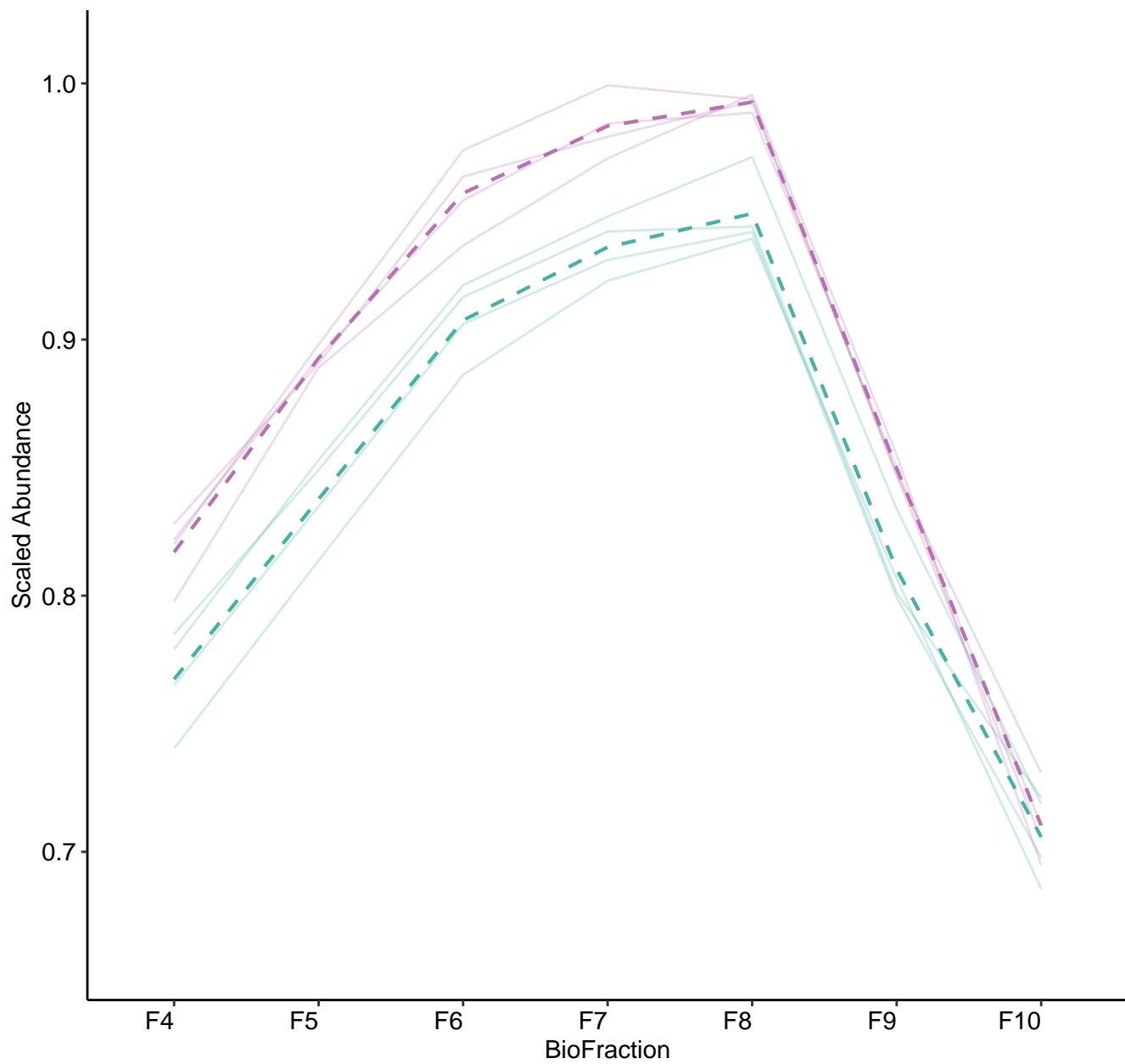
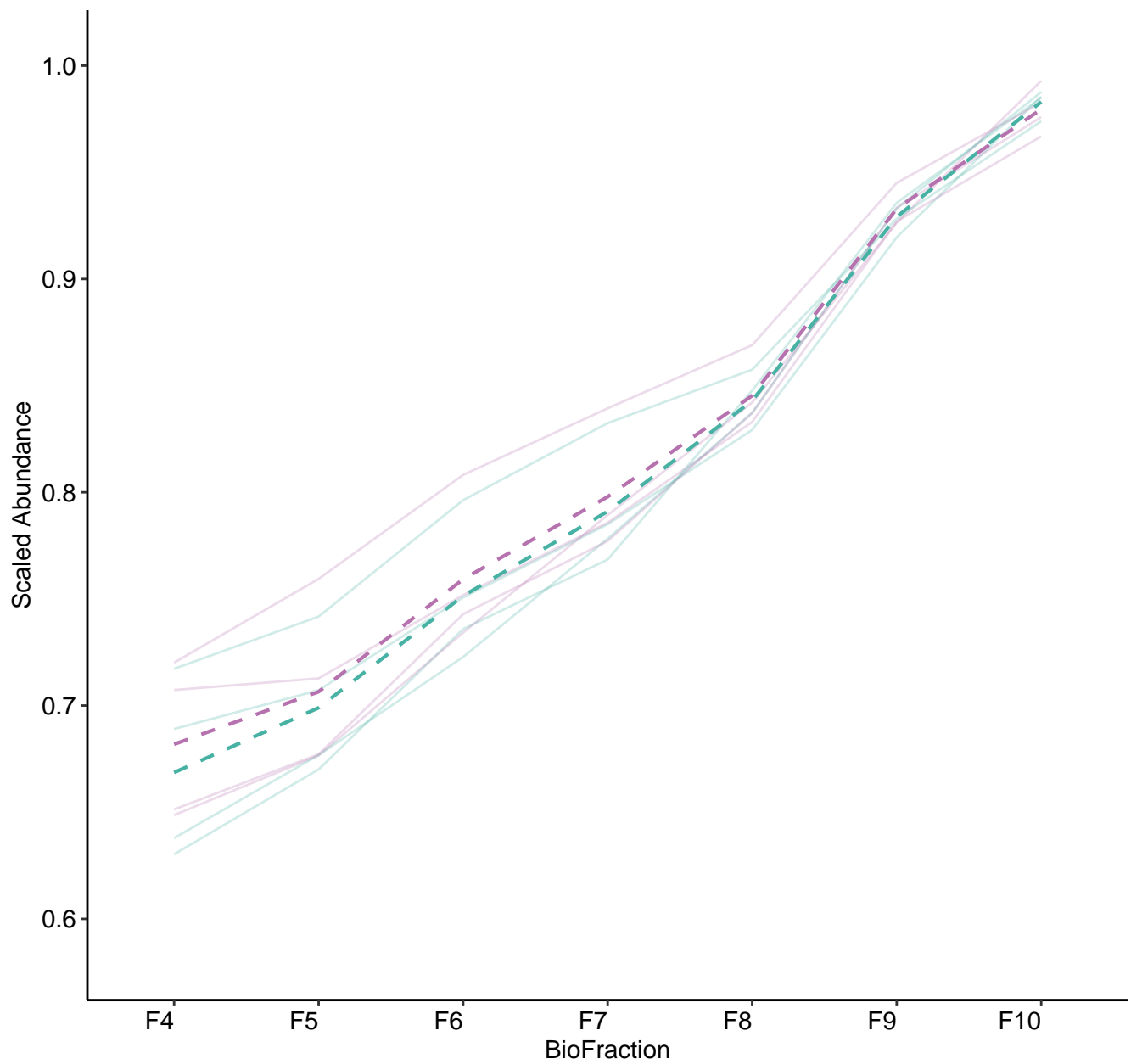


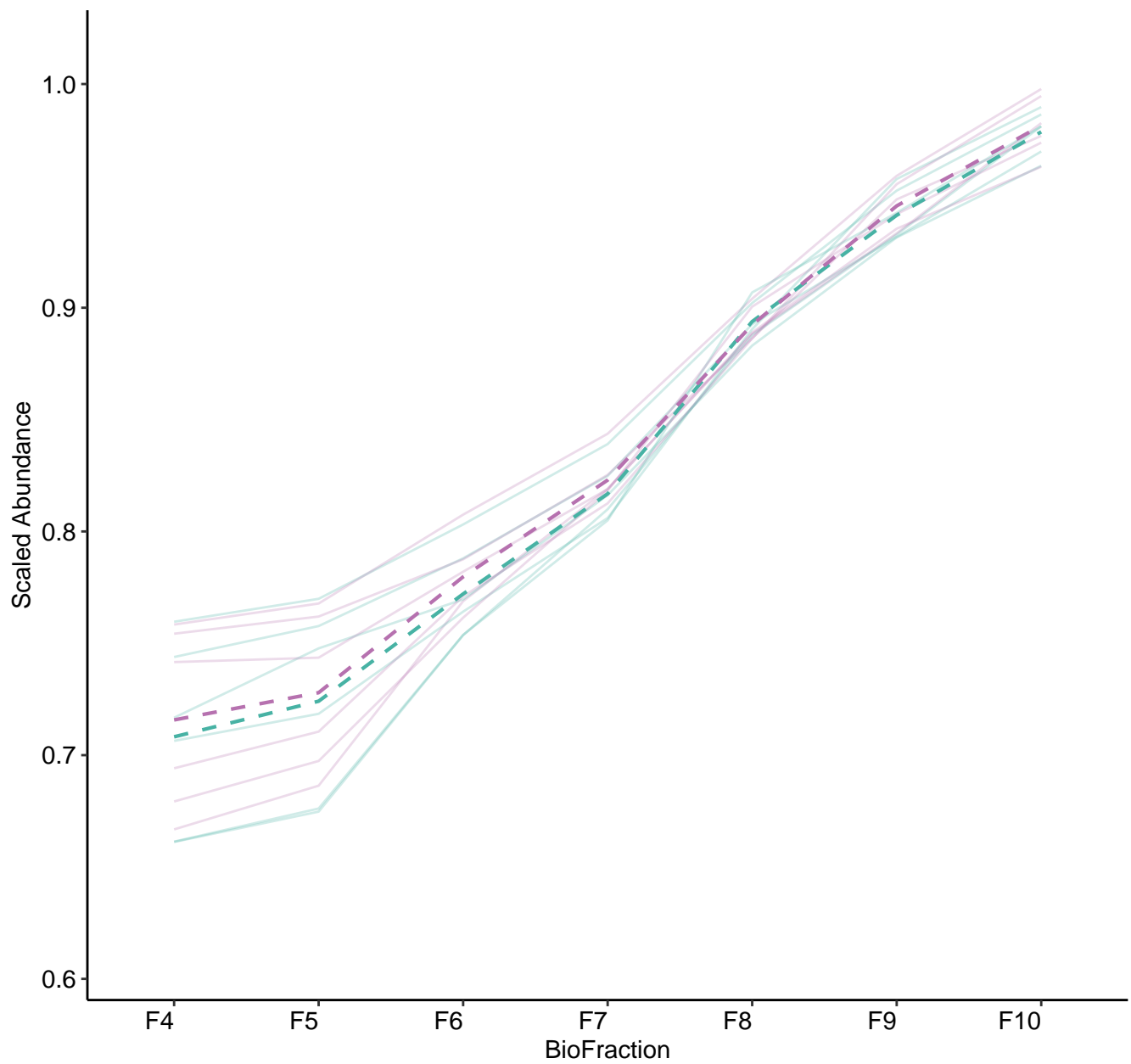
(ER)-localized multiprotein complex, Ig heavy chains associated (n = 4)
(R2.Fixef = 0.978)



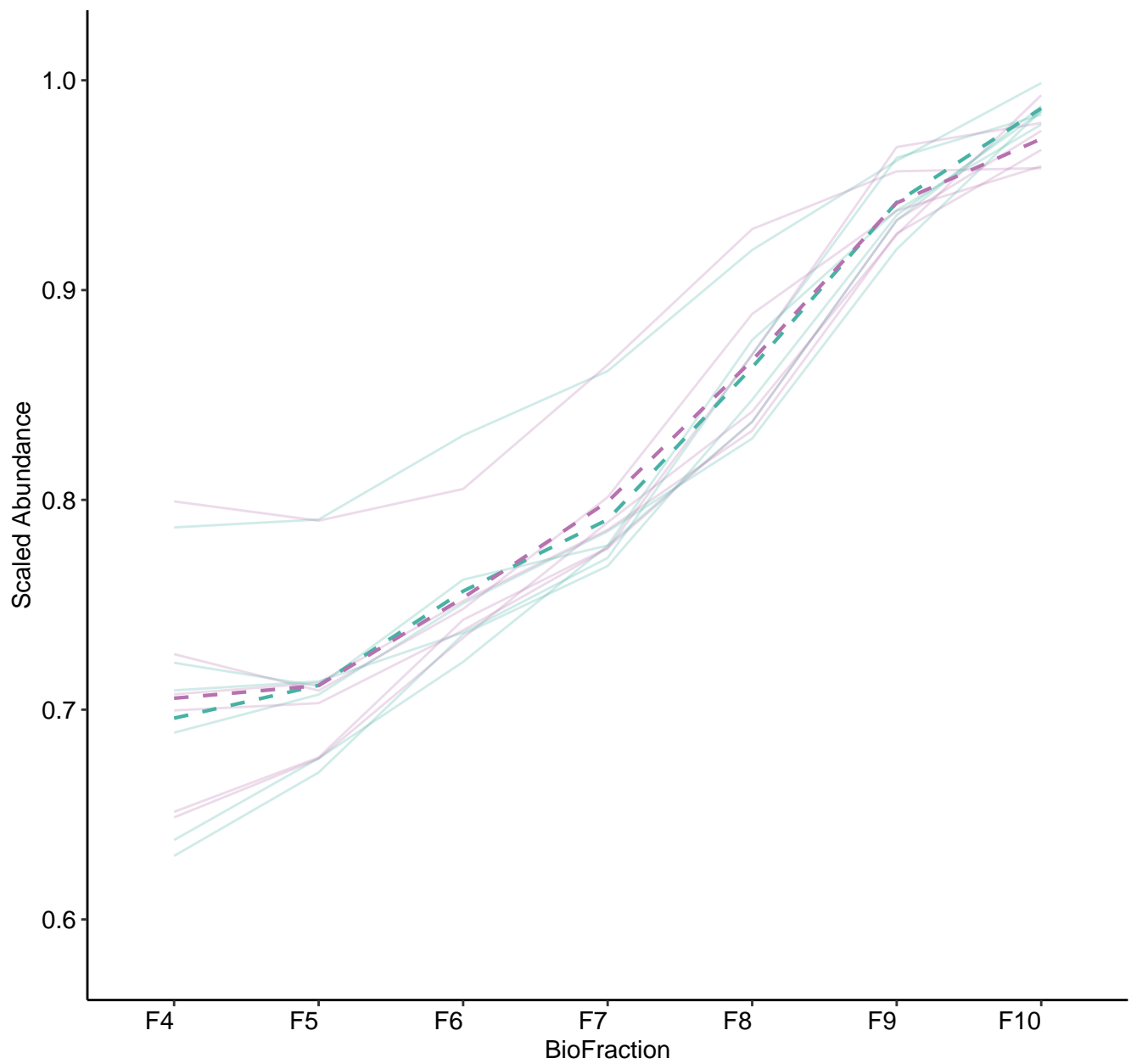
12S U11 snRNP (n = 4)
(R2.Fixef = 0.94)



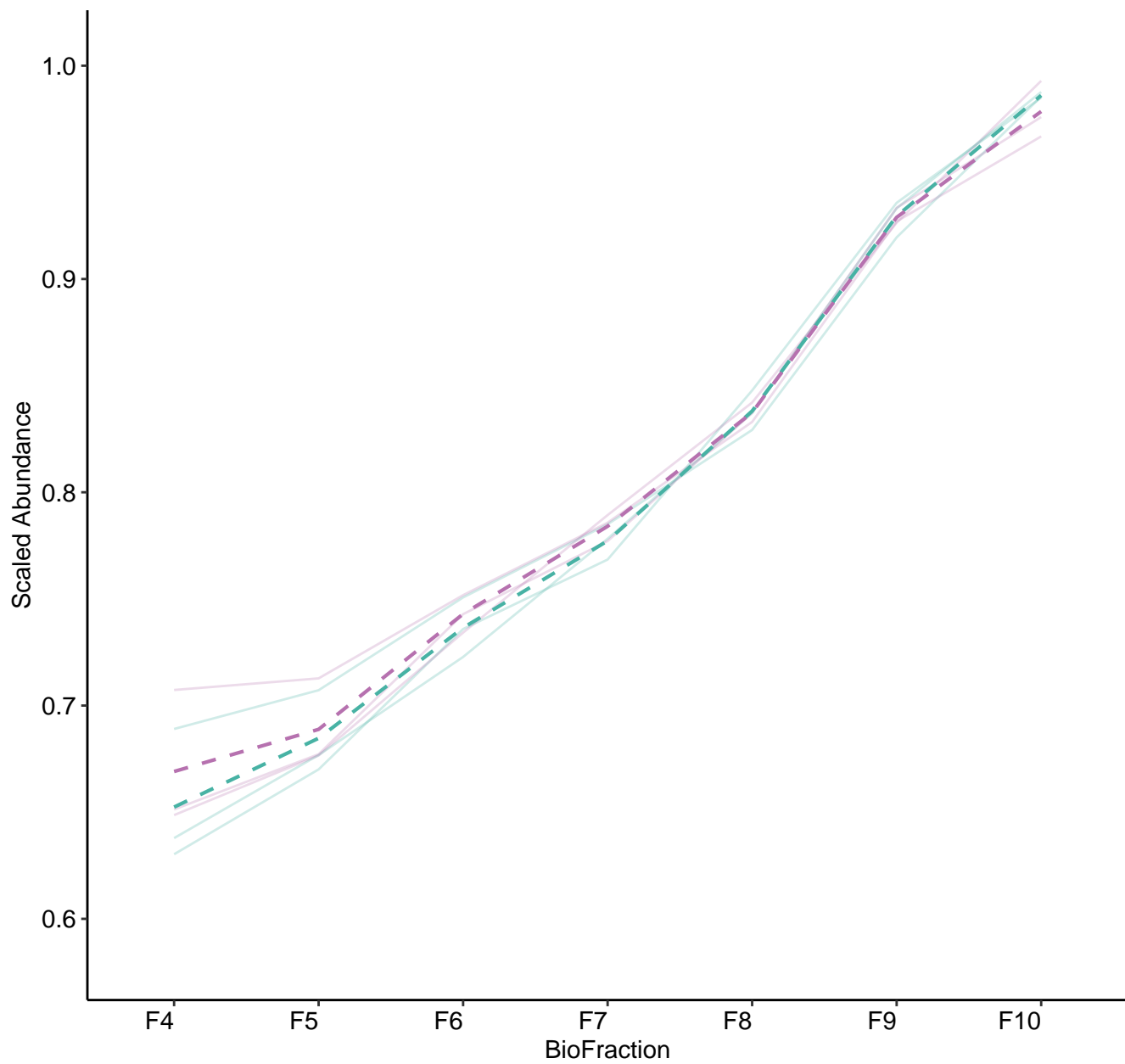
17S U2 snRNP (n = 6)
(R2.Fixef = 0.945)



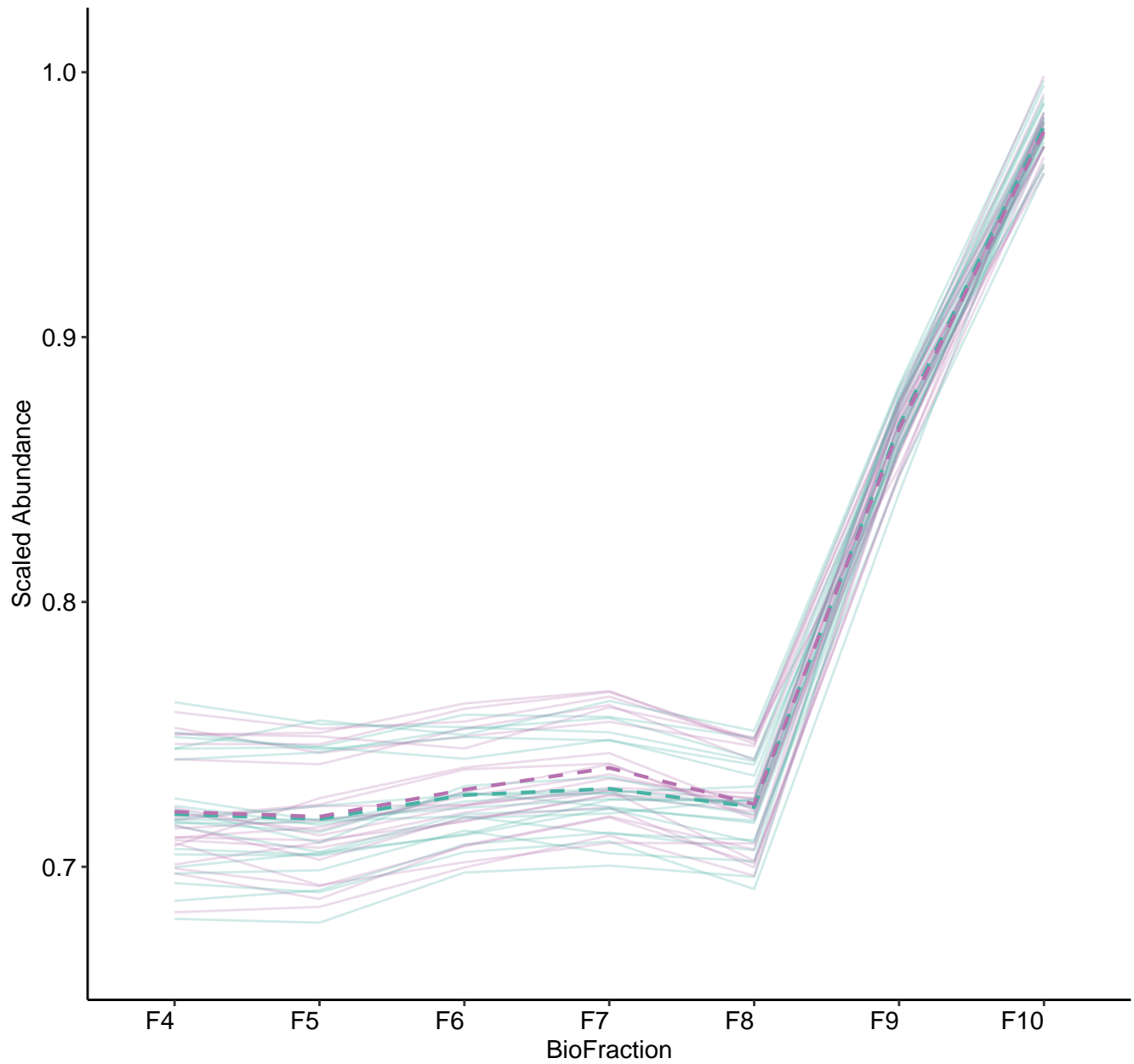
18S U11/U12 snRNP (n = 6)
(R2.Fixef = 0.892)



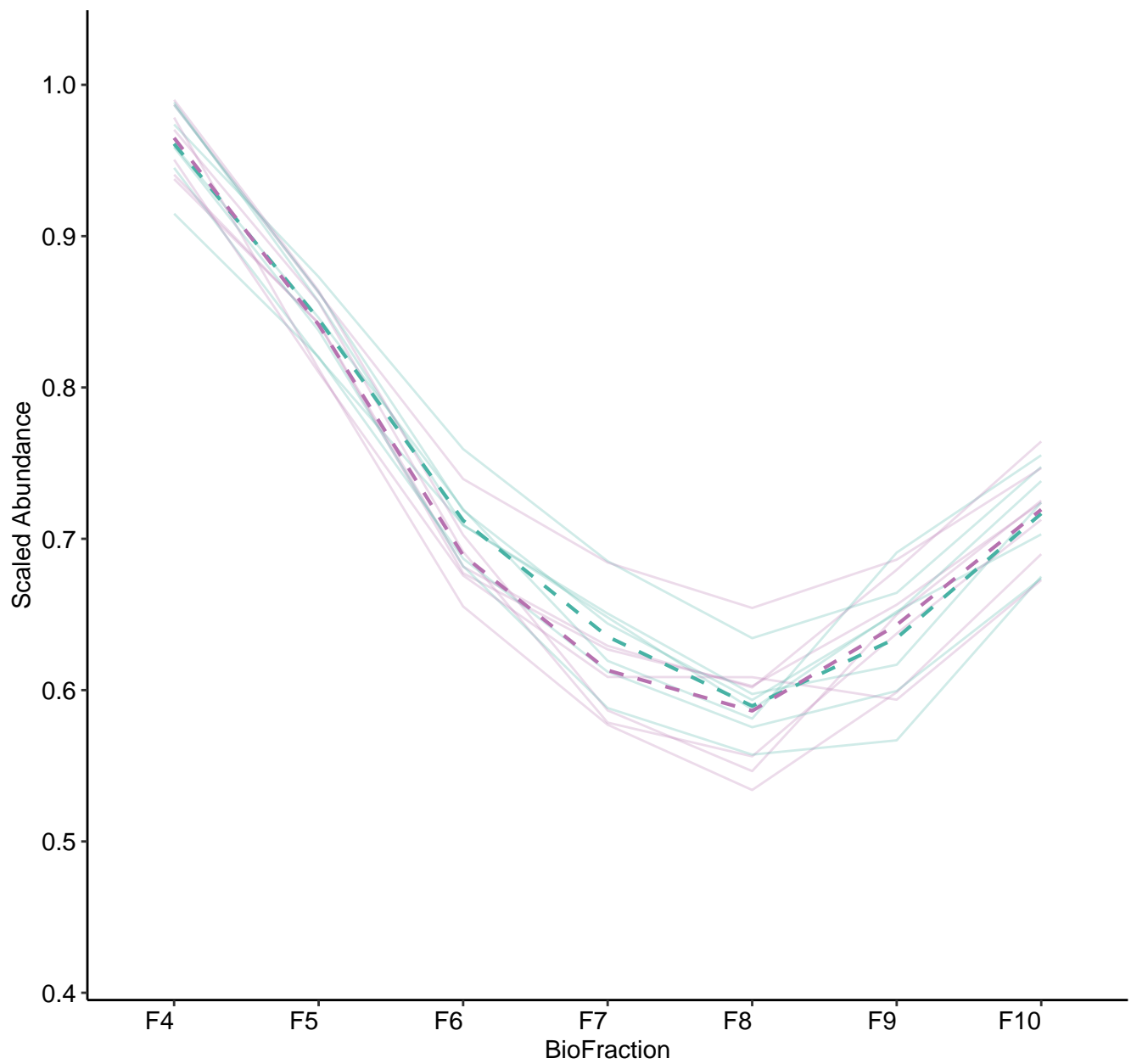
20S methylosome and RG-containing Sm protein complex (n = 3)
(R2.Fixef = 0.98)



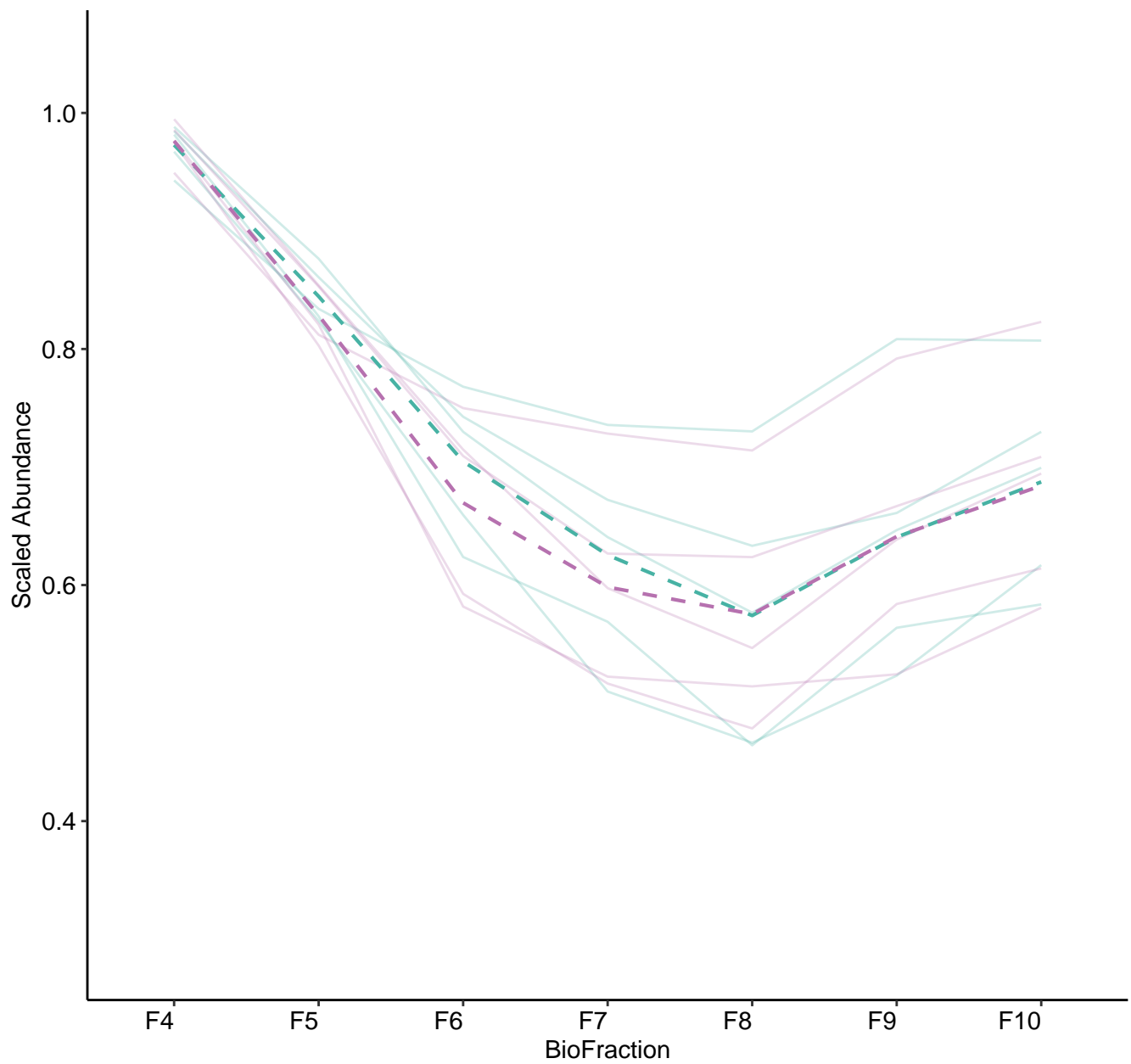
26S proteasome (n = 21)
(R2.Fixef = 0.968)



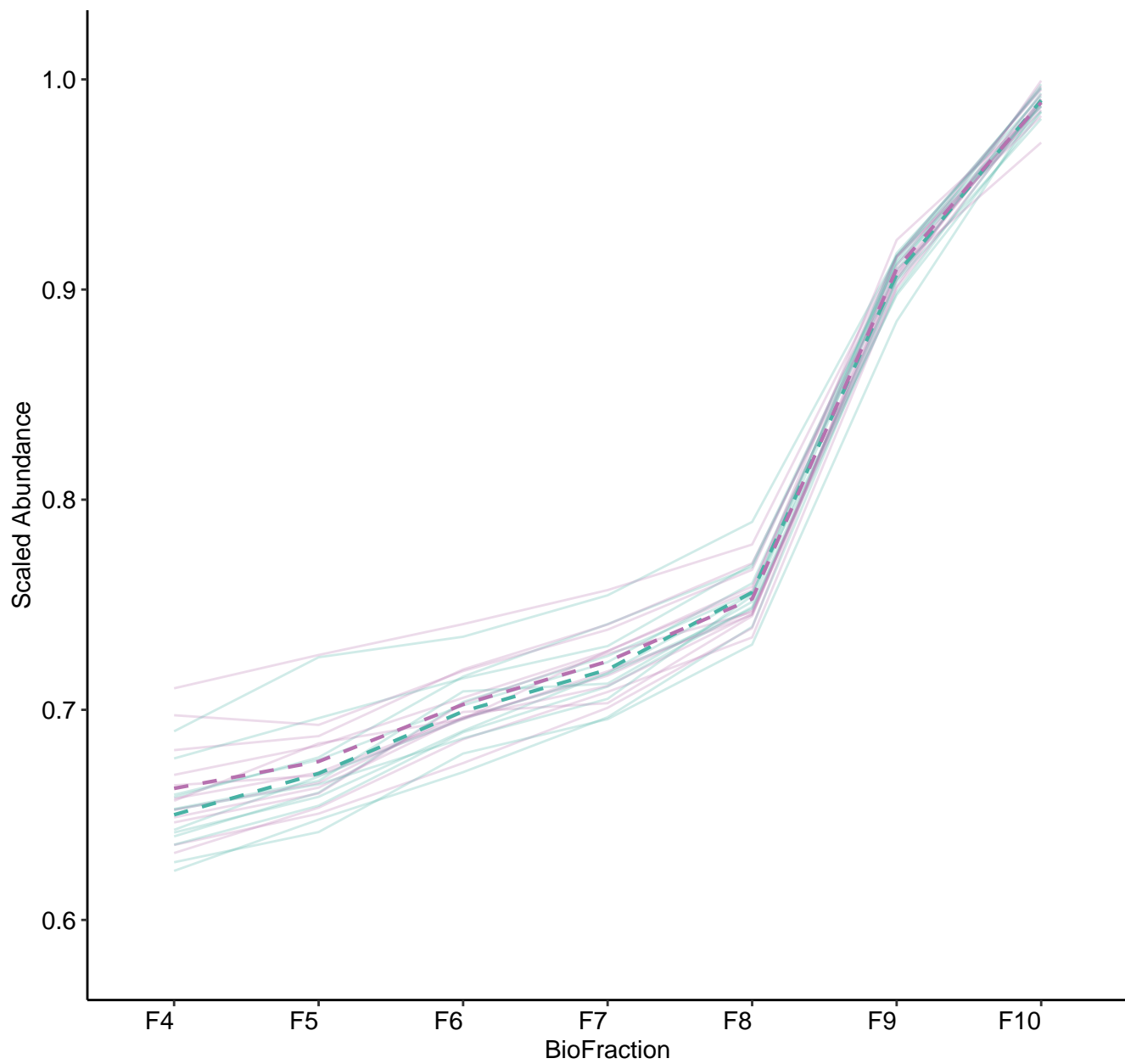
28S ribosomal subunit, mitochondrial (n = 7)
(R2.Fixef = 0.942)



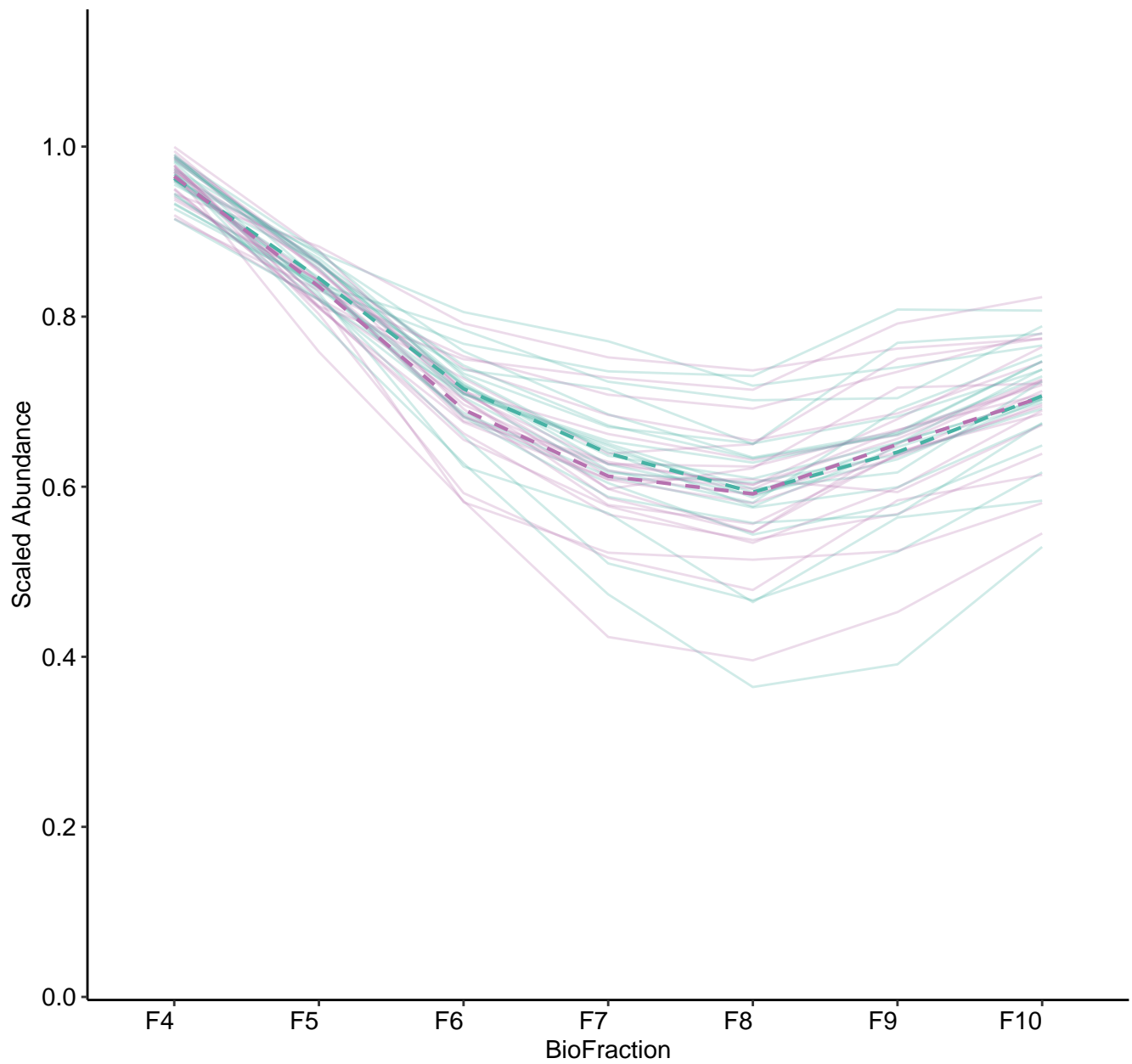
39S ribosomal subunit, mitochondrial (n = 5)
(R2.Fixef = 0.736)



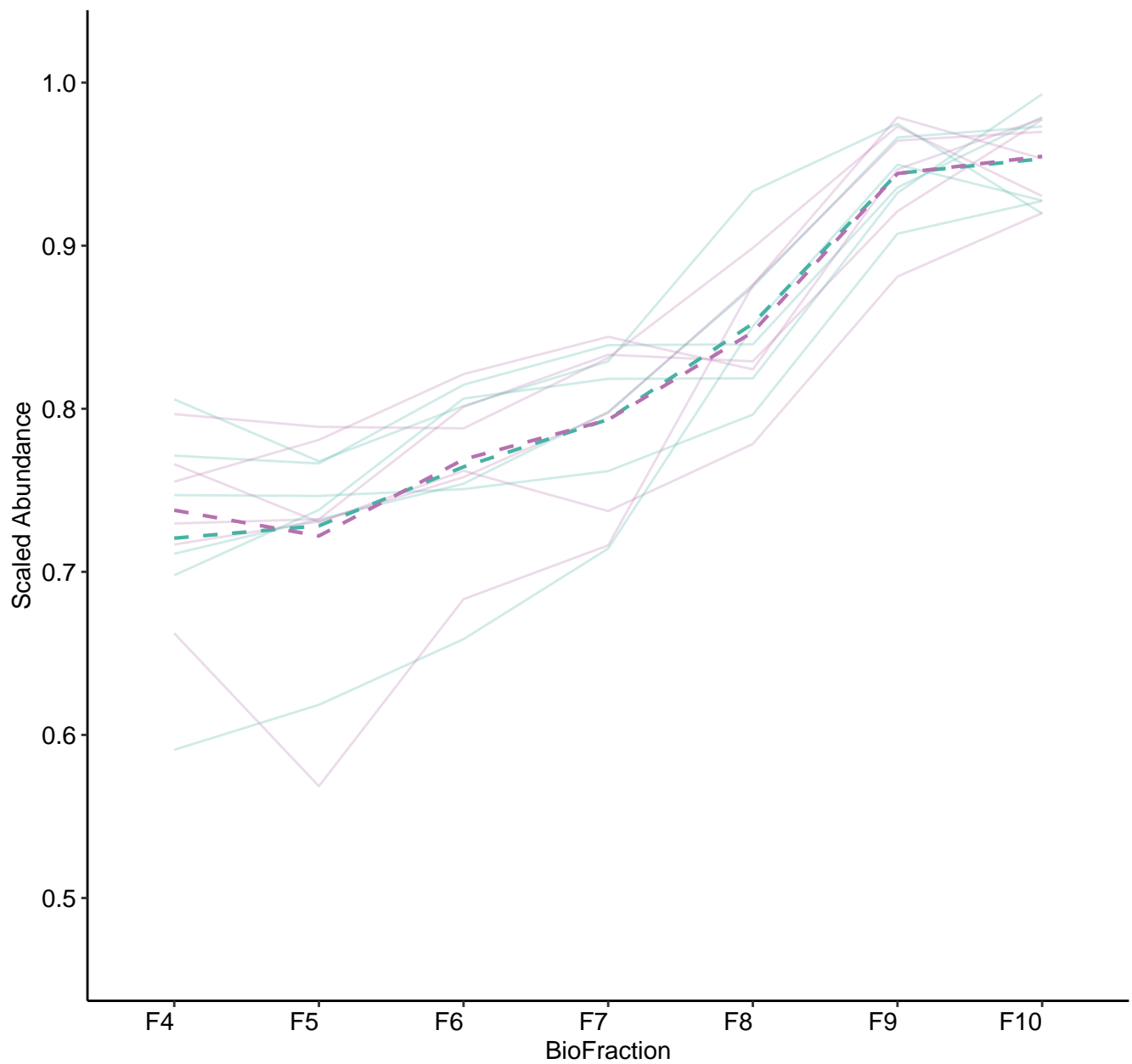
40S ribosomal subunit, cytoplasmic (n = 12)
(R2.Fixef = 0.981)



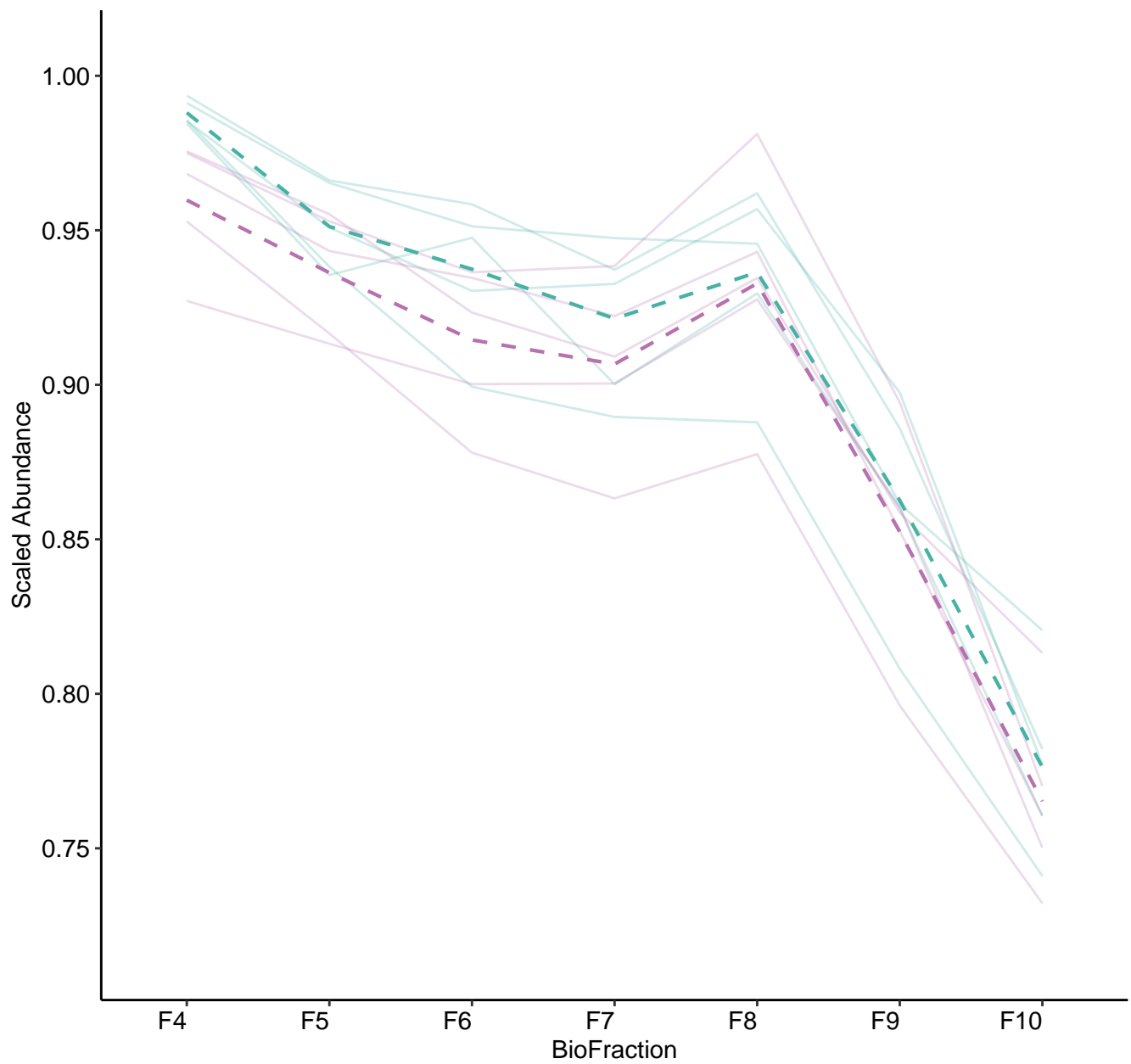
55S ribosome, mitochondrial (n = 22)
(R2.Fixef = 0.8)



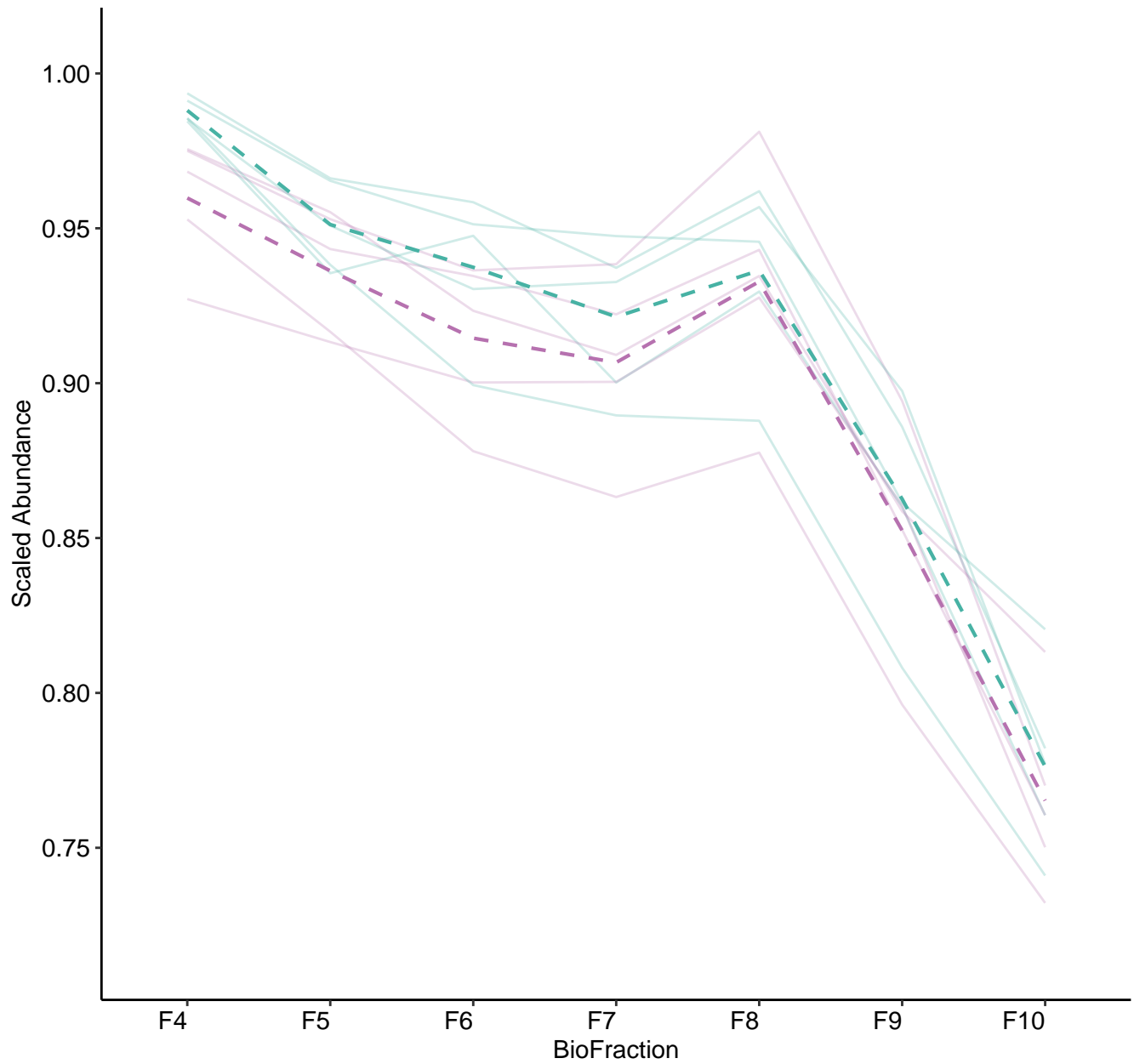
ALL-1 supercomplex (n = 6)
(R2.Fixef = 0.756)



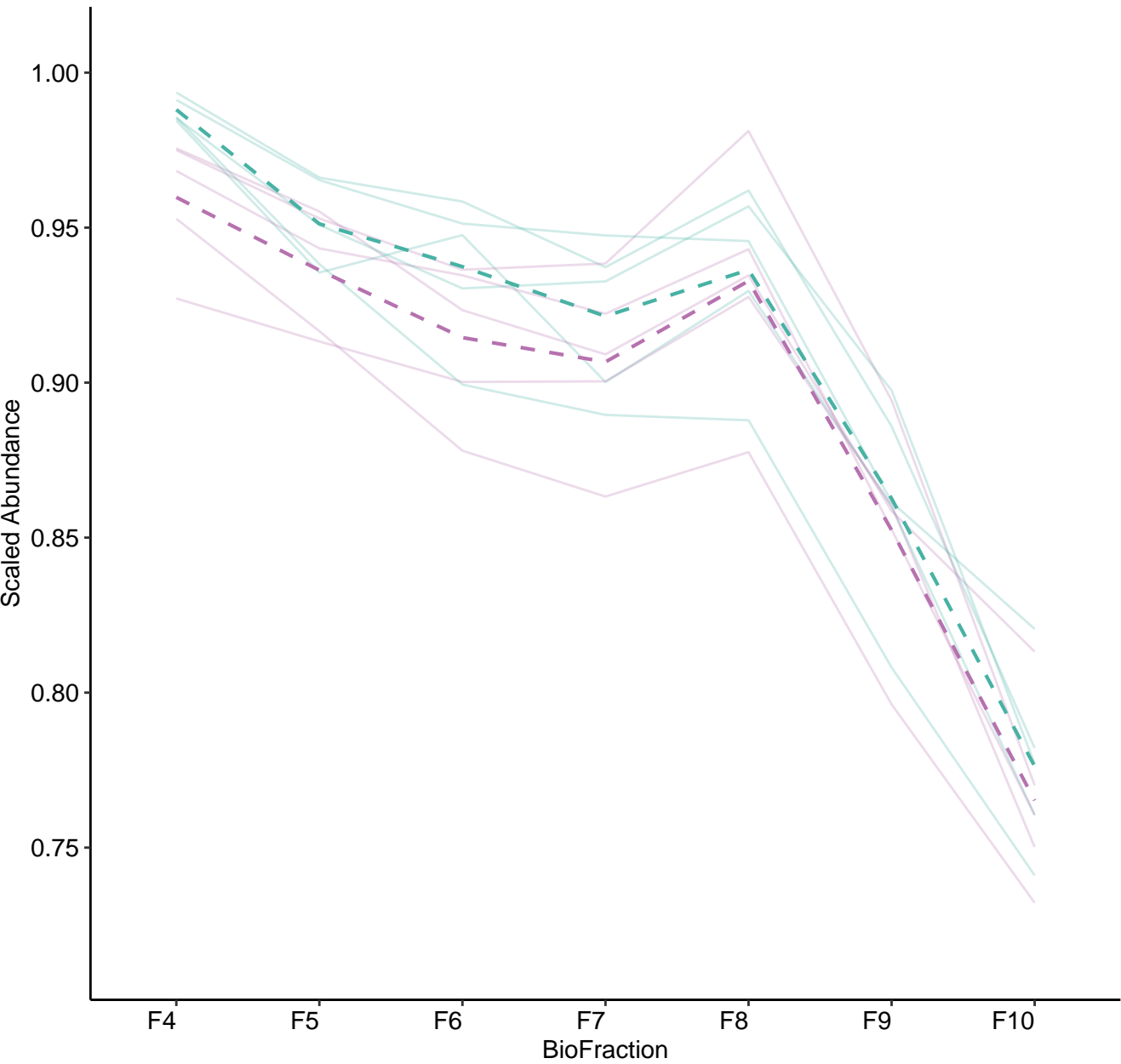
AMPA receptor complex (anti-GluA1-a) (n = 5)
(R2.Fixef = 0.85)



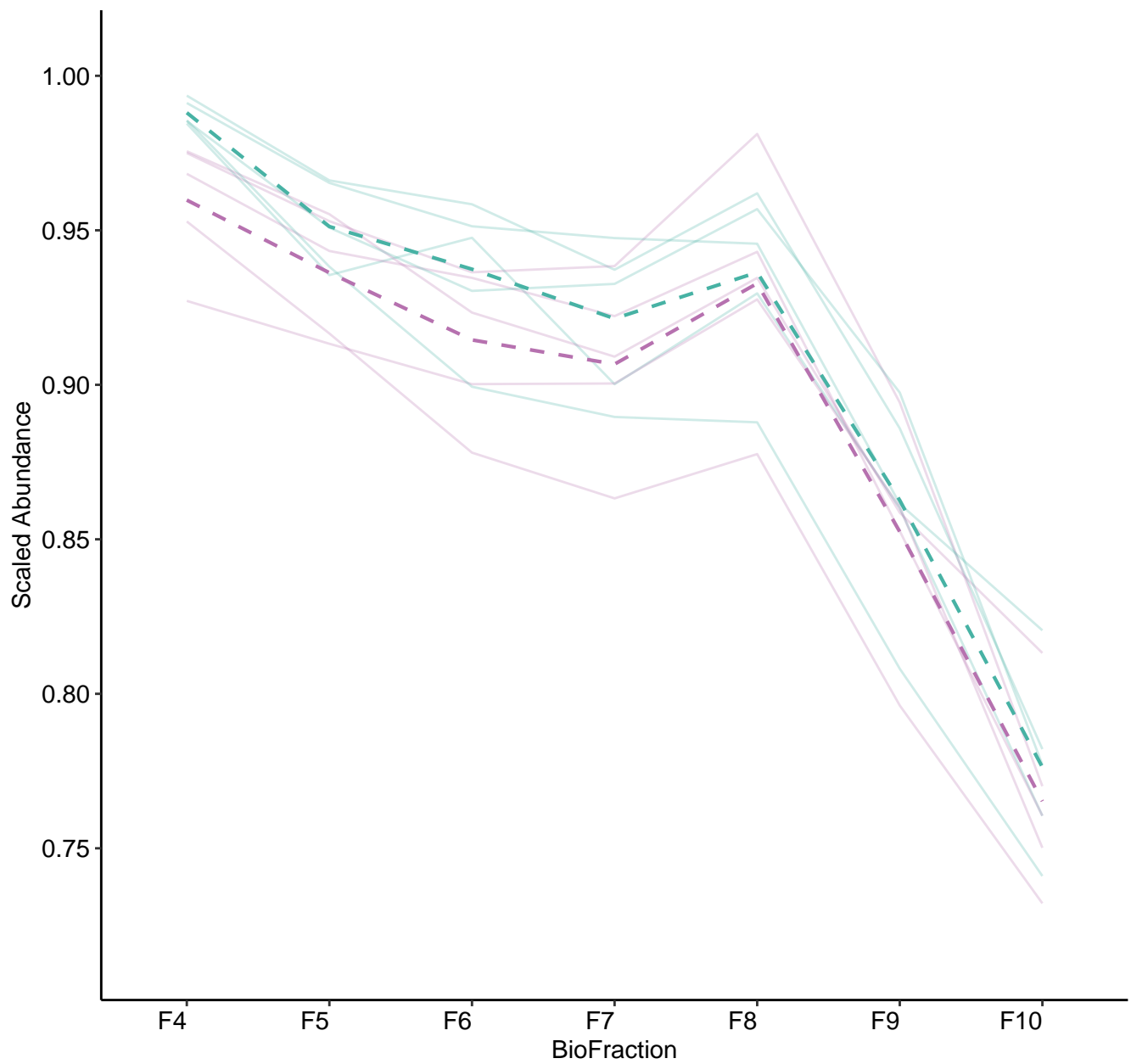
AMPA receptor complex (anti-GluA1-b) (n = 5)
(R2.Fixef = 0.85)



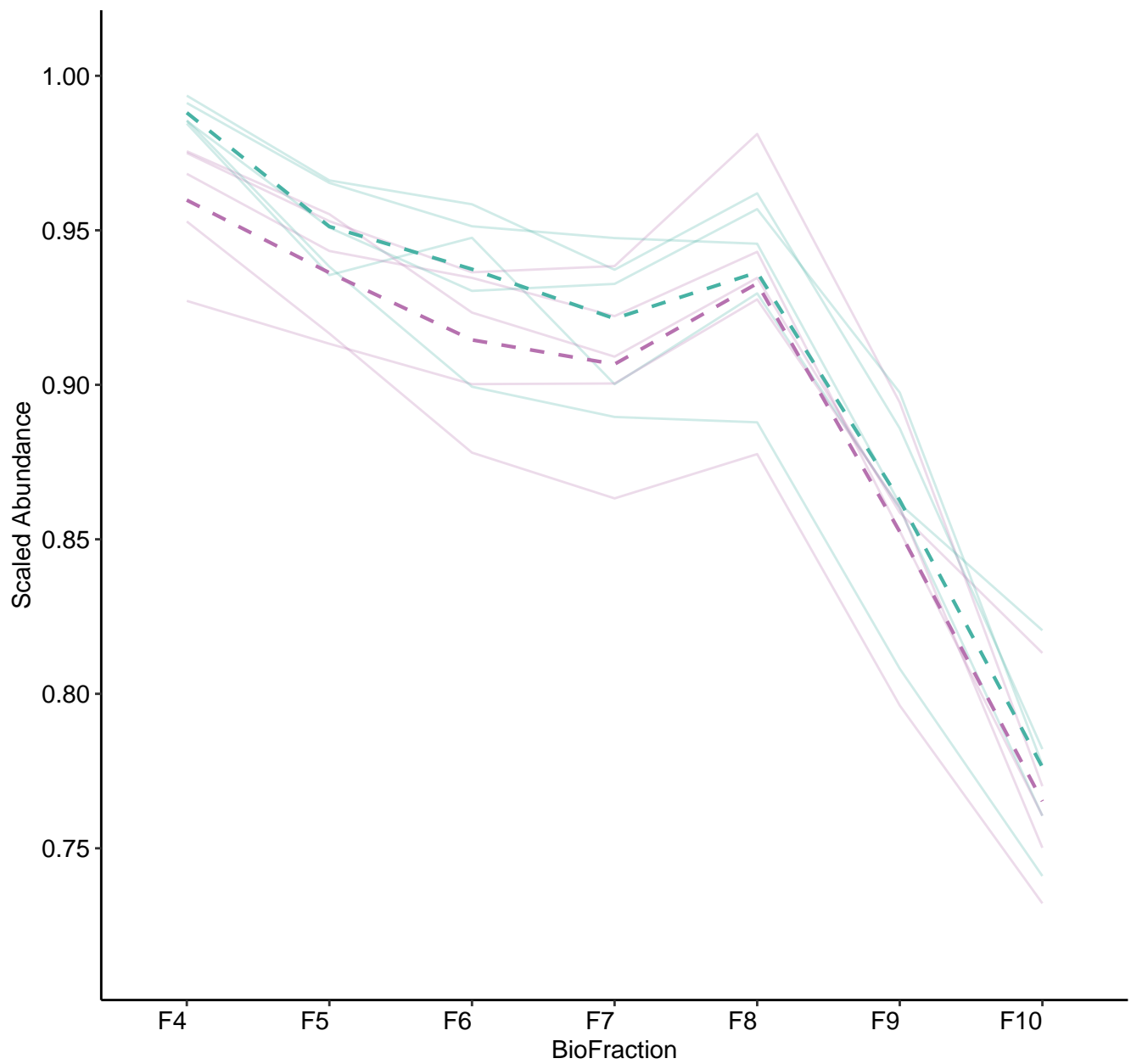
AMPA receptor complex (anti-GluA1-c) (n = 5)
(R2.Fixef = 0.85)



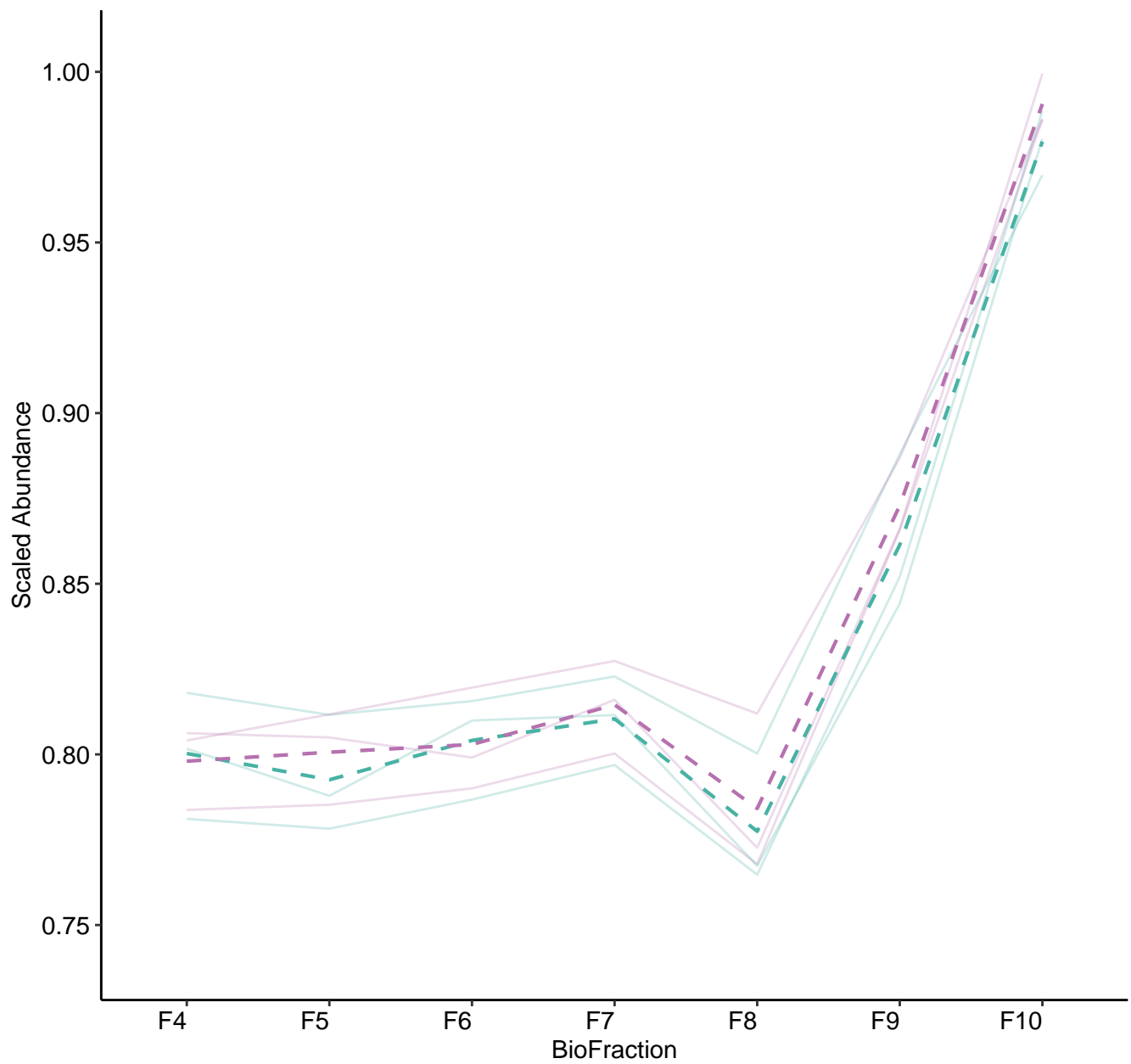
AMPA receptor complex (anti-GluA2-a) (n = 5)
(R2.Fixef = 0.85)



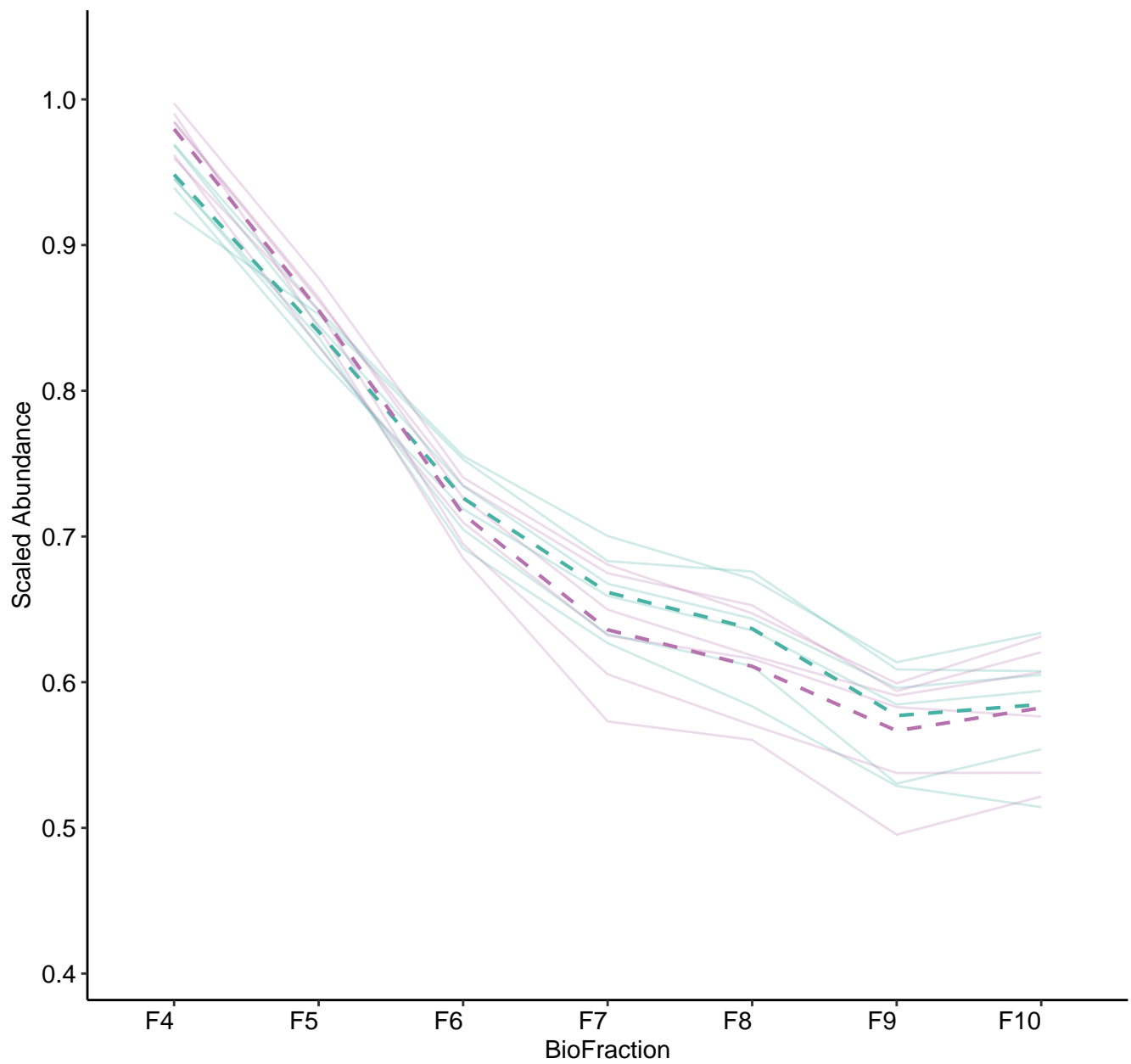
AMPA receptor complex (anti-GluA2-b) (n = 5)
(R2.Fixef = 0.85)



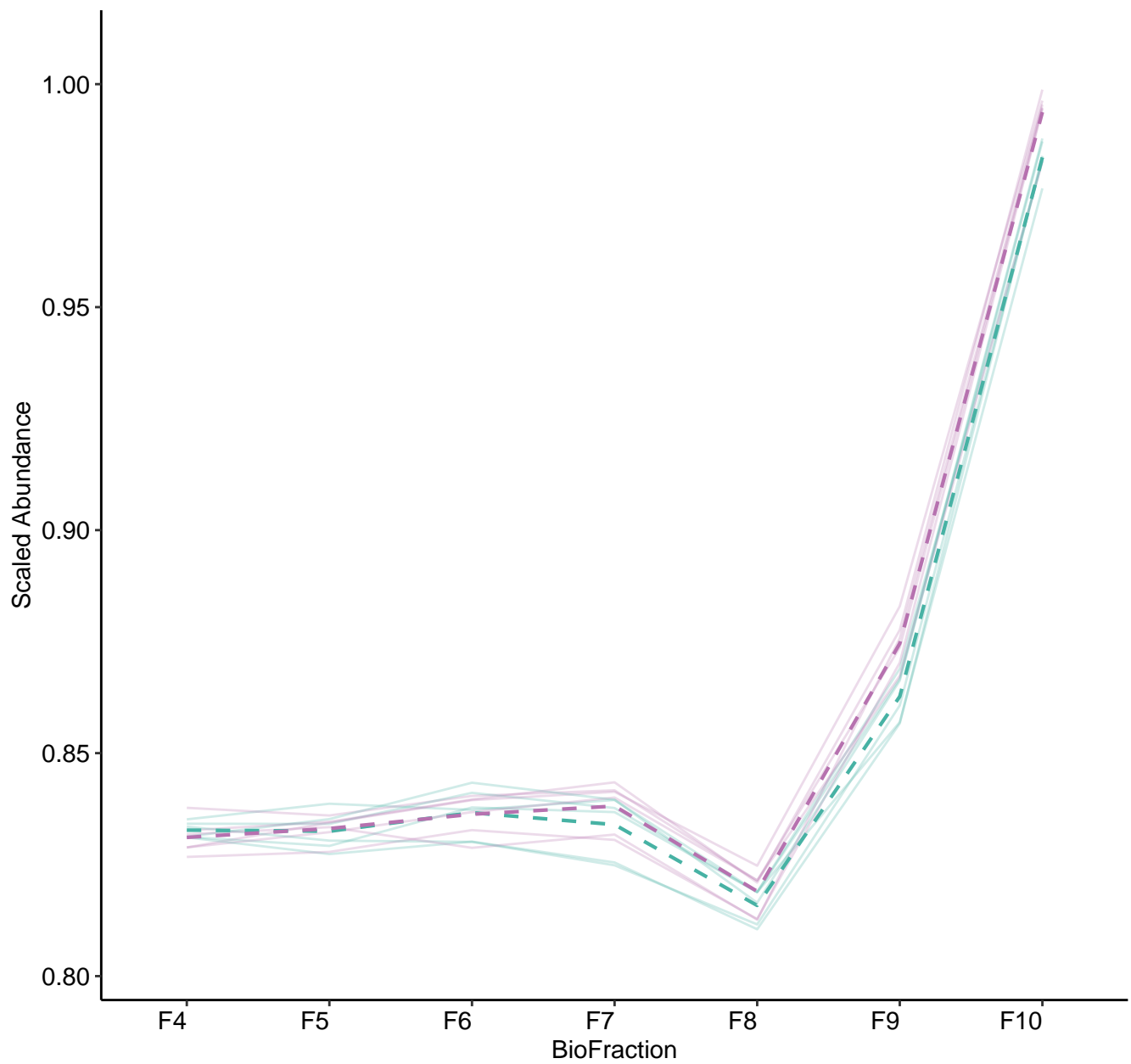
Anaphase-promoting complex (n = 3)
(R2.Fixef = 0.946)



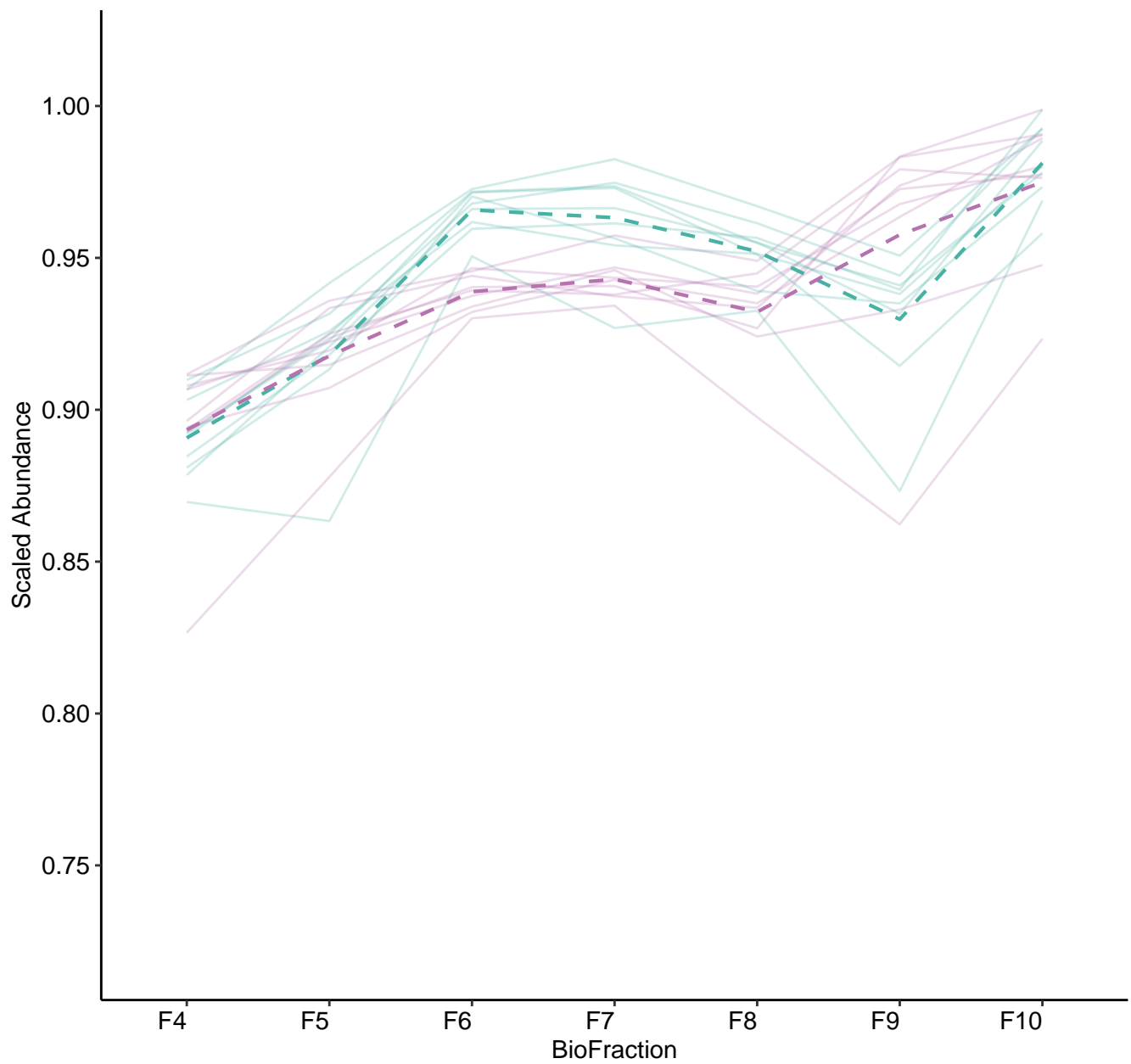
ATP synthasome (n = 6)
(R2.Fixef = 0.948)



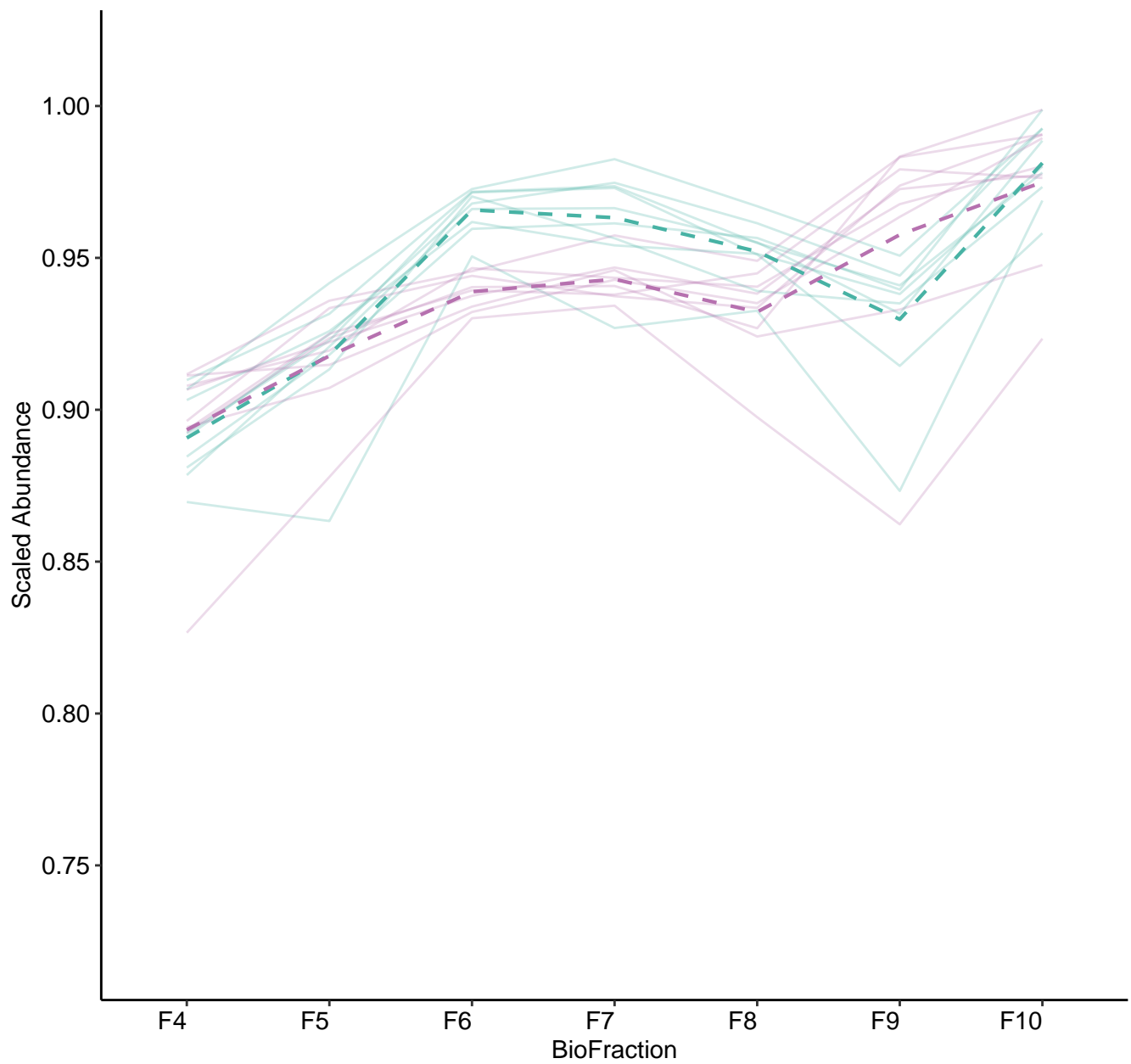
BBS-chaperonin complex (n = 6)
(R2.Fixef = 0.993)



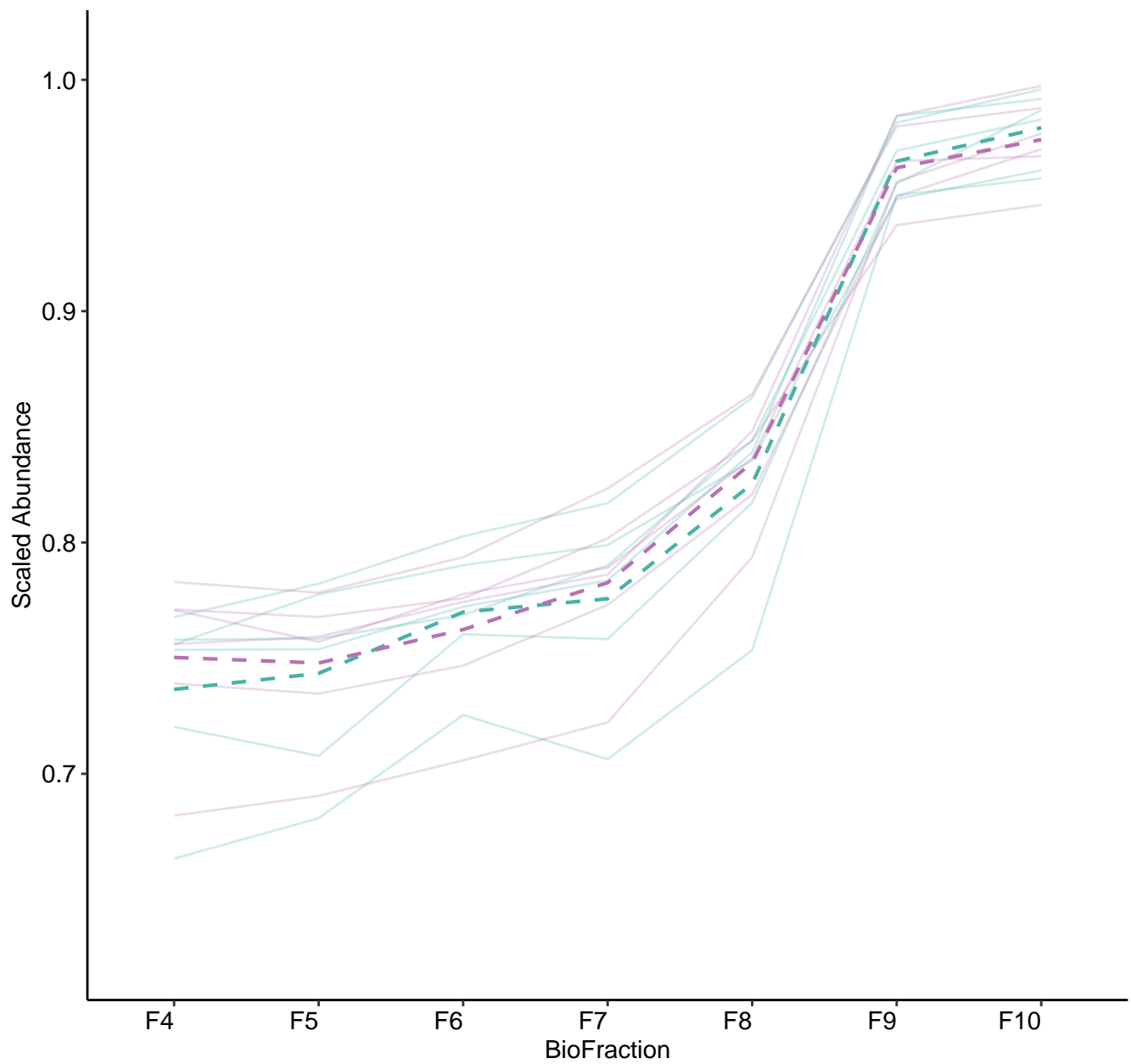
CCC complex (n = 9)
(R2.Fixef = 0.666)



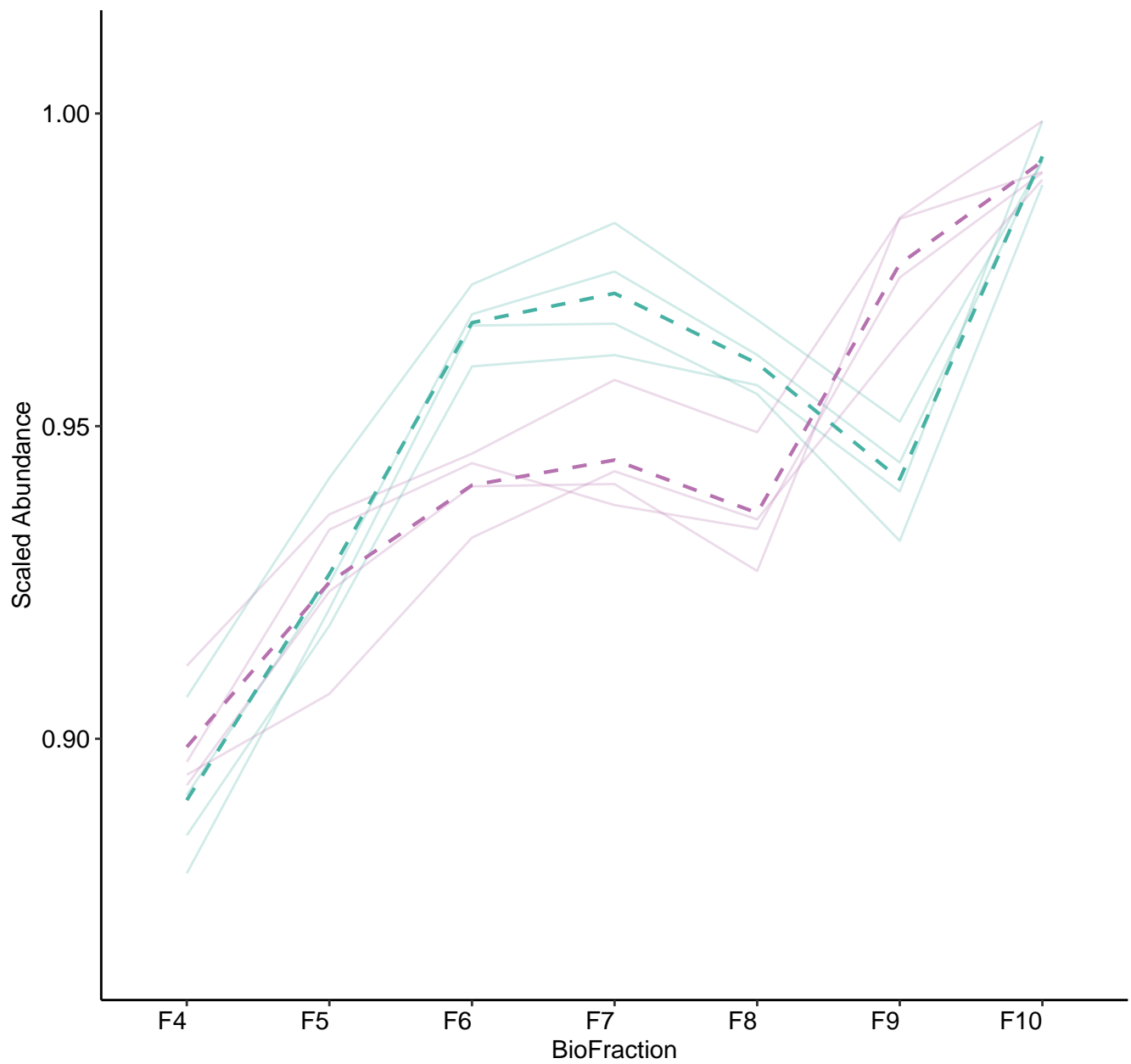
CCC-Wash (WASH1, FAM21C) complex (n = 9)
(R2.Fixef = 0.666)



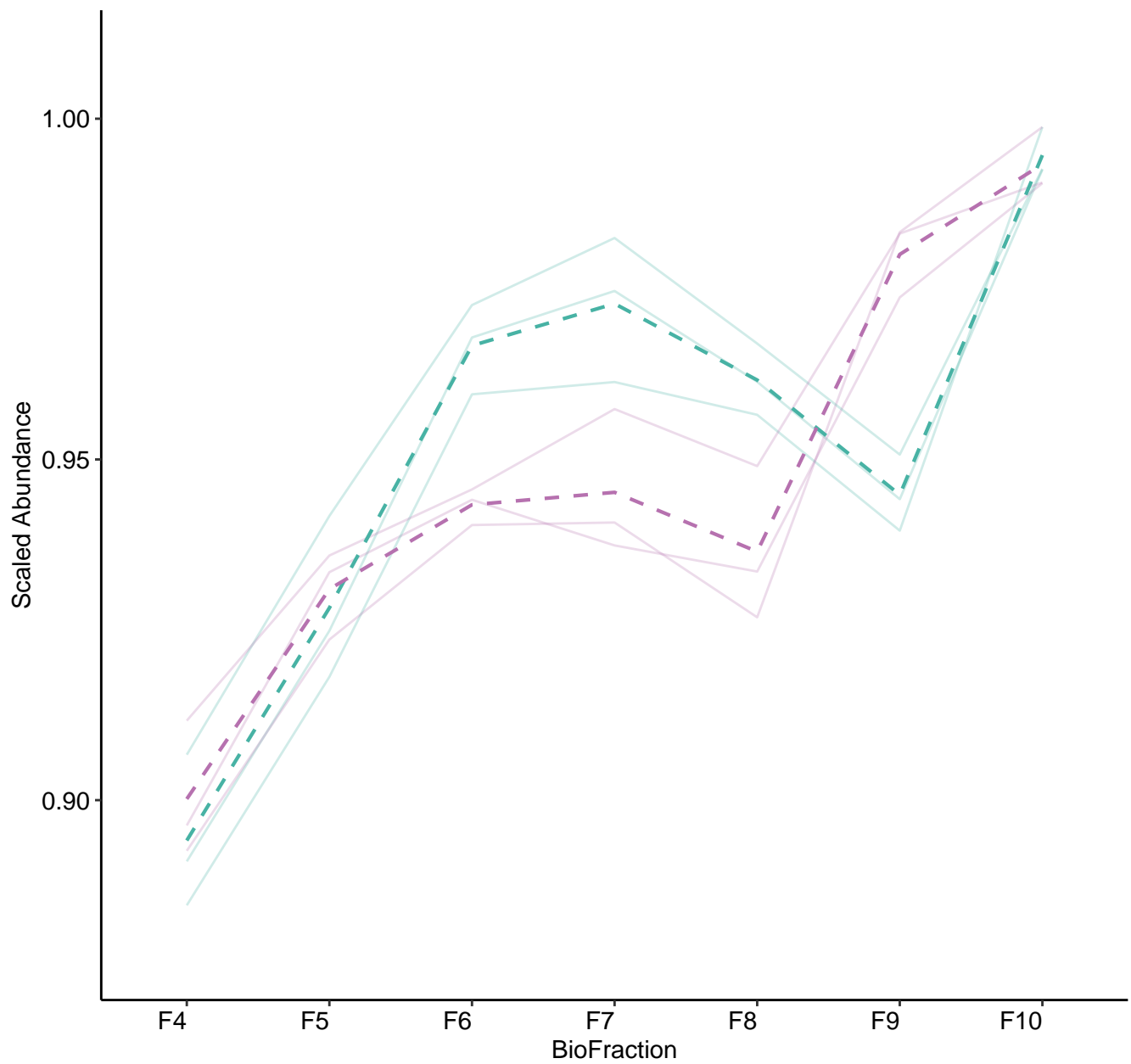
CCR4-NOT-CNOT7-CNOT6L complex (n = 6)
(R2.Fixef = 0.903)



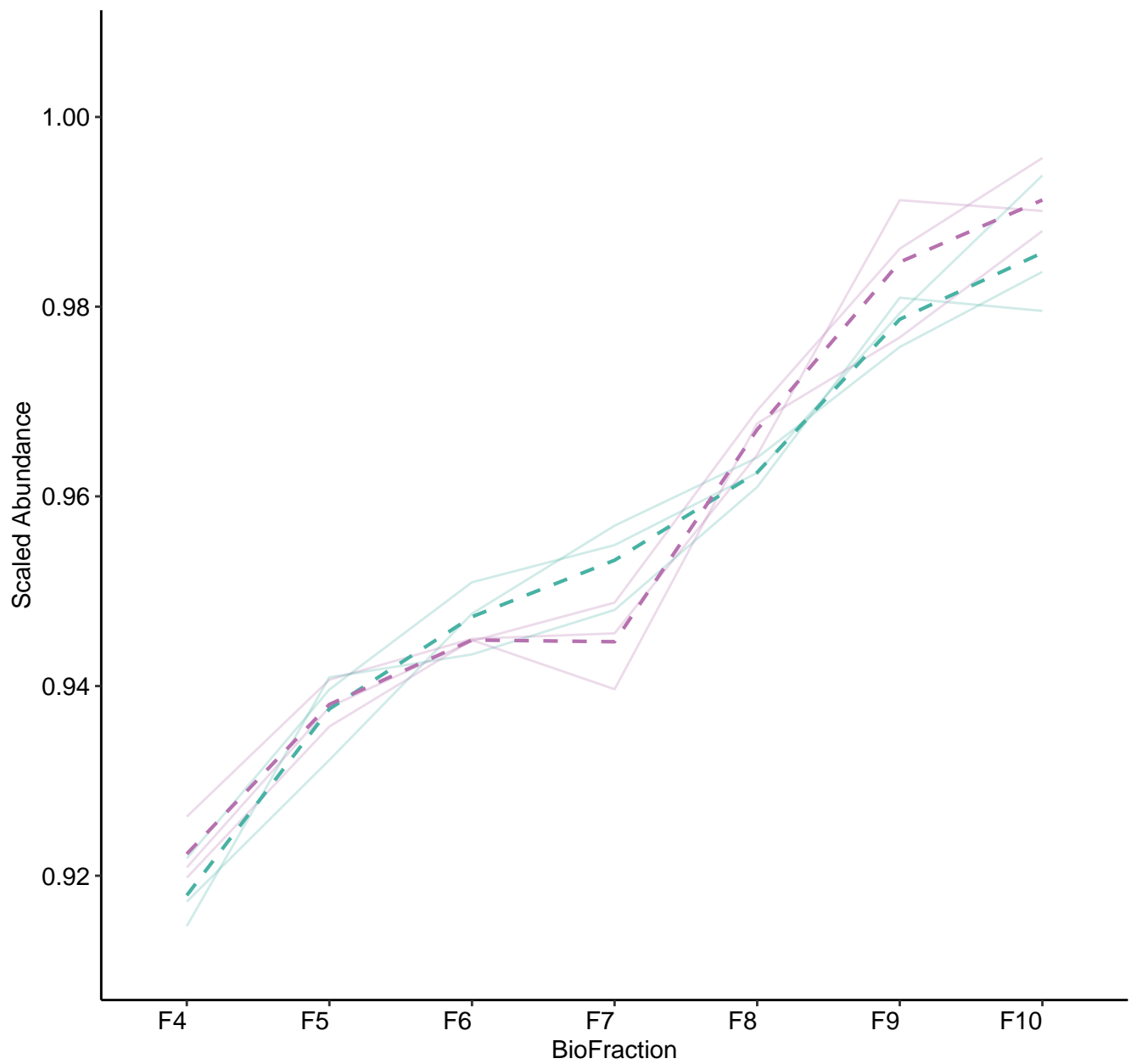
COMMD1-CCDC22-CCDC93-C16orf62 complex (n = 4)
(R2.Fixef = 0.926)



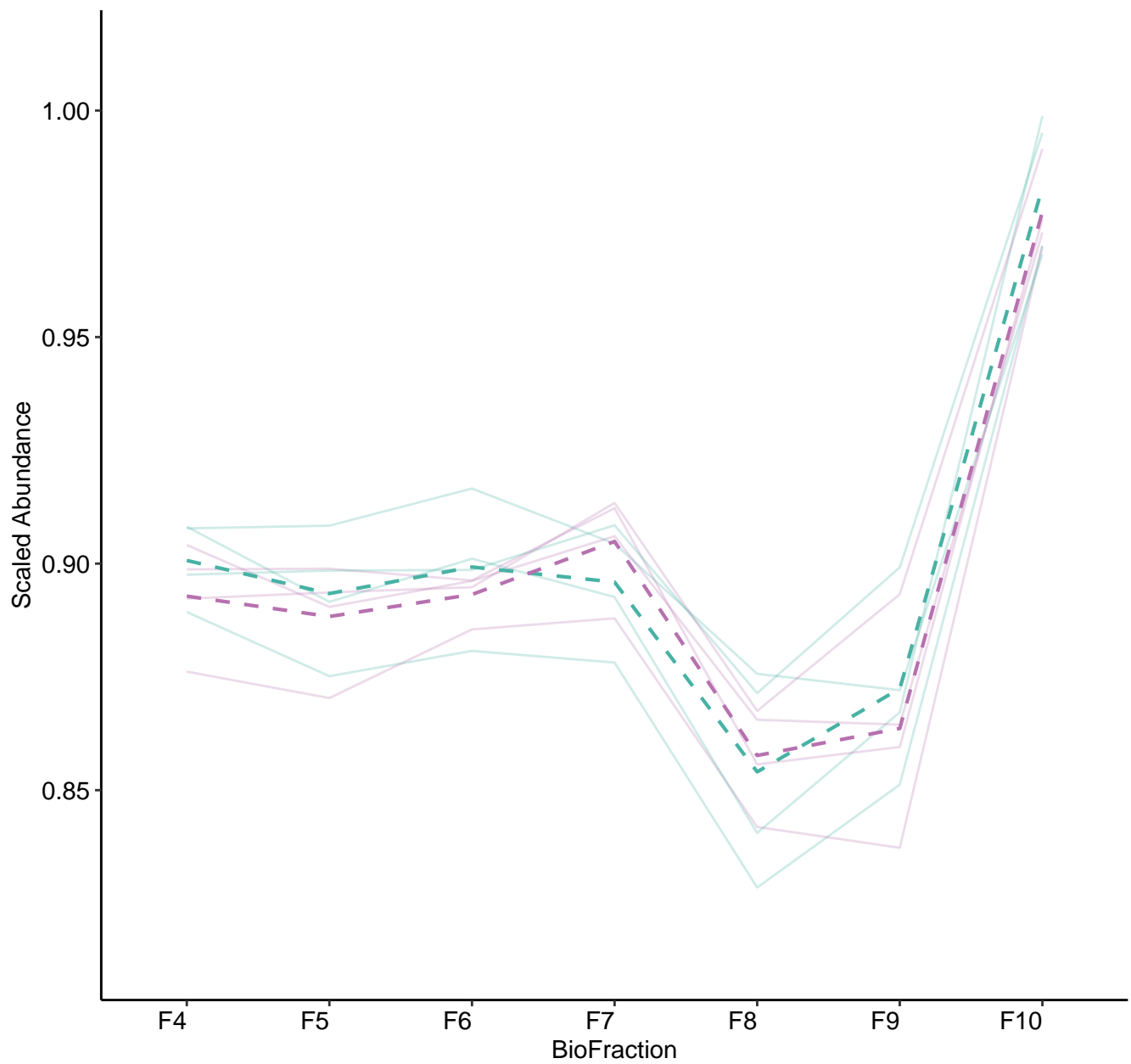
COMMD1-CCDC93-C16orf62 complex (n = 3)
(R2.Fixef = 0.93)



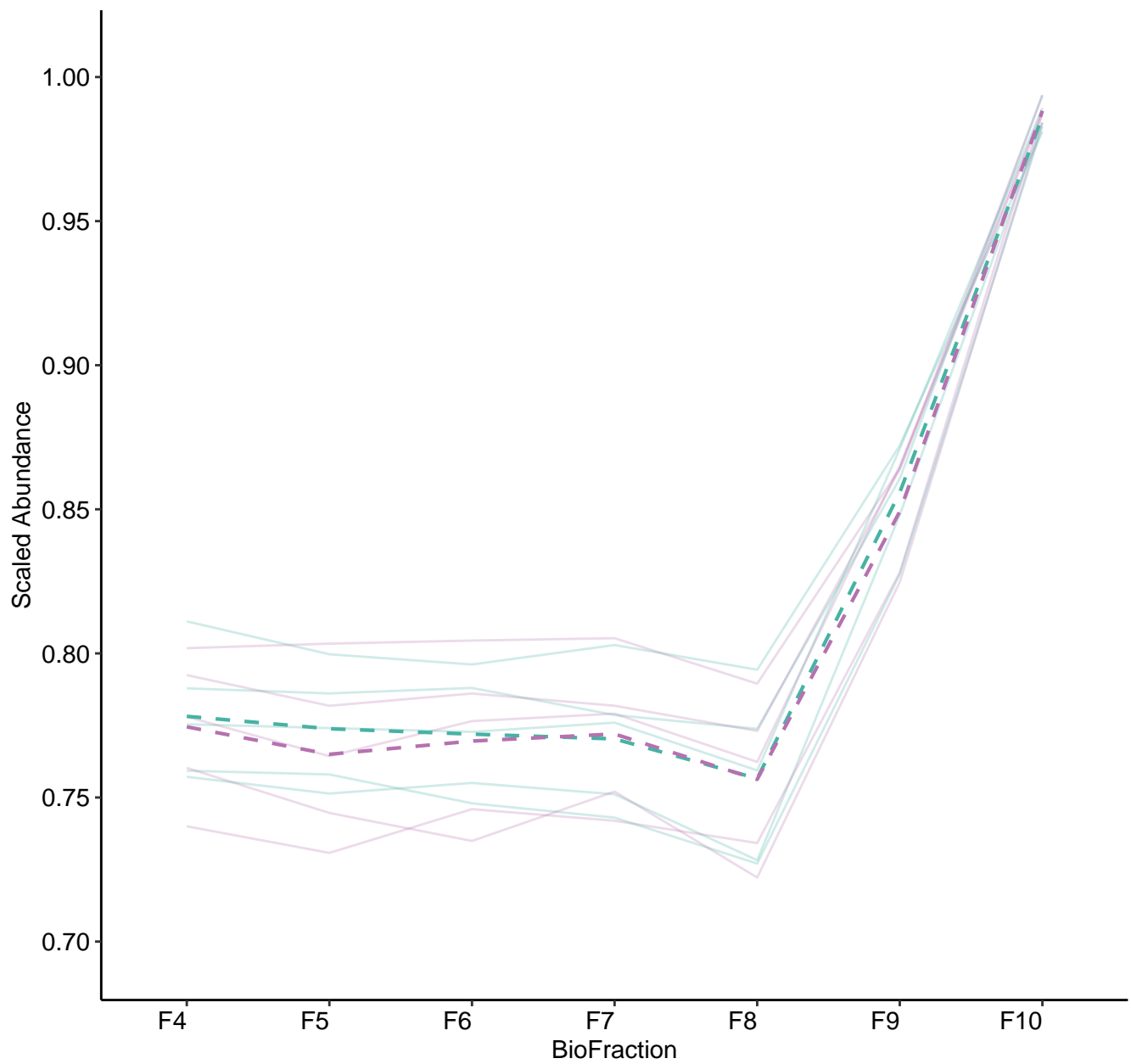
CORVET complex (n = 3)
(R2.Fixef = 0.967)



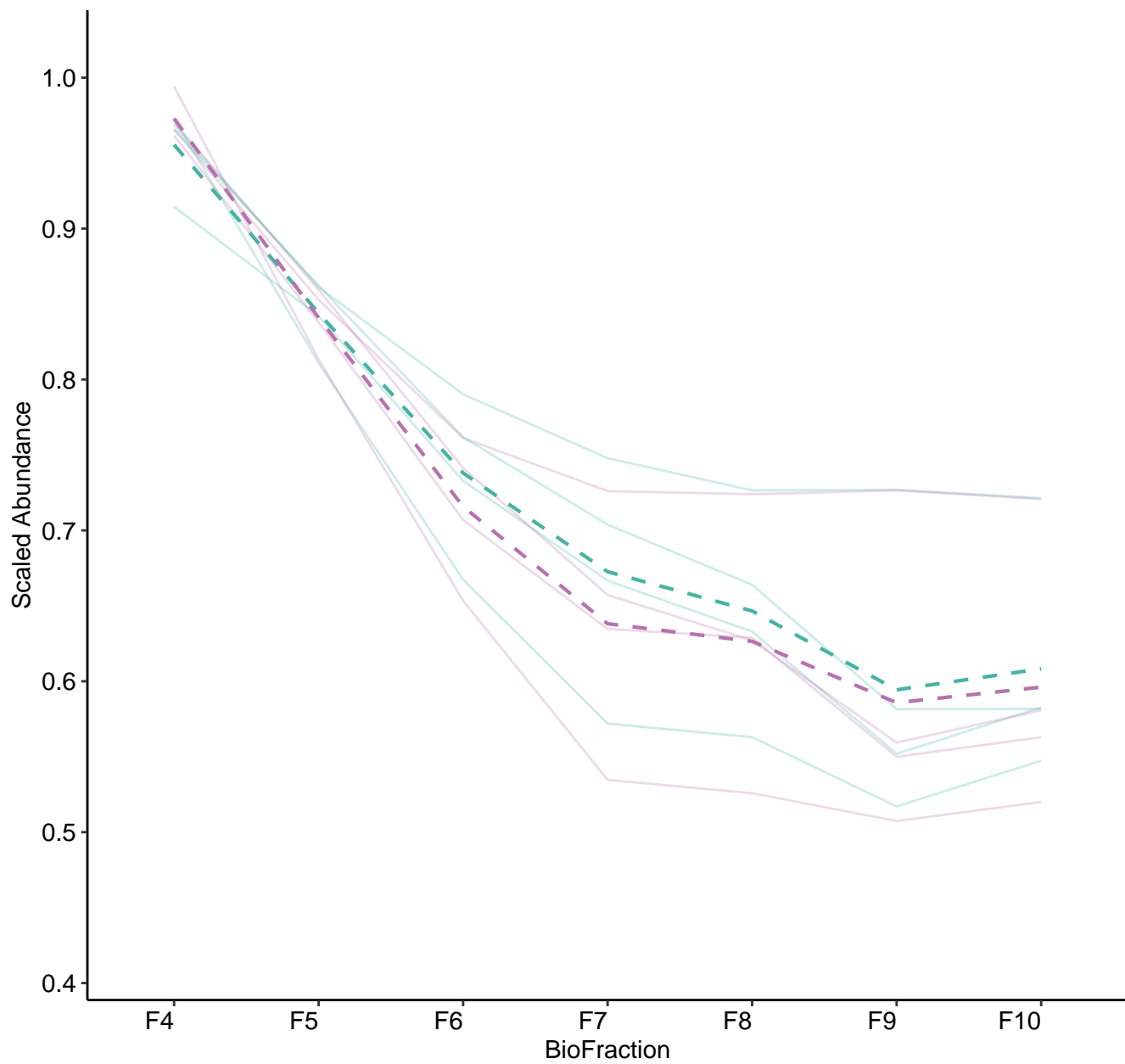
CSA-POLIIa complex (n = 4)
(R2.Fixef = 0.863)



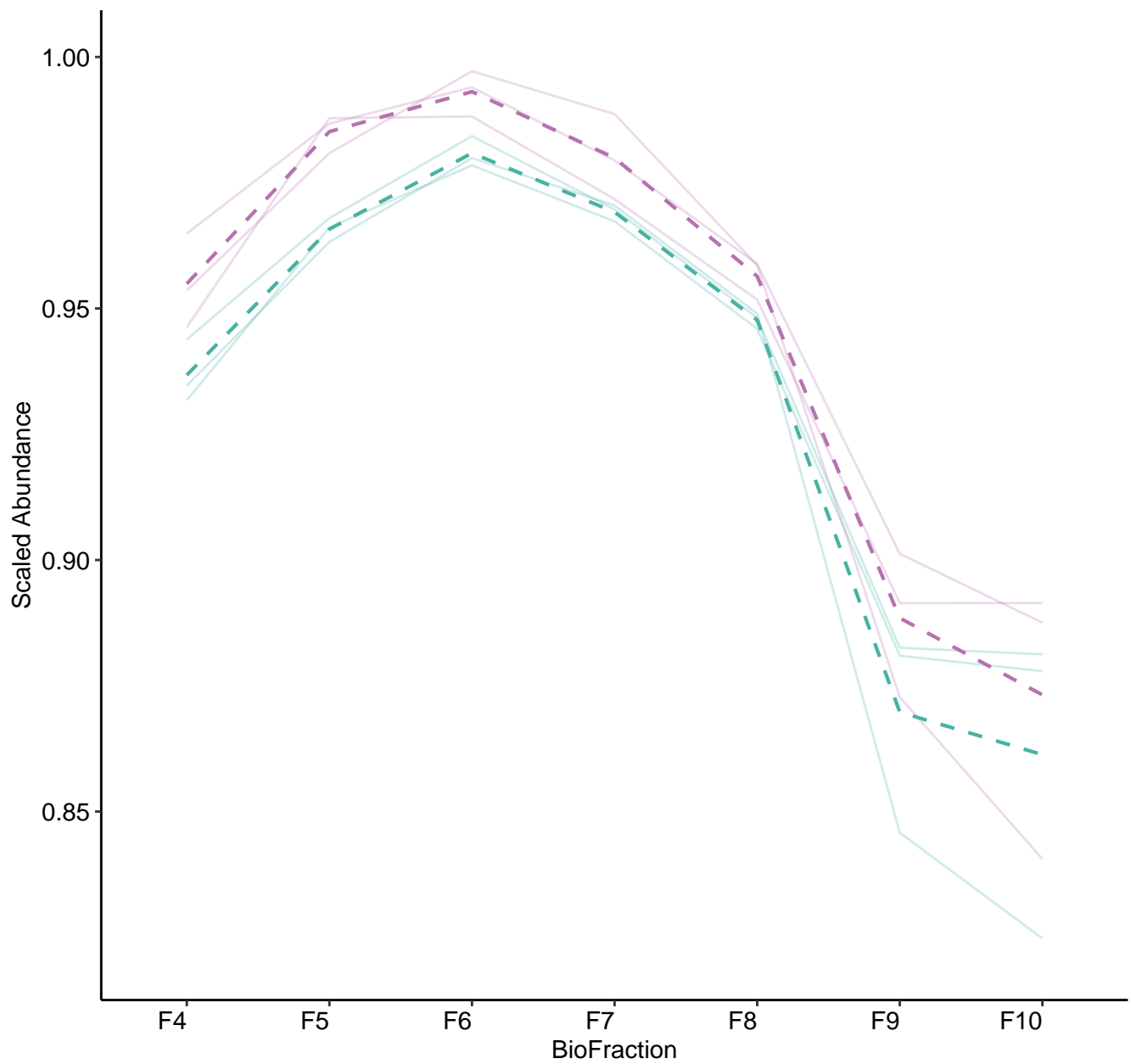
CTLH complex (n = 5)
(R2.Fixef = 0.922)



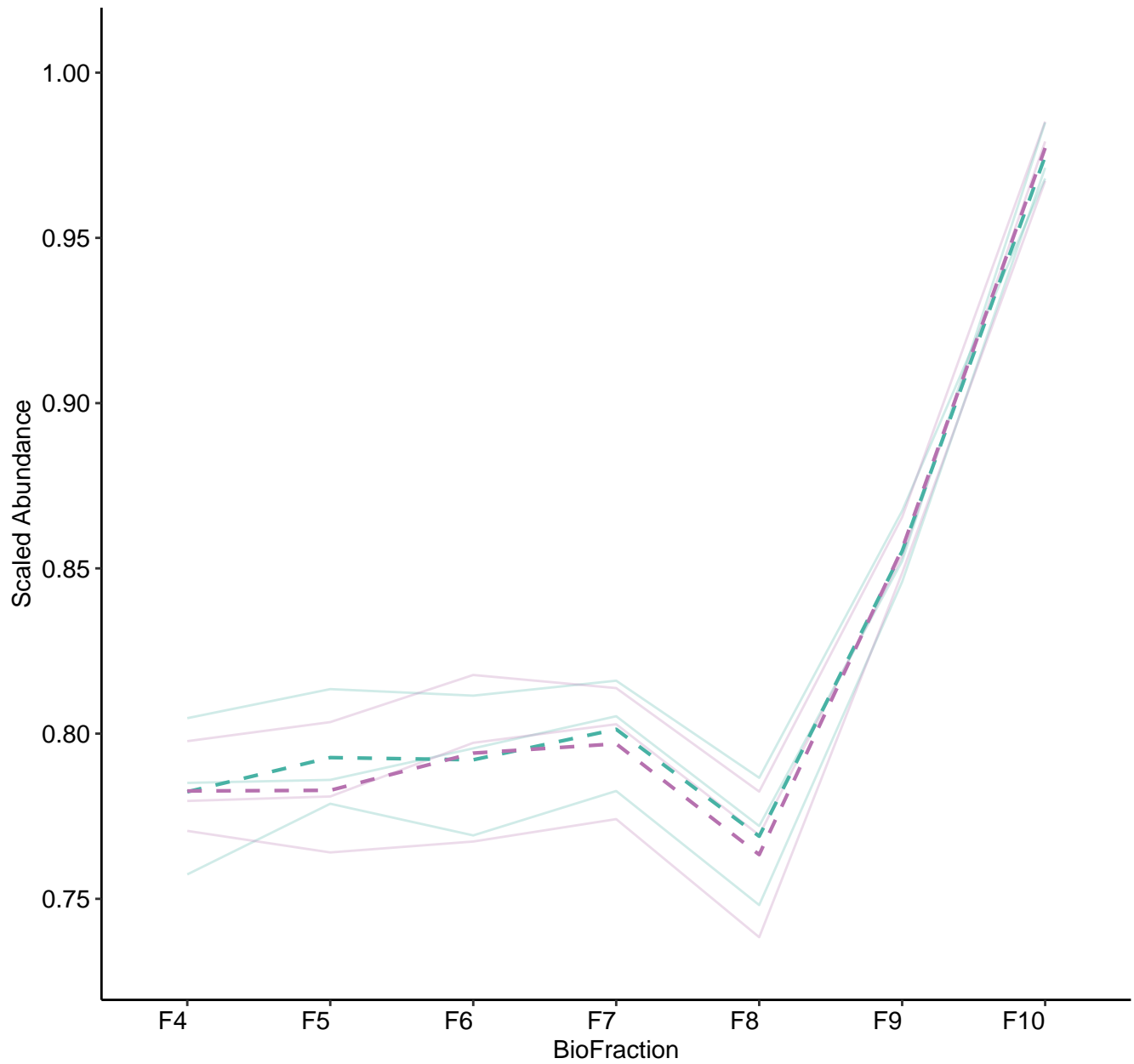
Cytochrome c oxidase, mitochondrial (n = 4)
(R2.Fixef = 0.796)



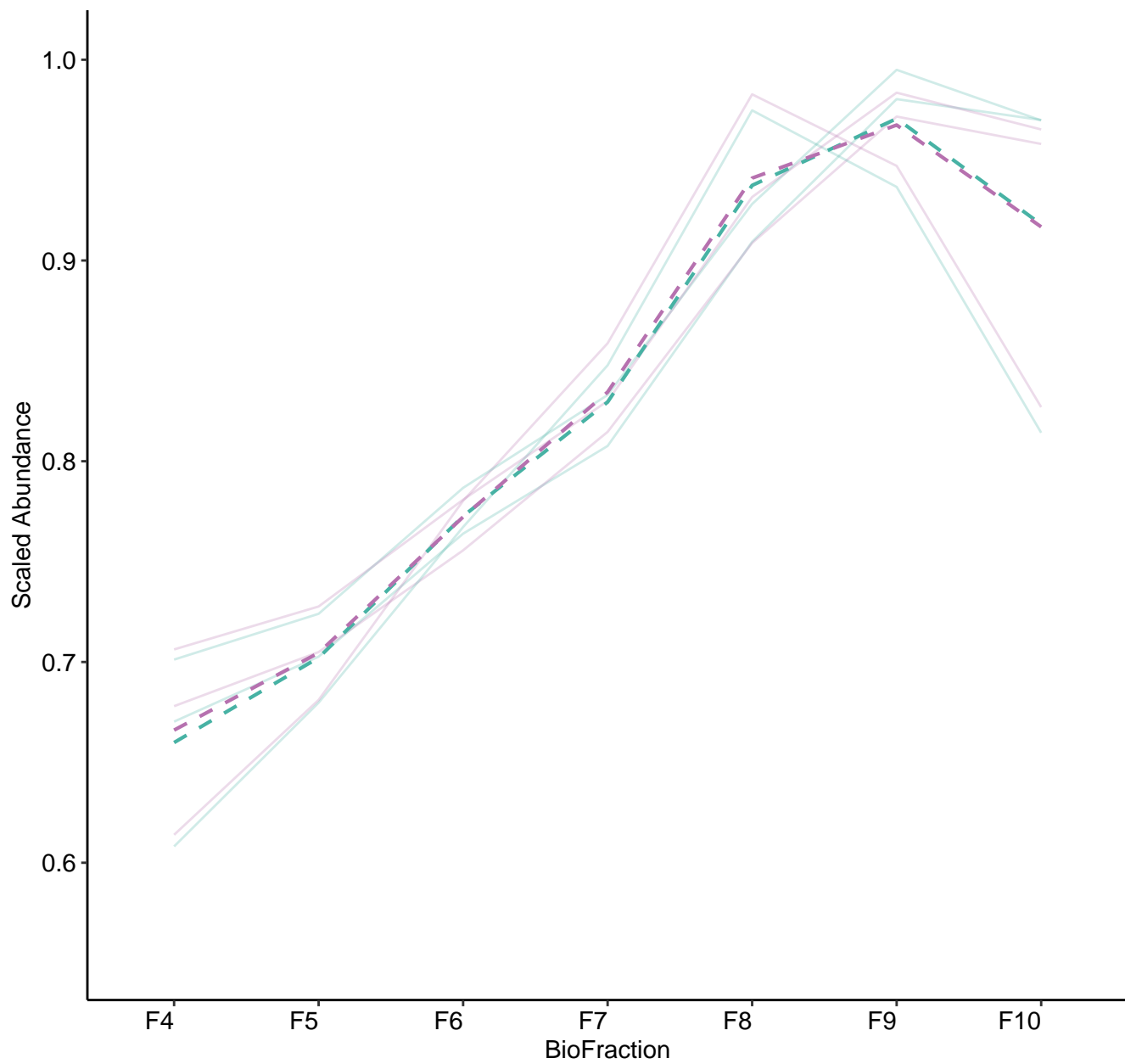
Dystrobrevin–syntrophin complex, brain-derived (n = 3)
(R2.Fixef = 0.915)



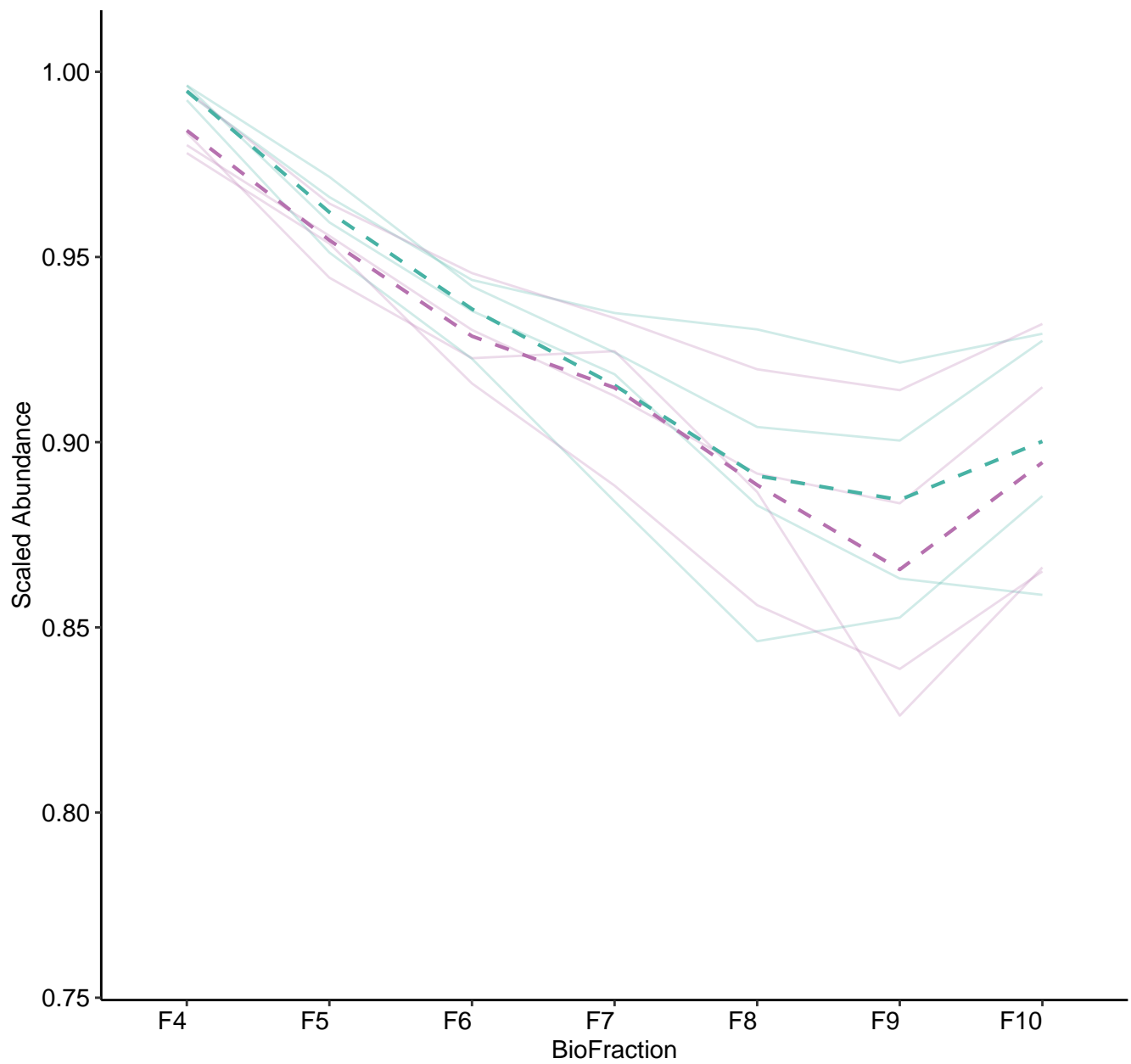
EIF2B1–EIF2B2–EIF2B3–EIF2B4–EIF2B5 complex (n = 3)
(R2.Fixef = 0.935)



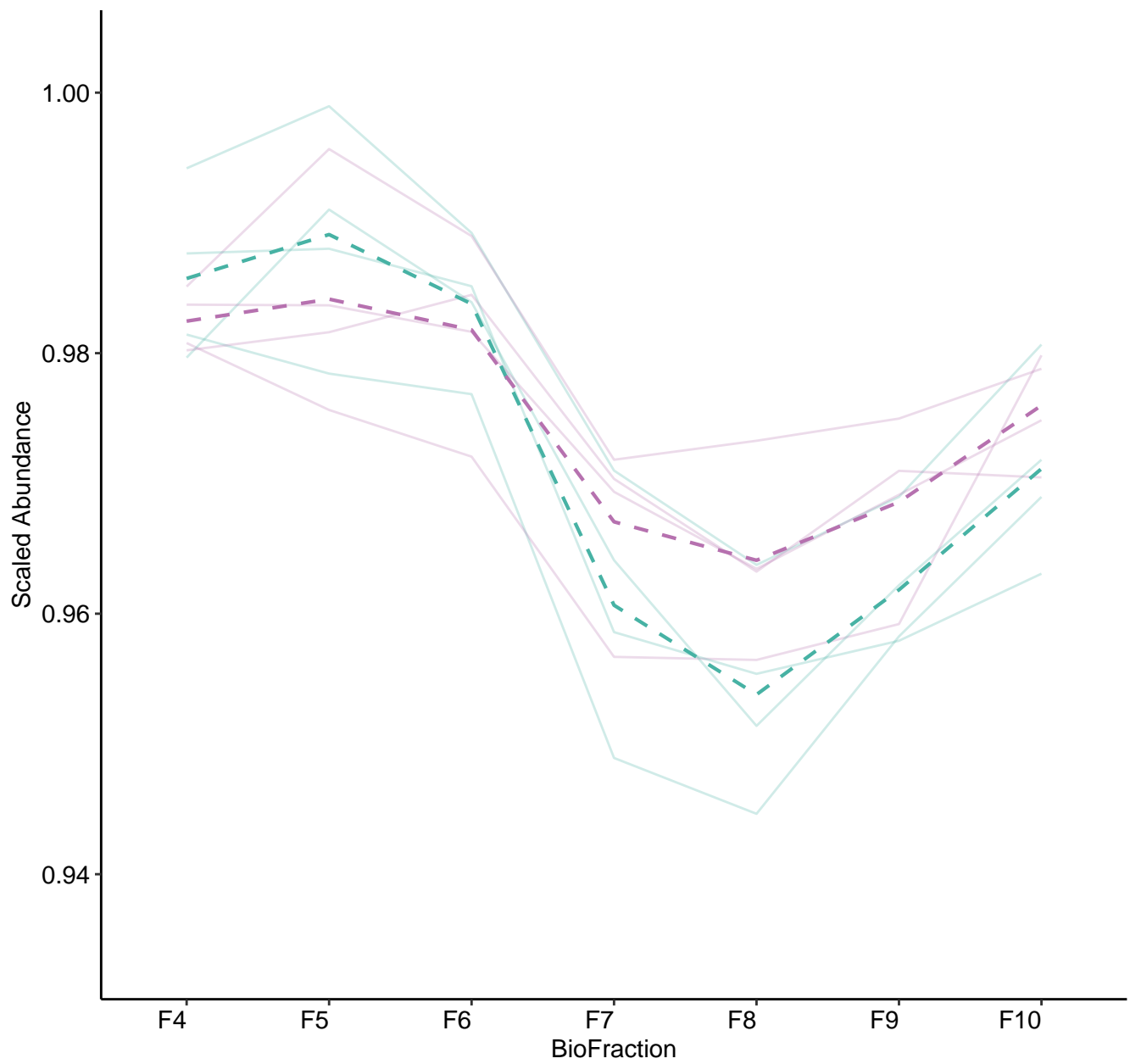
Emerin complex 52 (n = 3)
(R2.Fixef = 0.877)



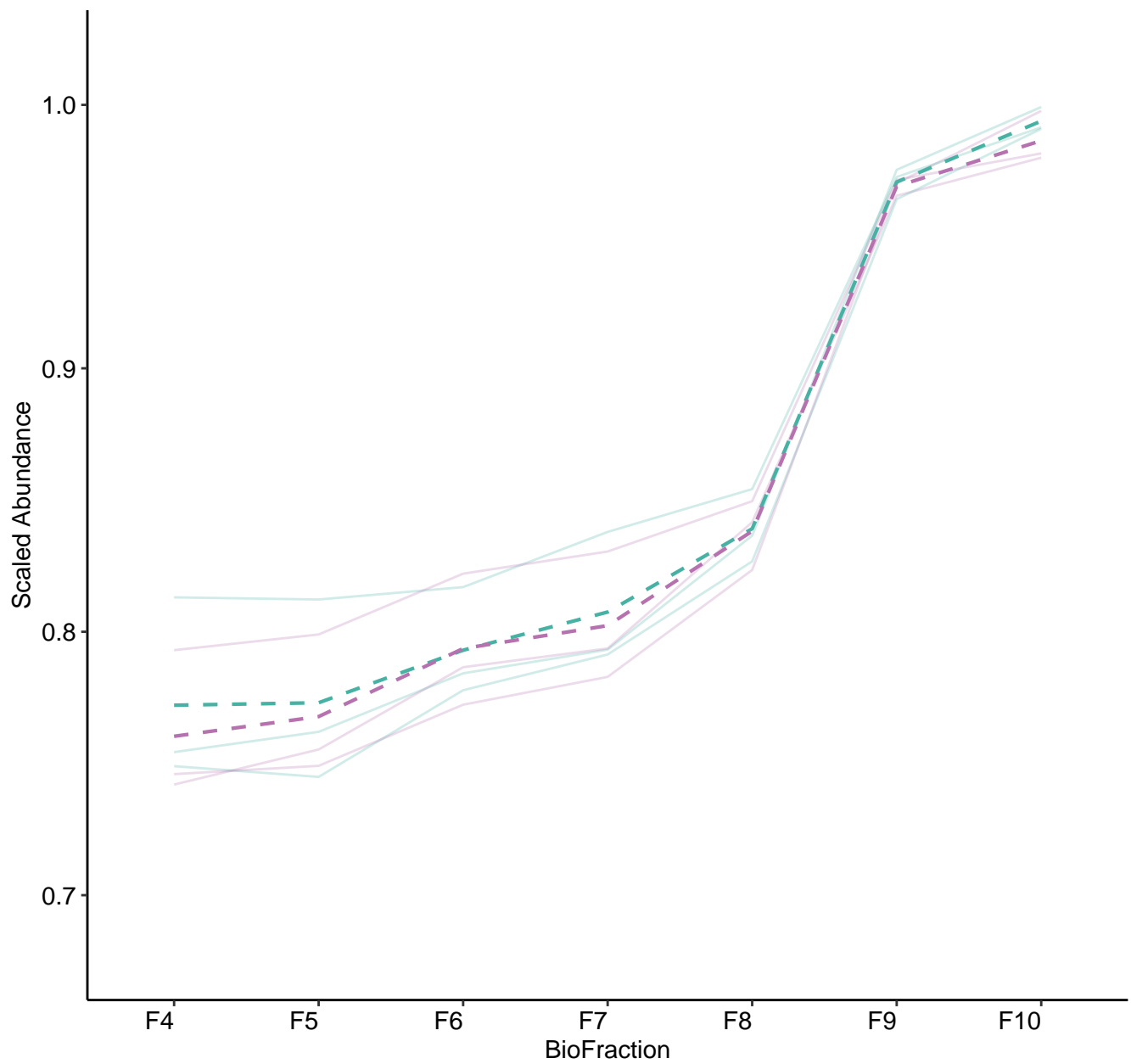
Endocytic coat complex (11 subunits) (n = 4)
(R2.Fixef = 0.712)



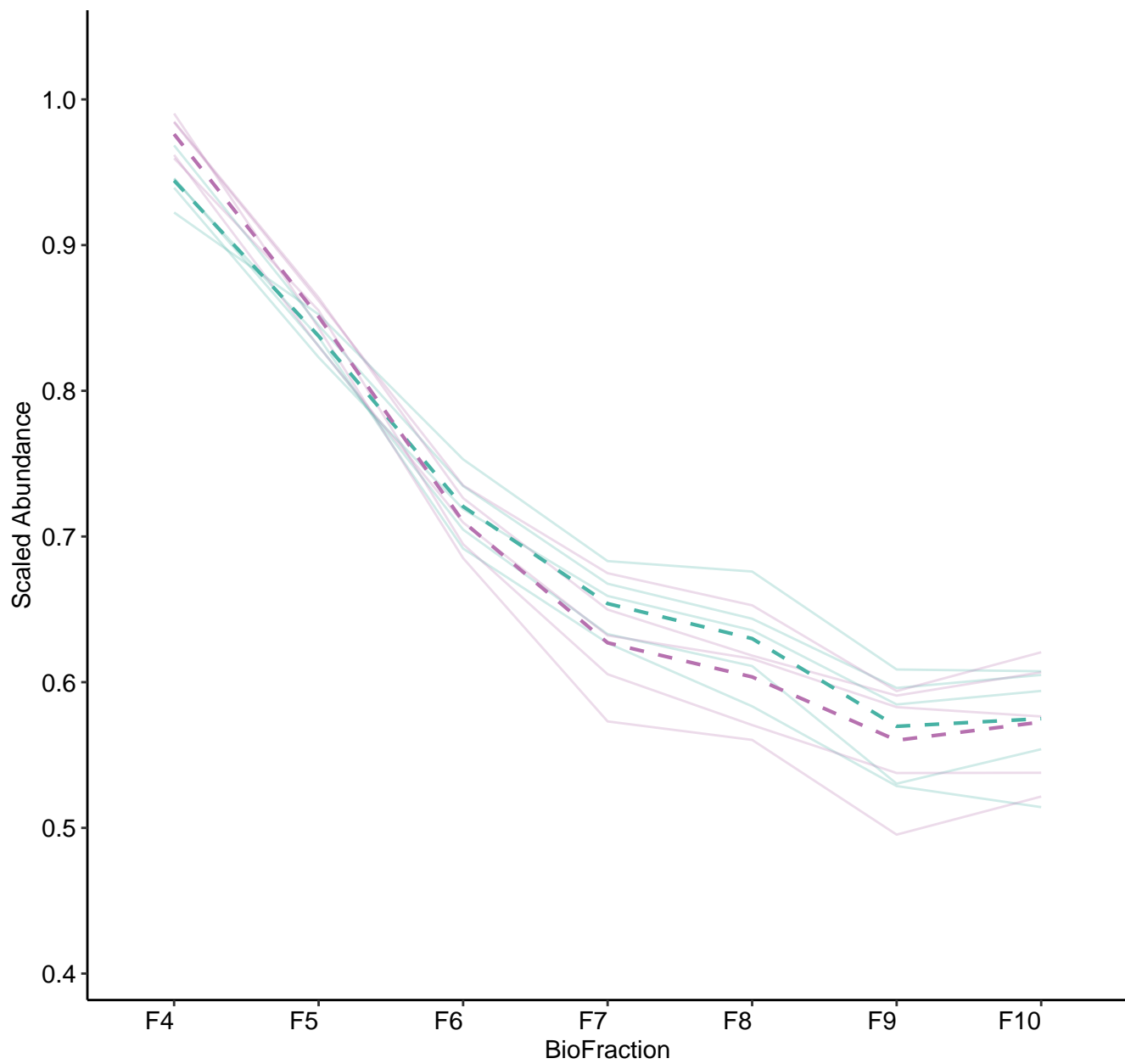
Exocyst complex (n = 4)
(R2.Fixef = 0.714)



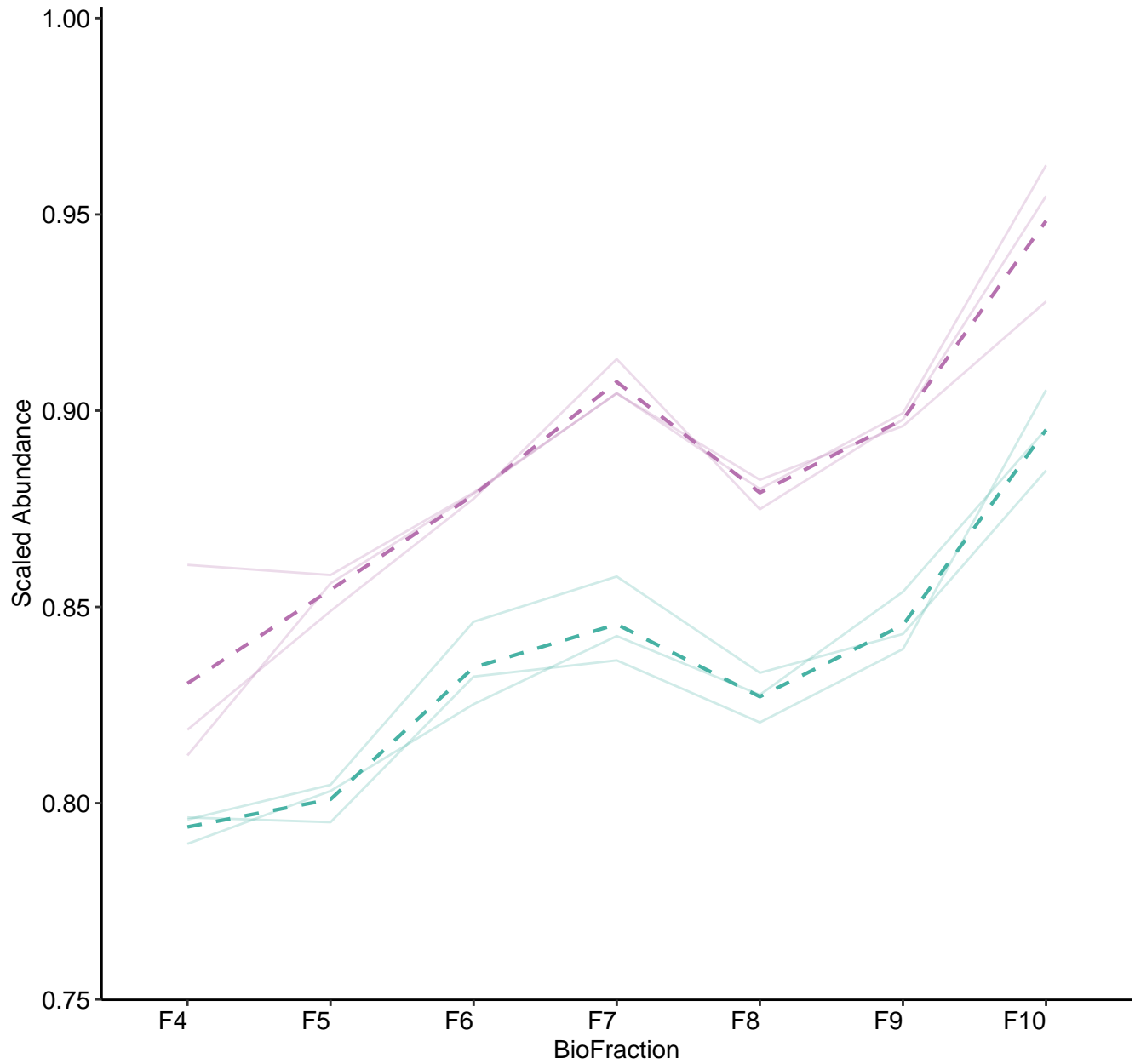
Exosome (n = 3)
(R2.Fixef = 0.939)



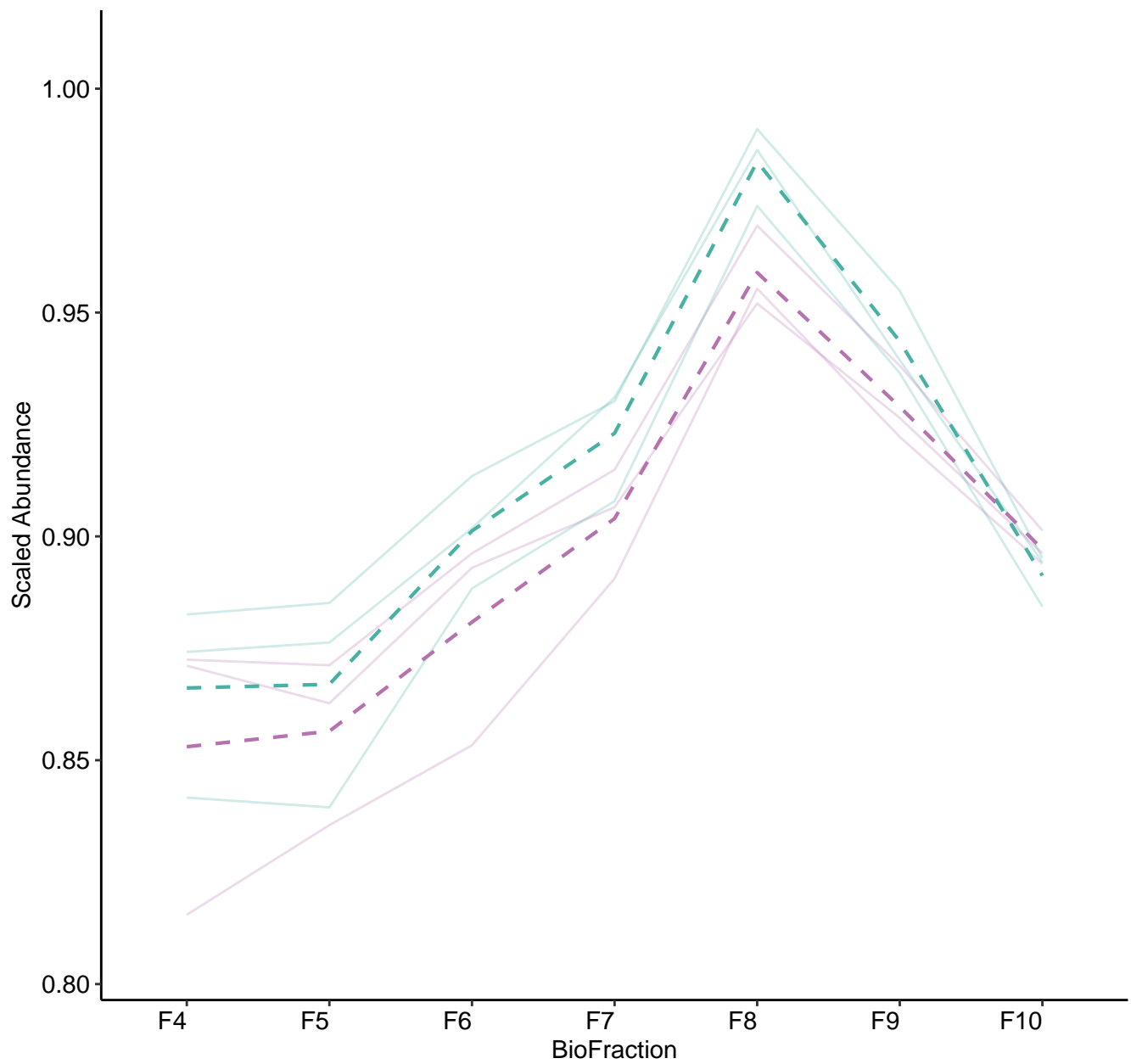
F0F1 ATP synthase, mitochondrial (n = 5)
(R2.Fixef = 0.953)



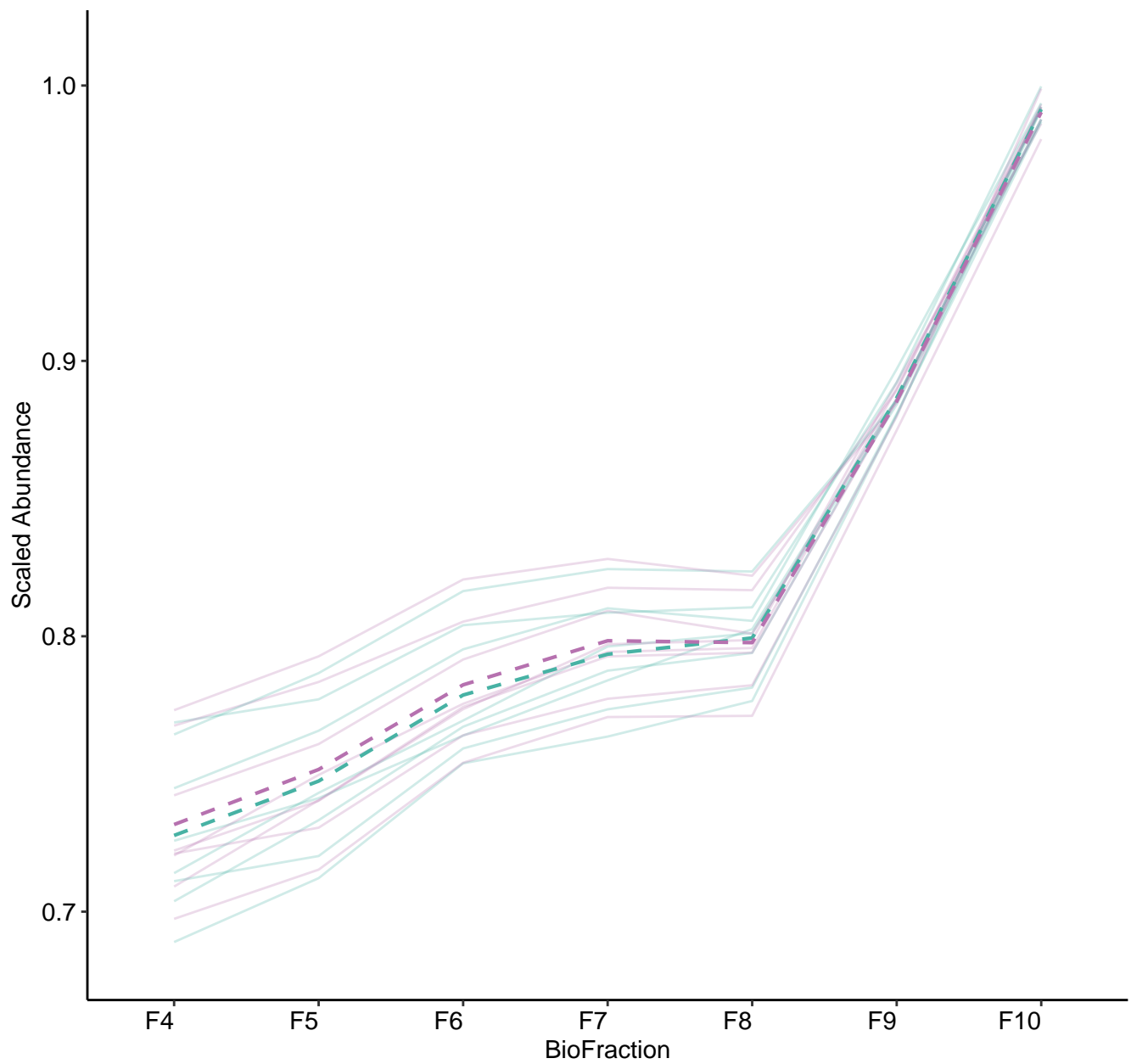
Fibrinogen complex (n = 3)
(R2.Fixef = 0.941)



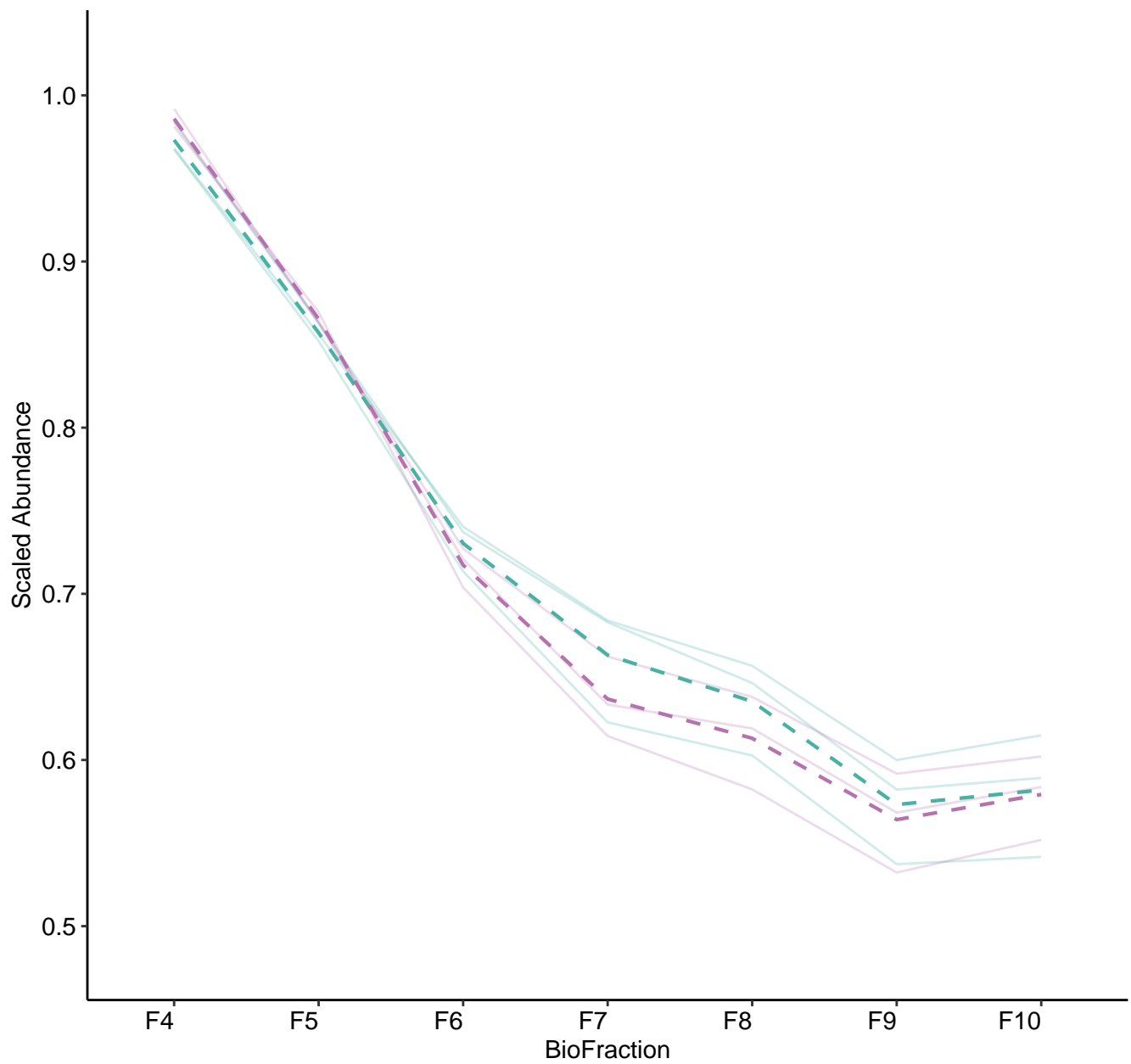
Gamma-BAR-AP1 complex (n = 3)
(R2.Fixef = 0.843)



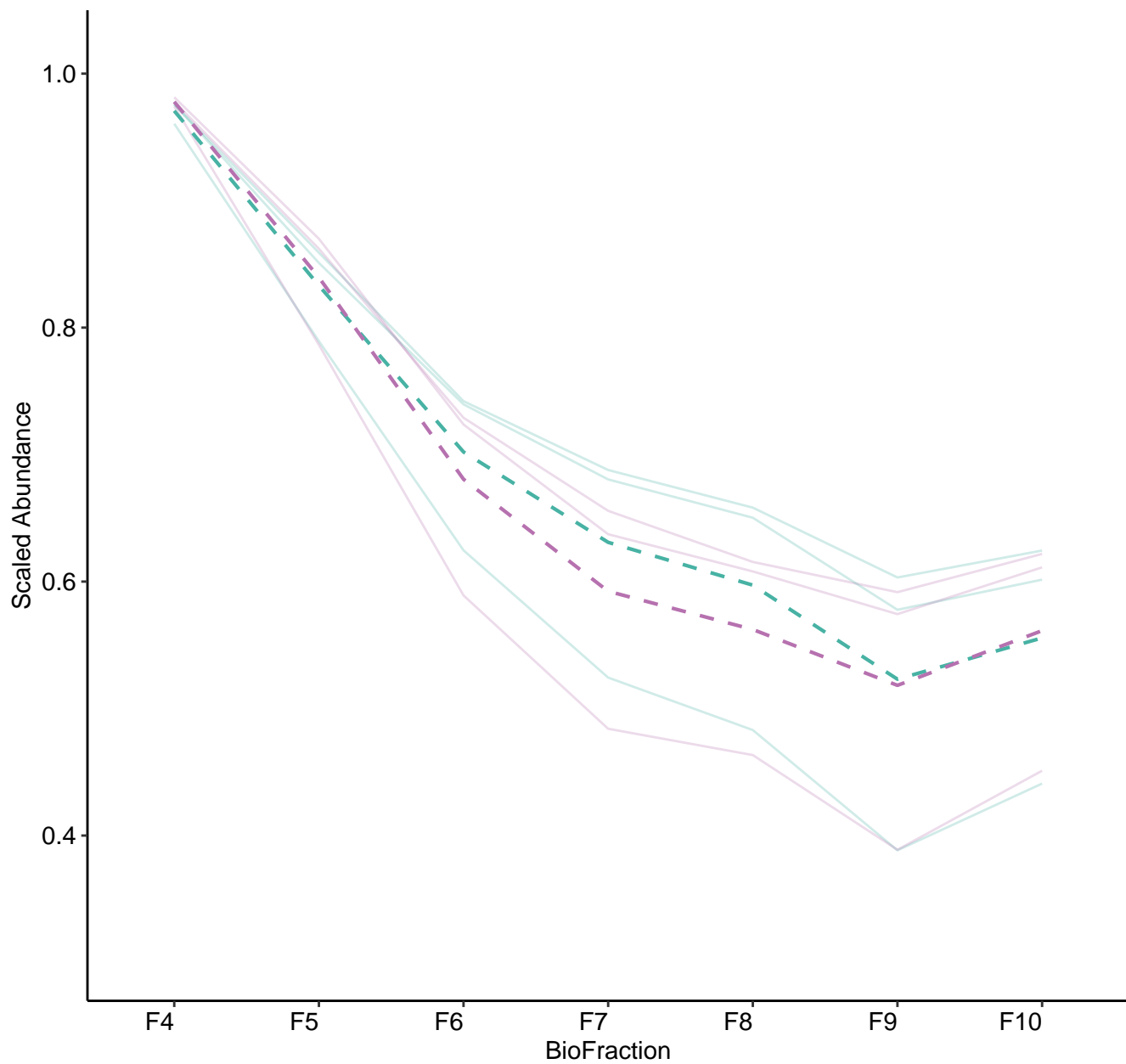
Multisynthetase complex (n = 8)
(R2.Fixef = 0.948)



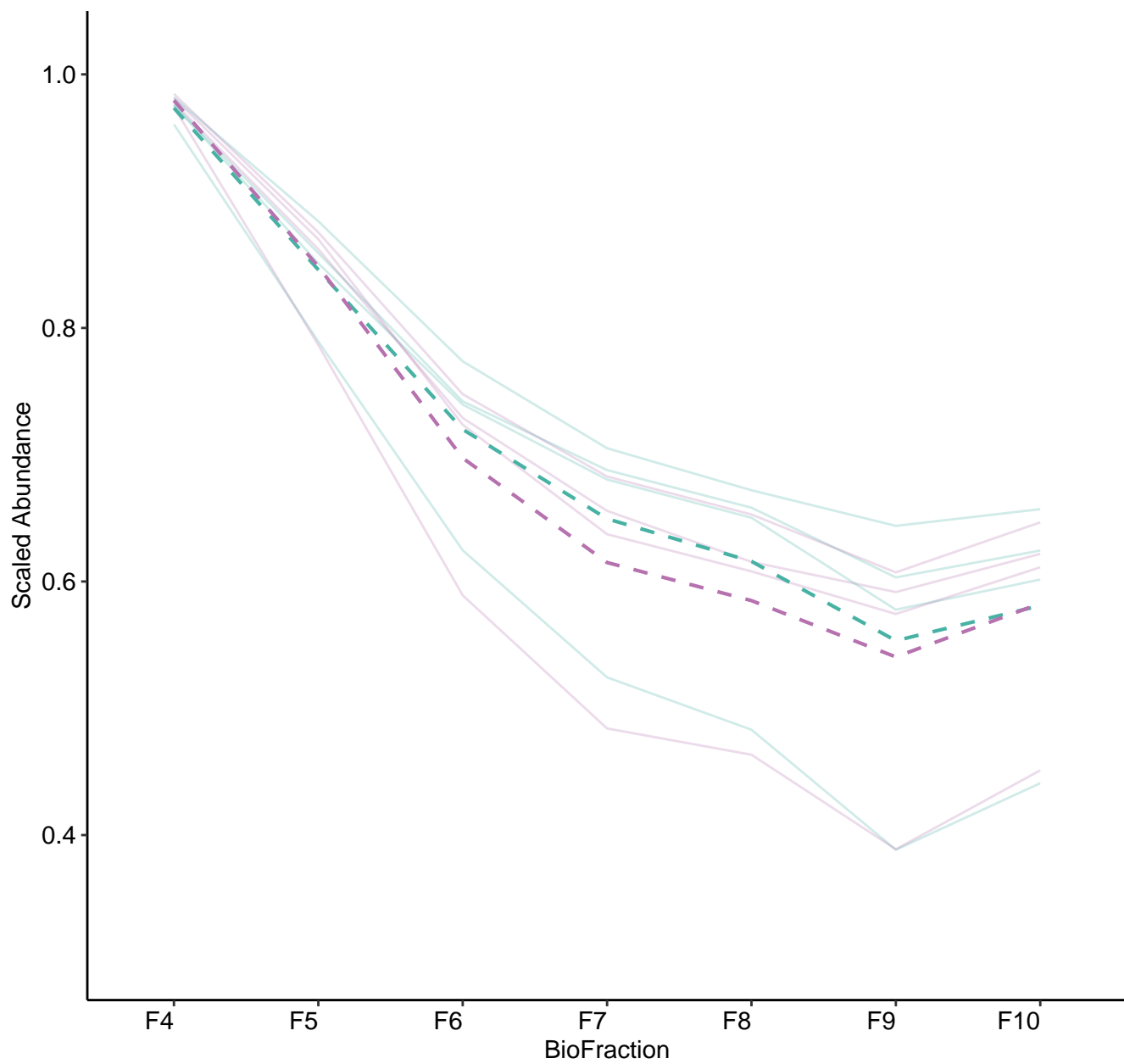
Respiratory chain complex I (early intermediate NDUF AF1 assembly), mitochondrial (n = 3)
(R2.Fixef = 0.974)



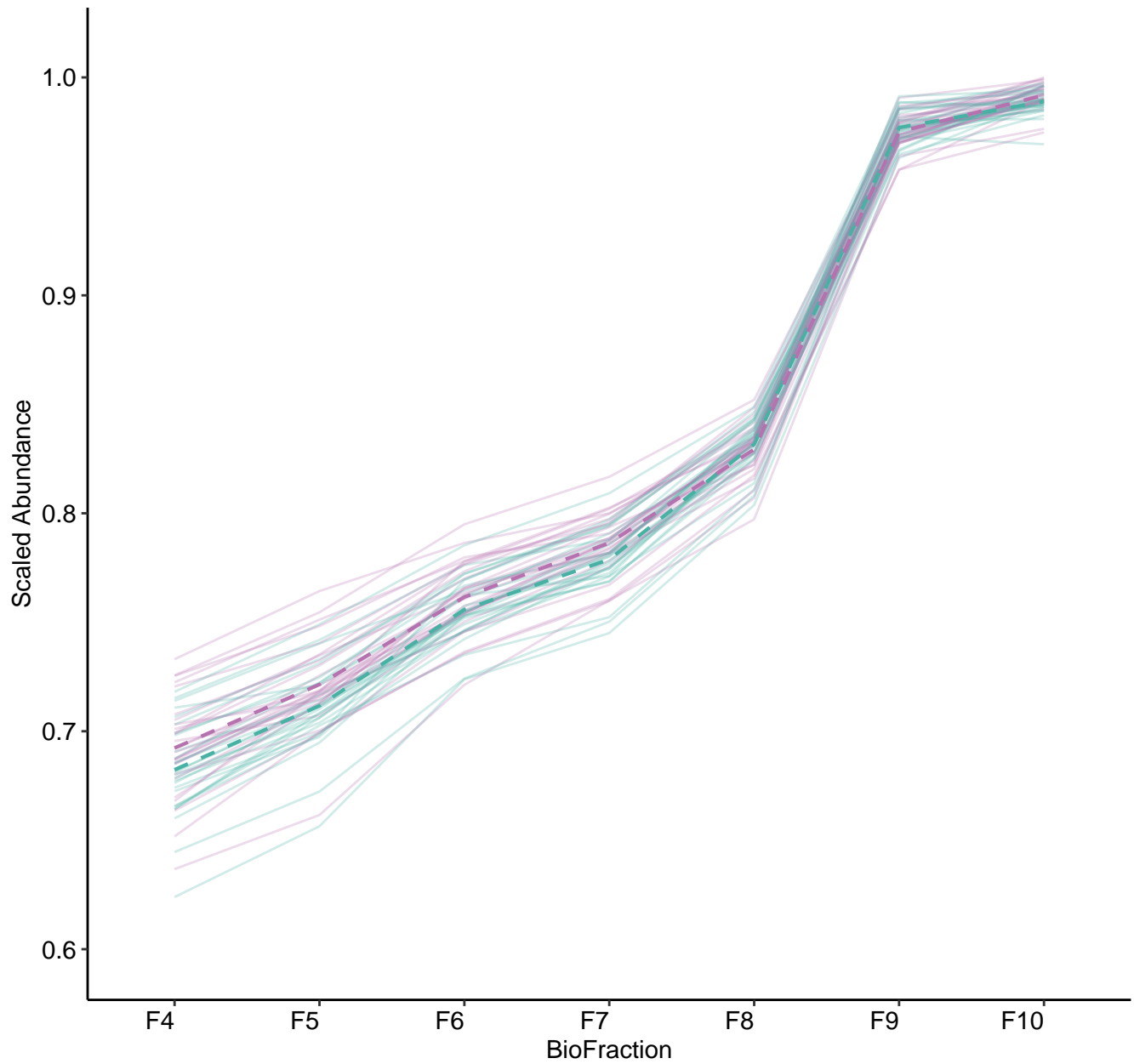
Respiratory chain complex I (intermediate V/380kD and VI/480kD), mitochondrial (n = 3)
(R2.Fixef = 0.783)



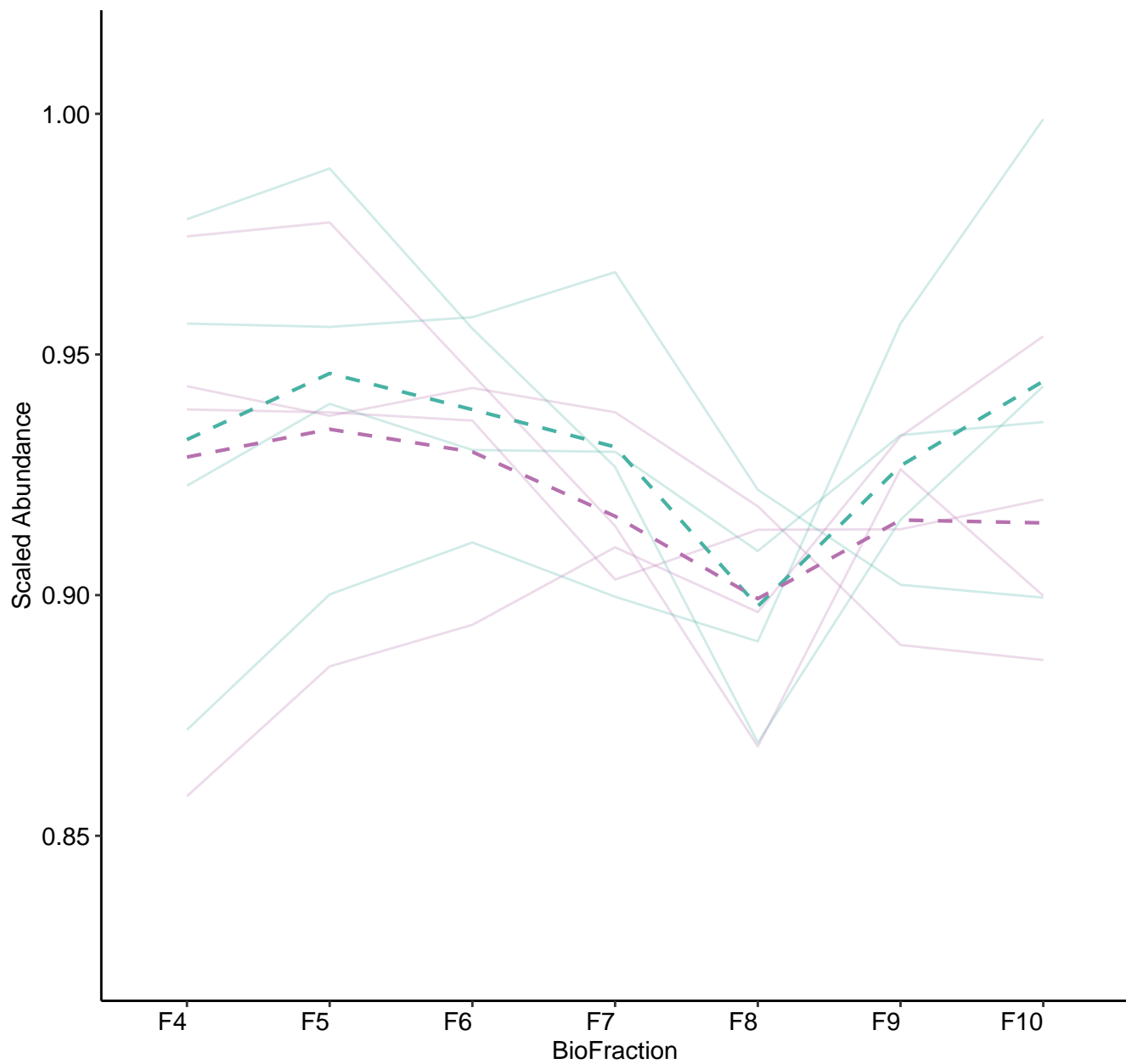
Respiratory chain complex I (intermediate VII/650kD), mitochondrial (n = 4)
(R2.Fixef = 0.787)



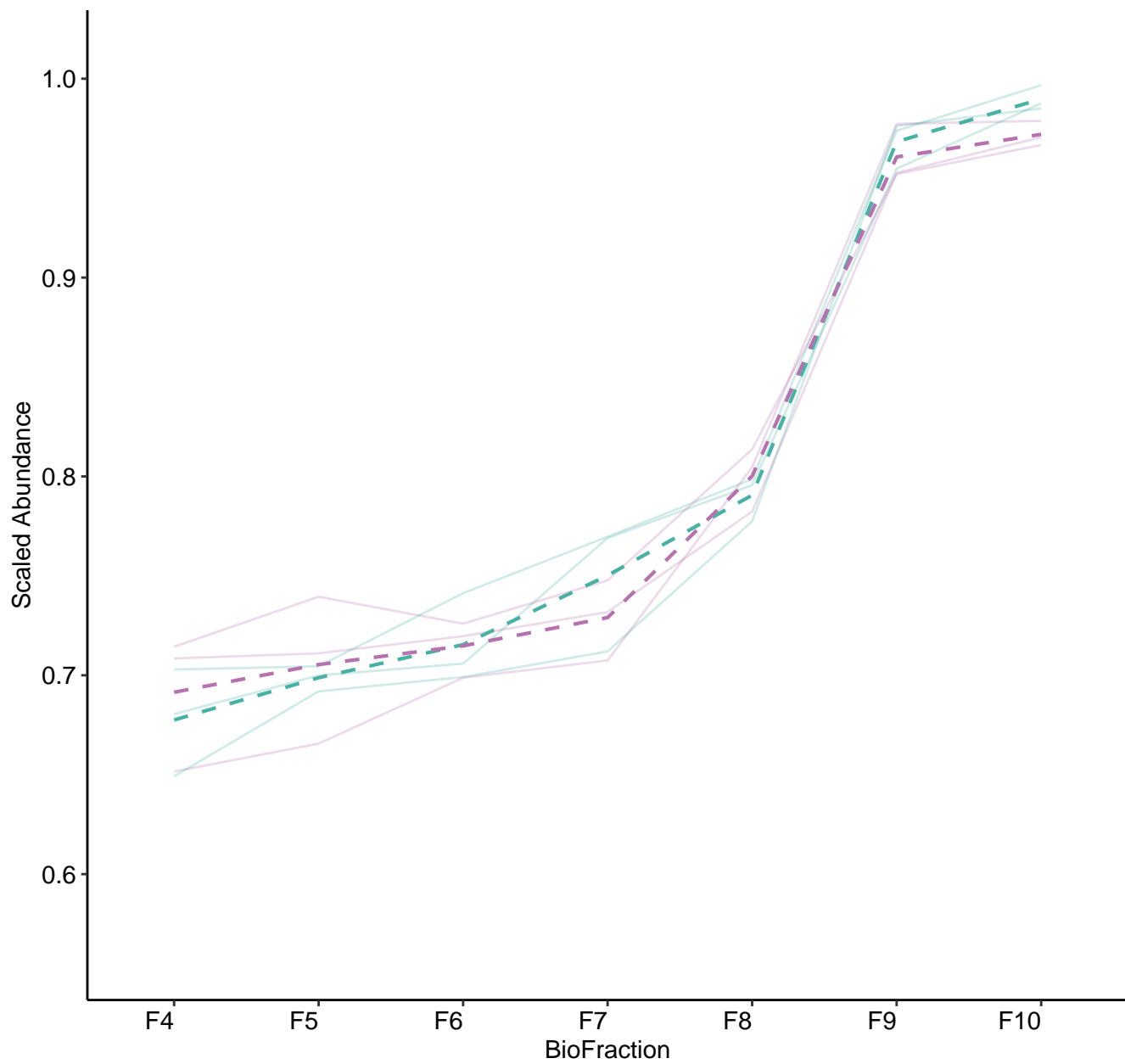
Ribosome, cytoplasmic (n = 27)
(R2.Fixef = 0.982)



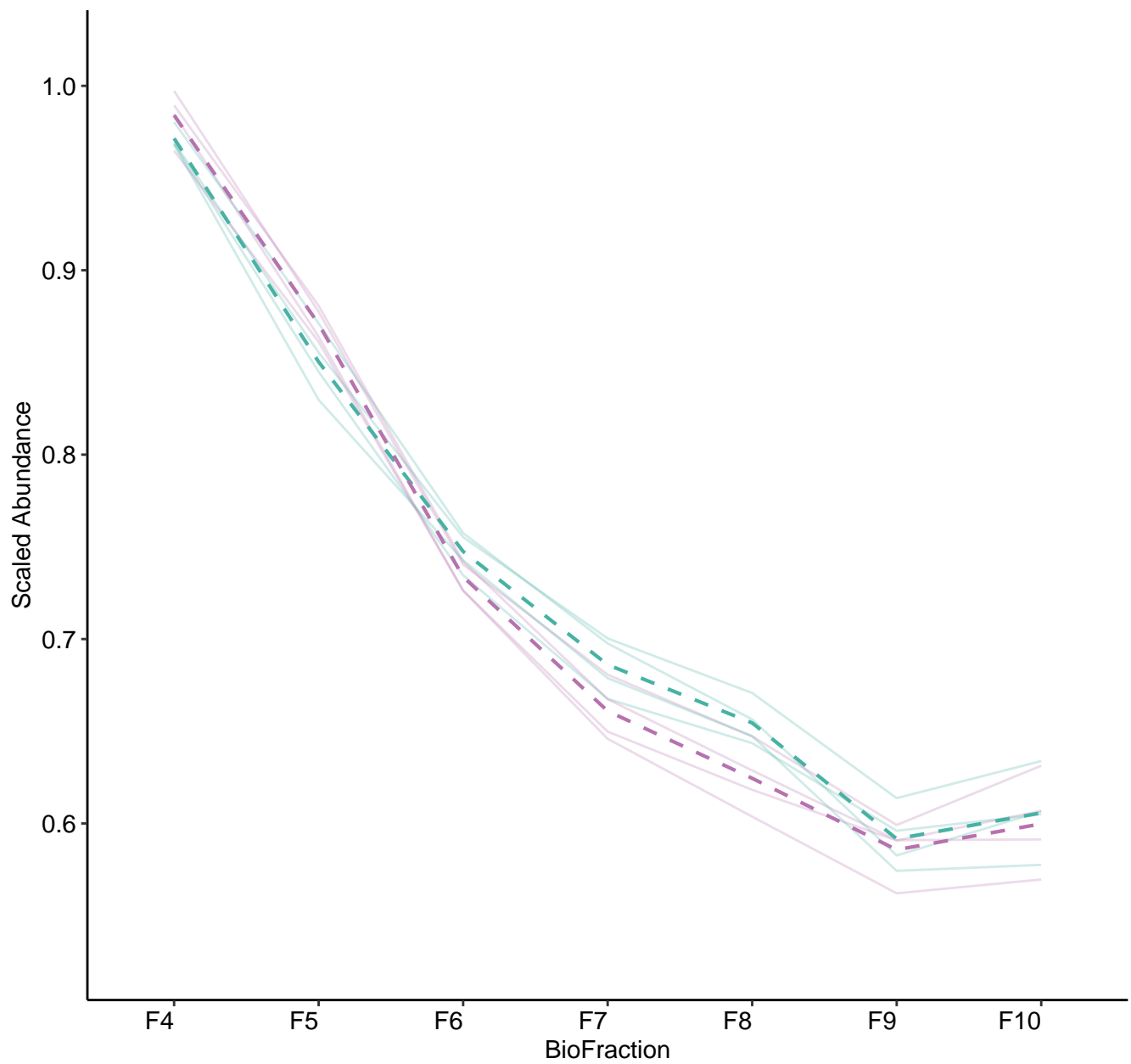
RICH1/AMOT polarity complex, Flag-Rich1 precipitated (n = 4)
(R2.Fixef = 0.174)



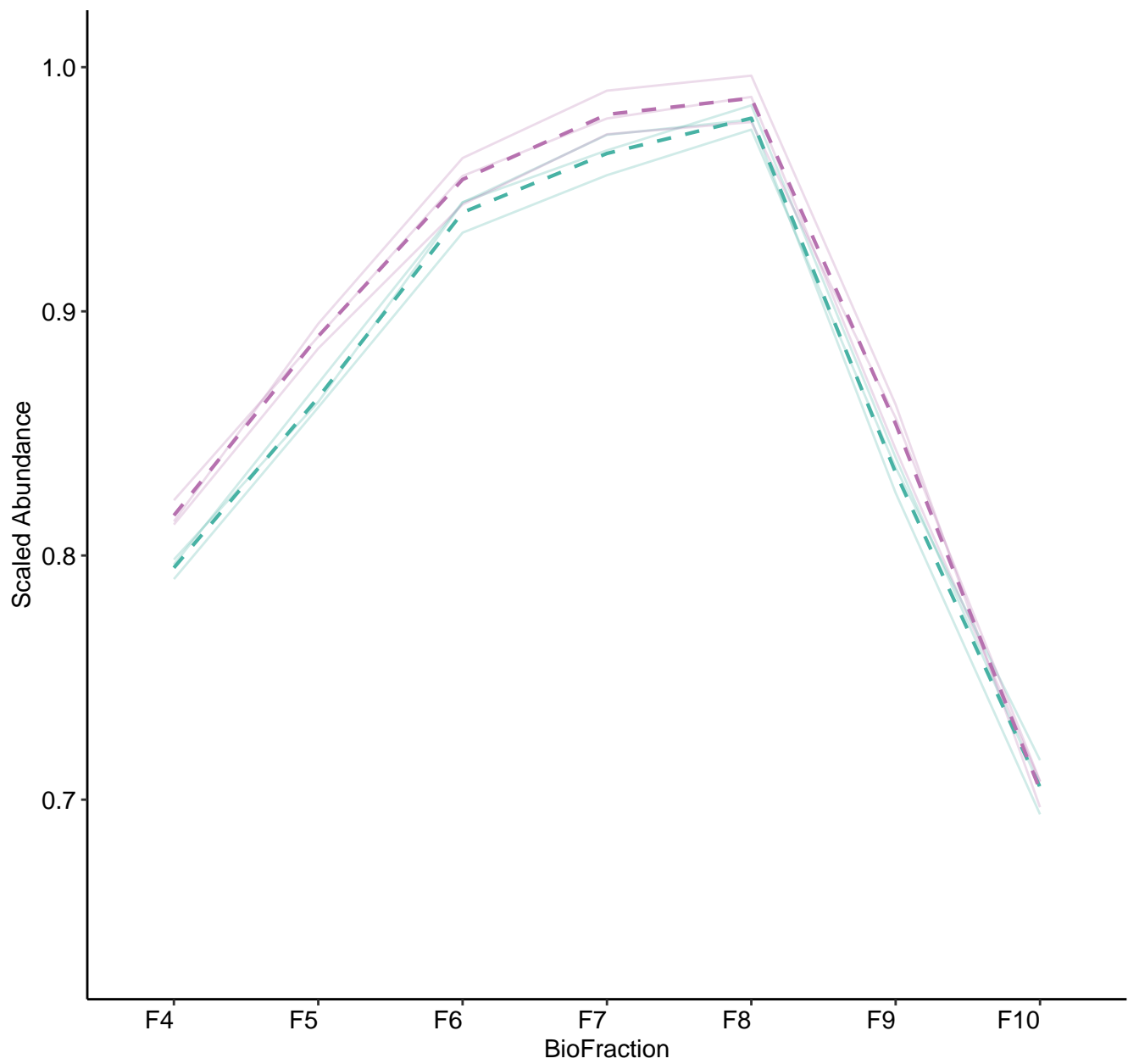
RNA polymerase II complex, chromatin structure modifying (n = 3)
(R2.Fixef = 0.968)



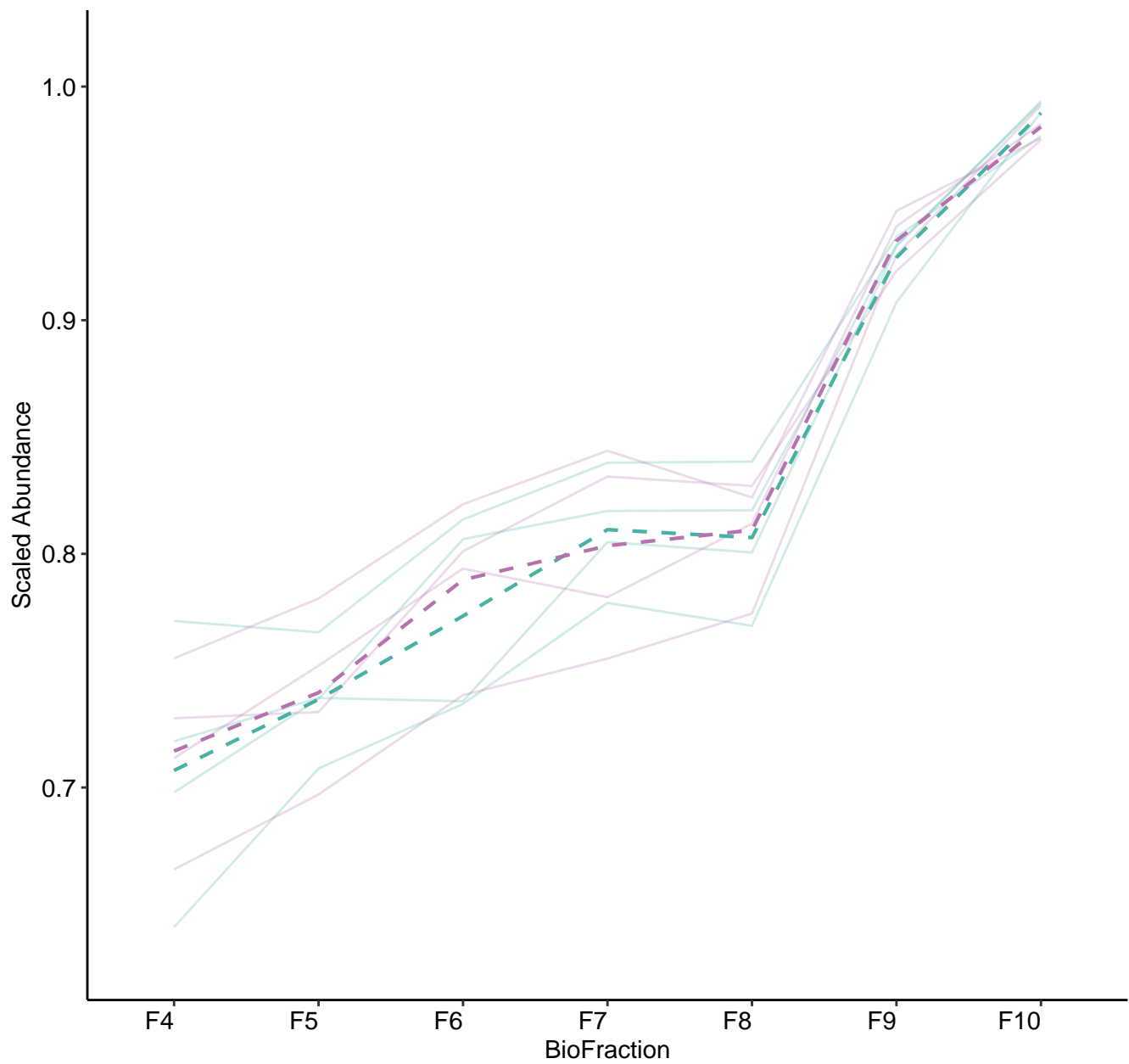
SDH-mABC1-PIC-ANT-ATPase complex (n = 4)
(R2.Fixef = 0.986)



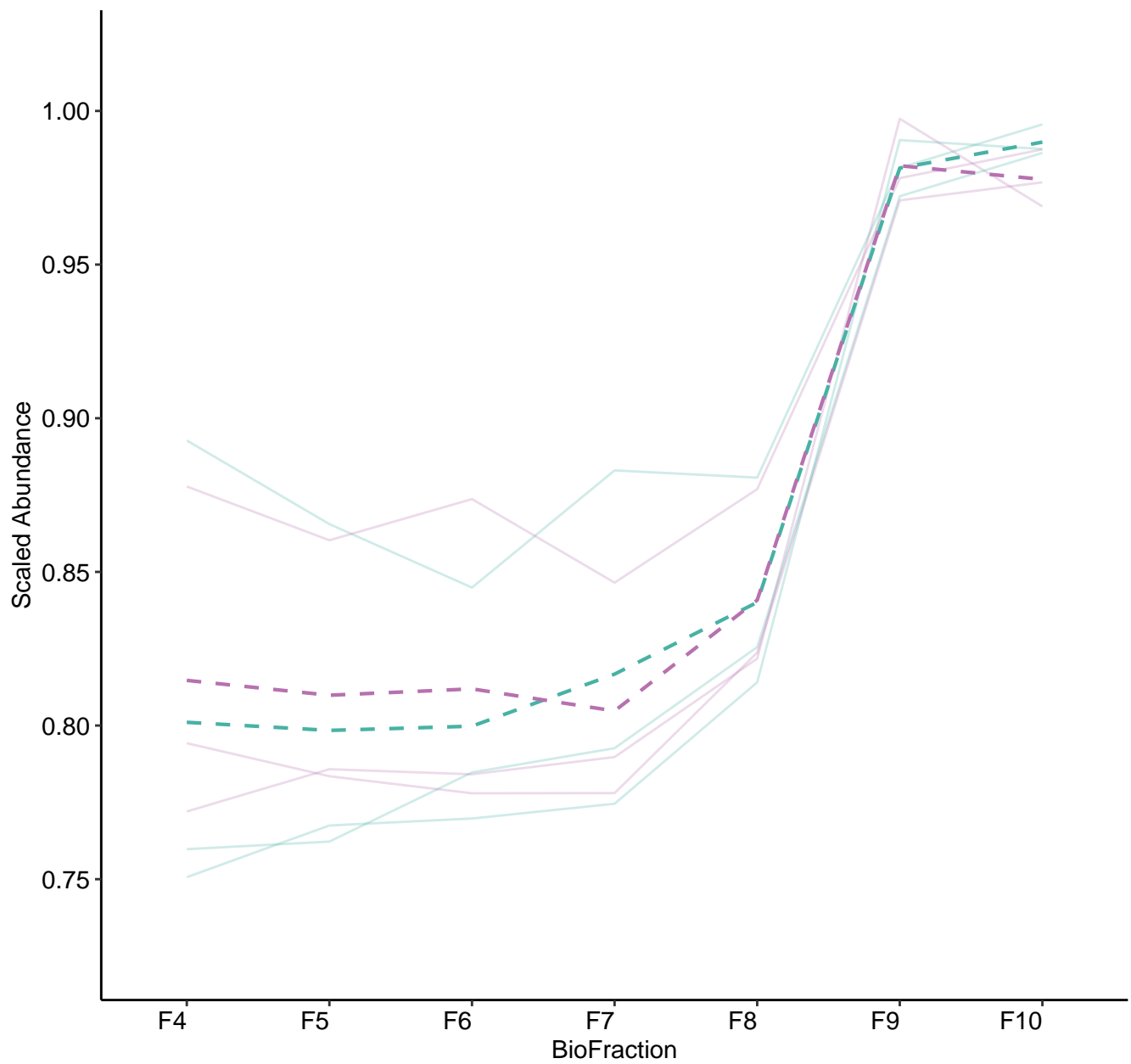
SELK multiprotein complex (n = 3)
(R2.Fixef = 0.994)



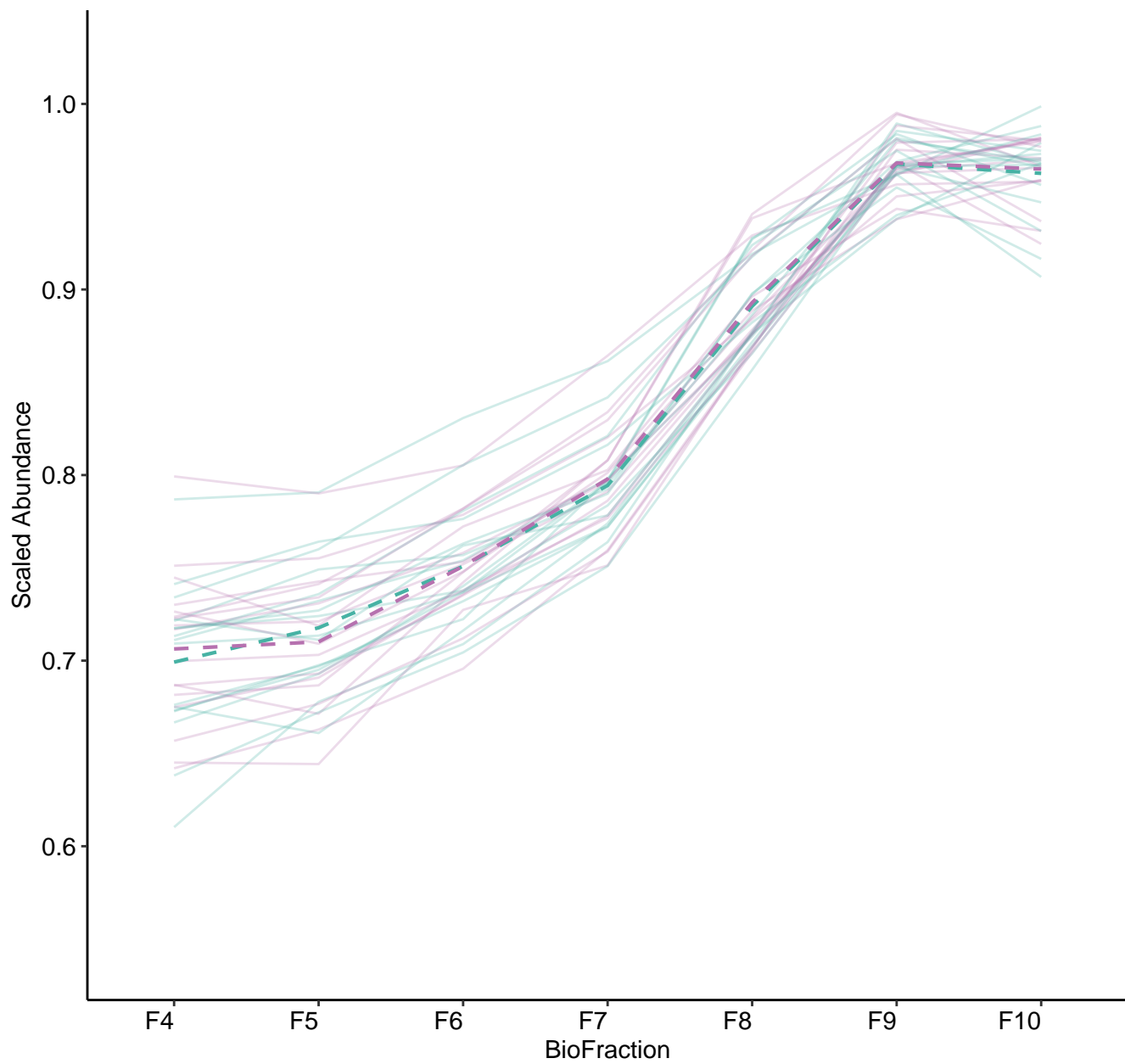
SNF2h-cohesin-NuRD complex (n = 4)
(R2.Fixef = 0.9)



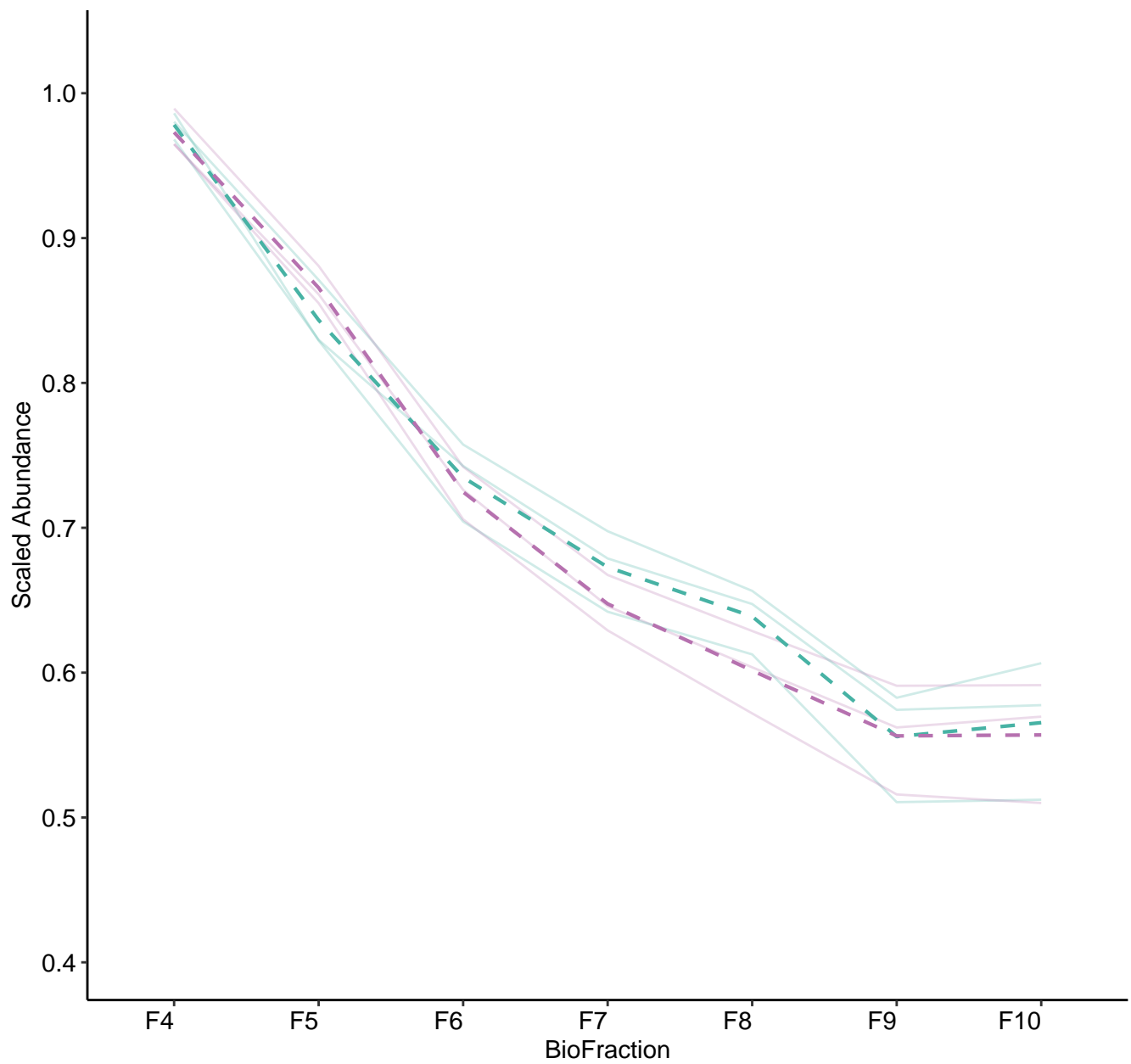
SOG complex (n = 3)
(R2.Fixef = 0.764)



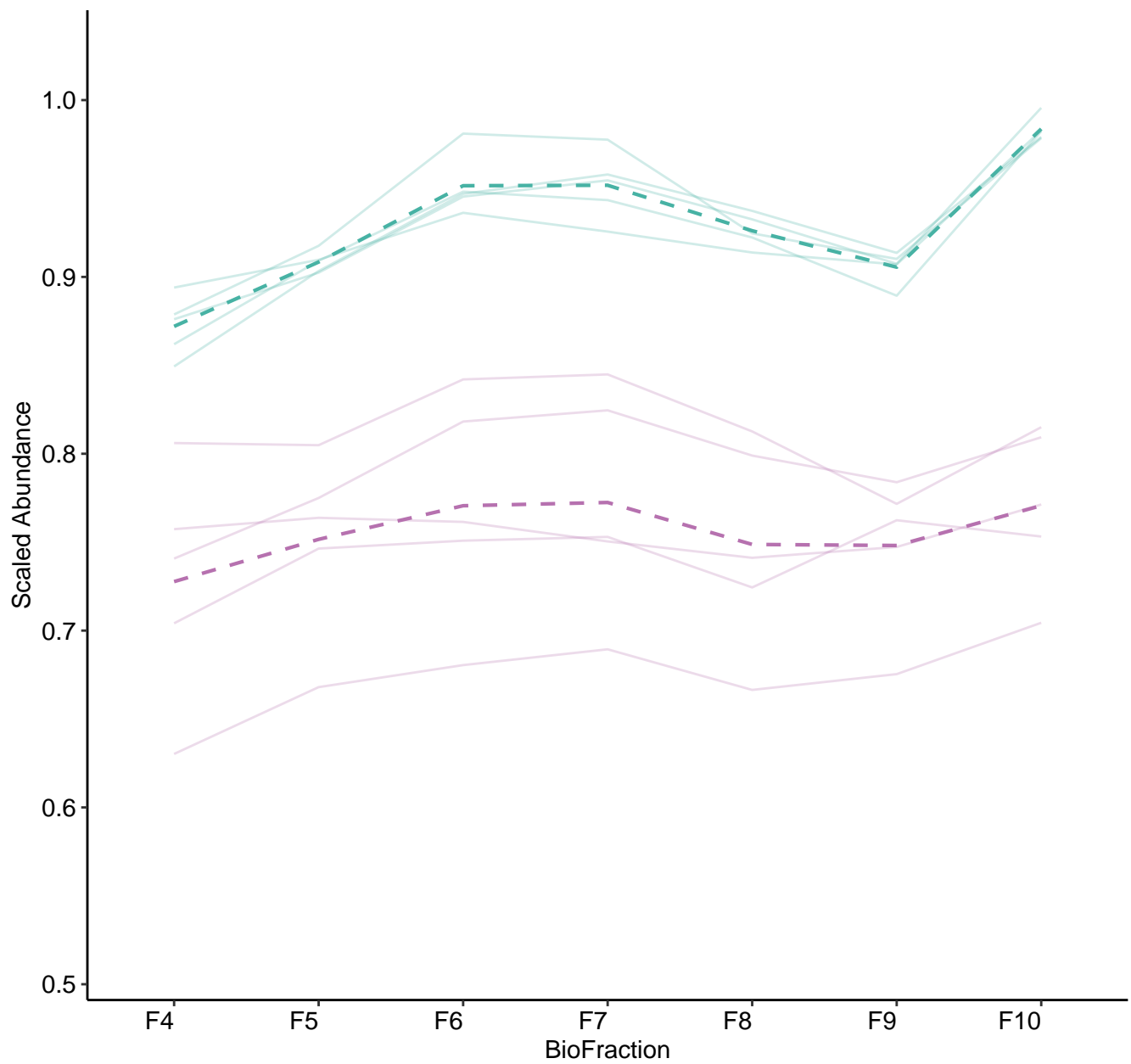
Spliceosome (n = 17)
(R2.Fixef = 0.927)



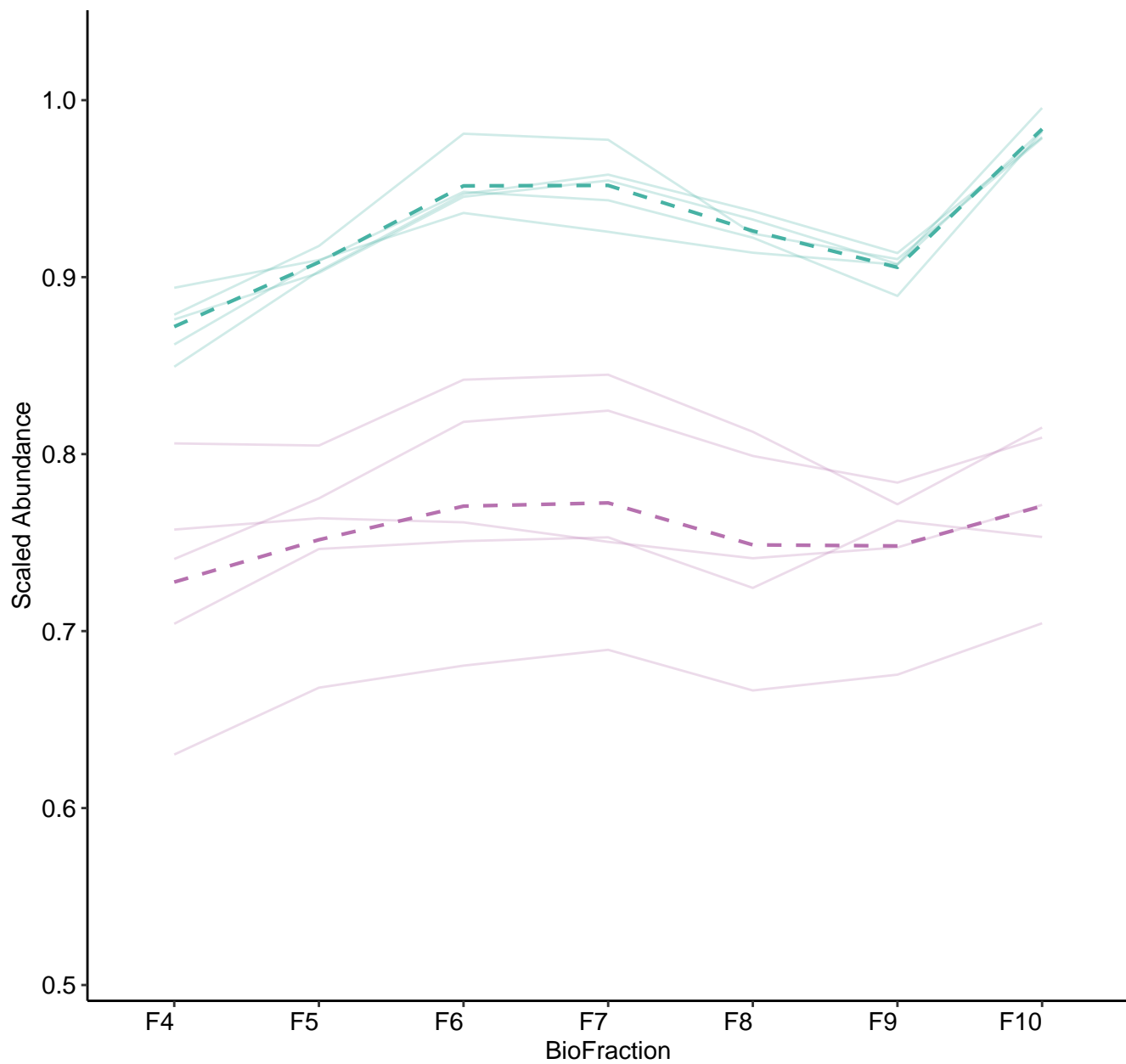
Succinate dehydrogenase complex II, mitochondrial (n = 3)
(R2.Fixef = 0.963)



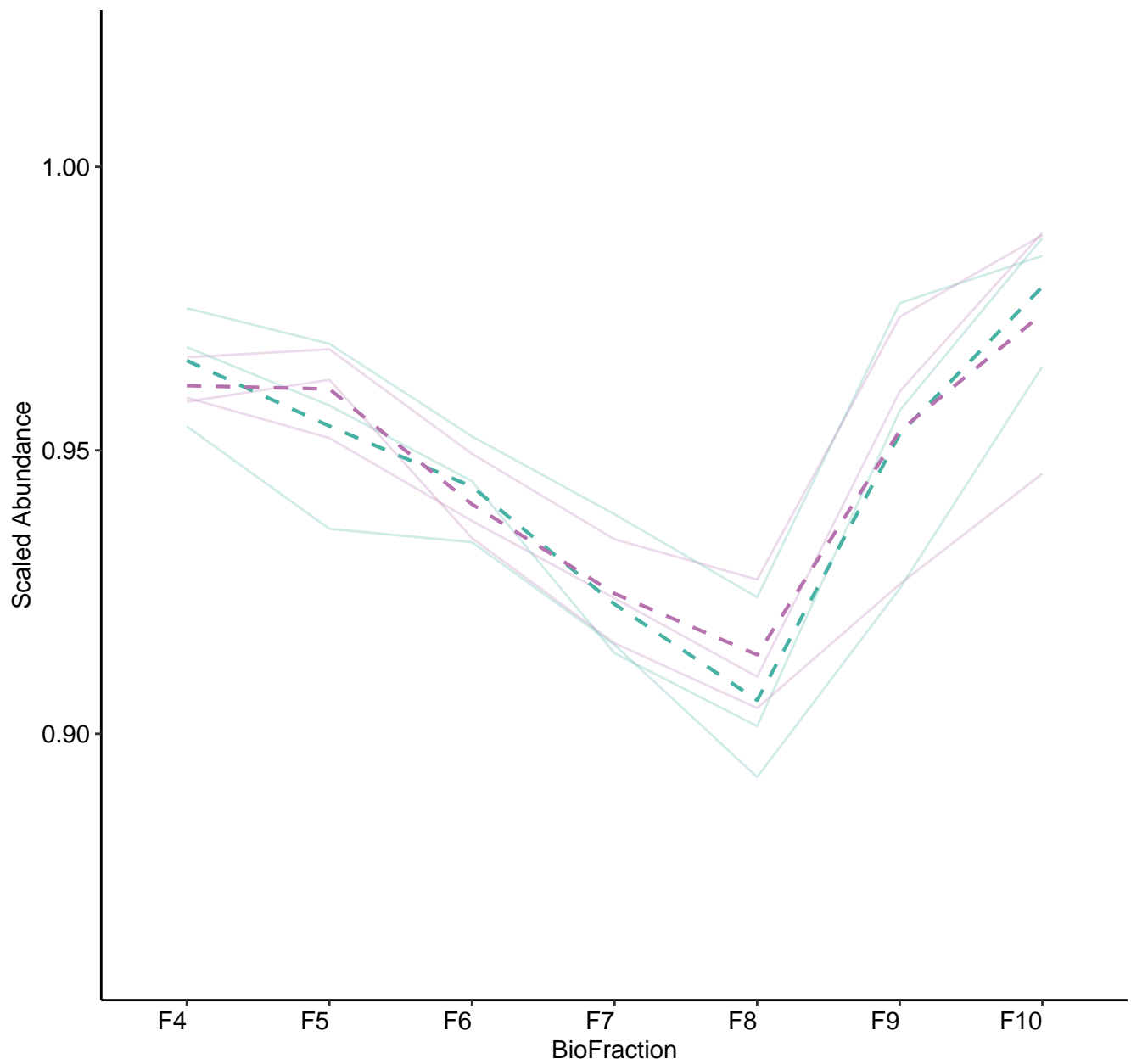
WASH complex (n = 5)
(R2.Fixef = 0.832)



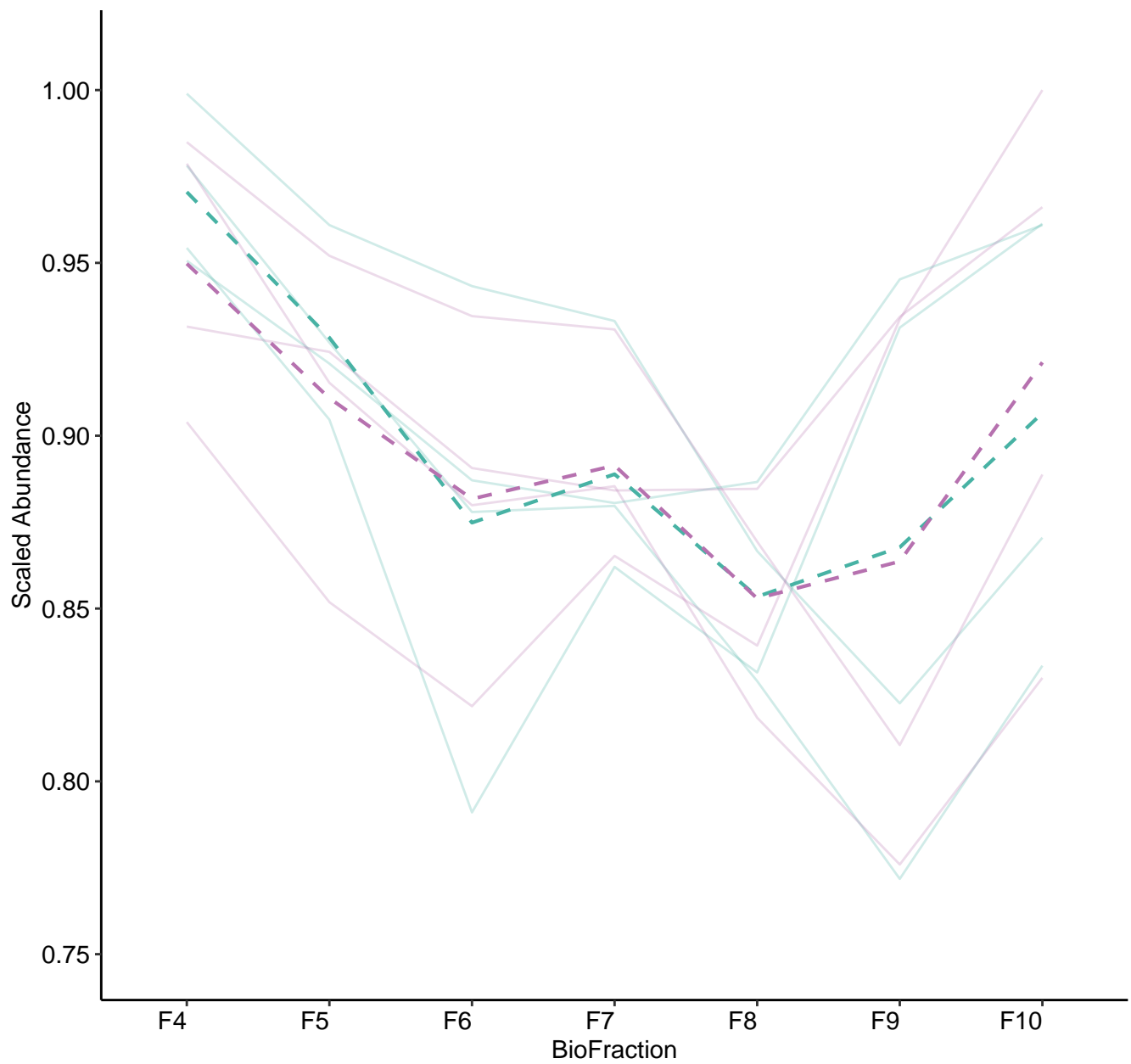
WASH-CAPZ α /beta complex (n = 5)
(R2.Fixef = 0.832)



Wave-2 complex (n = 3)
(R2.Fixef = 0.671)



WDR41-(C9orf72-SMCR8)-(FIP200-ULK1-ATG13-ATG101) complex (n = 4)
(R2.Fixef = 0.31)



RAVE (n = 3)
(R2.Fixef = 0.387)

