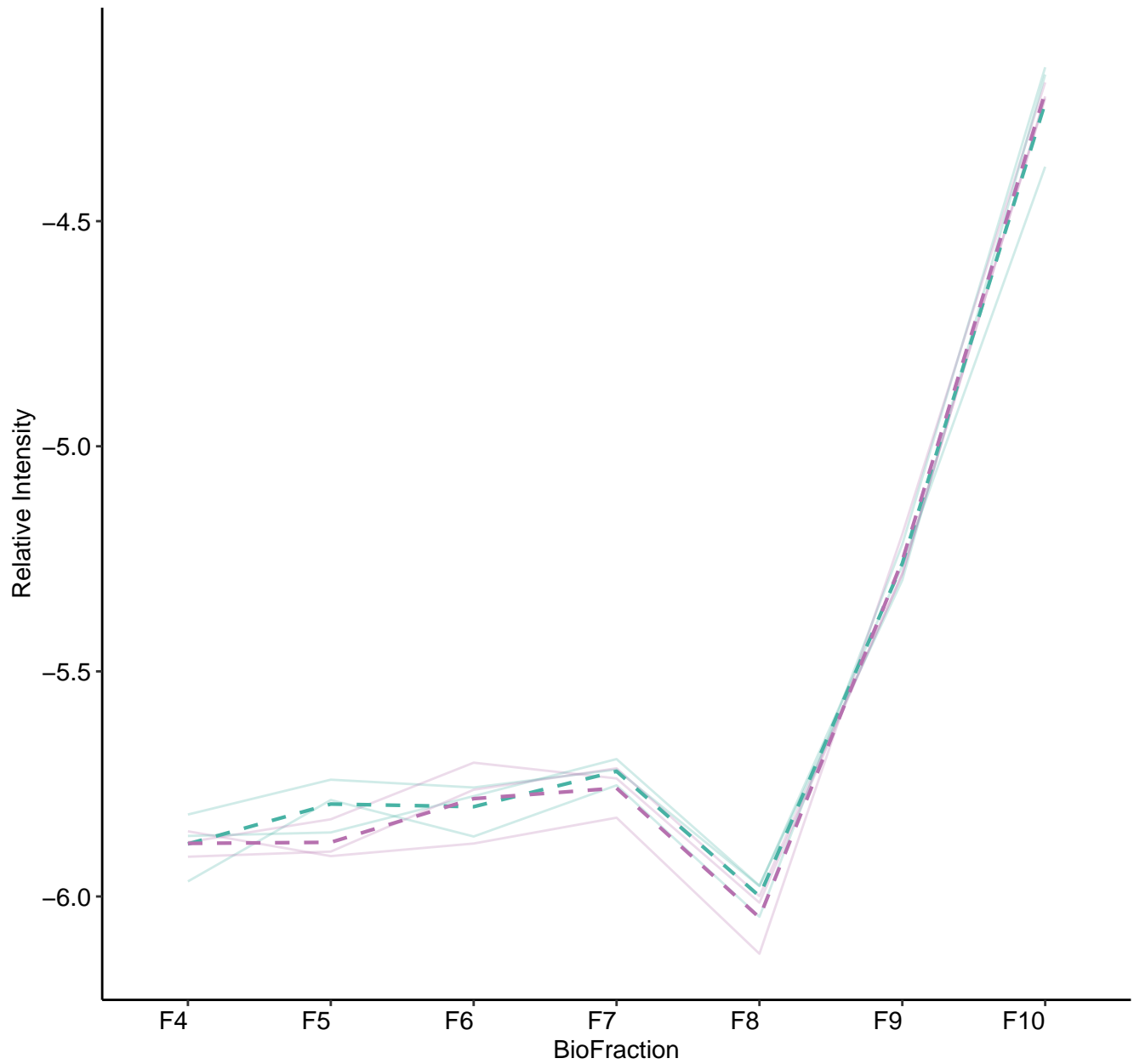
Cytochrome c oxidase, mitochondrial (n = 4)
(R2.Fixef = 0.872)



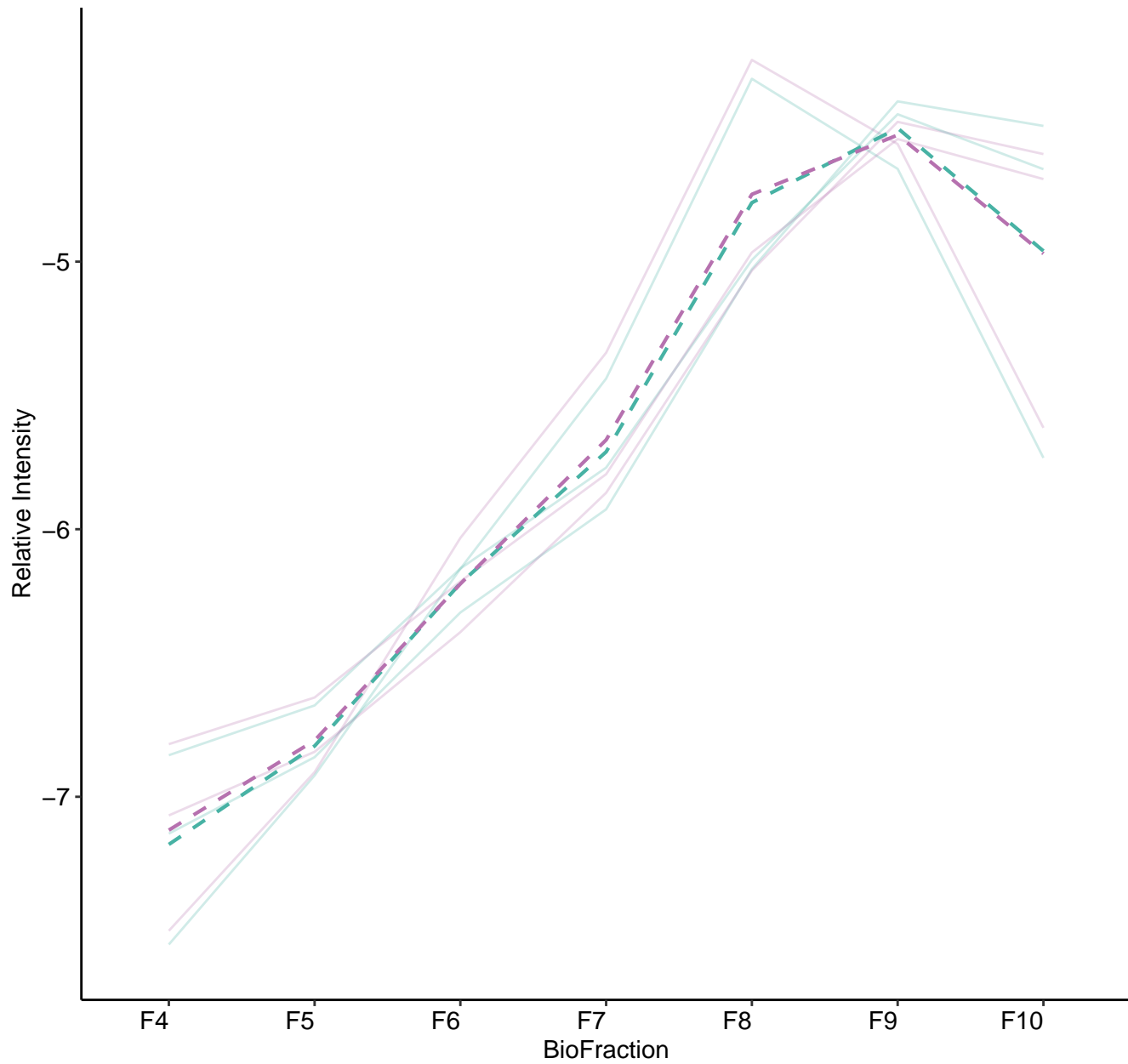
Dystrobrevin-syntrophin complex, brain-derived (n = 3)
(R2.Fixef = 0.877)



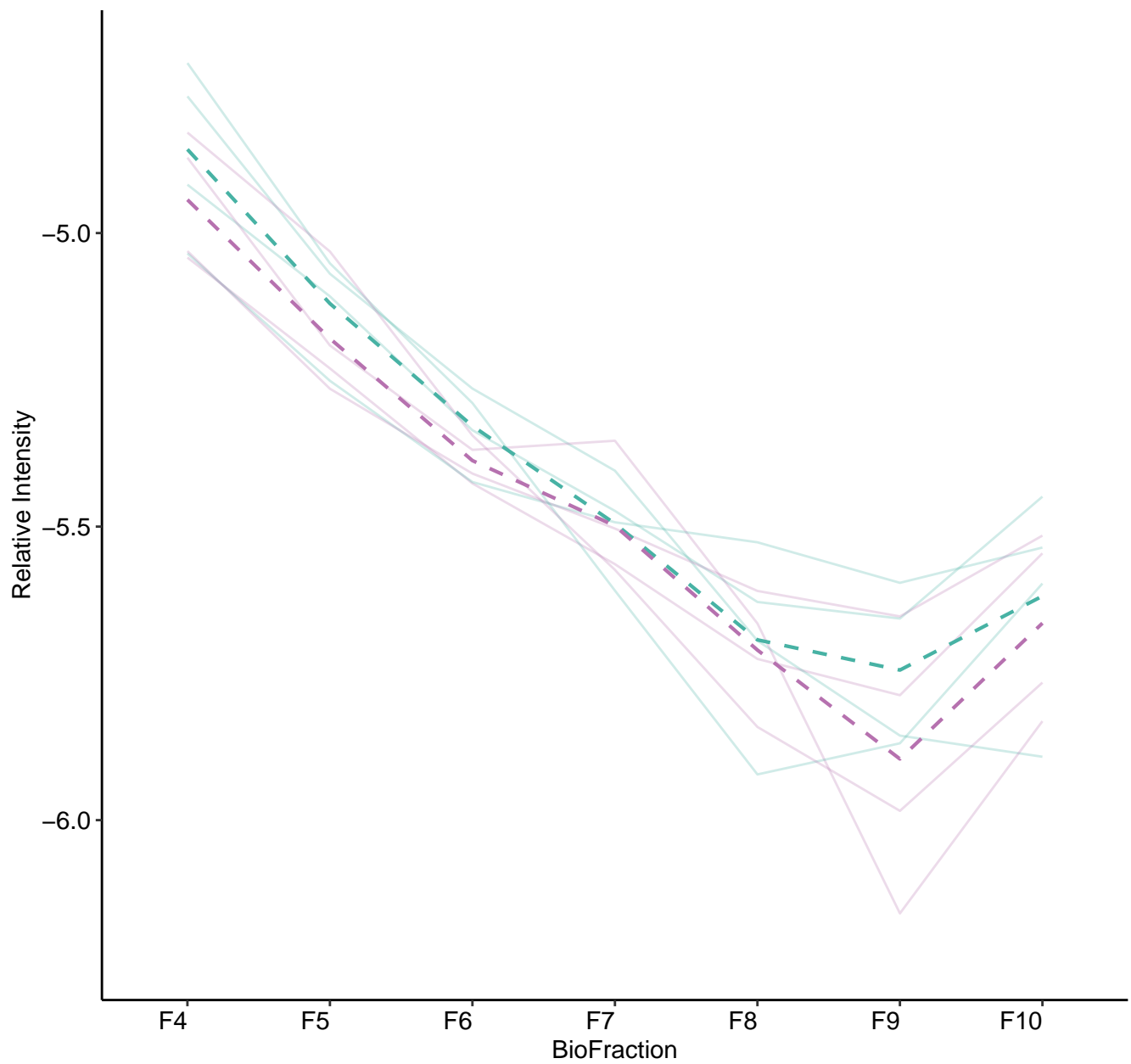
EIF2B1–EIF2B2–EIF2B3–EIF2B4–EIF2B5 complex (n = 3)
(R2.Fixef = 0.989)



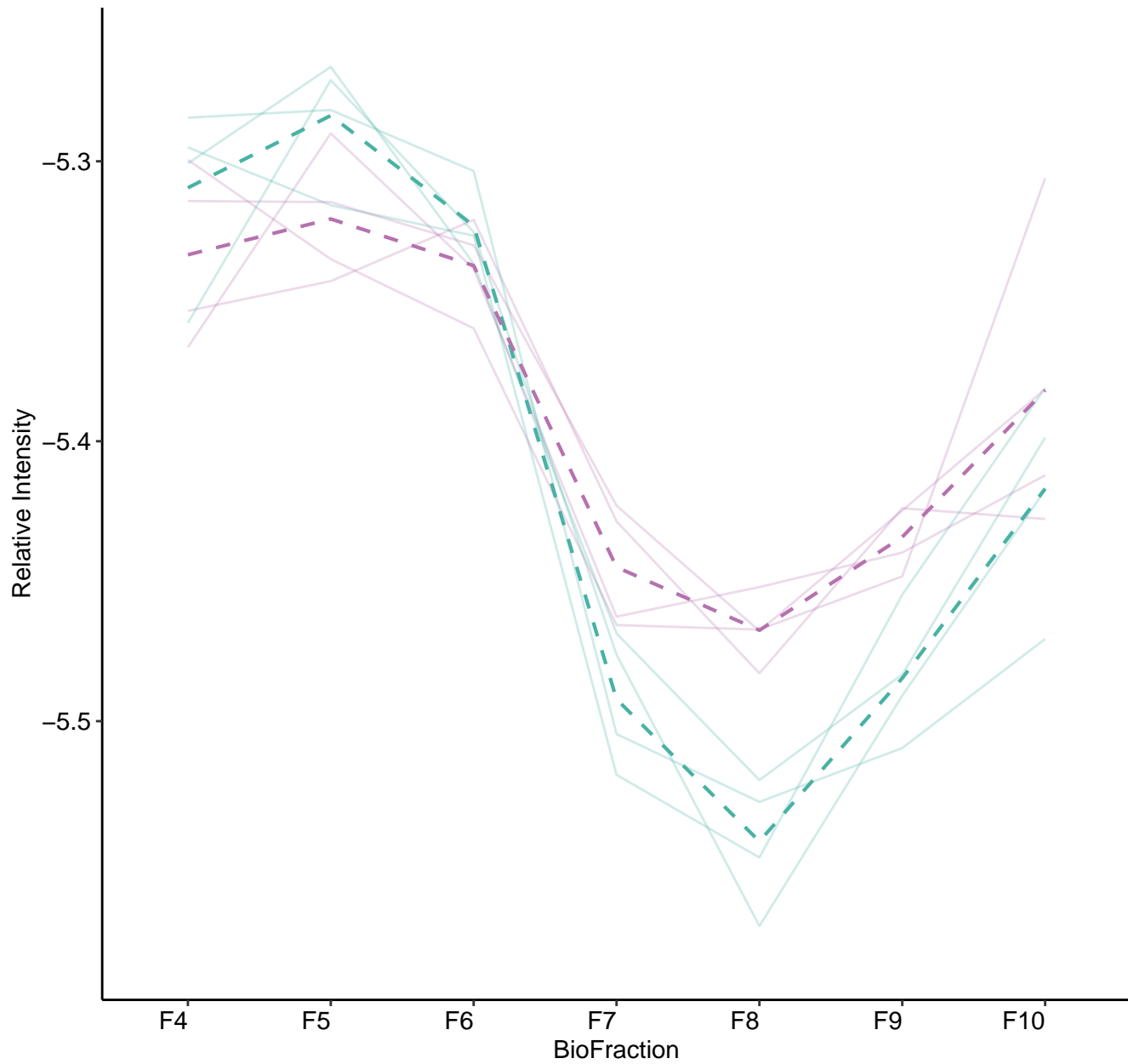
Emerin complex 52 (n = 3)
(R2.Fixef = 0.89)



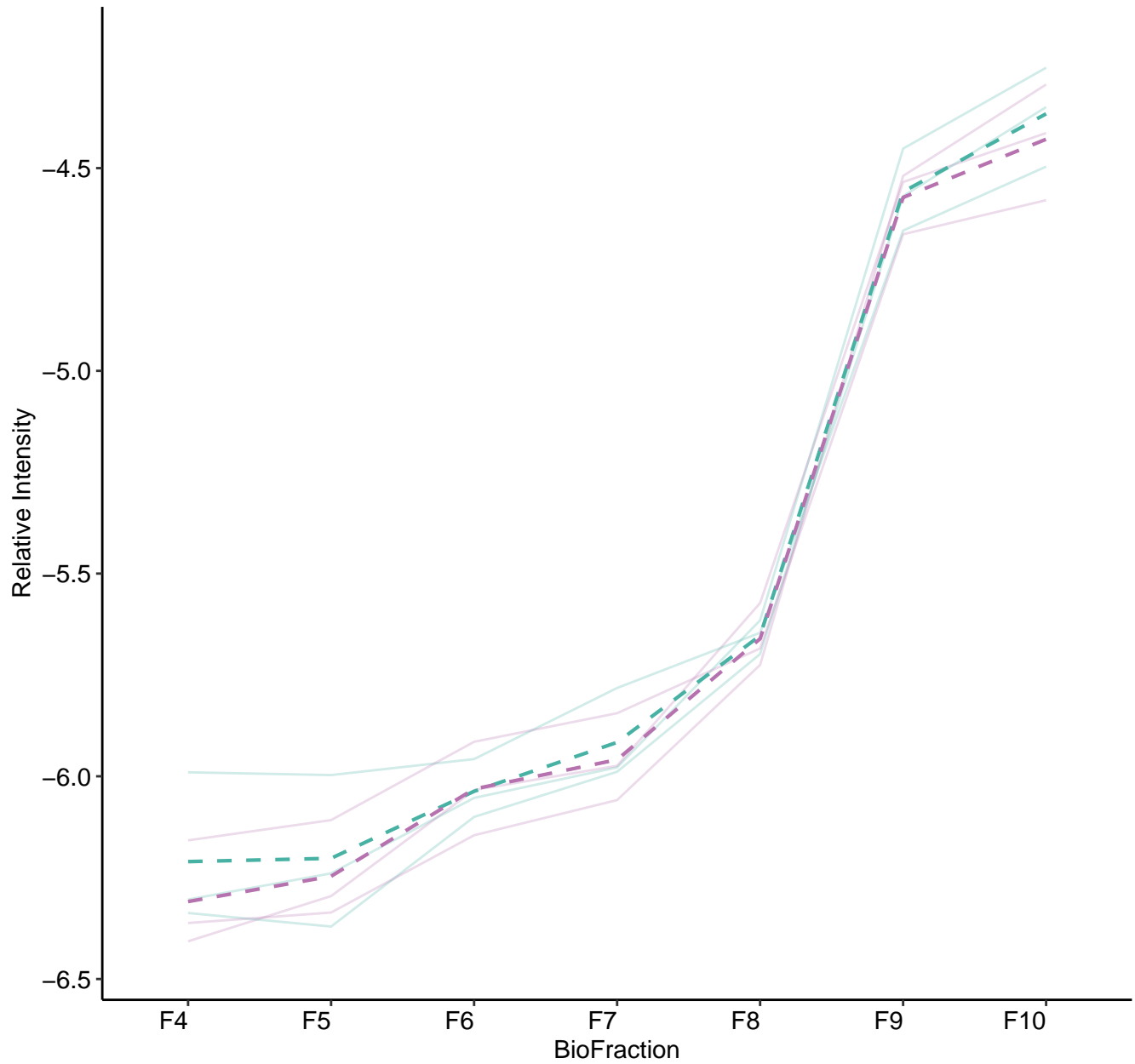
Endocytic coat complex (11 subunits) (n = 4)
(R2.Fixef = 0.844)



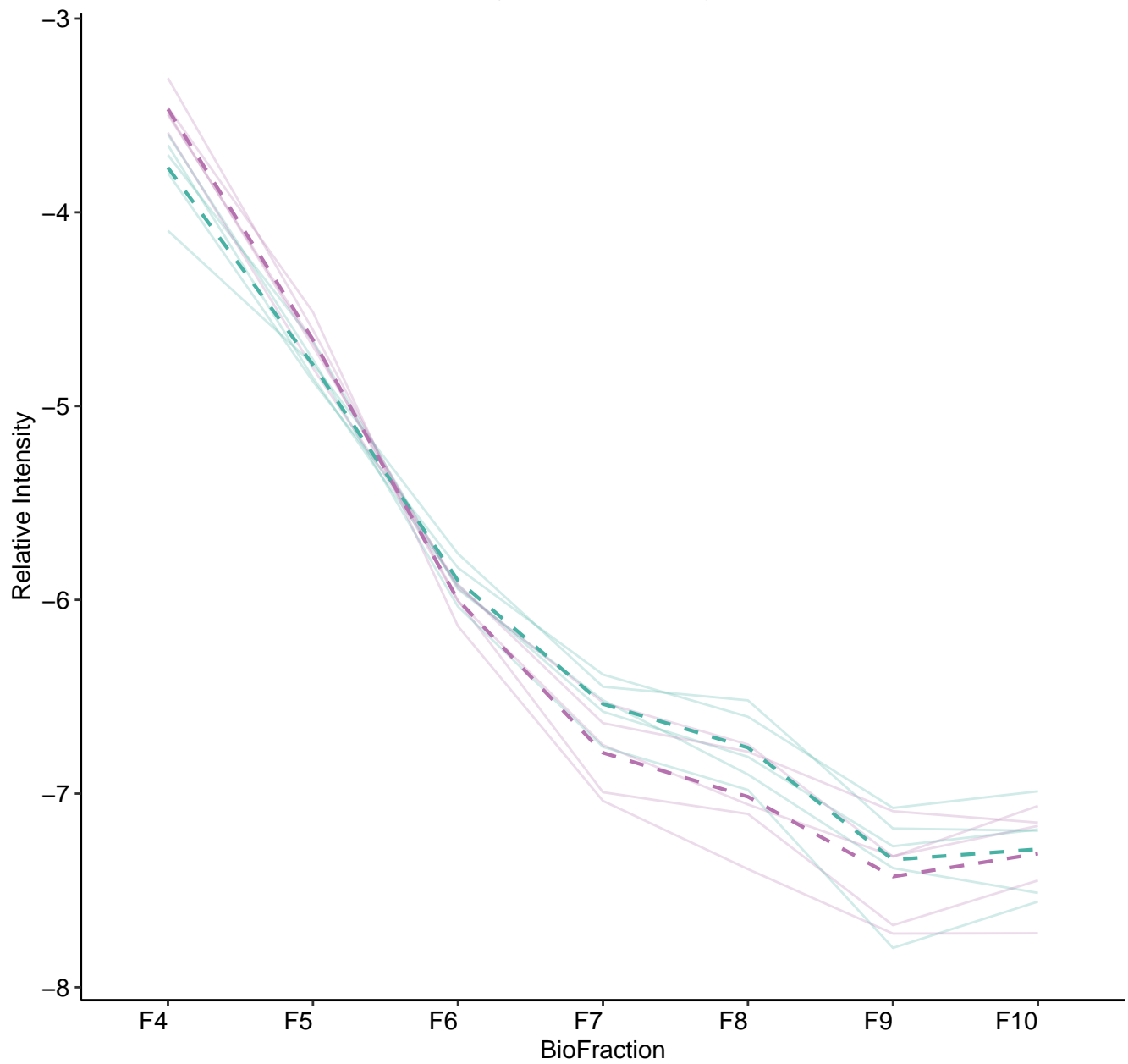
Exocyst complex (n = 4)
(R2.Fixef = 0.894)



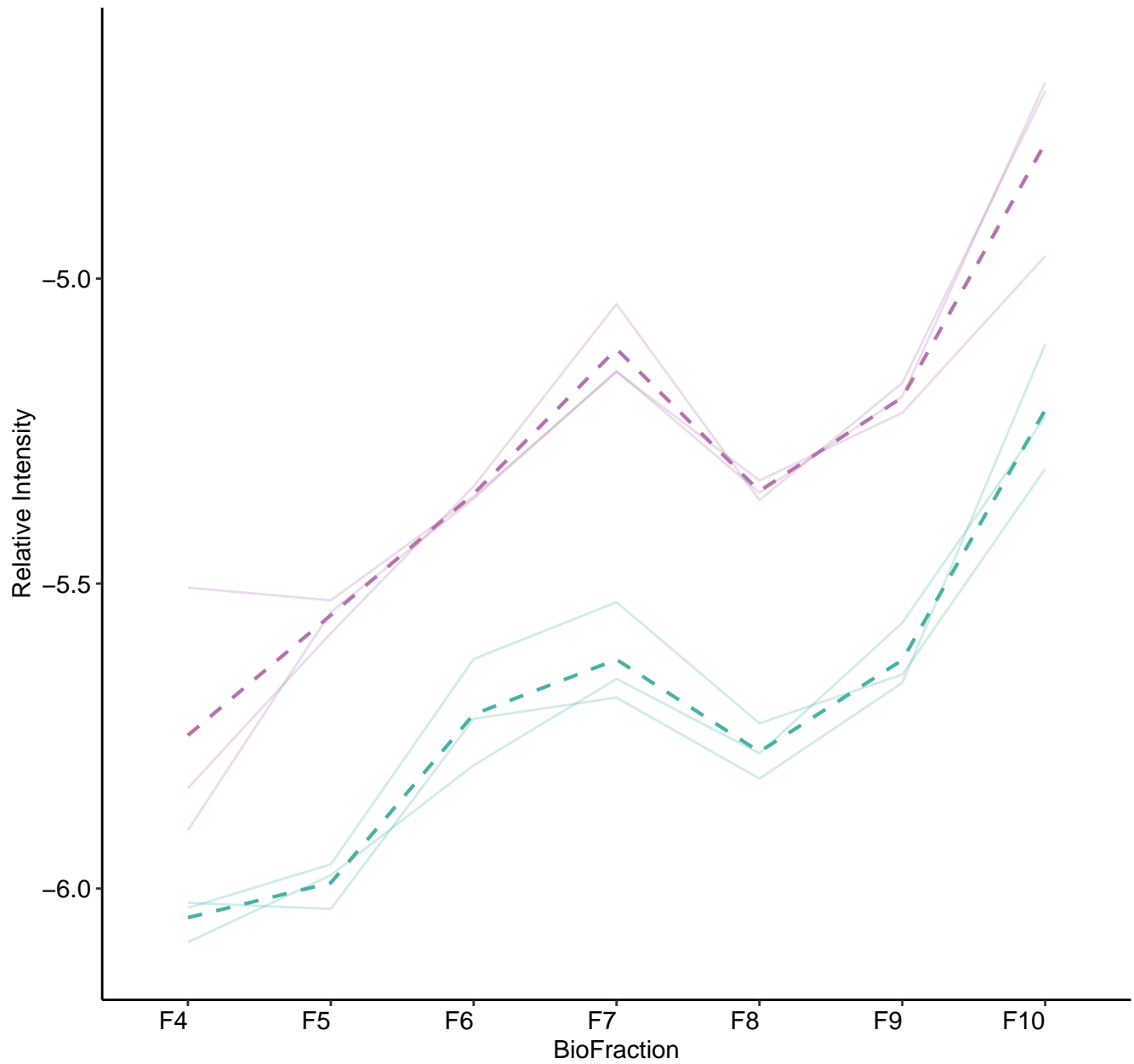
Exosome (n = 3)
(R2.Fixef = 0.973)



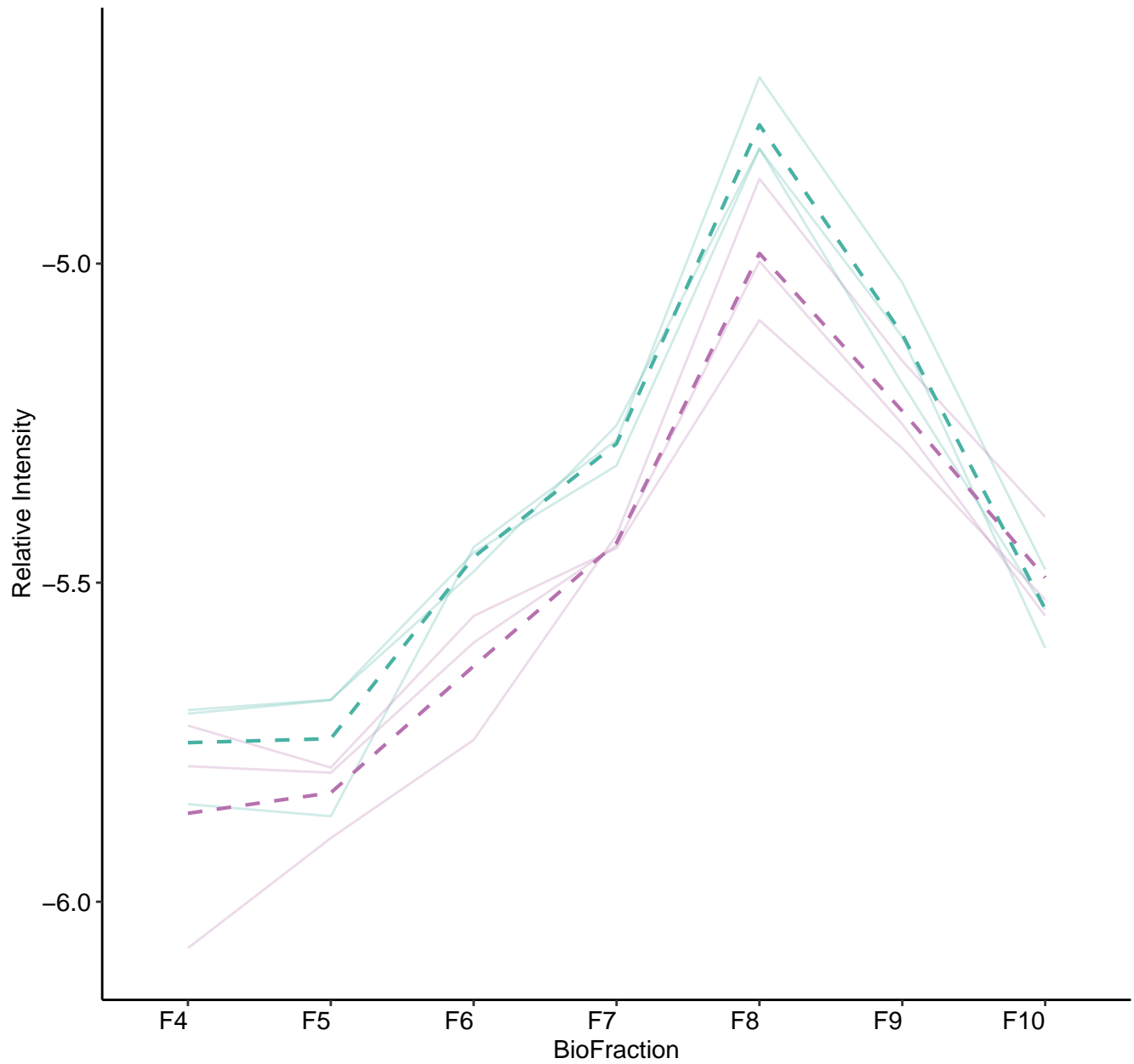
F0F1 ATP synthase, mitochondrial (n = 5)
(R2.Fixef = 0.978)



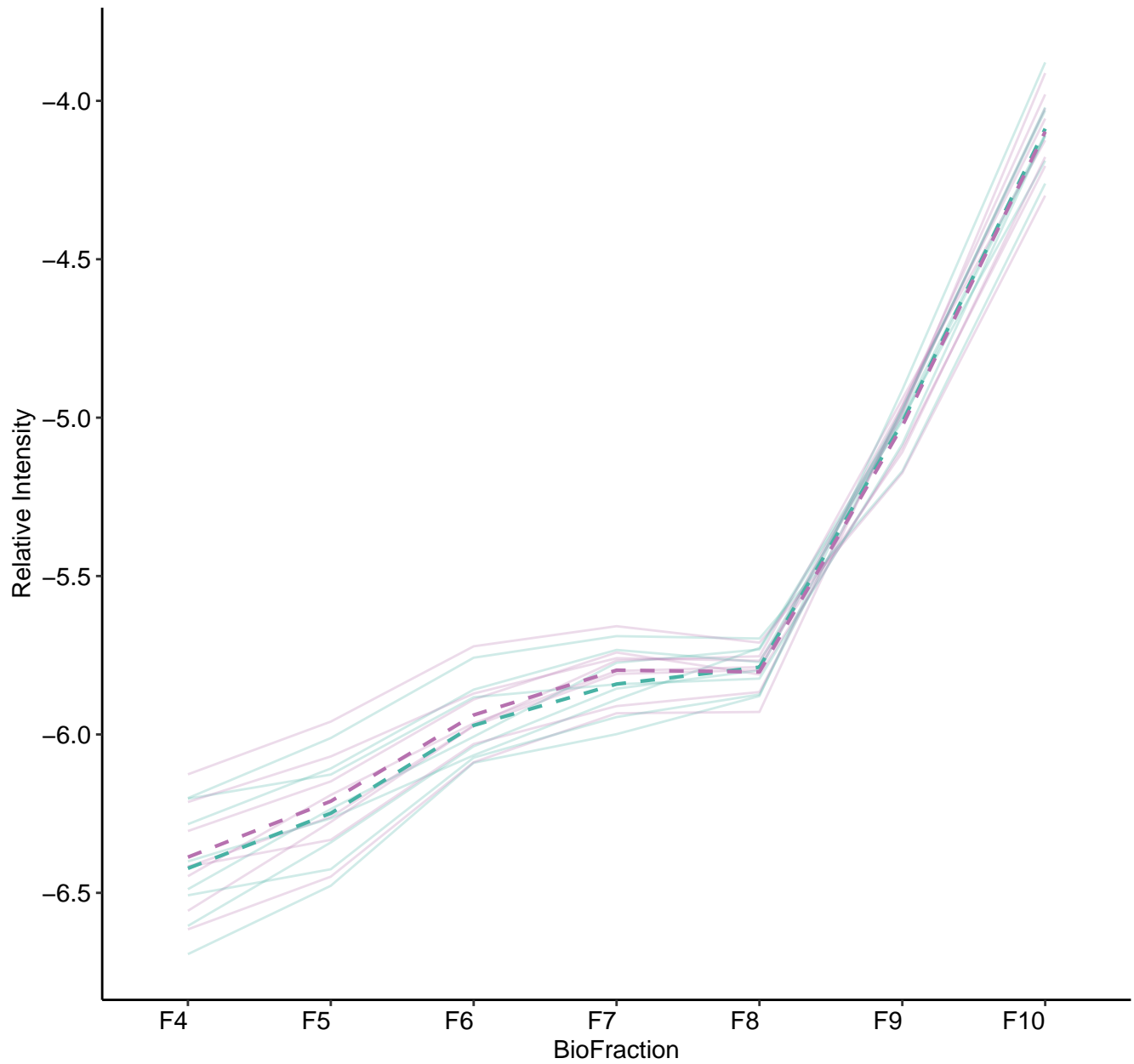
Fibrinogen complex (n = 3)
(R2.Fixef = 0.939)



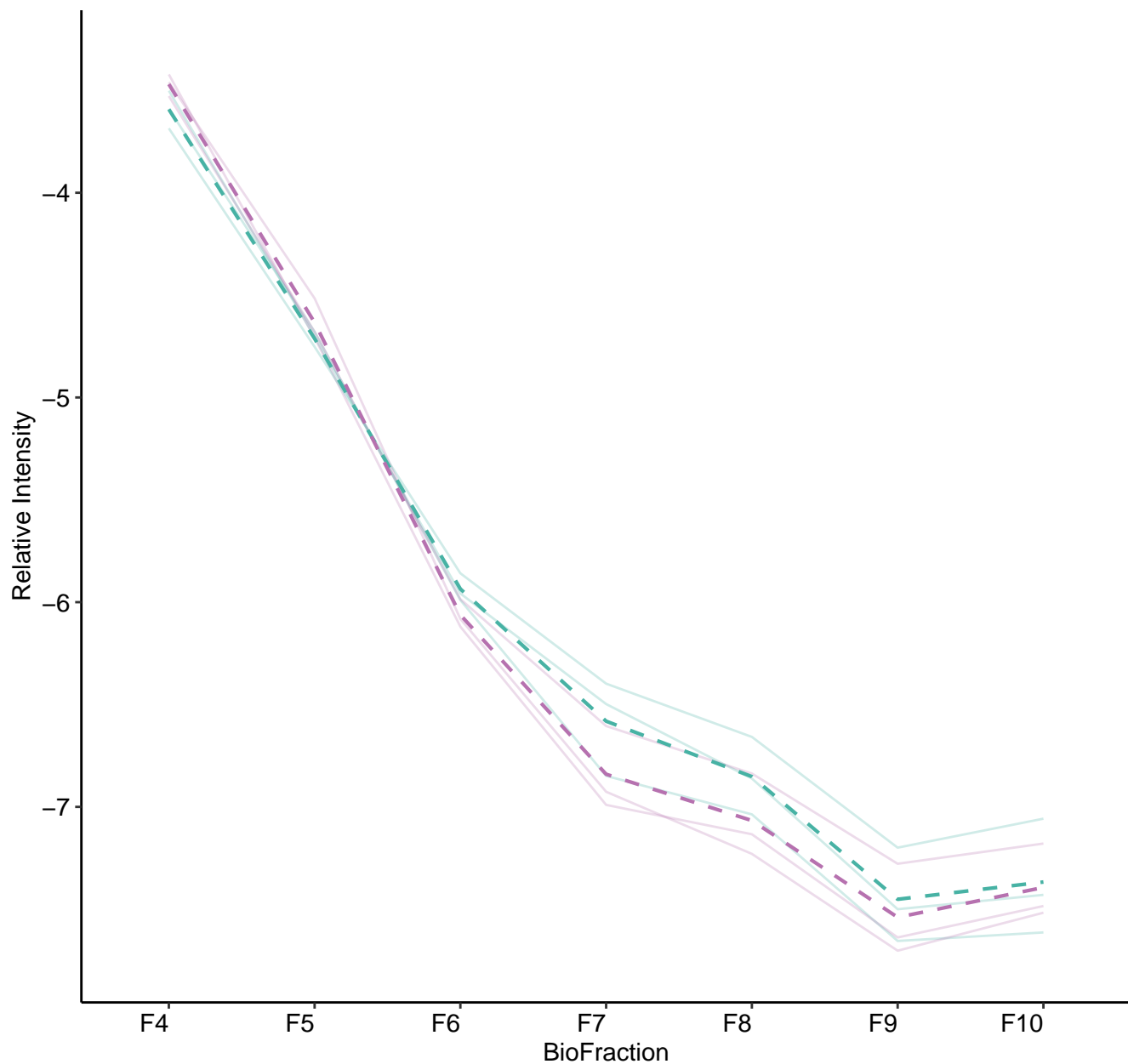
Gamma-BAR-AP1 complex (n = 3)
(R2.Fixef = 0.93)



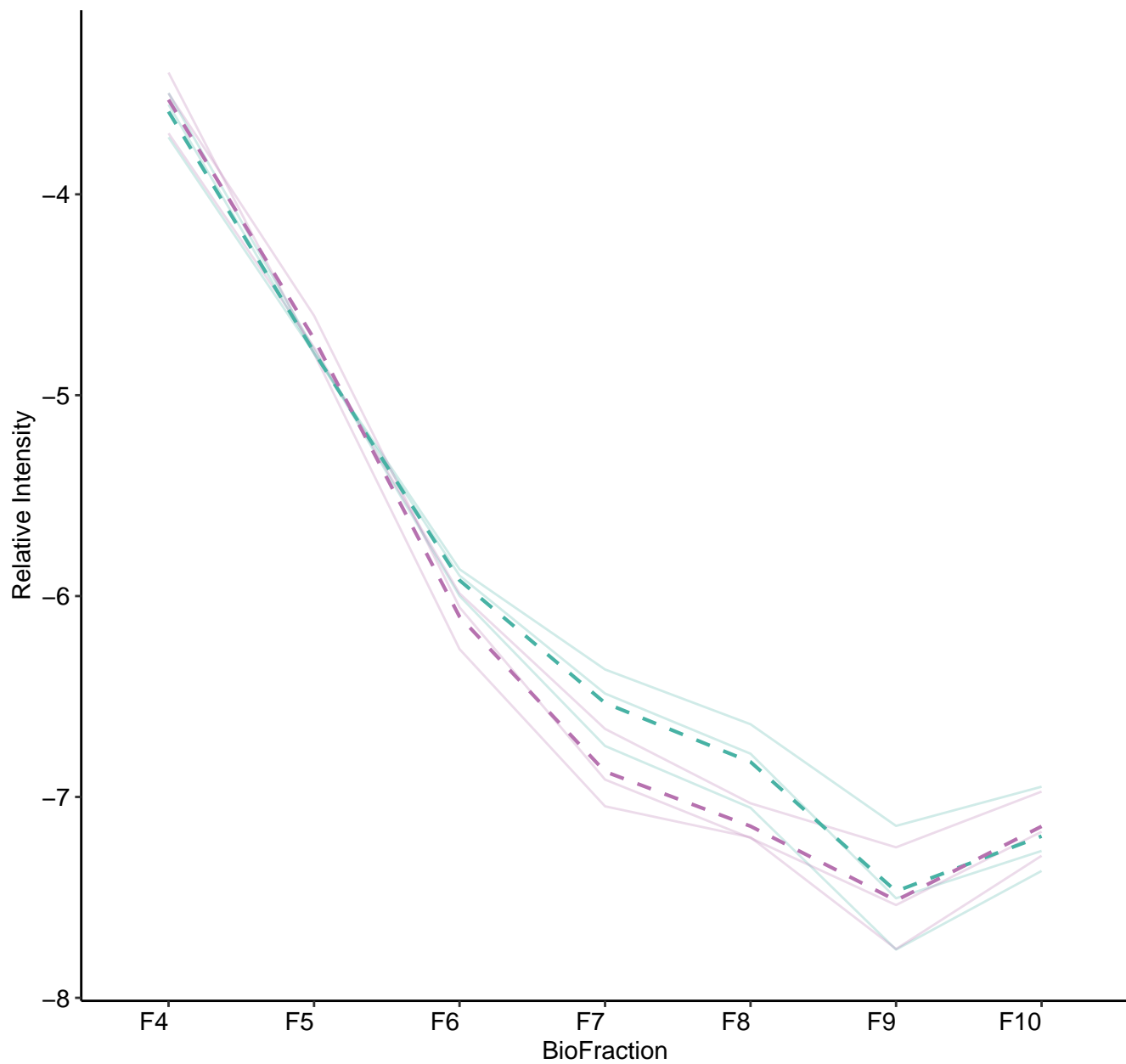
Multisynthetase complex (n = 8)
(R2.Fixef = 0.974)



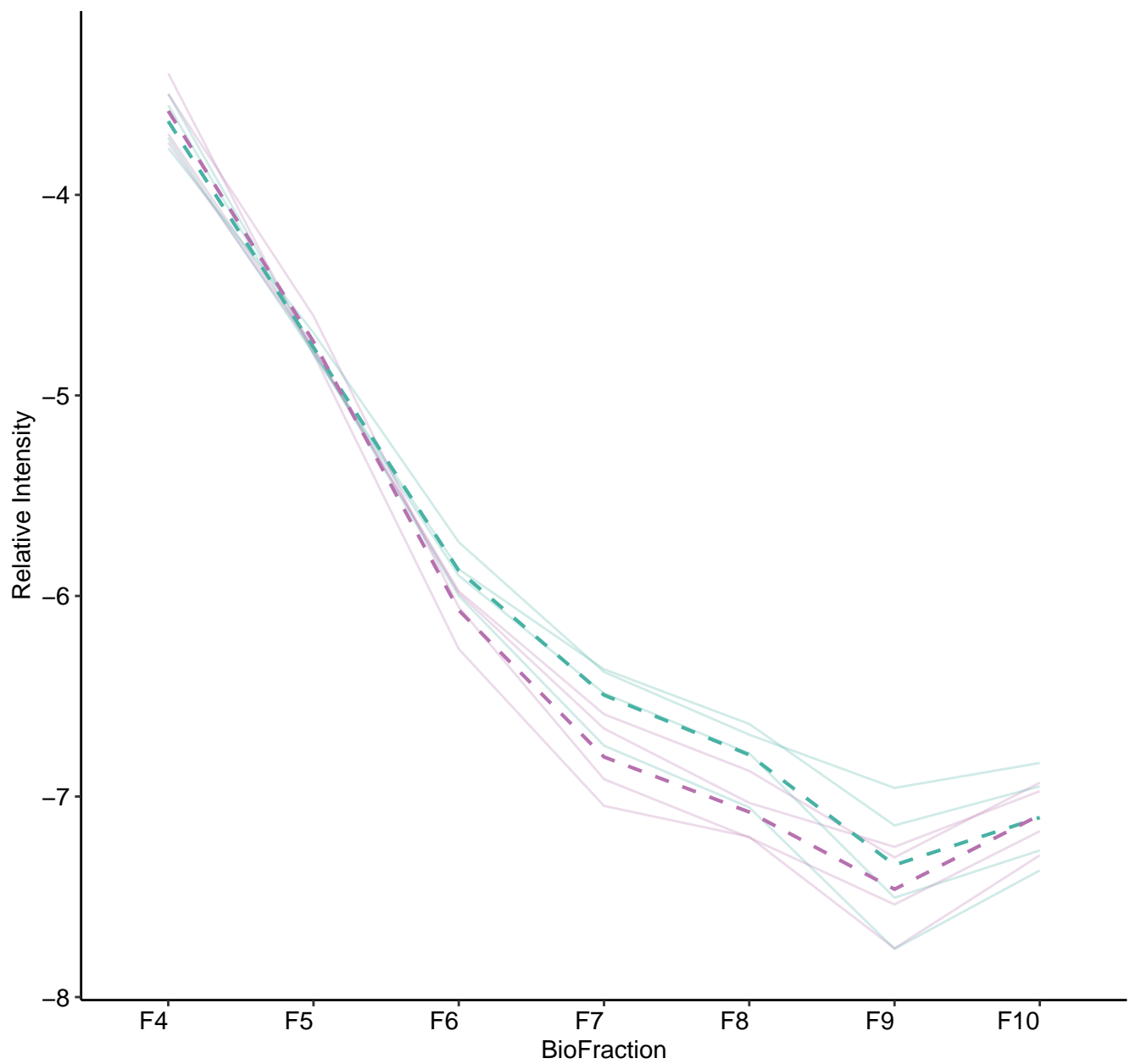
Respiratory chain complex I (early intermediate NDUFAF1 assembly), mitochondrial (n = 3)
(R2.Fixef = 0.985)



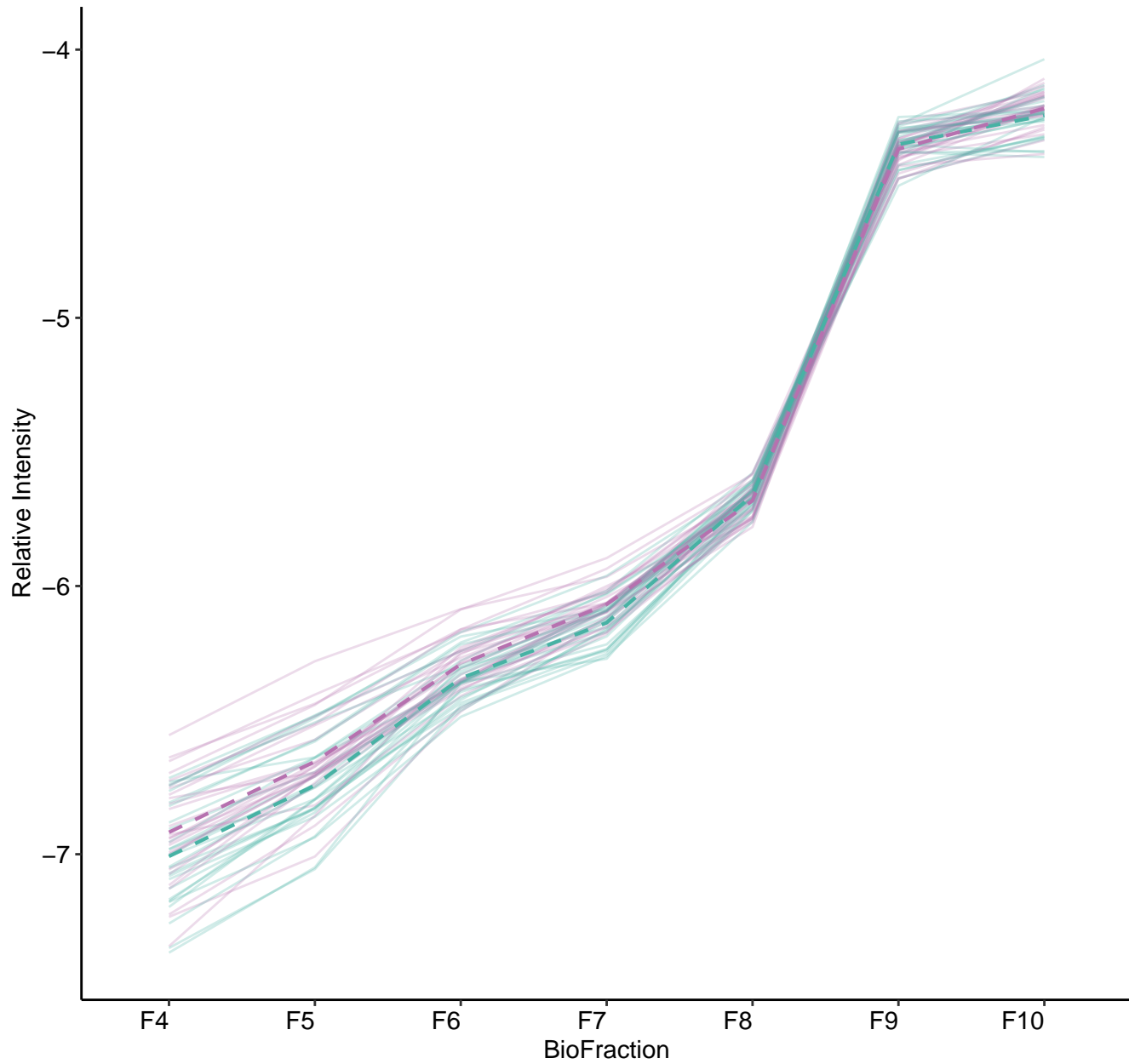
Respiratory chain complex I (intermediate V/380kD and VI/480kD), mitochondrial (n = 3)
(R2.Fixef = 0.983)



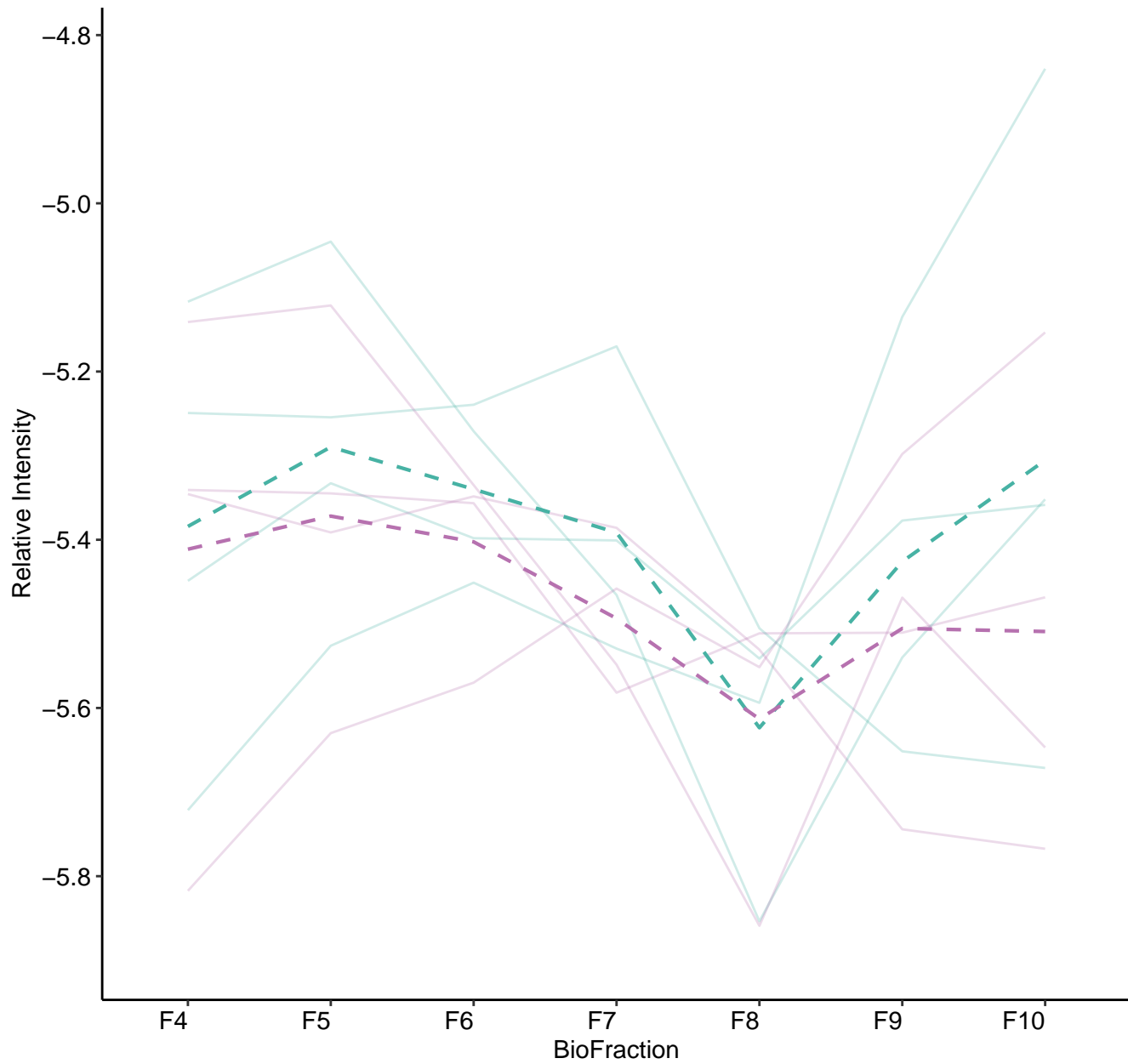
Respiratory chain complex I (intermediate VII/650kD), mitochondrial (n = 4)
(R2.Fixef = 0.98)



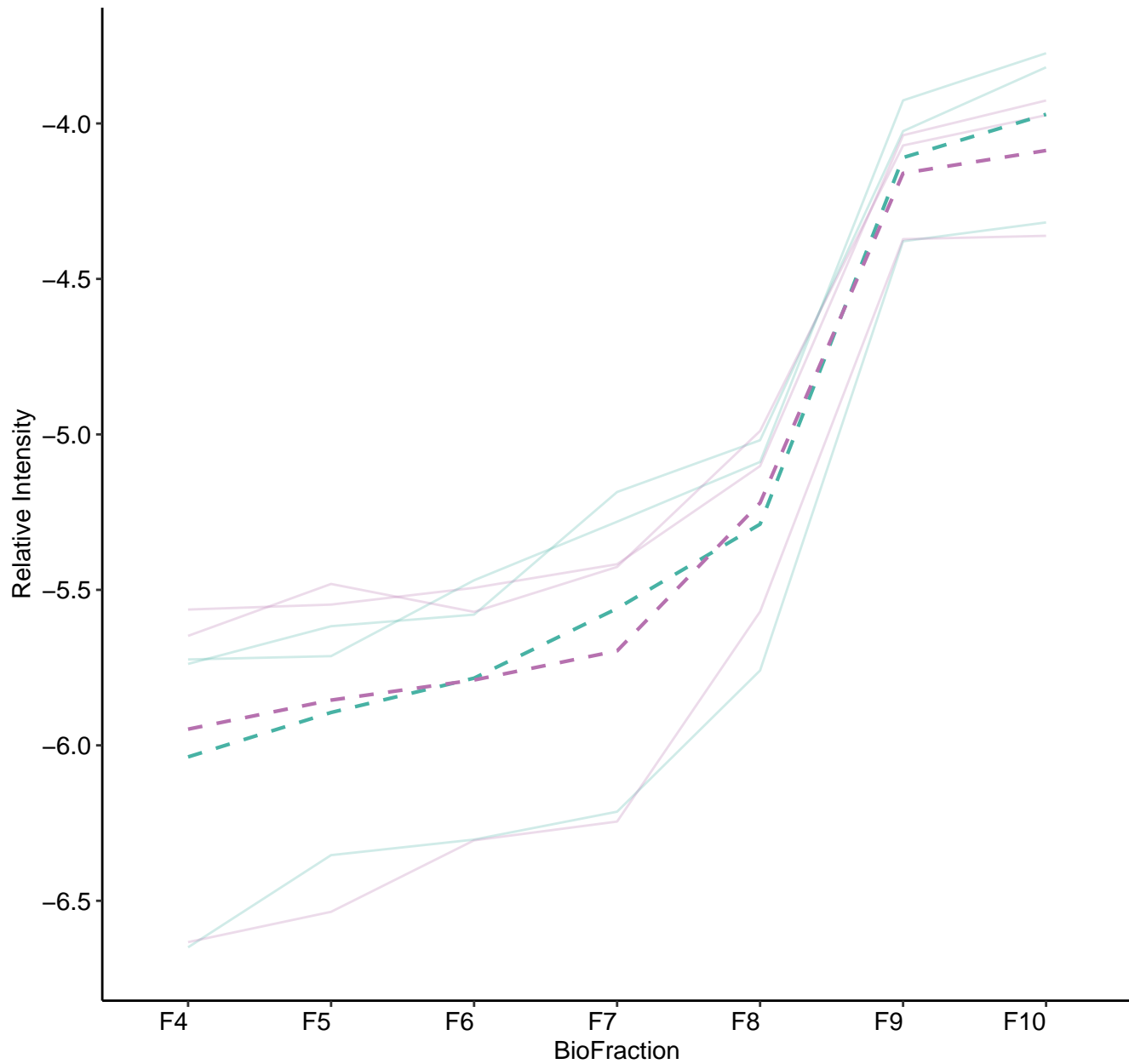
Ribosome, cytoplasmic (n = 27)
(R2.Fixef = 0.988)



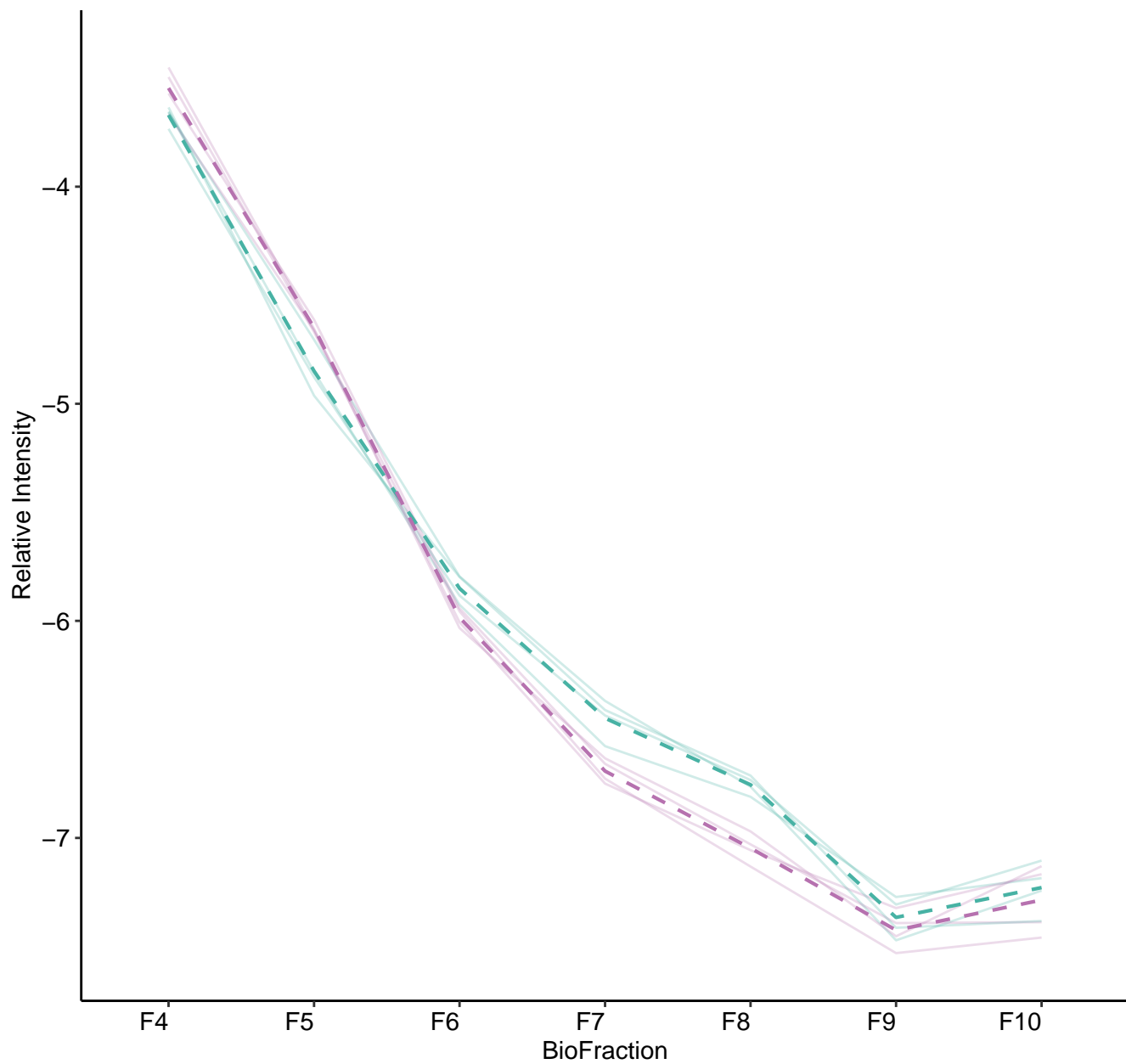
RICH1/AMOT polarity complex, Flag-Rich1 precipitated (n = 4)
(R2.Fixef = 0.186)



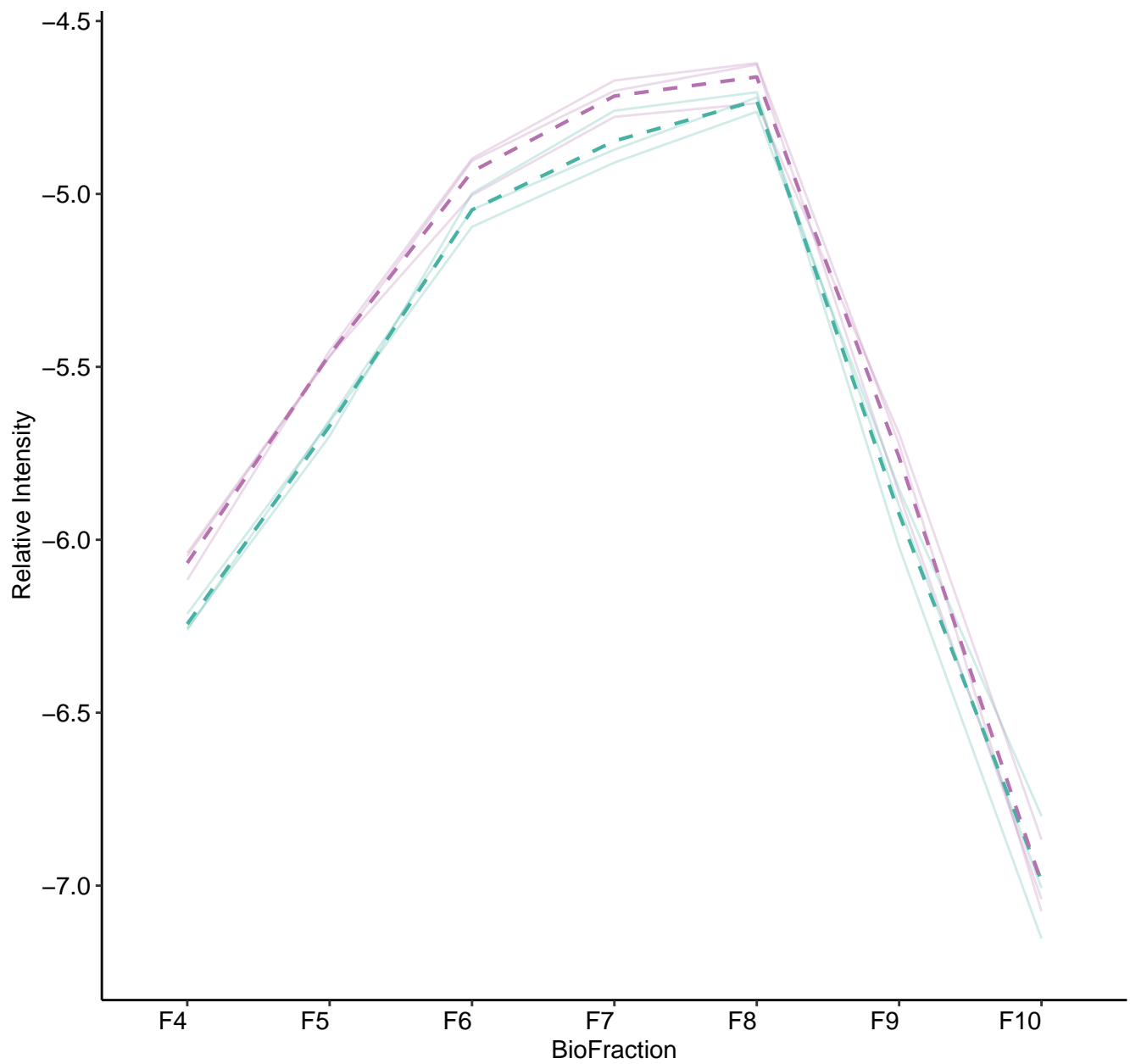
RNA polymerase II complex, chromatin structure modifying (n = 3)
(R2.Fixef = 0.765)



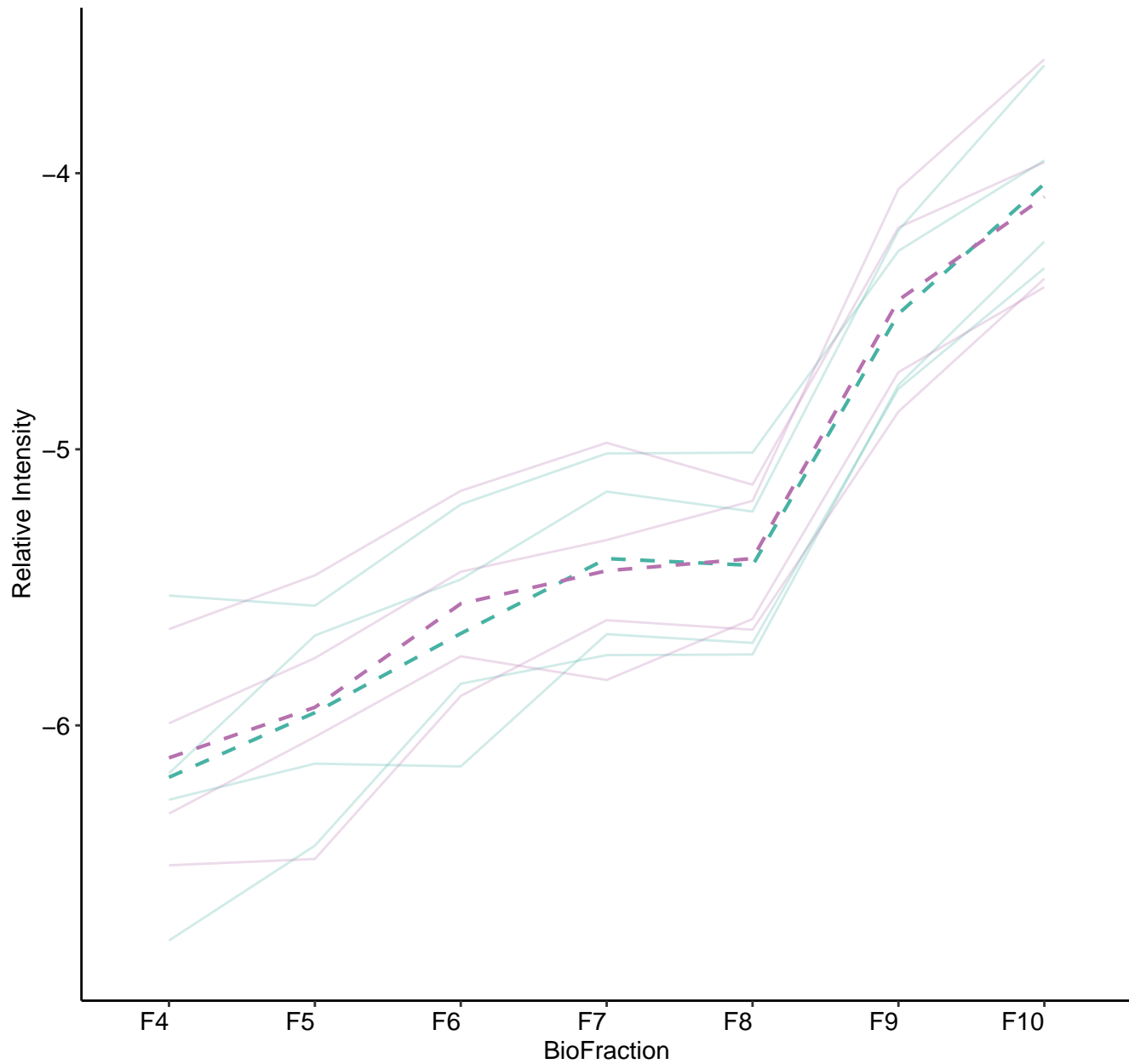
SDH-mABC1-PIC-ANT-ATPase complex (n = 4)
(R2.Fixef = 0.996)



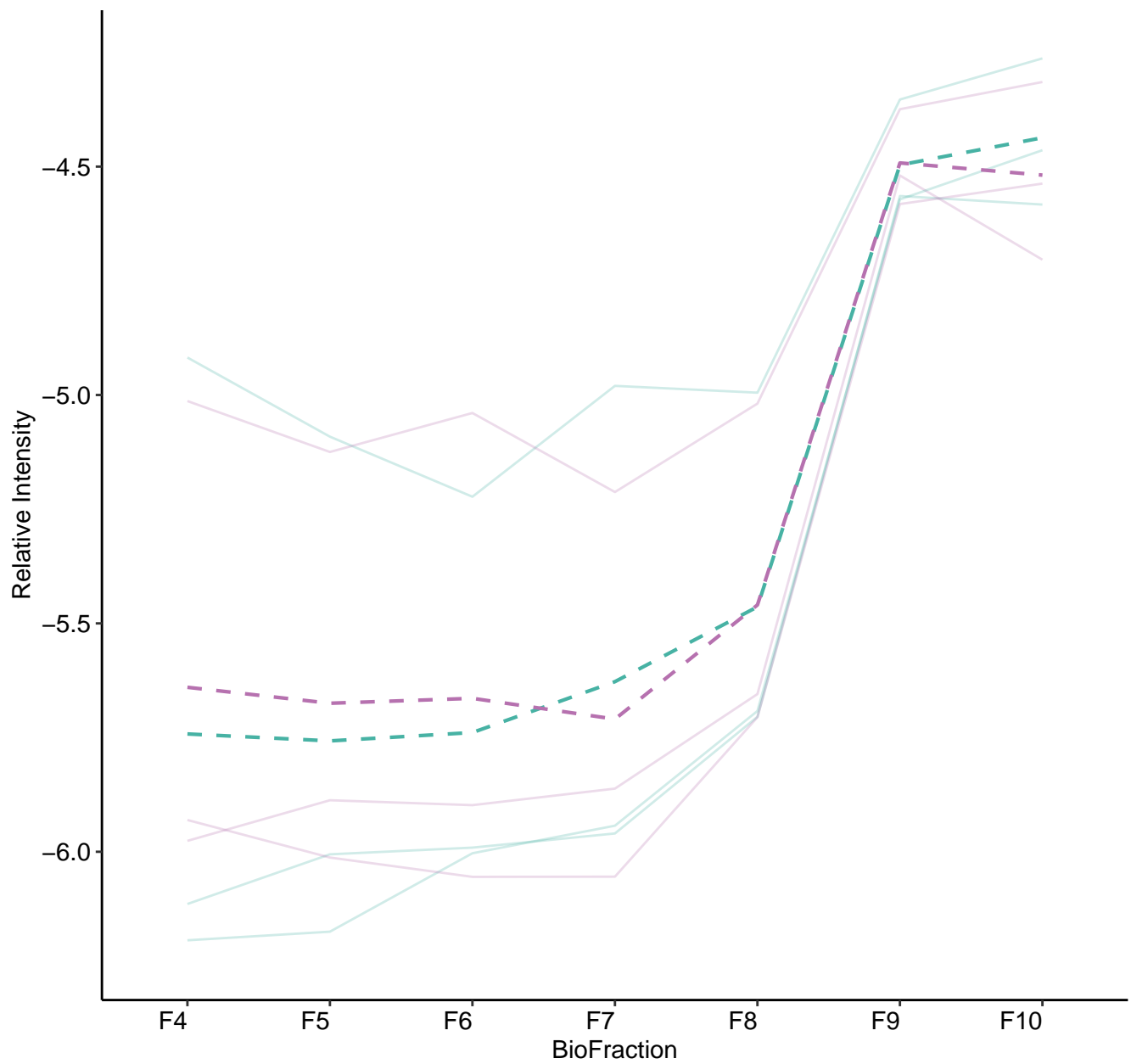
SELK multiprotein complex (n = 3)
(R2.Fixef = 0.99)



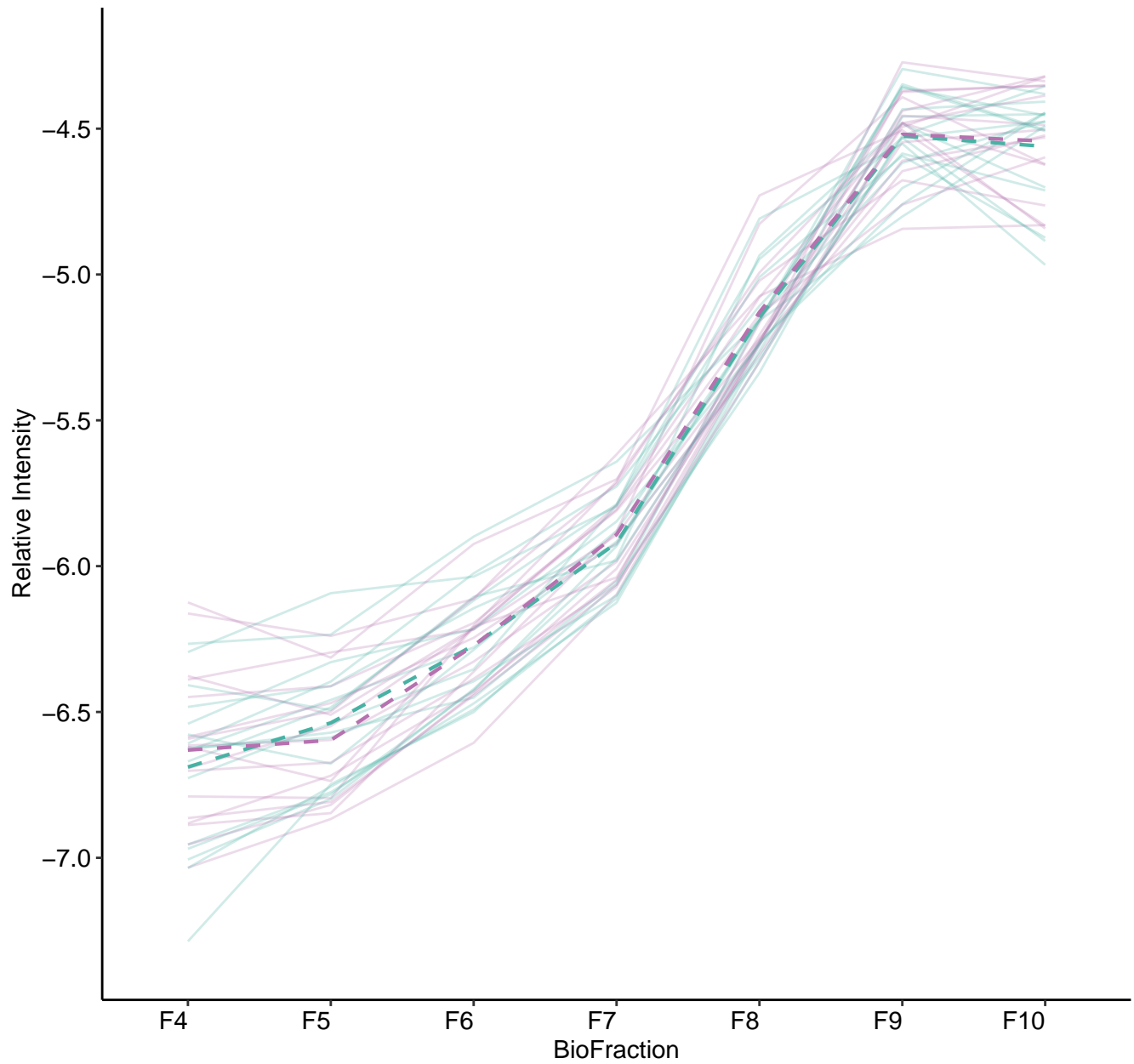
SNF2h-cohesin-NuRD complex (n = 4)
(R2.Fixef = 0.777)



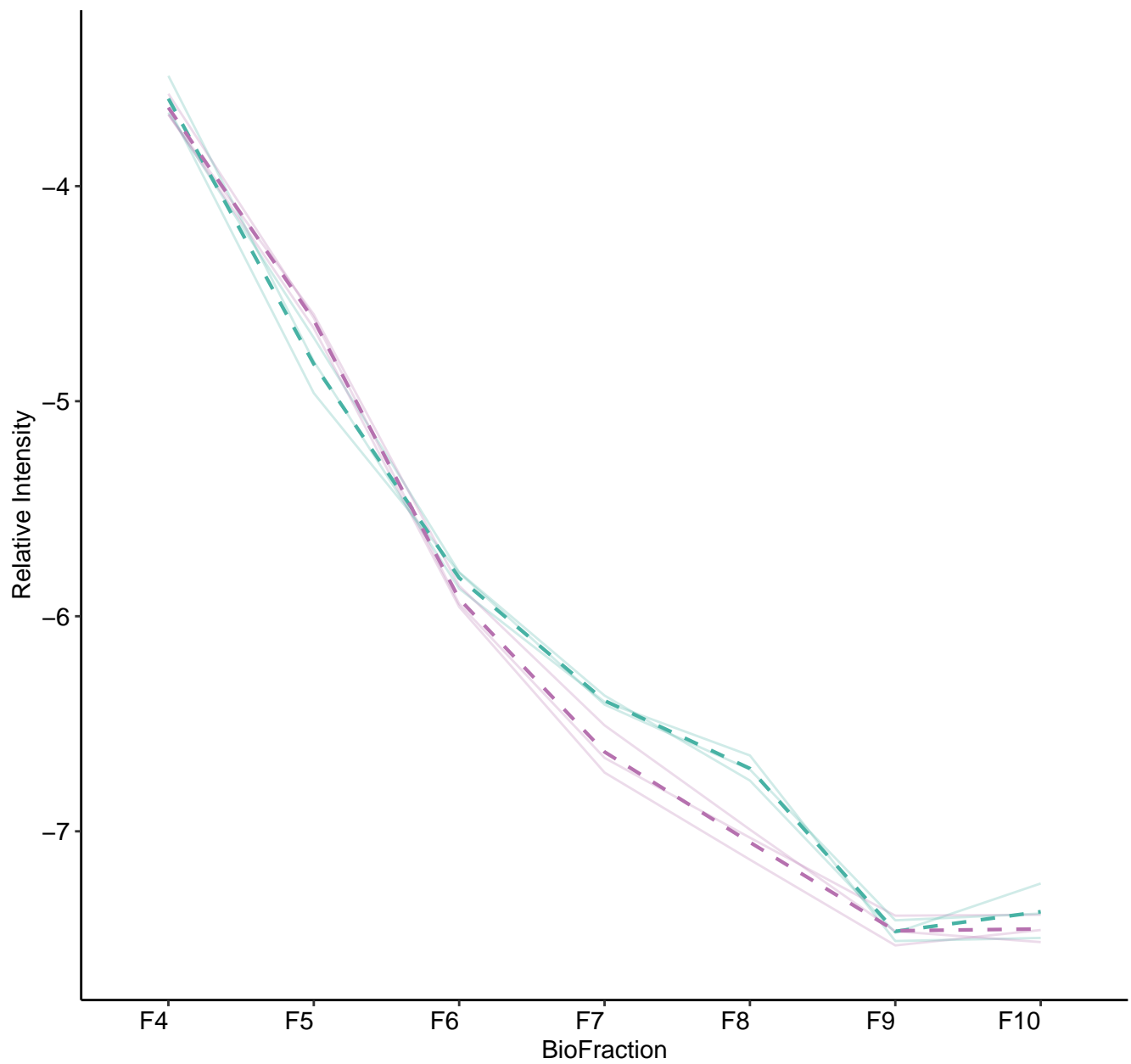
SOG complex (n = 3)
(R2.Fixef = 0.593)



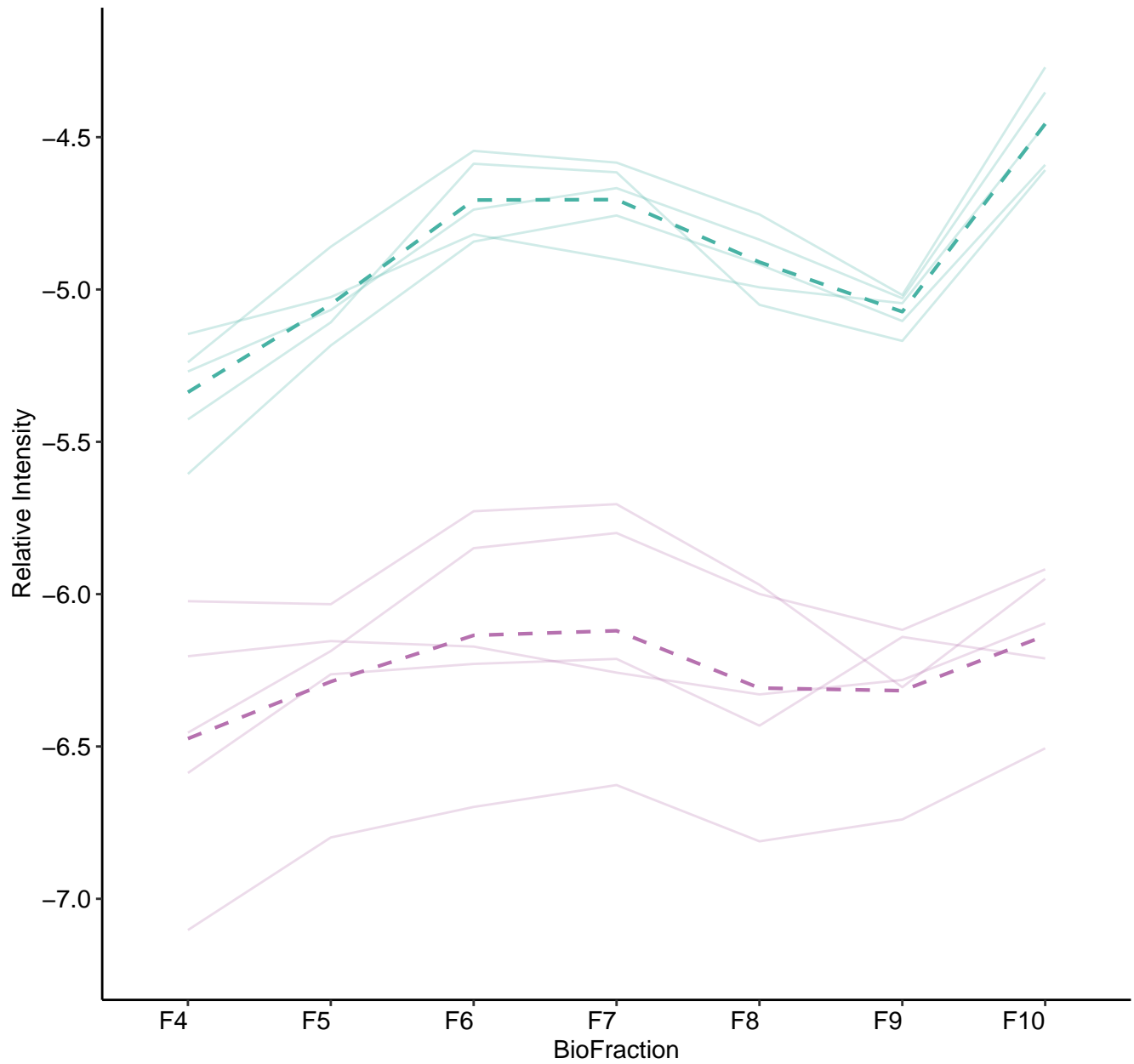
Spliceosome (n = 17)
(R2.Fixef = 0.953)



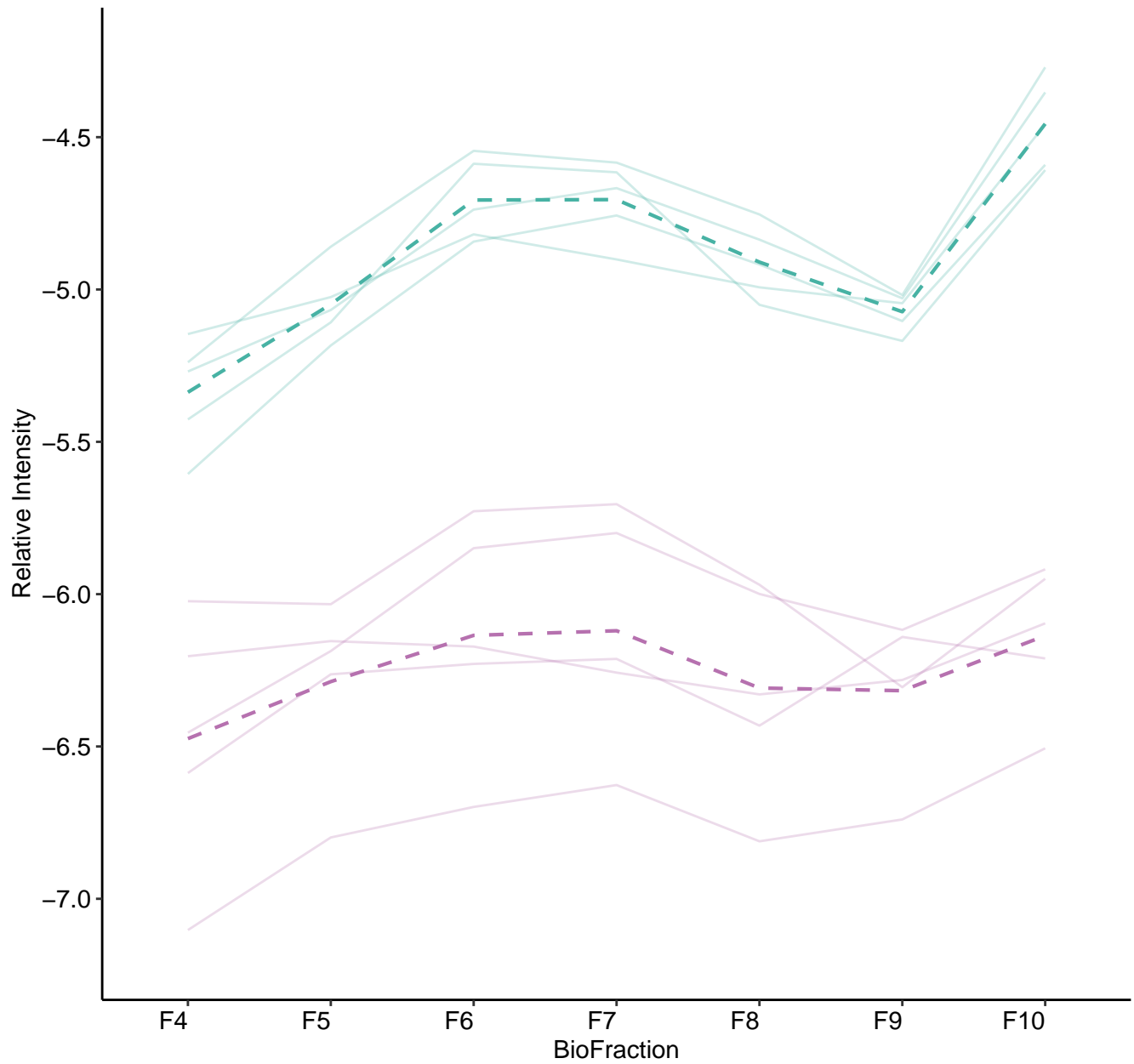
Succinate dehydrogenase complex II, mitochondrial (n = 3)
(R2.Fixef = 0.997)



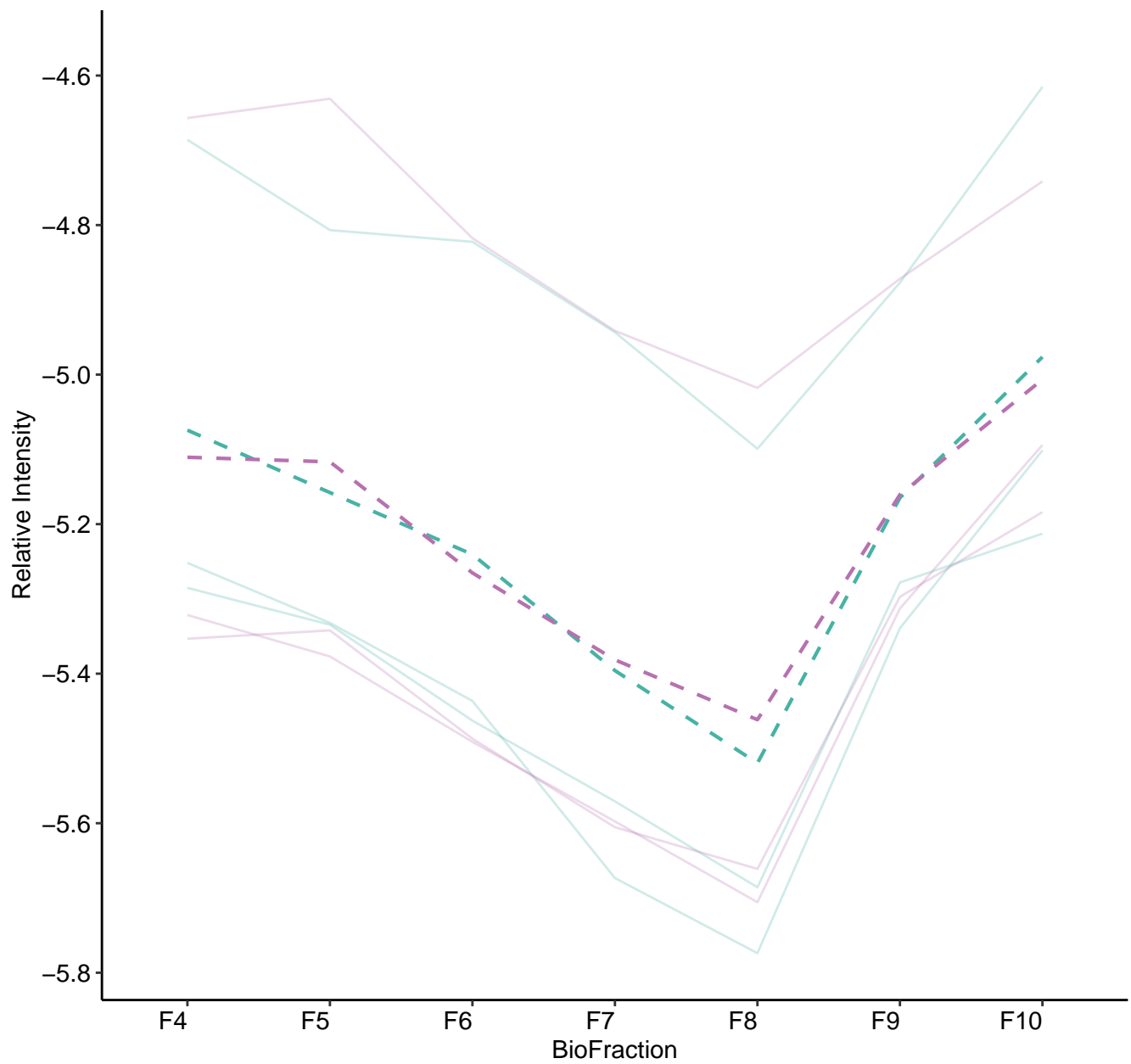
WASH complex (n = 5)
(R2.Fixef = 0.889)



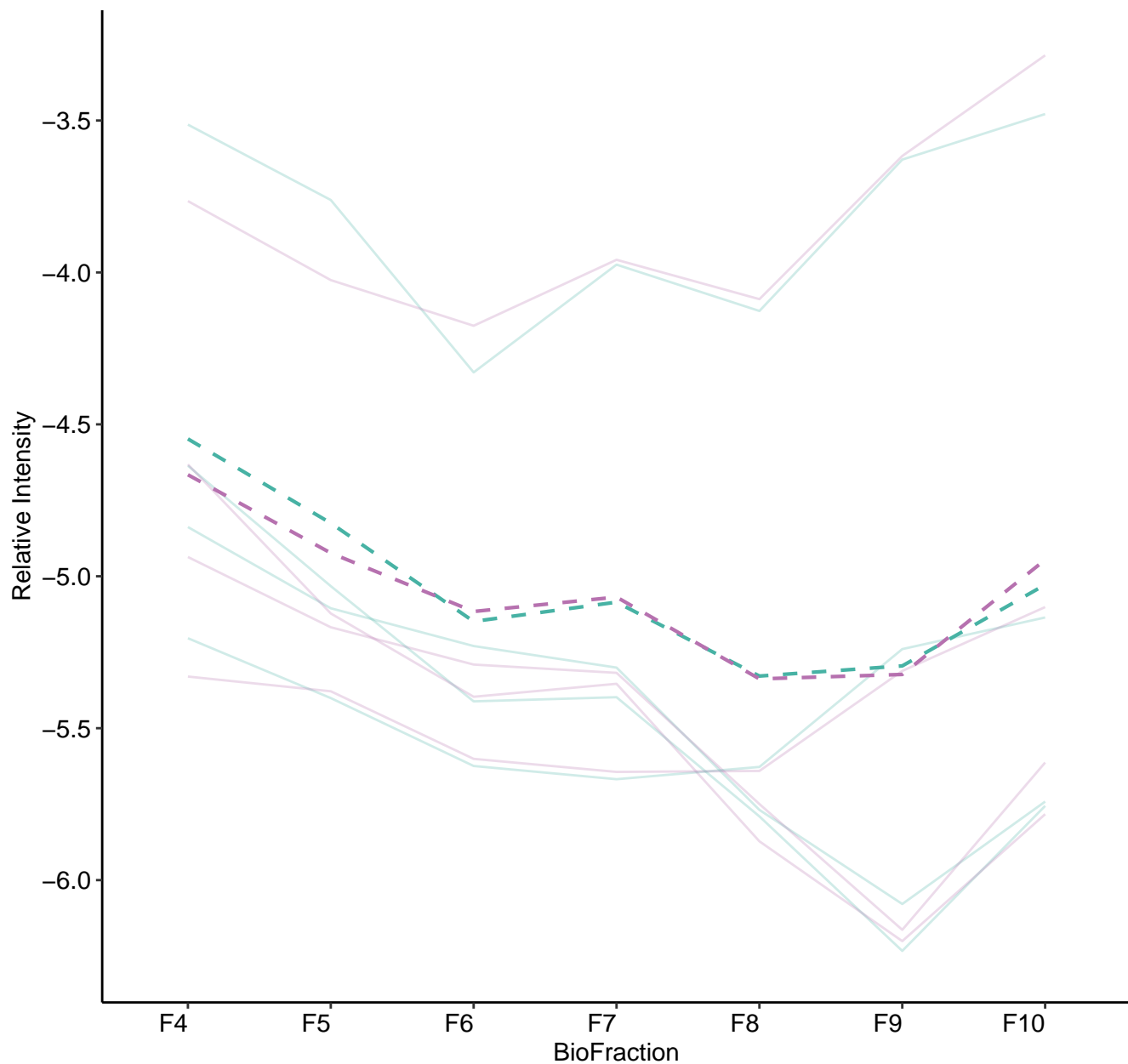
WASH-CAPZalpa/beta complex (n = 5)
(R2.Fixef = 0.889)



Wave-2 complex (n = 3)
(R2.Fixef = 0.182)



WDR41-(C9orf72-SMCR8)-(FIP200-ULK1-ATG13-ATG101) complex (n = 4)
(R2.Fixef = 0.071)



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

