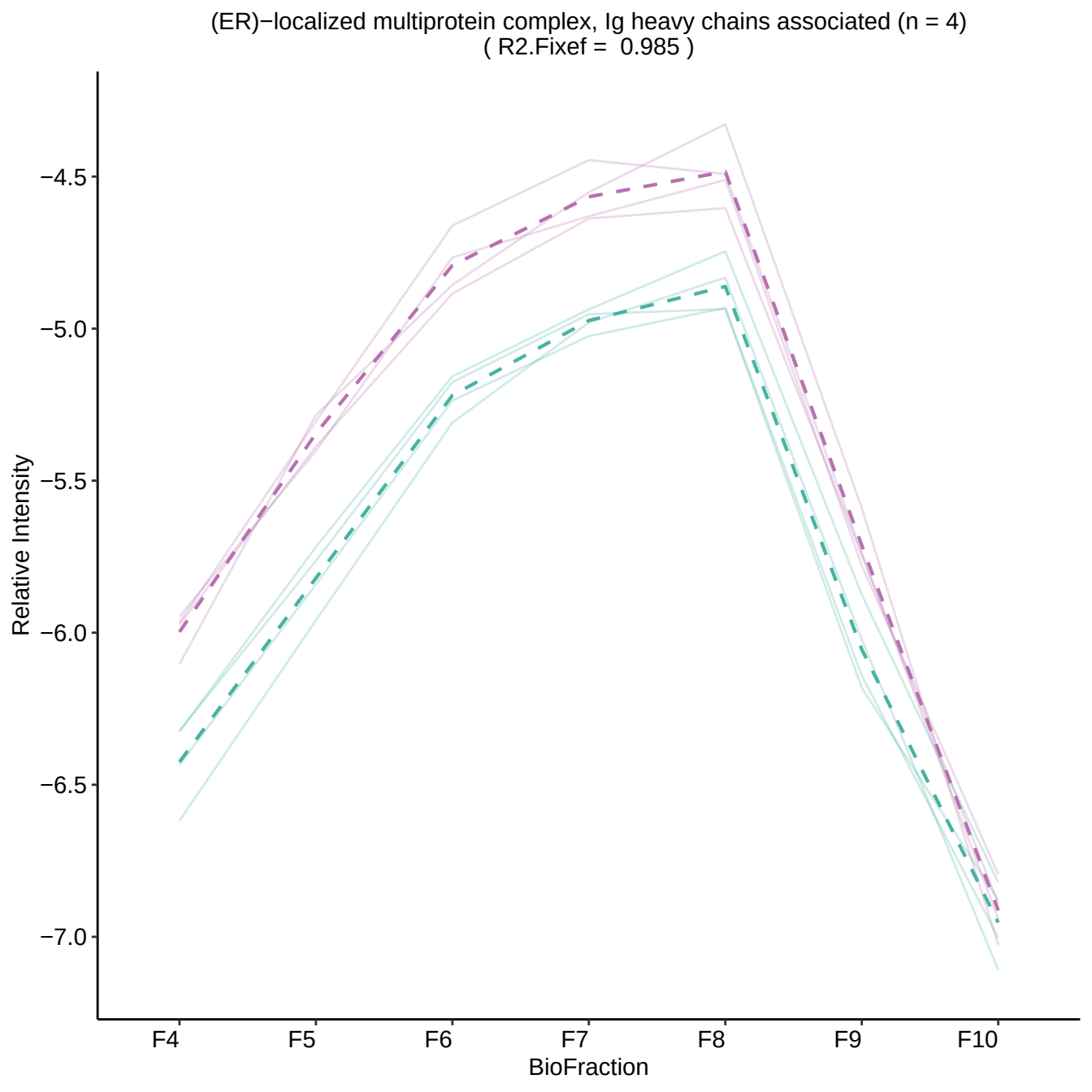
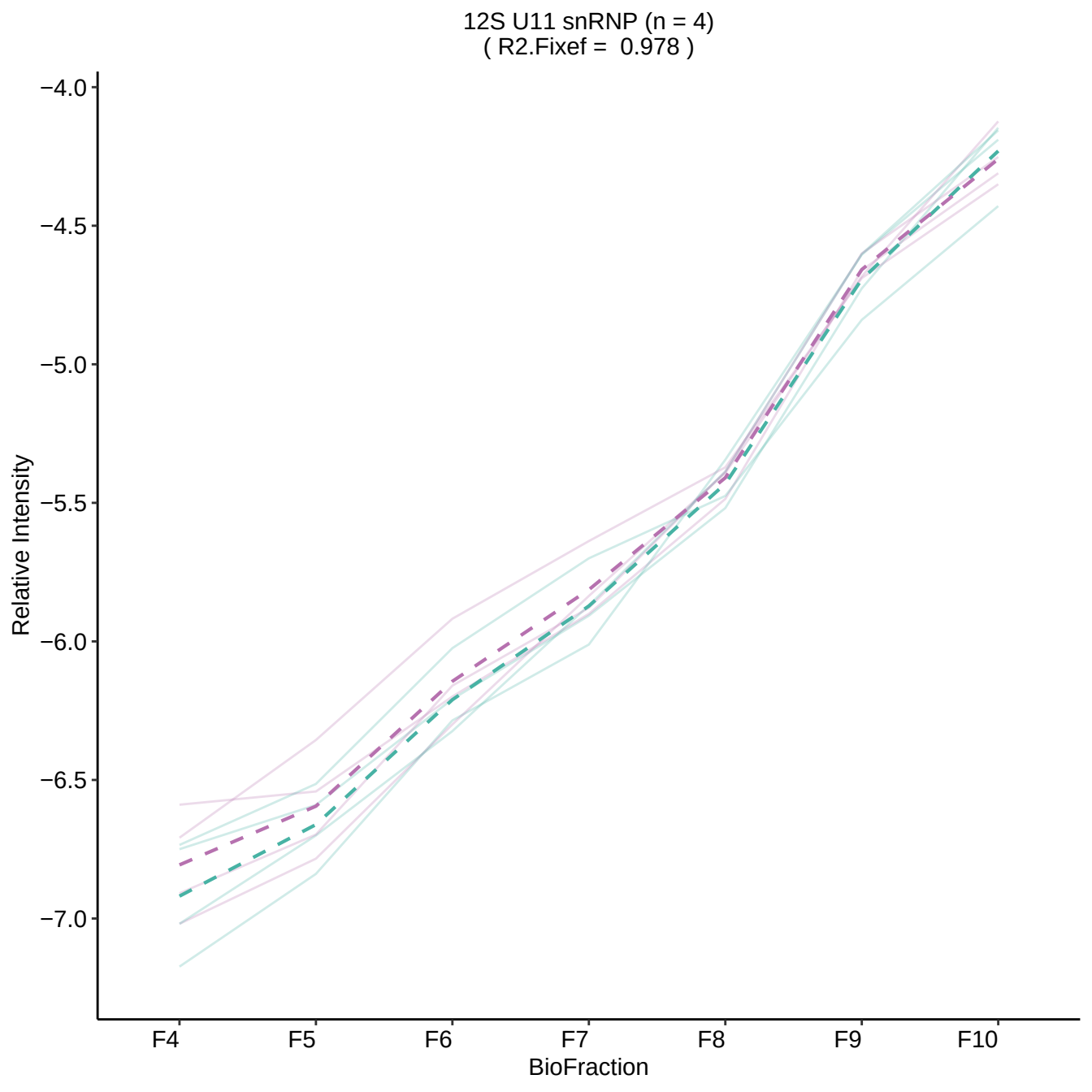
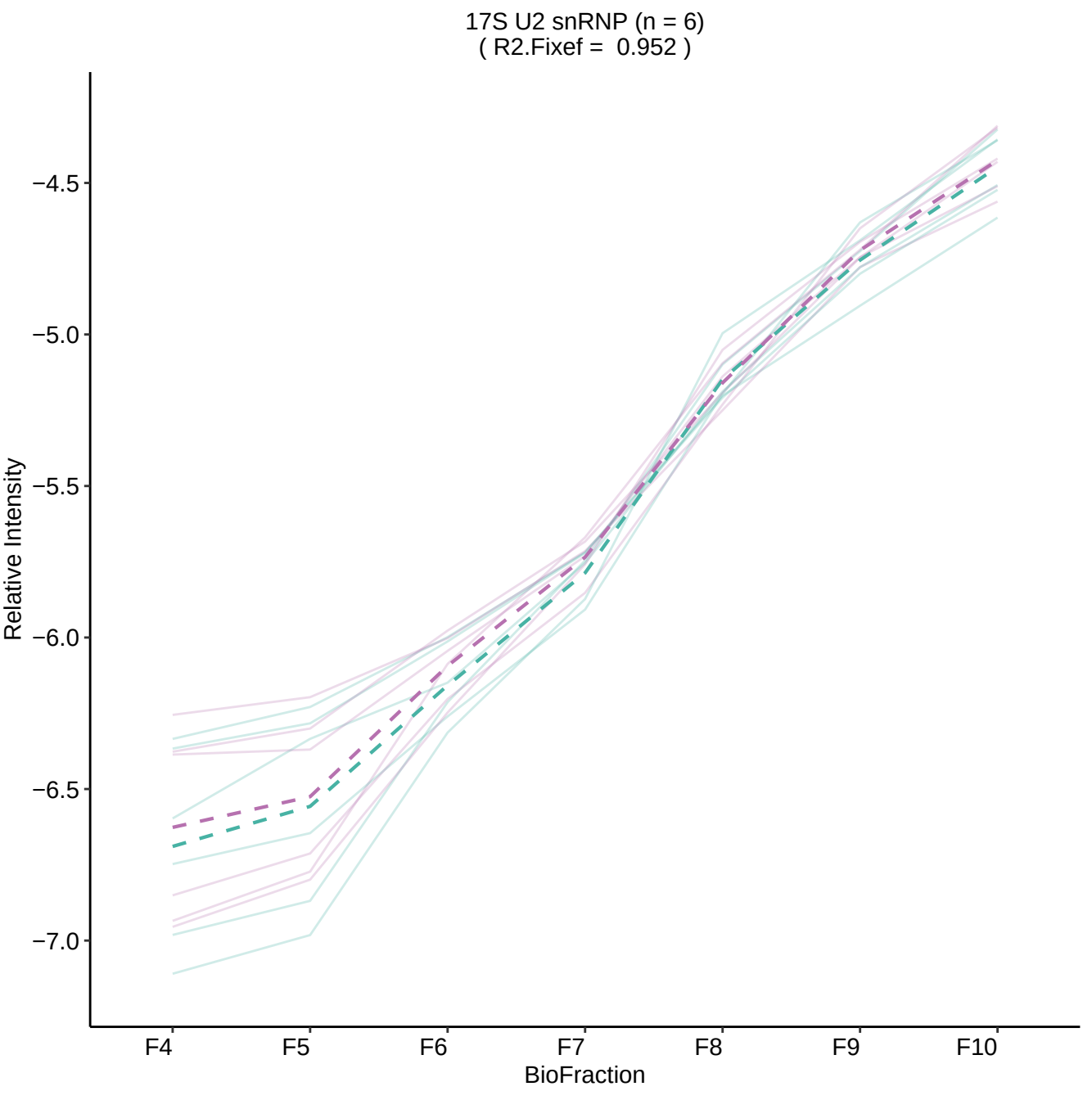
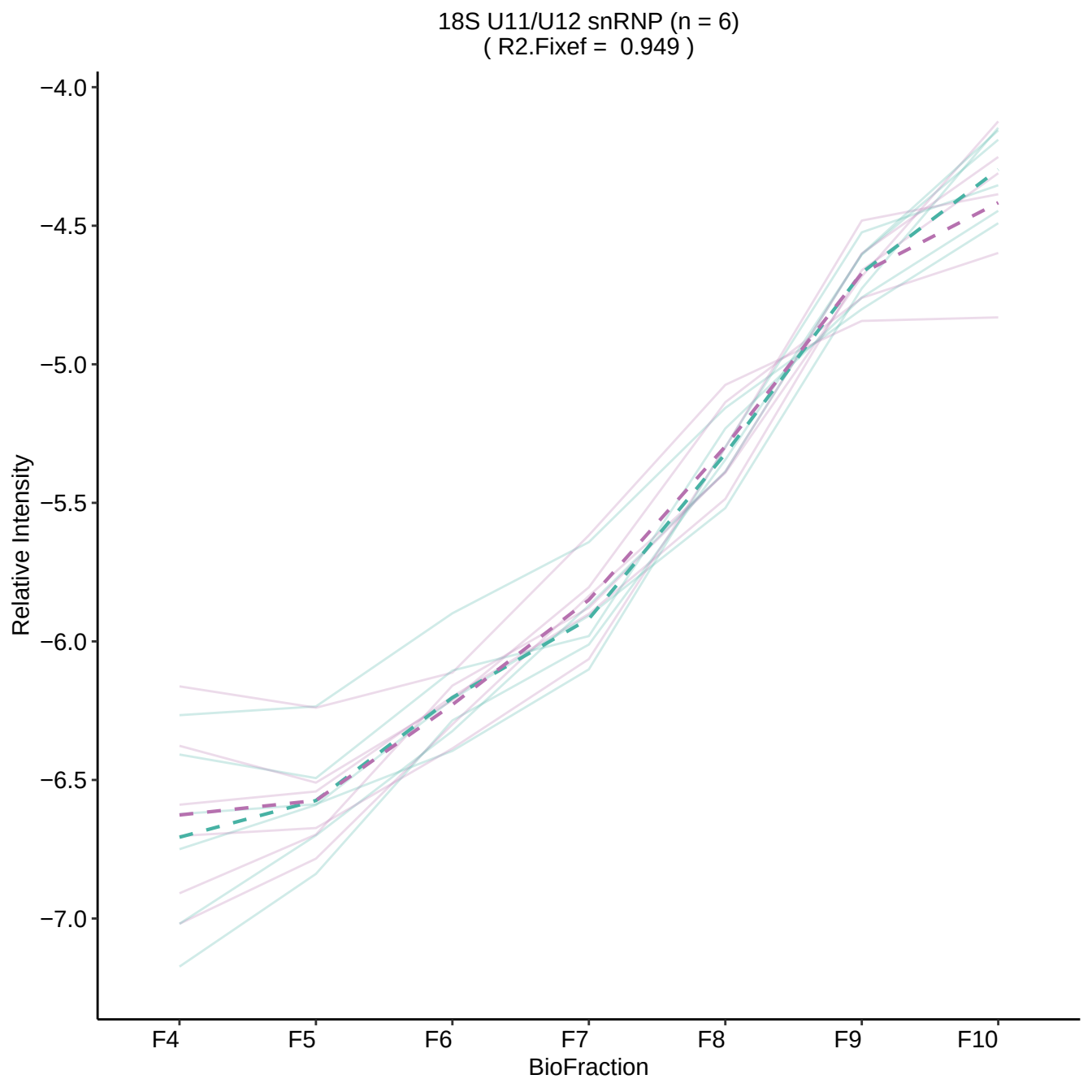
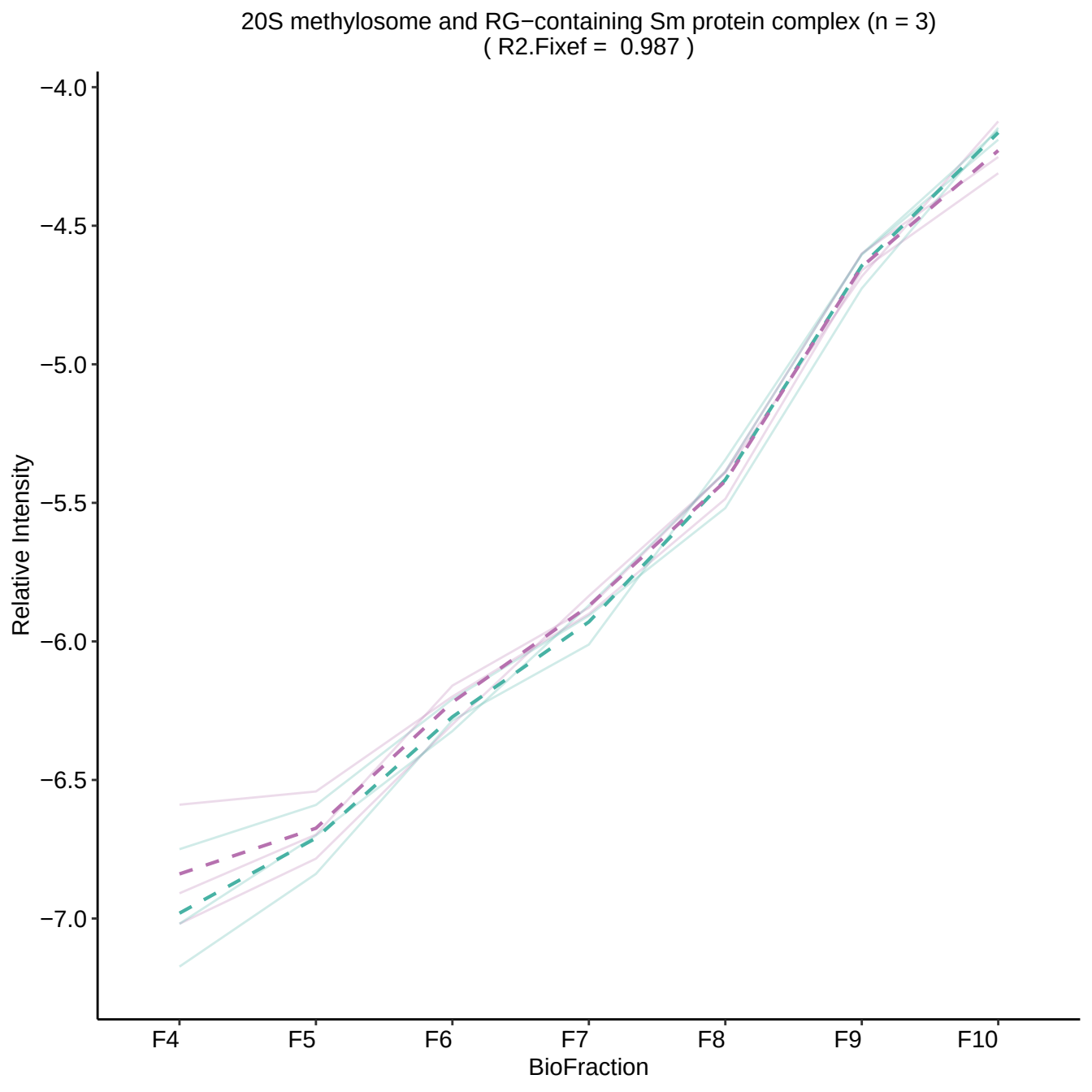
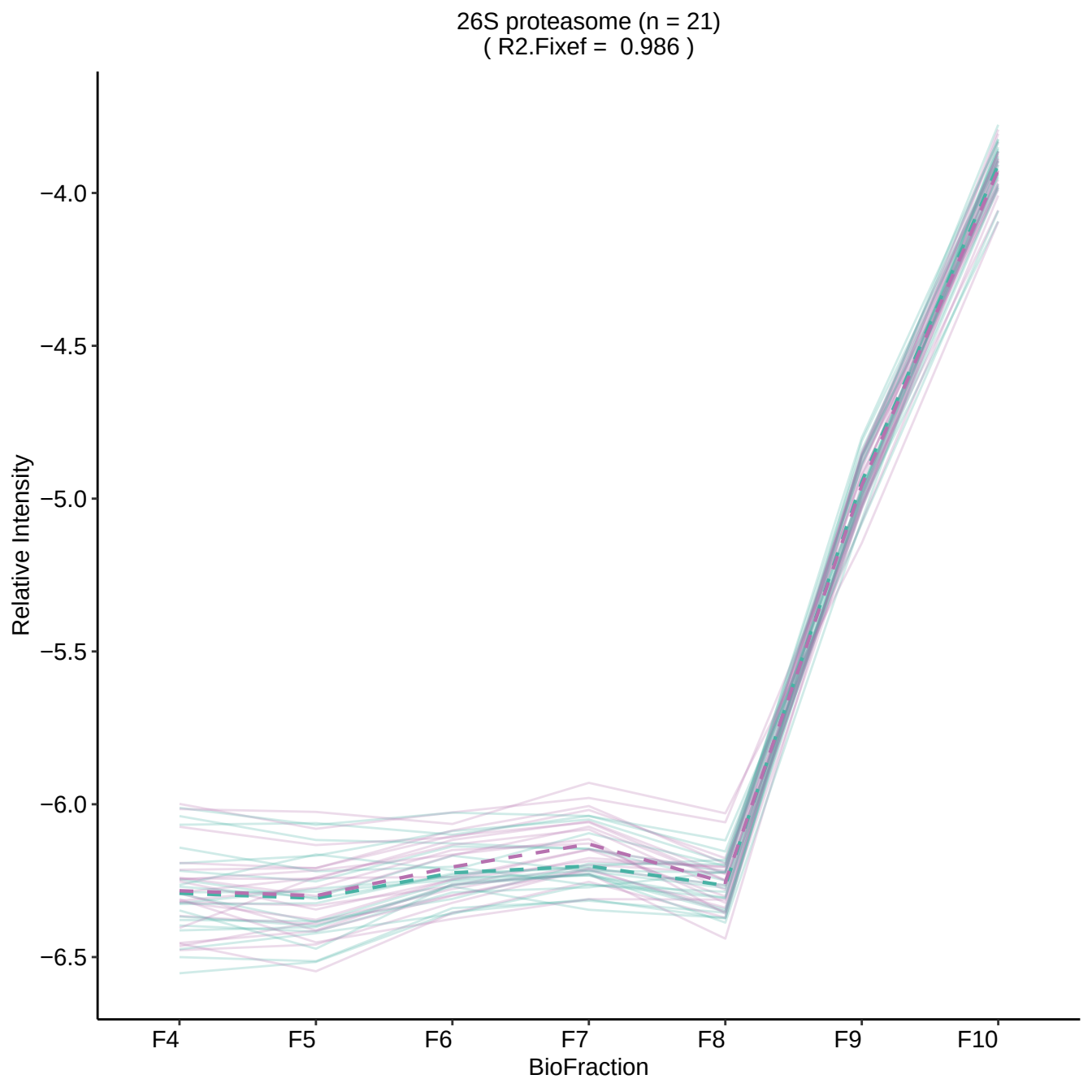

(ER)−localized multiprotein complex, Ig heavy chains associated (n = 4)
(R2.Fixef = 0.985)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

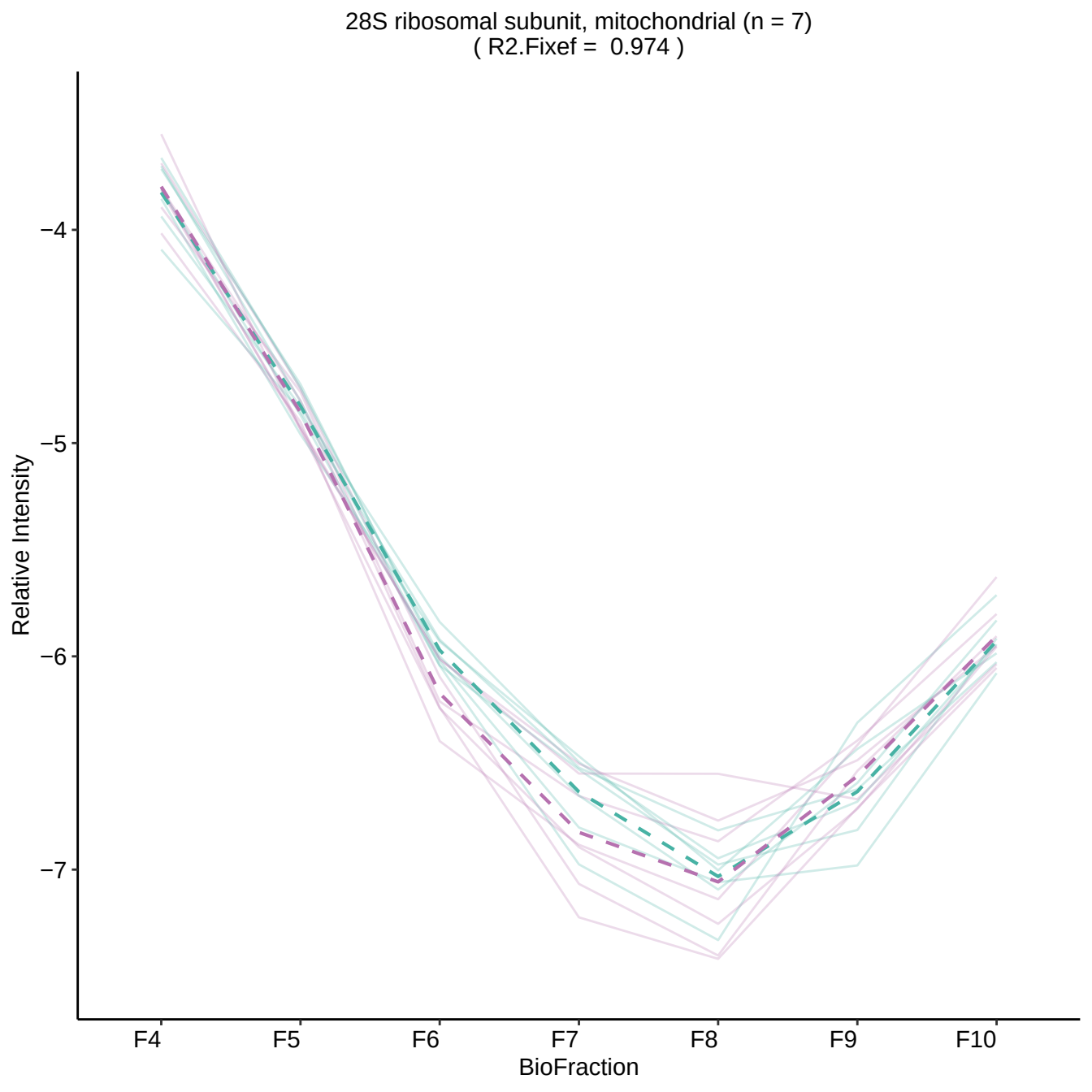

12S U11 snRNP (n = 4)
(R2.Fixef = 0.978)
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

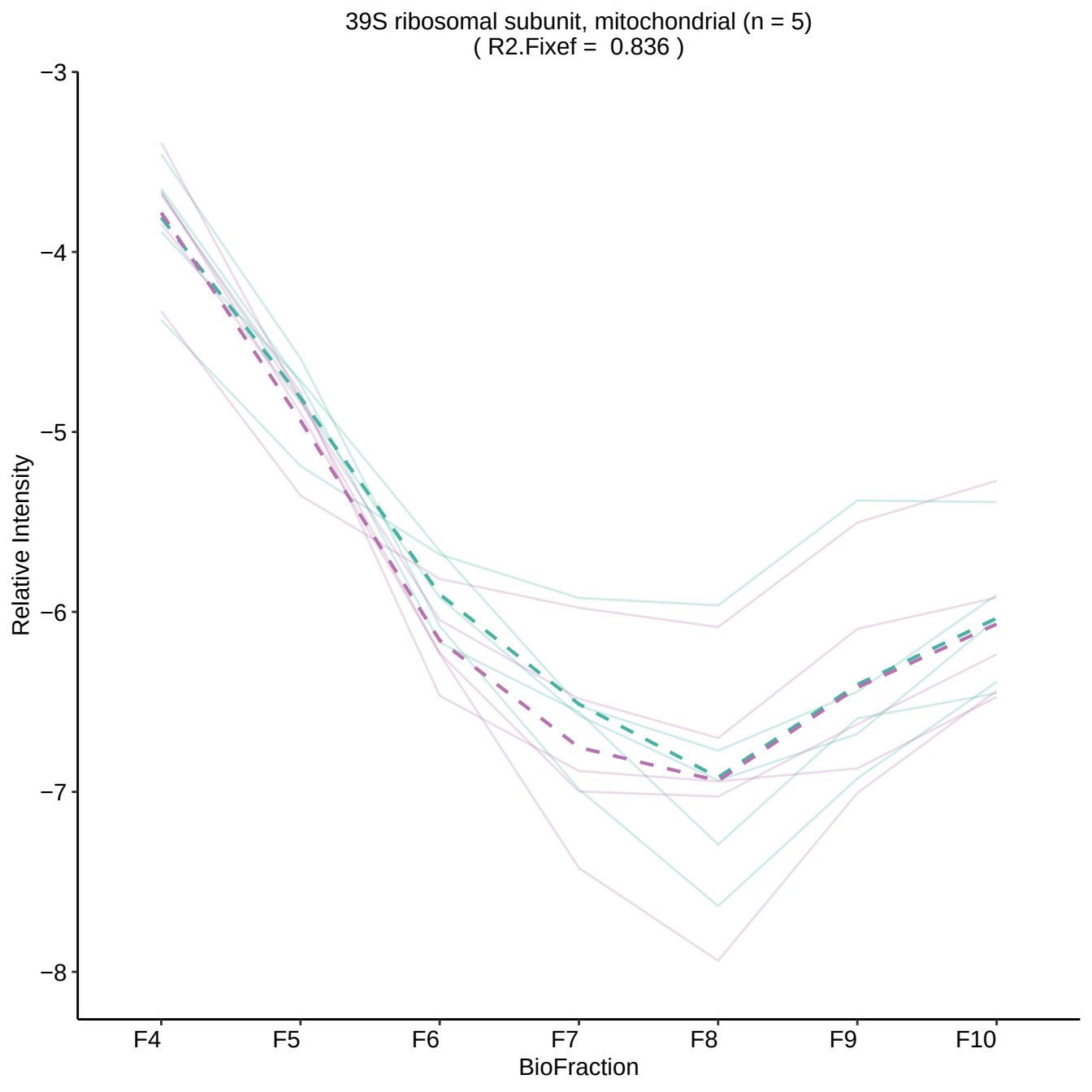

17S U2 snRNP (n = 6)
(R2.Fixef = 0.952)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

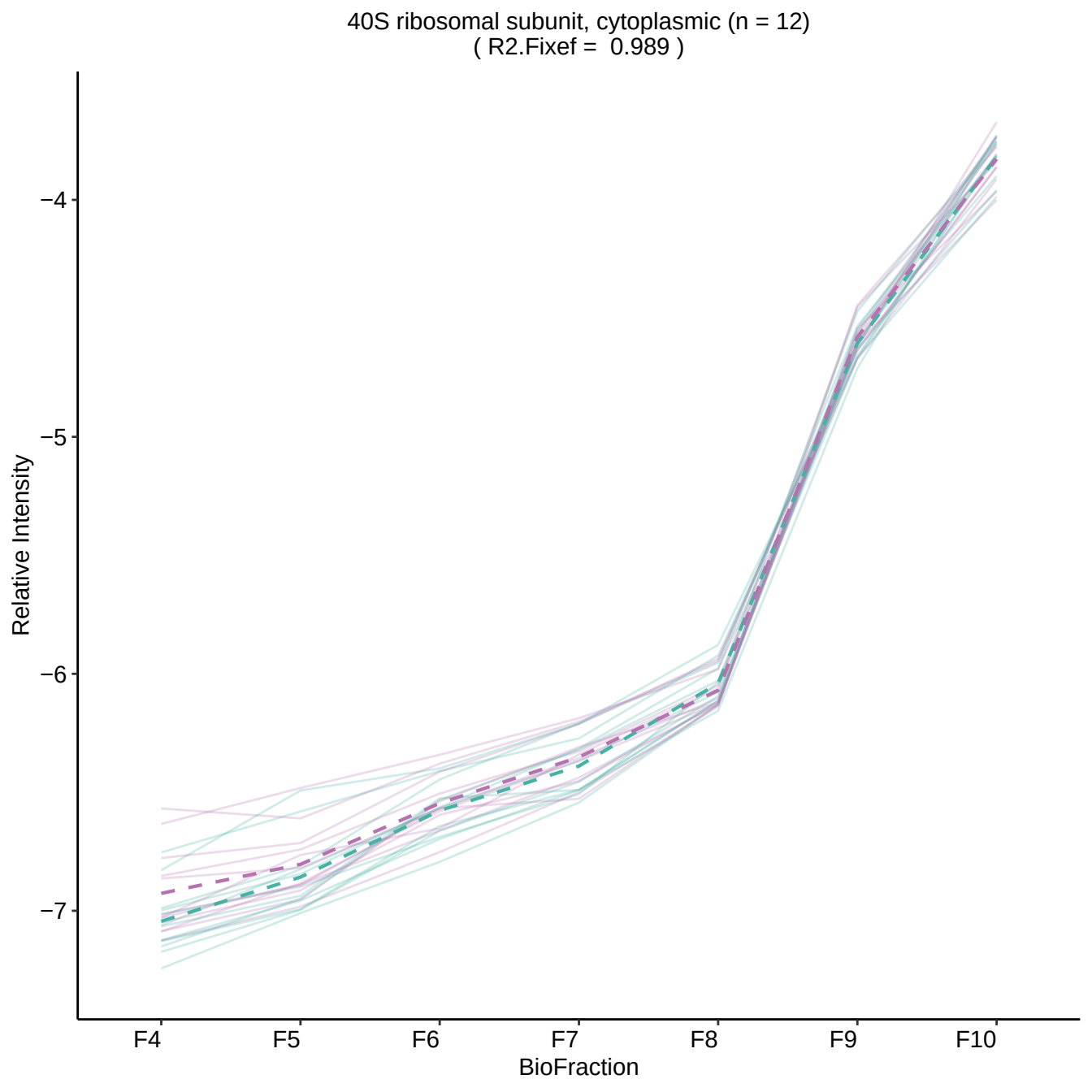

18S U11/U12 snRNP (n = 6)
(R2.Fixef = 0.949)
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

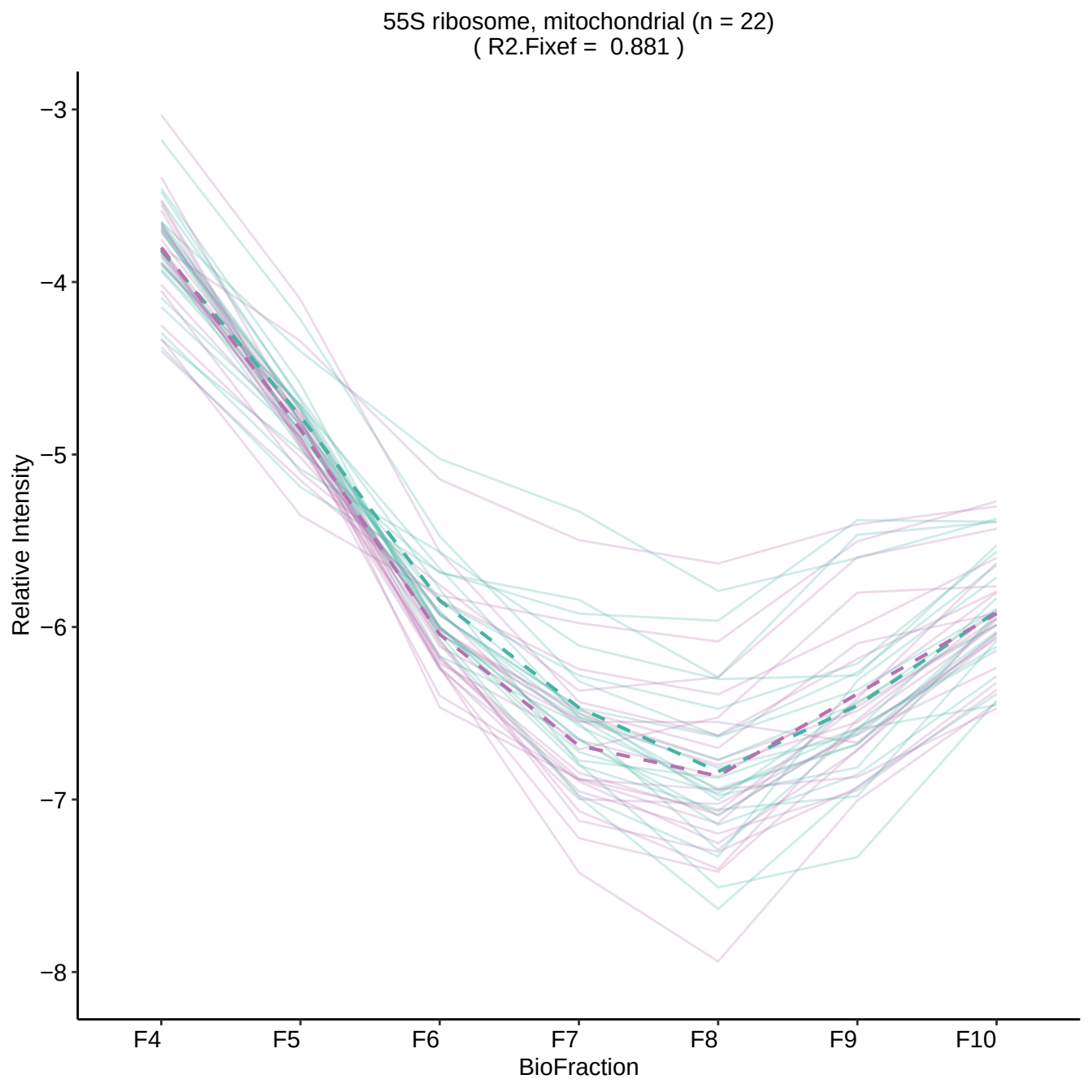

20S methylosome and RG−containing Sm protein complex (n = 3)
(R2.Fixef = 0.987)
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

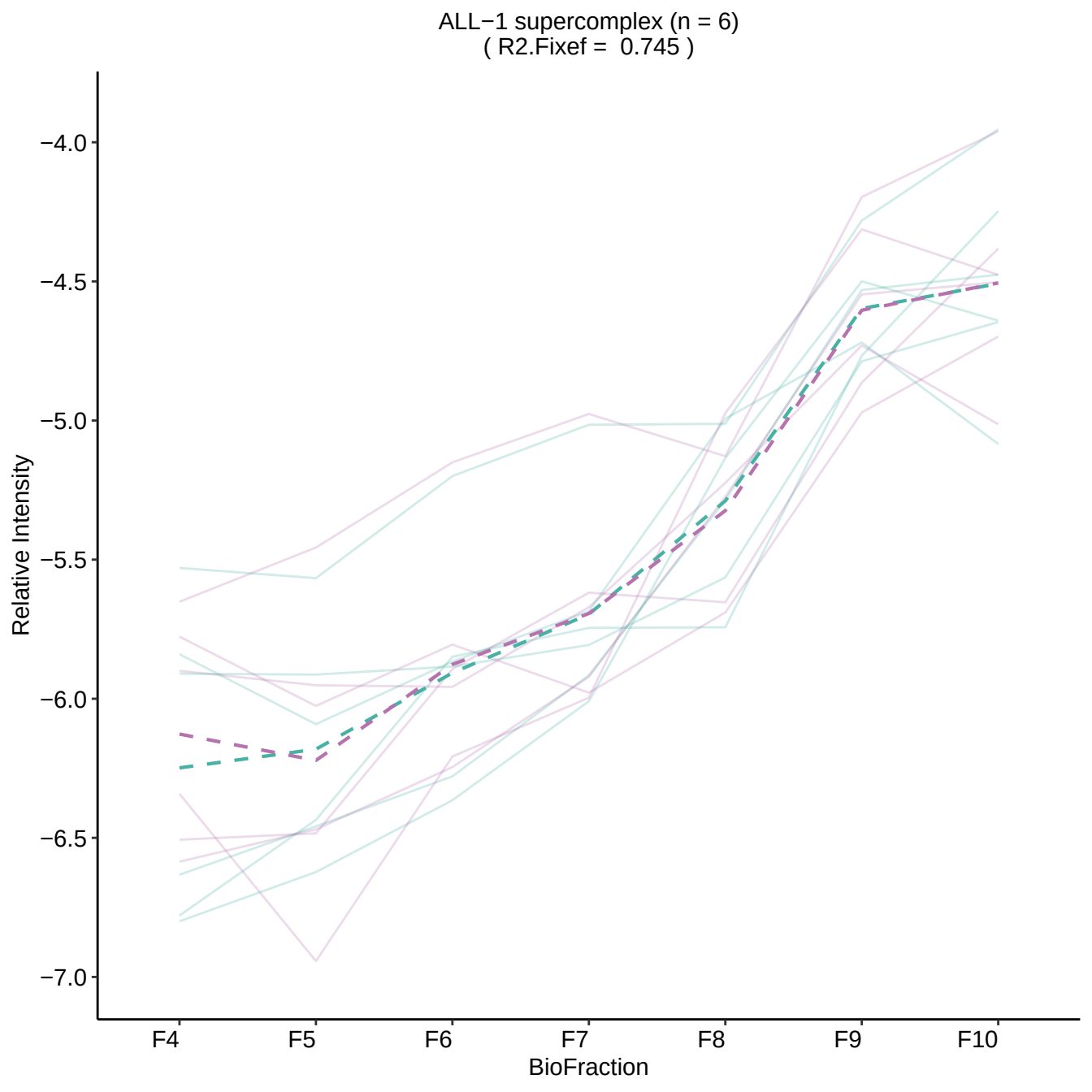

26S proteasome (n = 21)
(R2.Fixef = 0.986)
Relative Intensity
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
F4
F5
F6
F7
F8
F9
F10
BioFraction

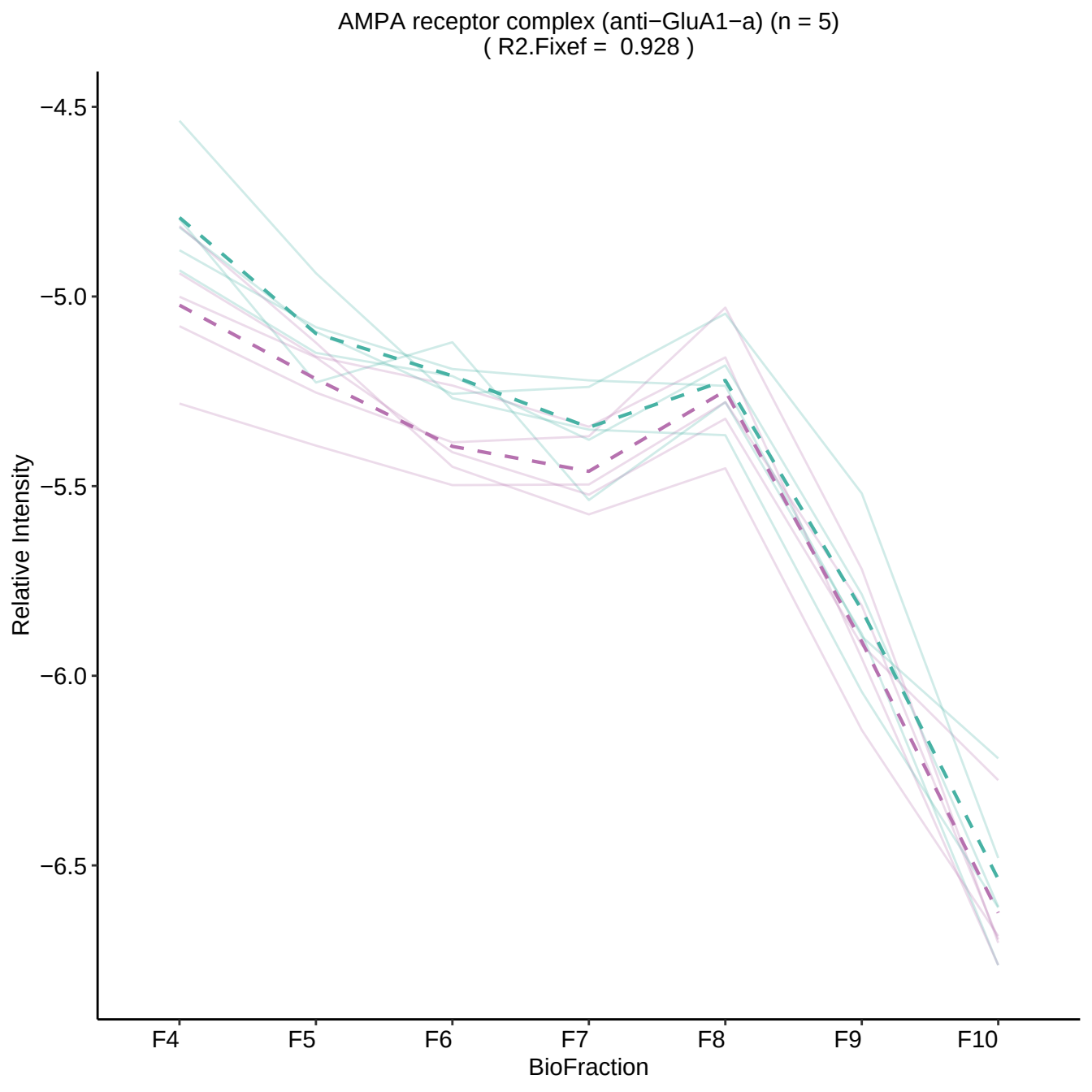

28S ribosomal subunit, mitochondrial (n = 7)
(R2.Fixef = 0.974)
Relative Intensity
−4
−5
−6
−7
F4
F5
F6
F7
F8
F9
F10
BioFraction

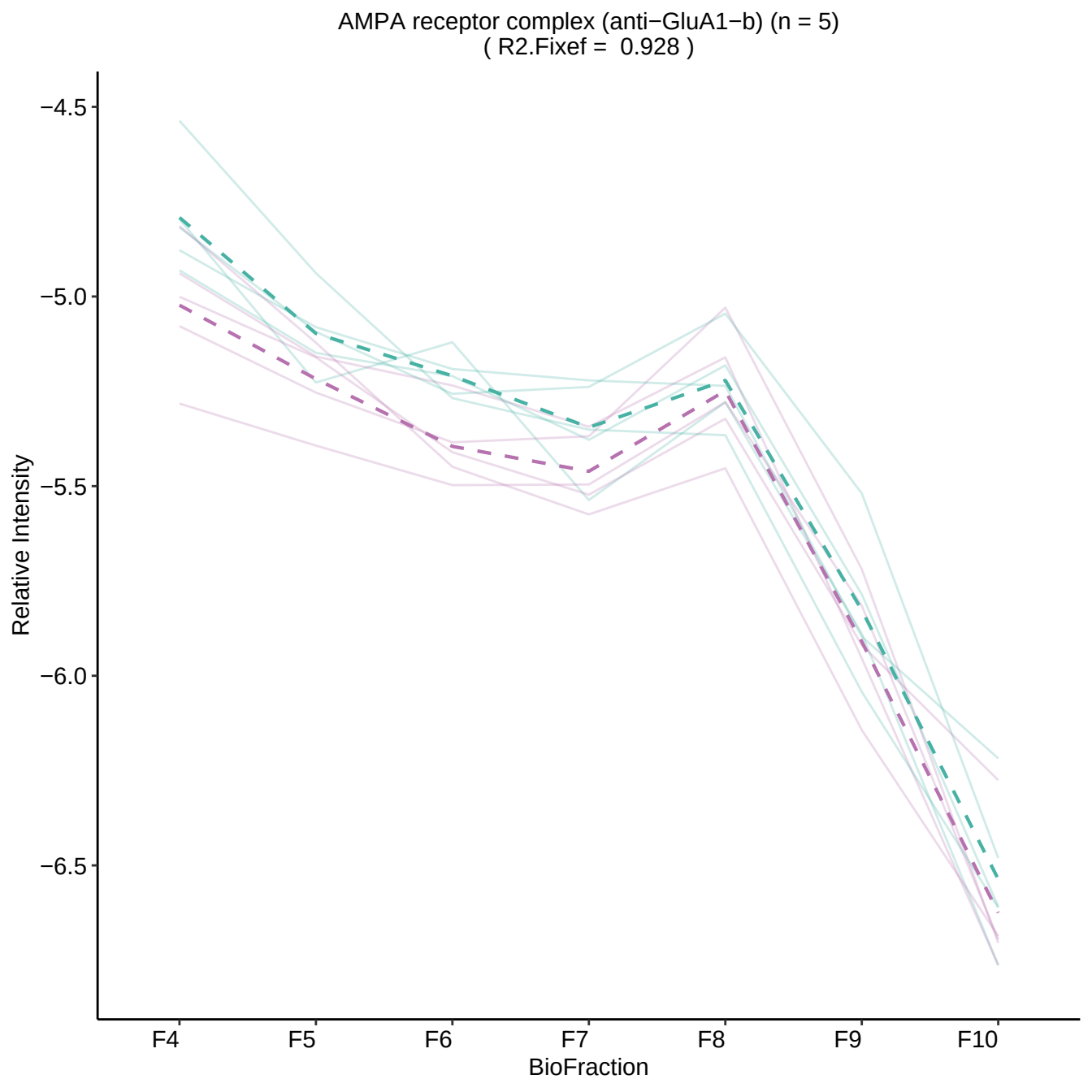

39S ribosomal subunit, mitochondrial (n = 5)
(R2.Fixef = 0.836)
−3
−4
−5
−6
−7
−8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

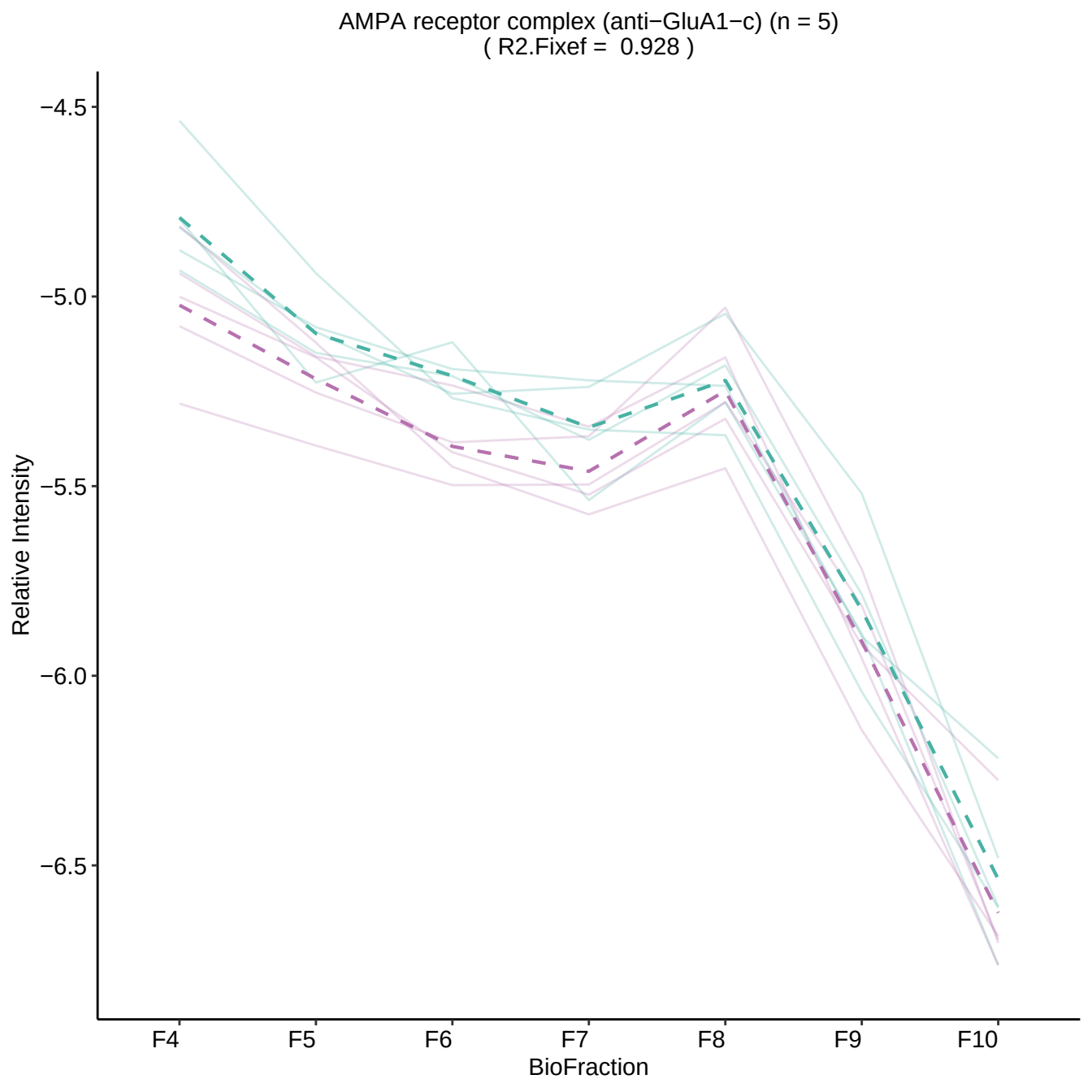

40S ribosomal subunit, cytoplasmic (n = 12)
(R2.Fixef = 0.989)
Relative Intensity
−4
−5
−6
−7
F4
F5
F6
F7
F8
F9
F10
BioFraction

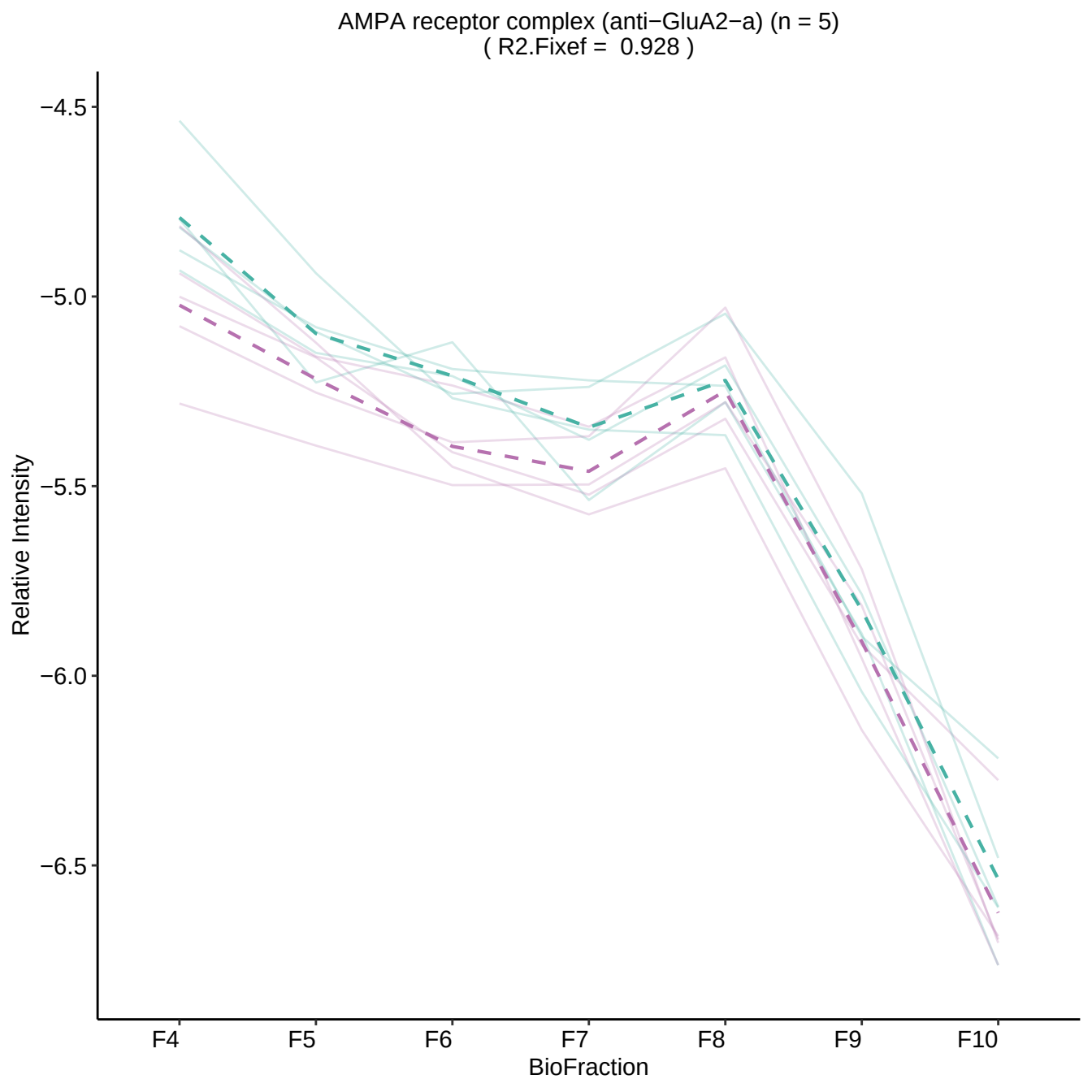

55S ribosome, mitochondrial (n = 22)
(R2.Fixef = 0.881)
−3
−4
−5
−6
−7
−8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

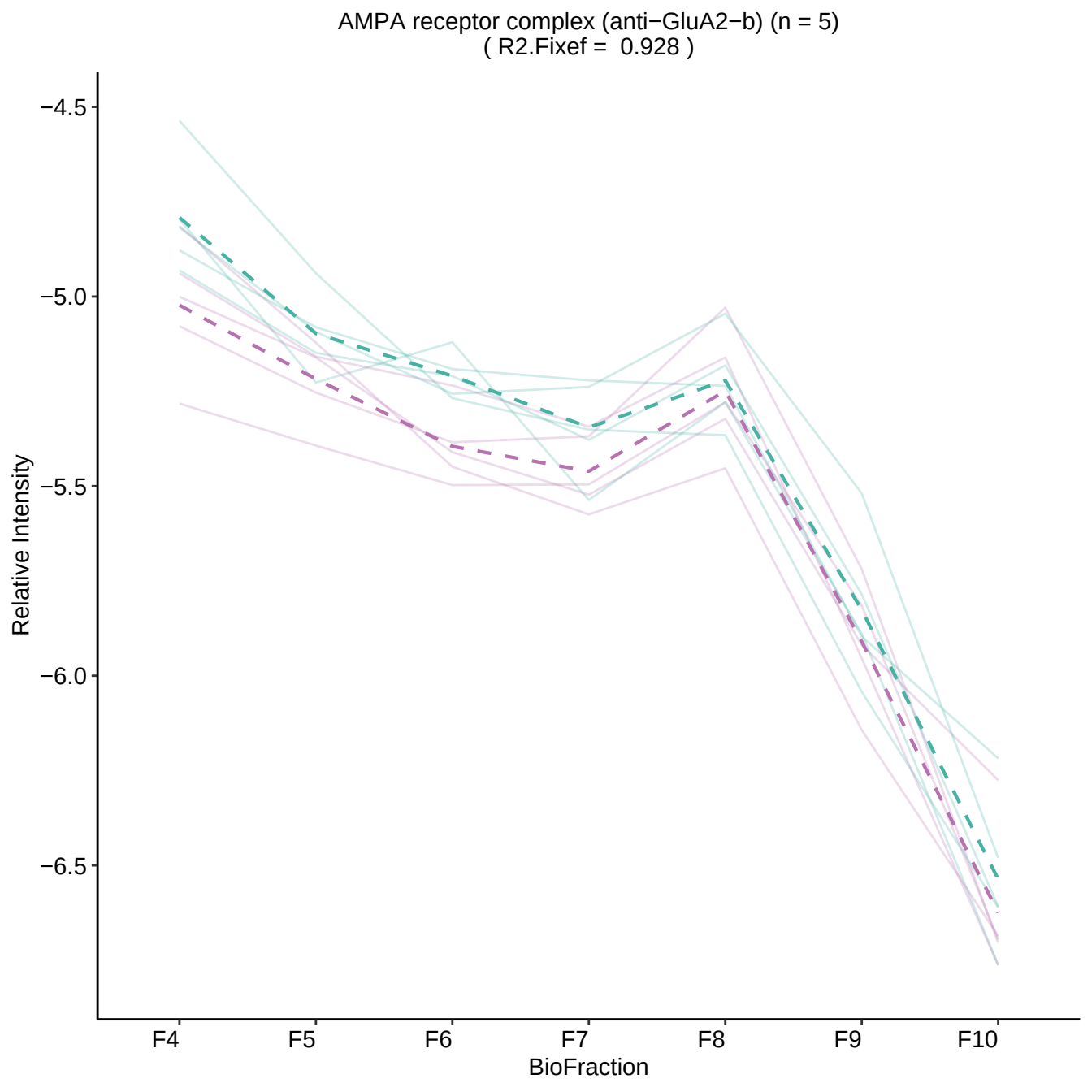

ALL−1 supercomplex (n = 6)
(R2.Fixef = 0.745)
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

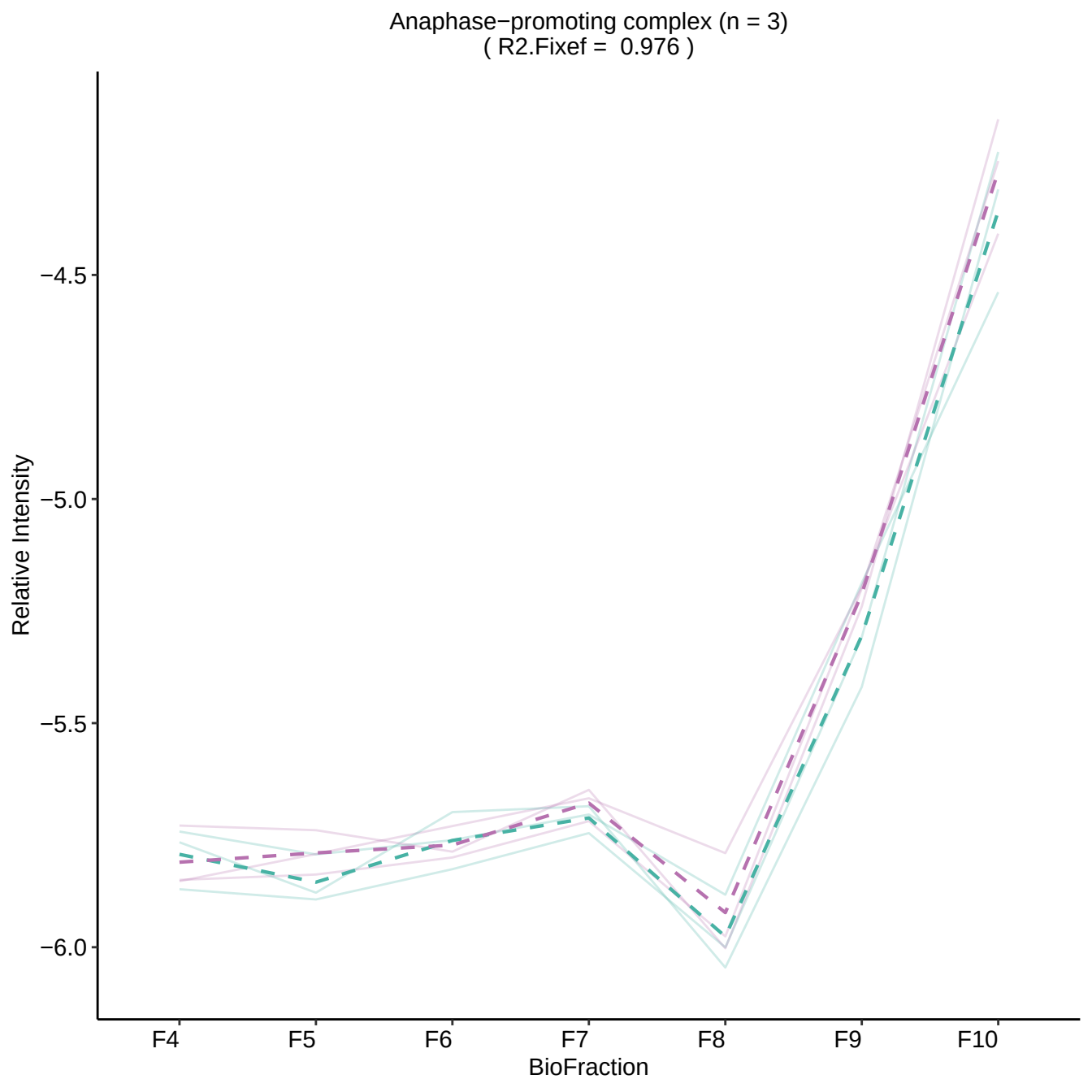

AMPA receptor complex (anti−GluA1−a) (n = 5)
(R2.Fixef = 0.928)
−4.5
−5.0
−5.5
−6.0
−6.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

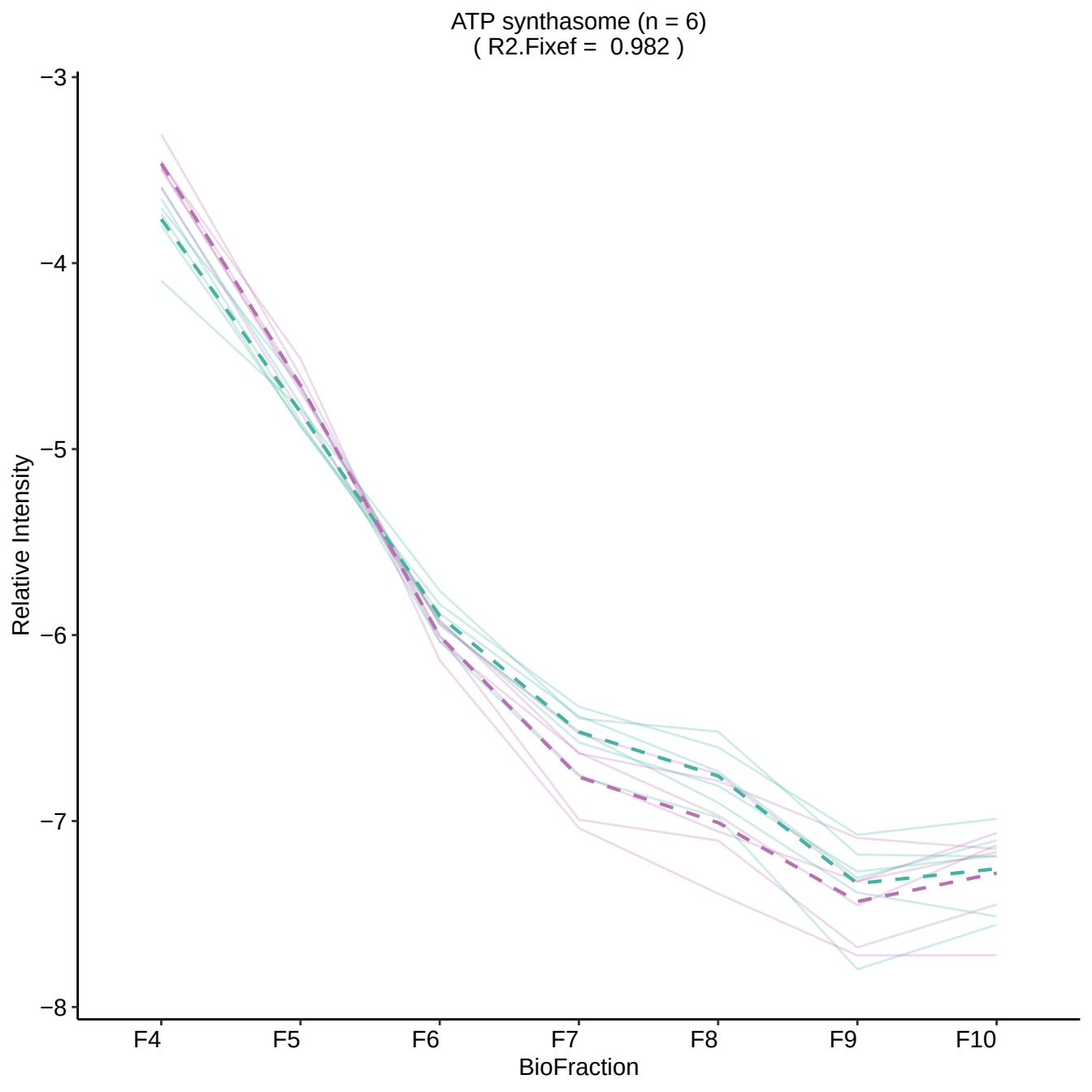

AMPA receptor complex (anti−GluA1−b) (n = 5)
(R2.Fixef = 0.928)
−4.5
−5.0
−5.5
−6.0
−6.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

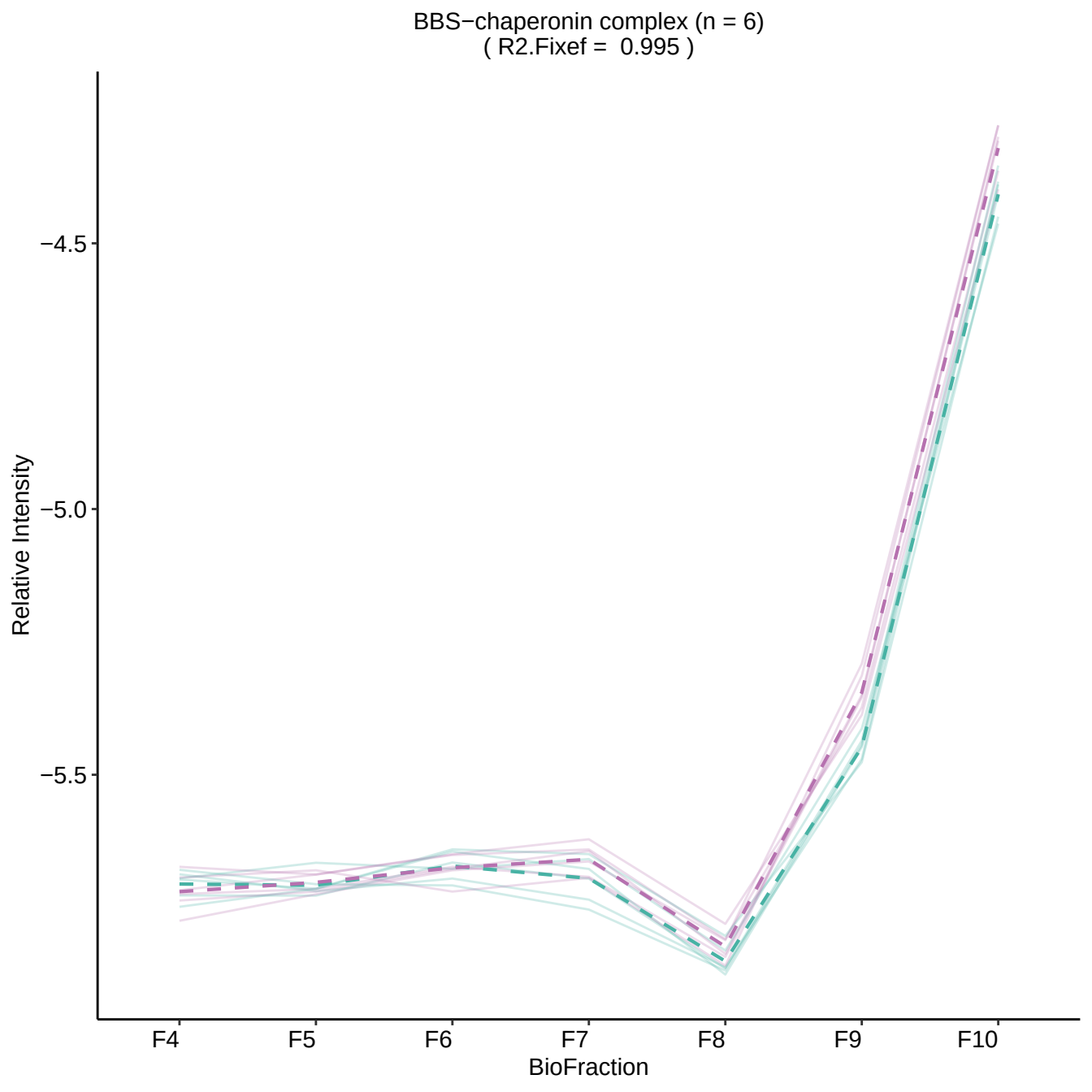

AMPA receptor complex (anti−GluA1−c) (n = 5)
(R2.Fixef = 0.928)
−4.5
−5.0
−5.5
−6.0
−6.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

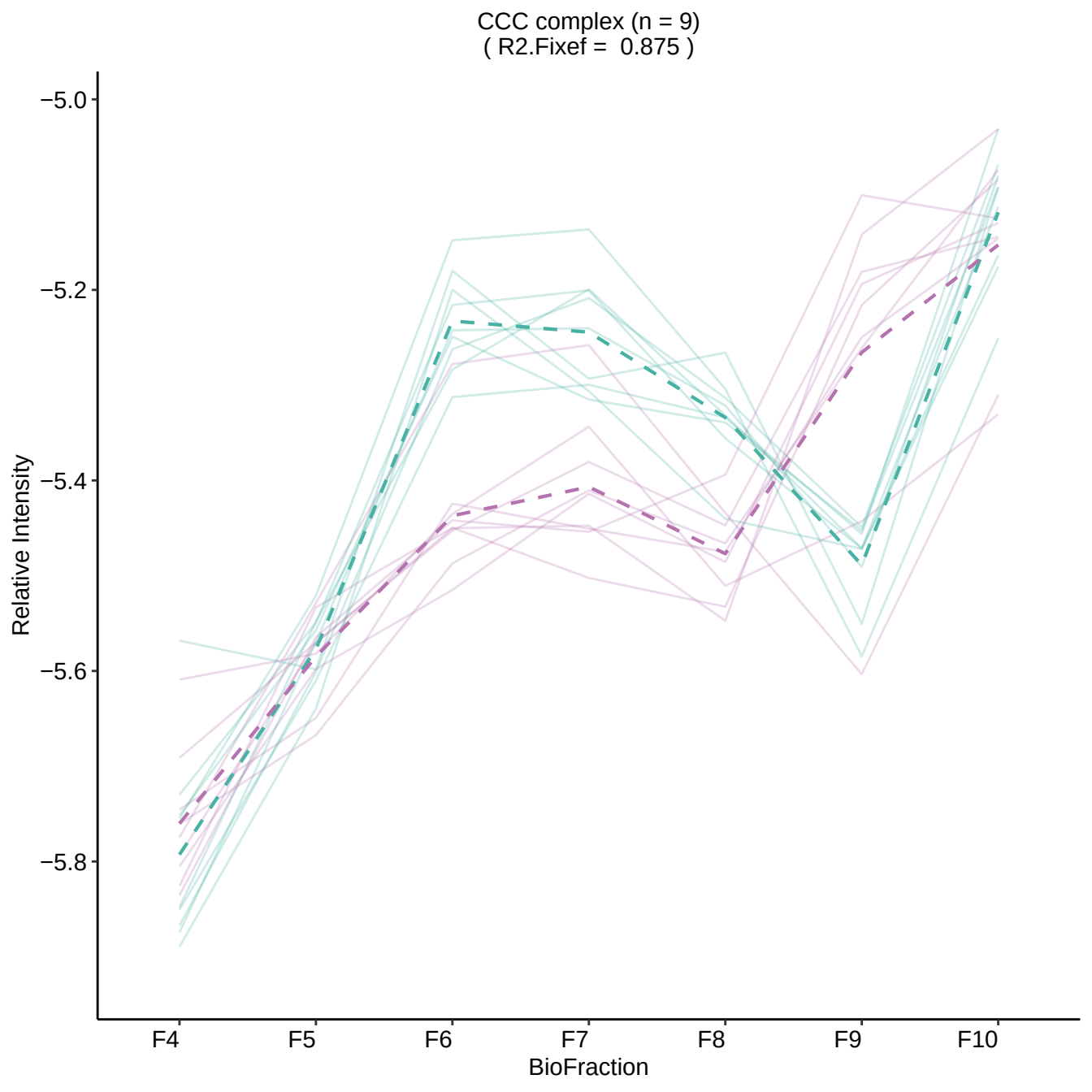

AMPA receptor complex (anti−GluA2−a) (n = 5)
(R2.Fixef = 0.928)
−4.5
−5.0
−5.5
−6.0
−6.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

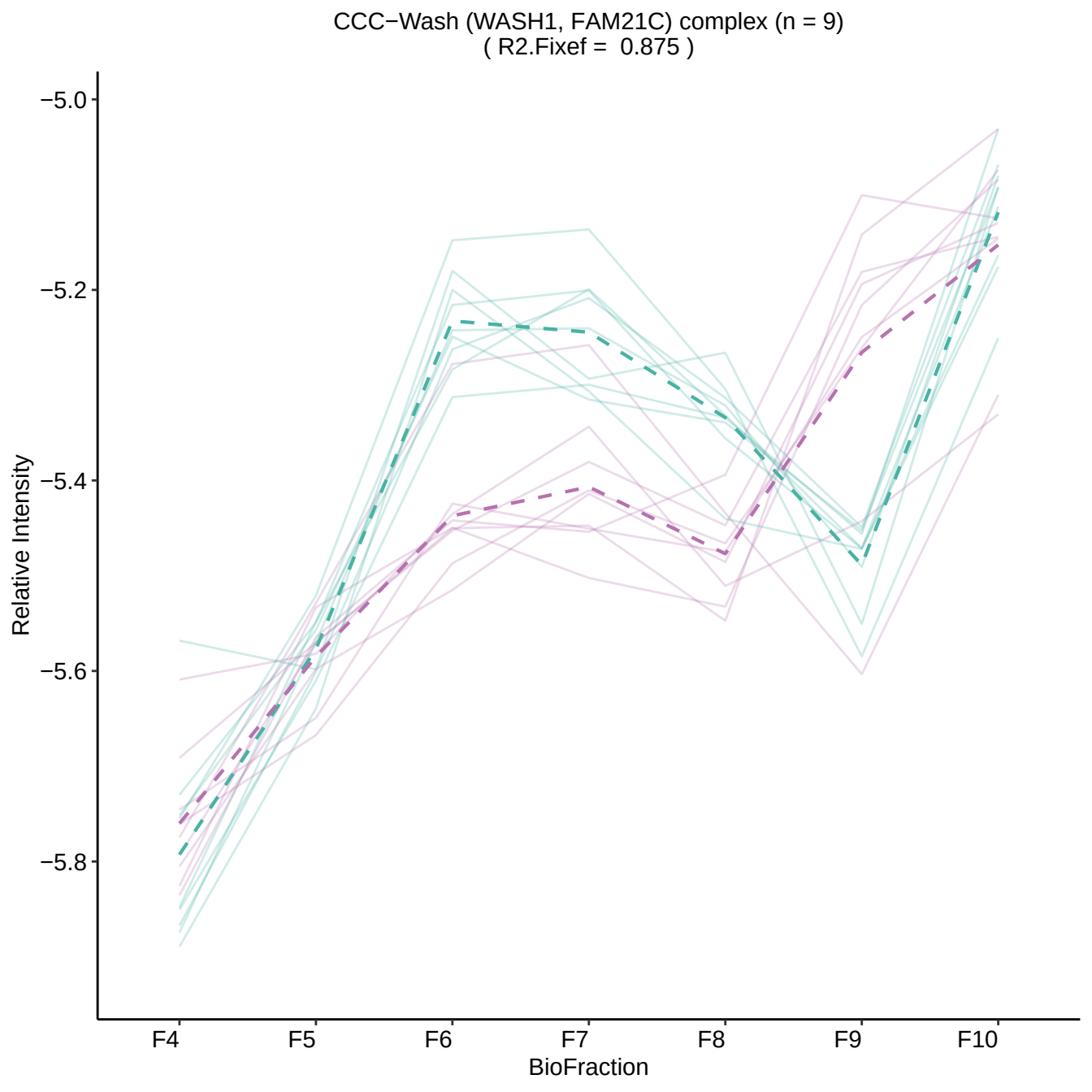

AMPA receptor complex (anti−GluA2−b) (n = 5)
(R2.Fixef = 0.928)
−4.5
−5.0
−5.5
−6.0
−6.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

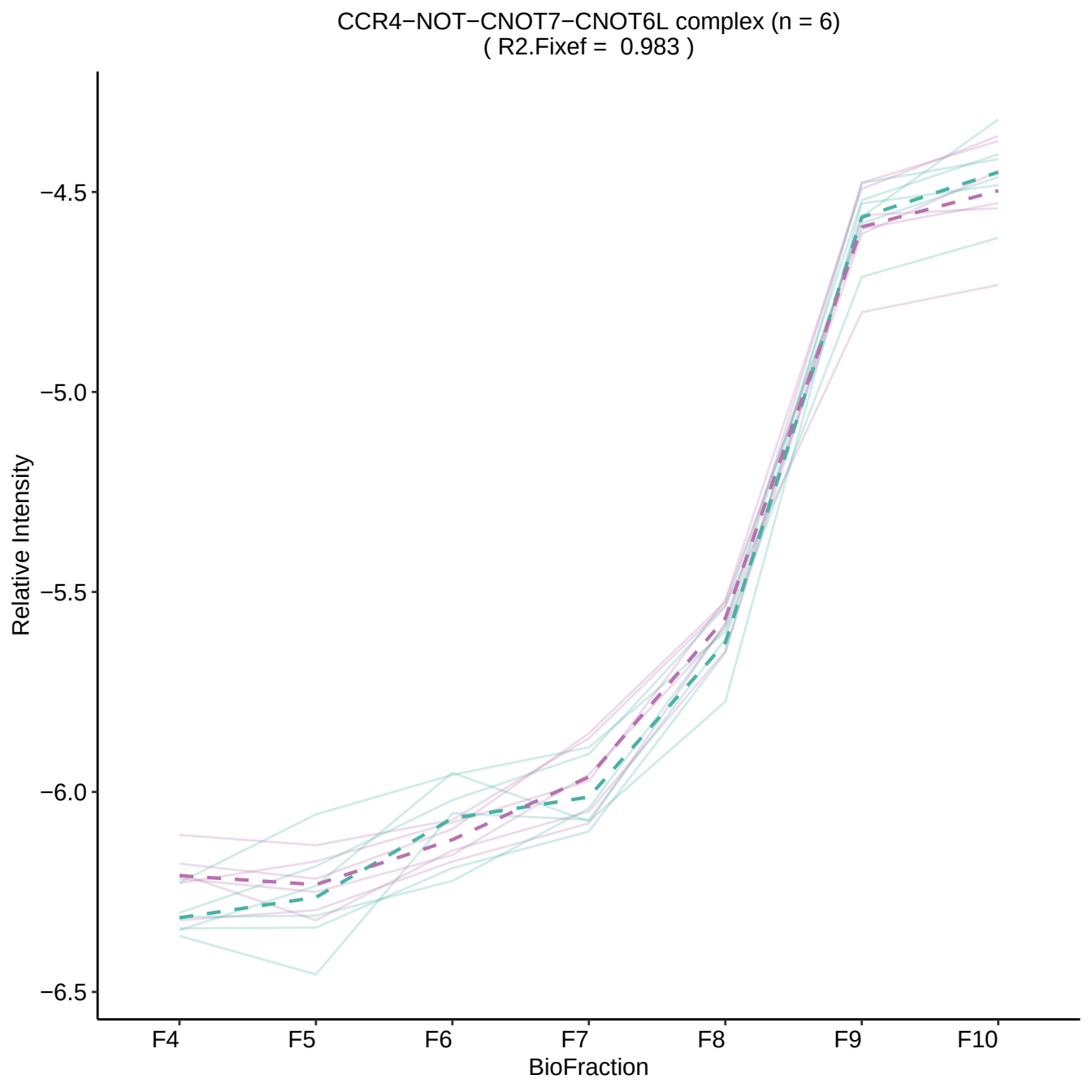

Anaphase−promoting complex (n = 3)
(R2.Fixef = 0.976)
−4.5
−5.0
−5.5
−6.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

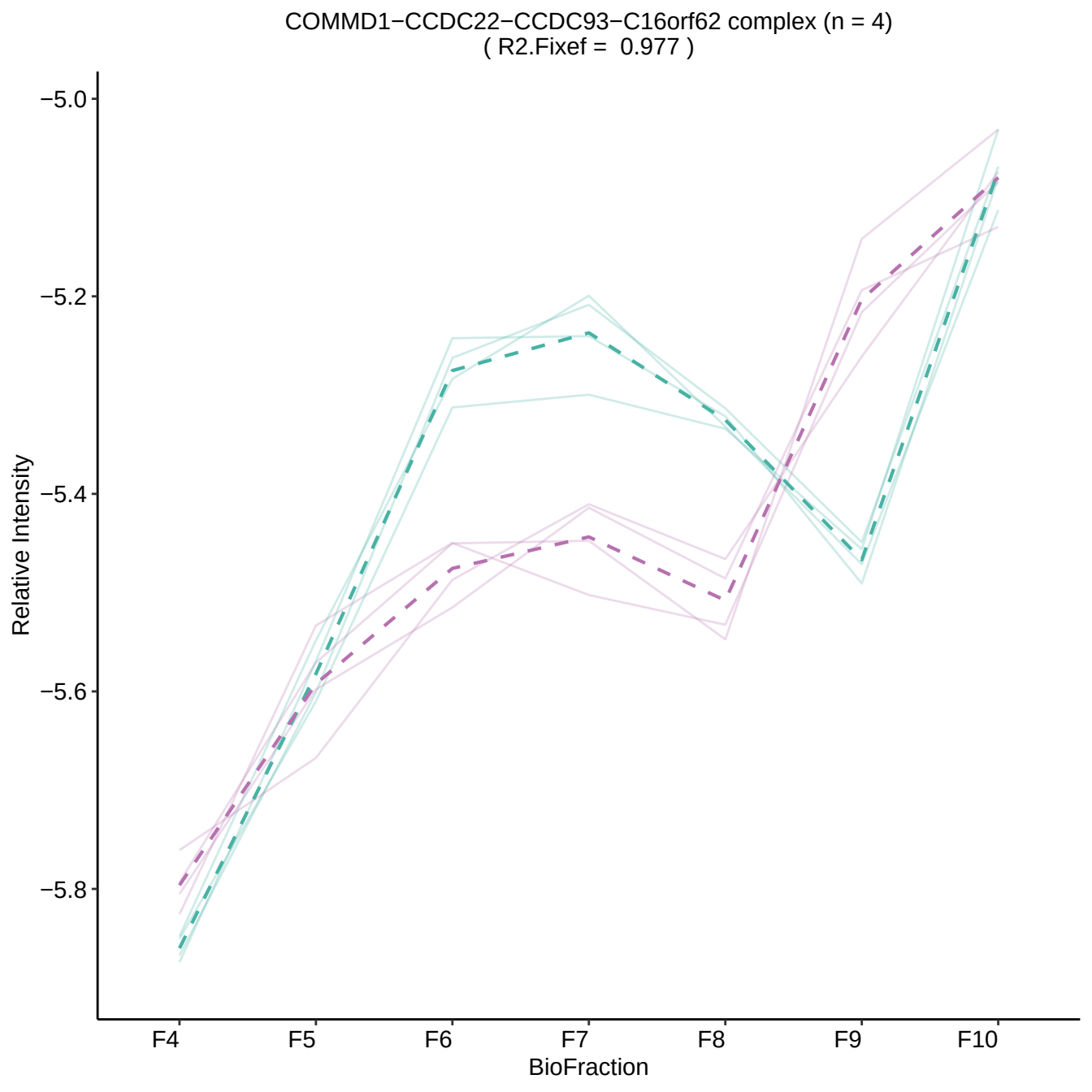

ATP synthasome (n = 6)
(R2.Fixef = 0.982)
−3
−4
−5
−6
−7
−8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

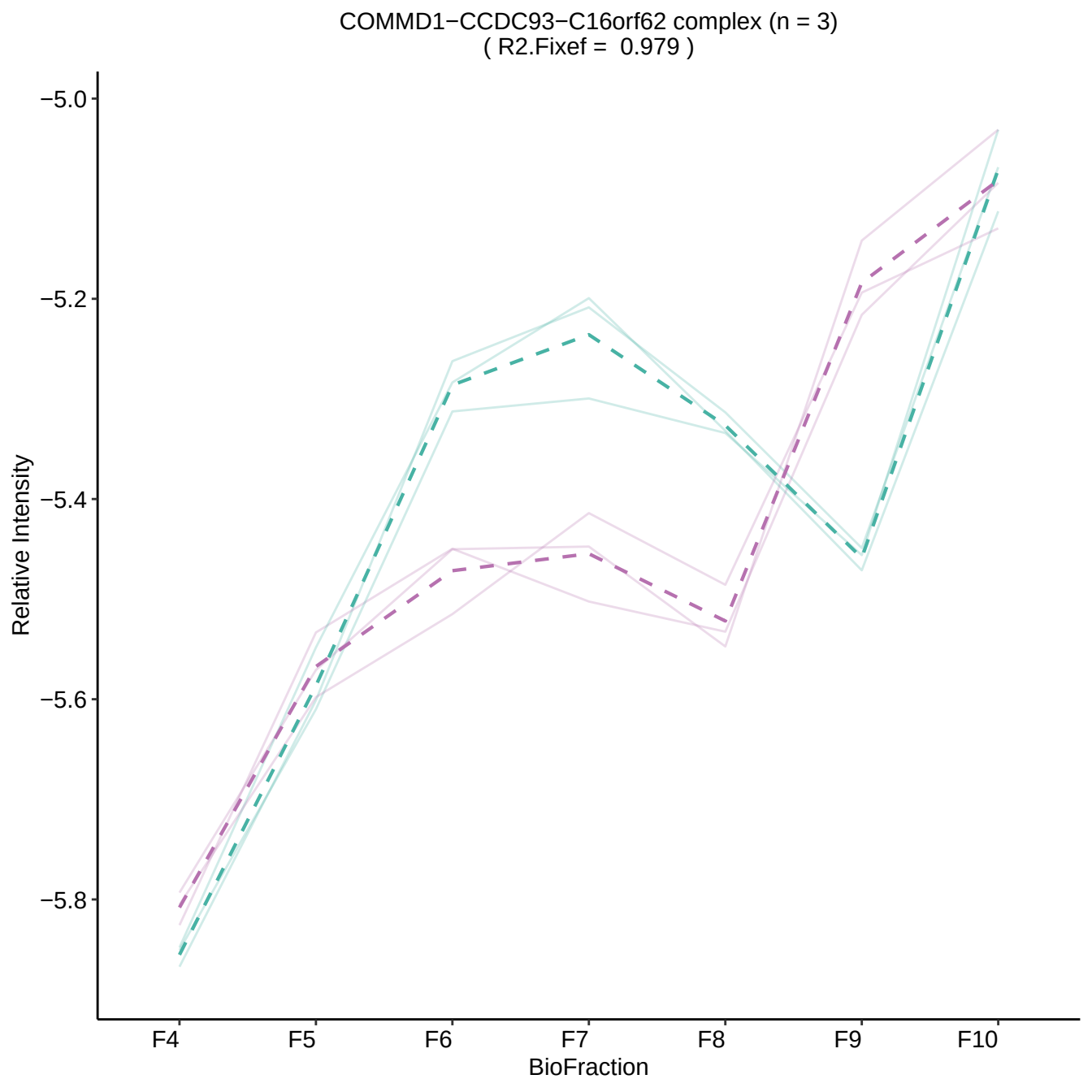

BBS−chaperonin complex (n = 6)
(R2.Fixef = 0.995)
Relative Intensity
−4.5
−5.0
−5.5
F4
F5
F6
F7
F8
F9
F10
BioFraction

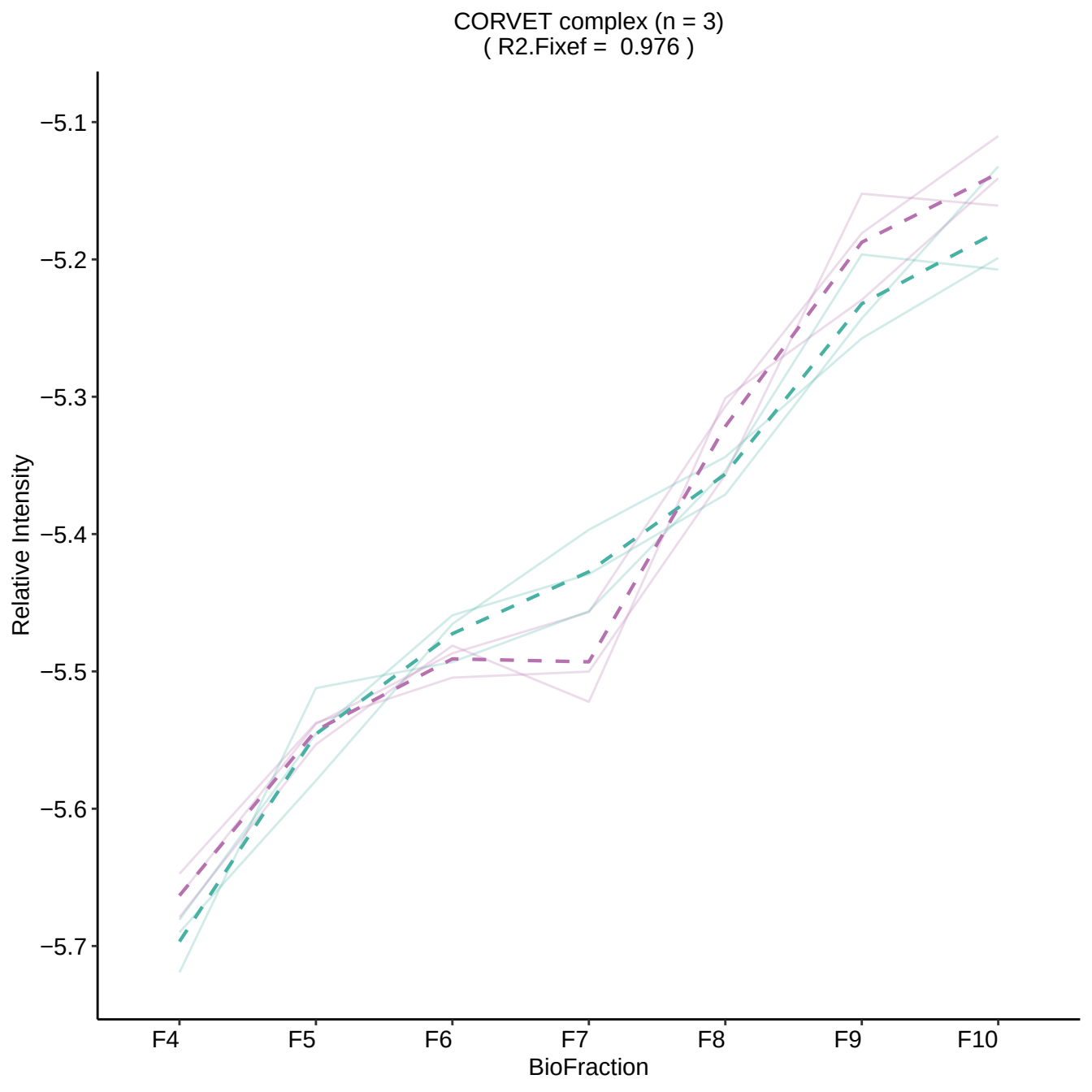

CCC complex (n = 9)
(R2.Fixef = 0.875)
−5.0
−5.2
−5.4
−5.6
−5.8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

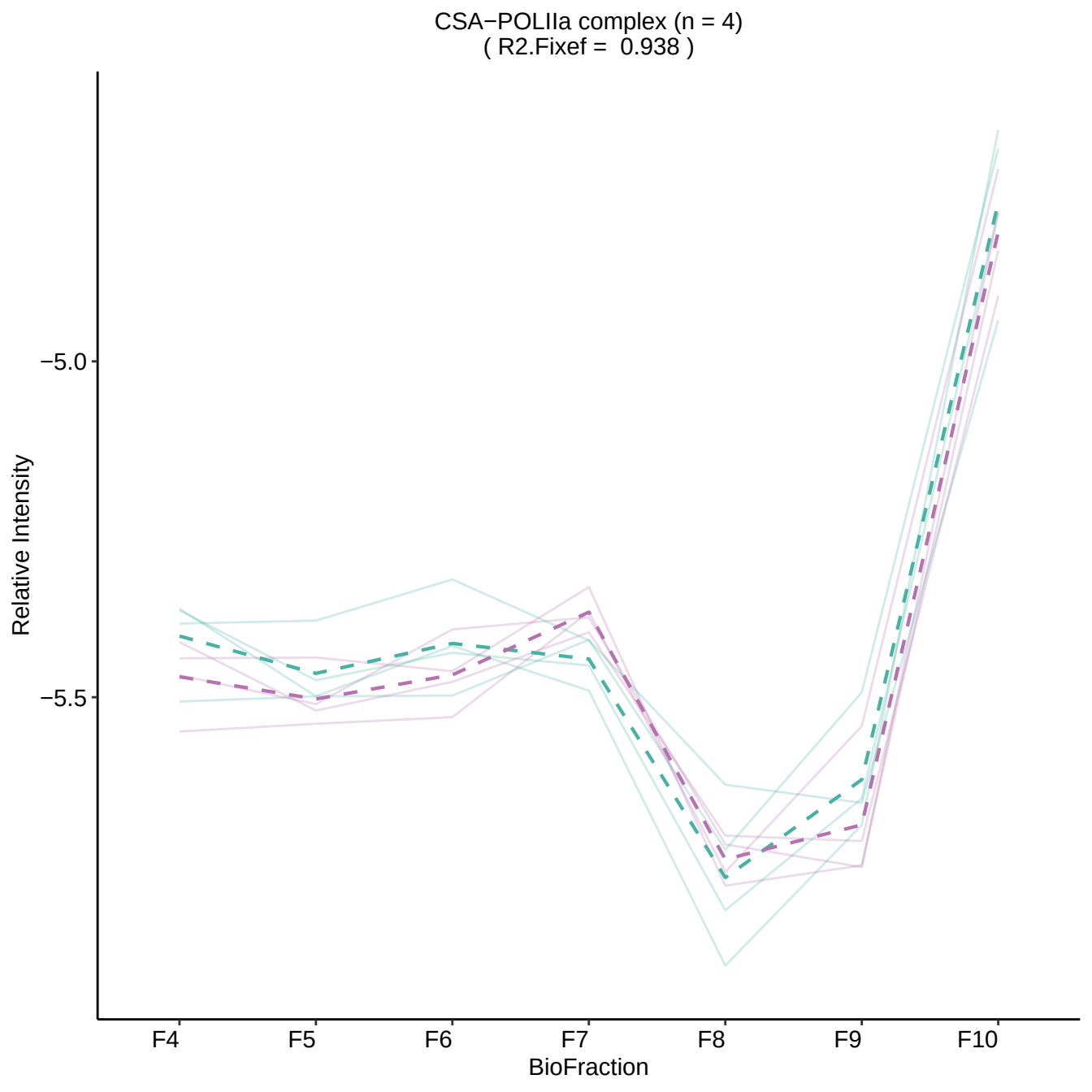

CCC−Wash (WASH1, FAM21C) complex (n = 9)
(R2.Fixef = 0.875)
−5.0
−5.2
−5.4
−5.6
−5.8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

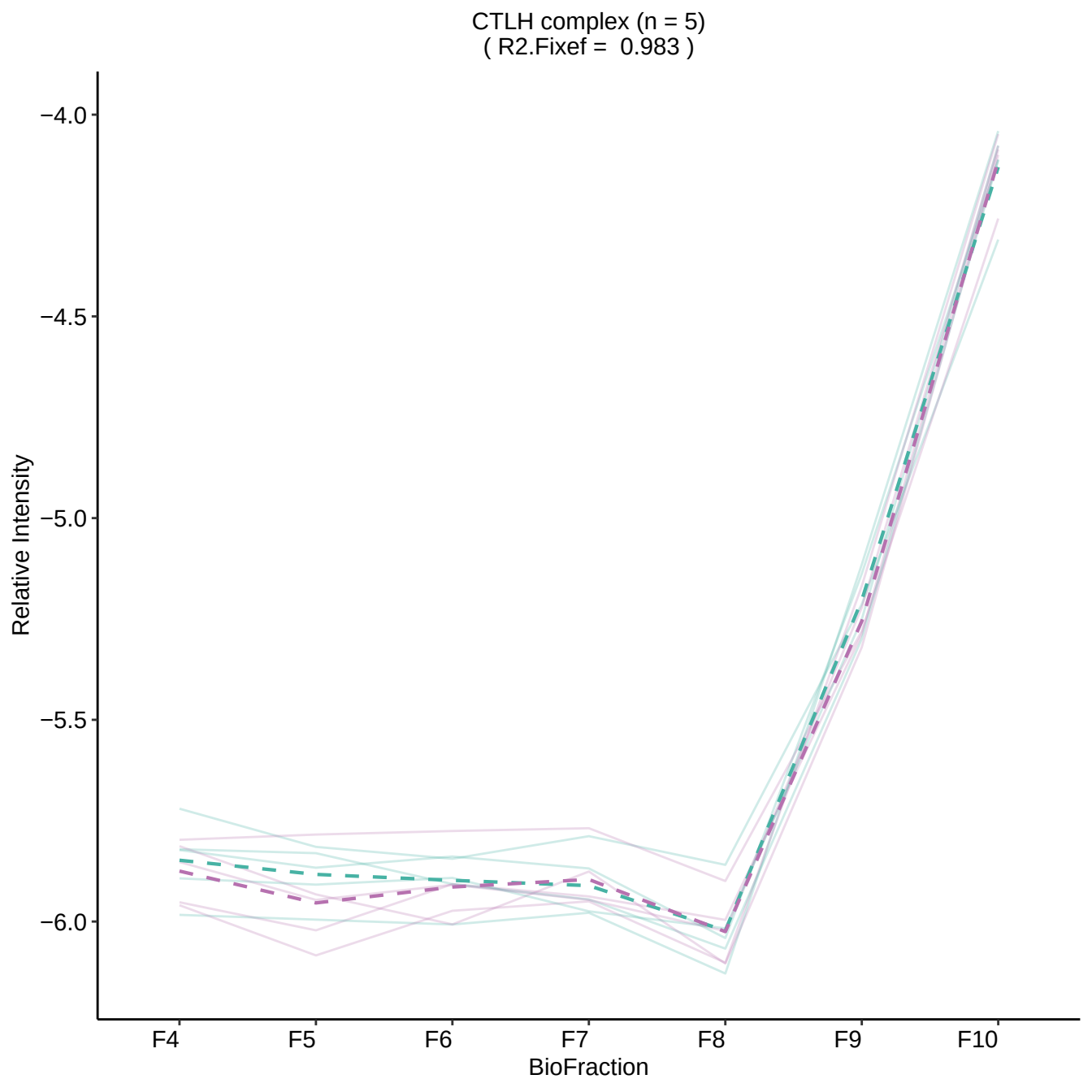

CCR4−NOT−CNOT7−CNOT6L complex (n = 6)
(R2.Fixef = 0.983)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
−6.5
F4
F5
F6
F7
F8
F9
F10
BioFraction

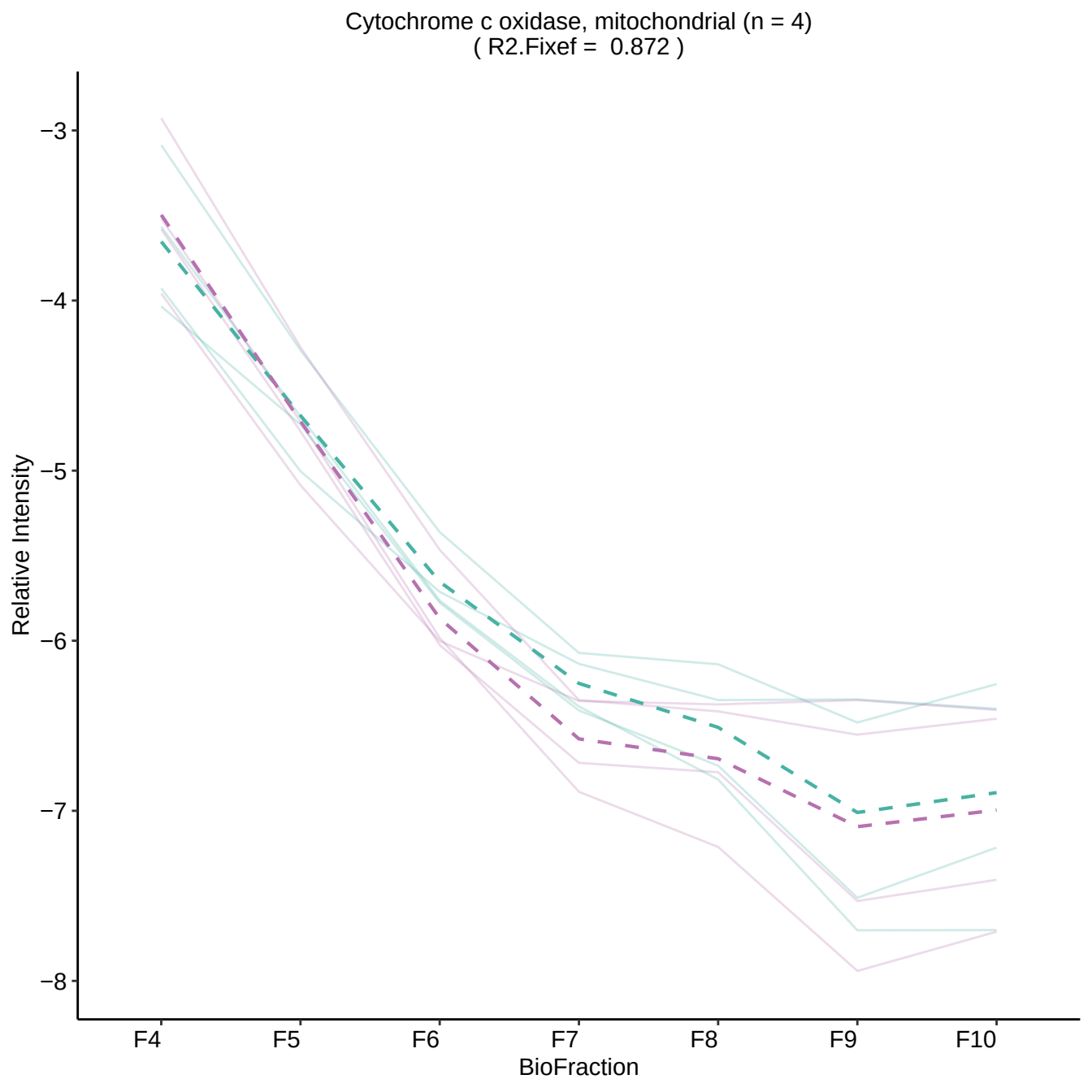

COMMD1−CCDC22−CCDC93−C16orf62 complex (n = 4)
(R2.Fixef = 0.977)
−5.0
−5.2
−5.4
−5.6
−5.8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

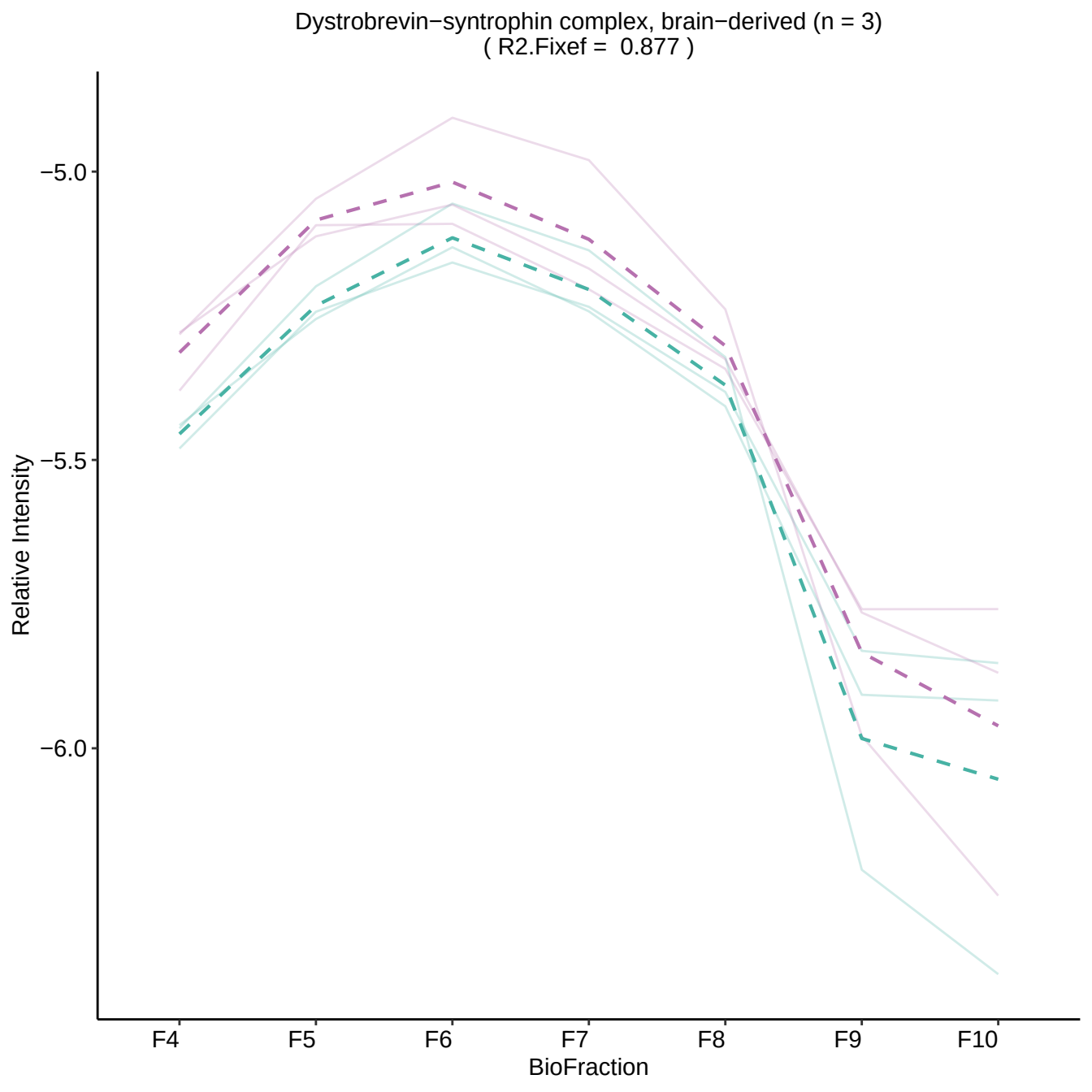

COMMD1−CCDC93−C16orf62 complex (n = 3)
(R2.Fixef = 0.979)
−5.0
−5.2
−5.4
−5.6
−5.8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

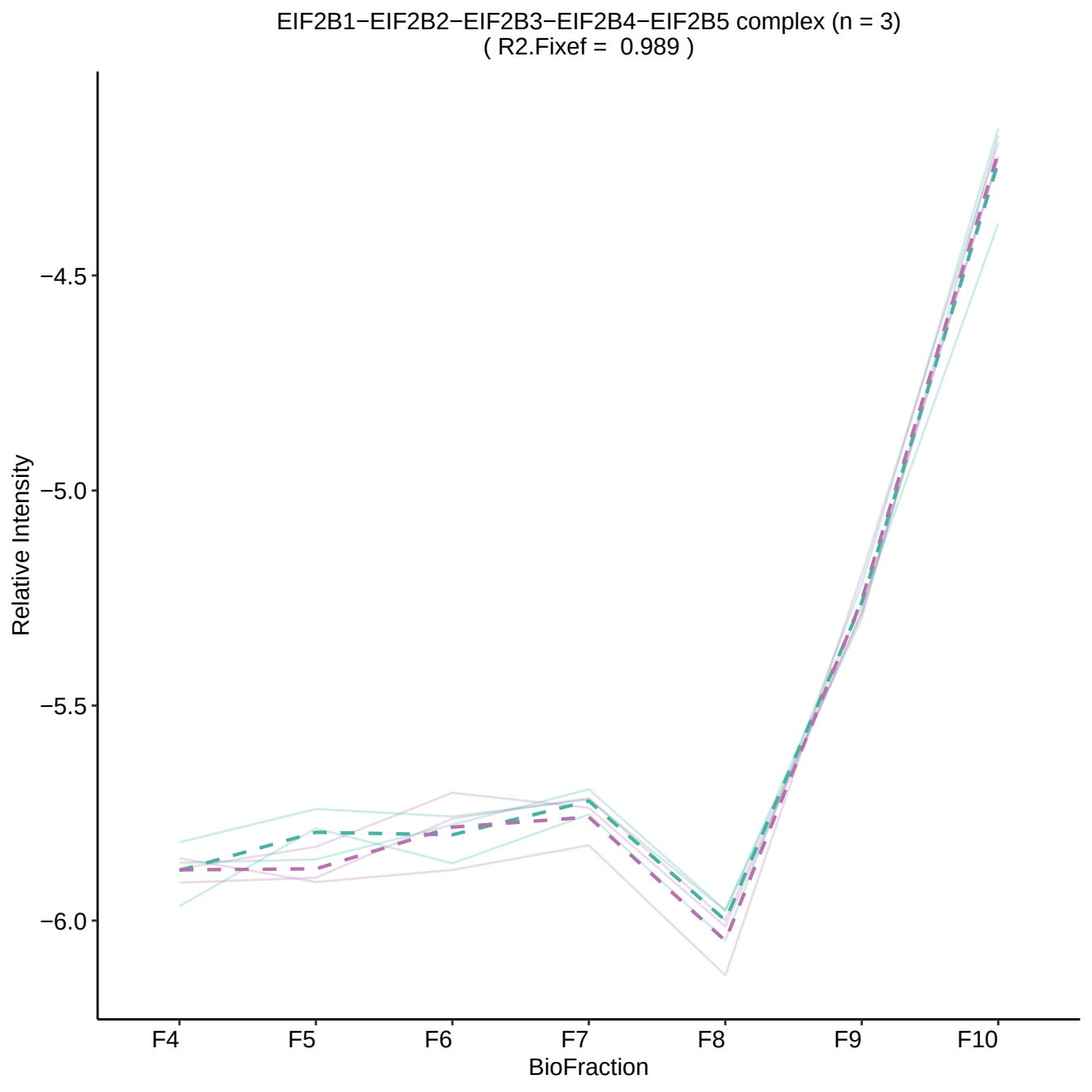

CORVET complex (n = 3)
(R2.Fixef = 0.976)
−5.1
−5.2
−5.3
−5.4
−5.5
−5.6
−5.7
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

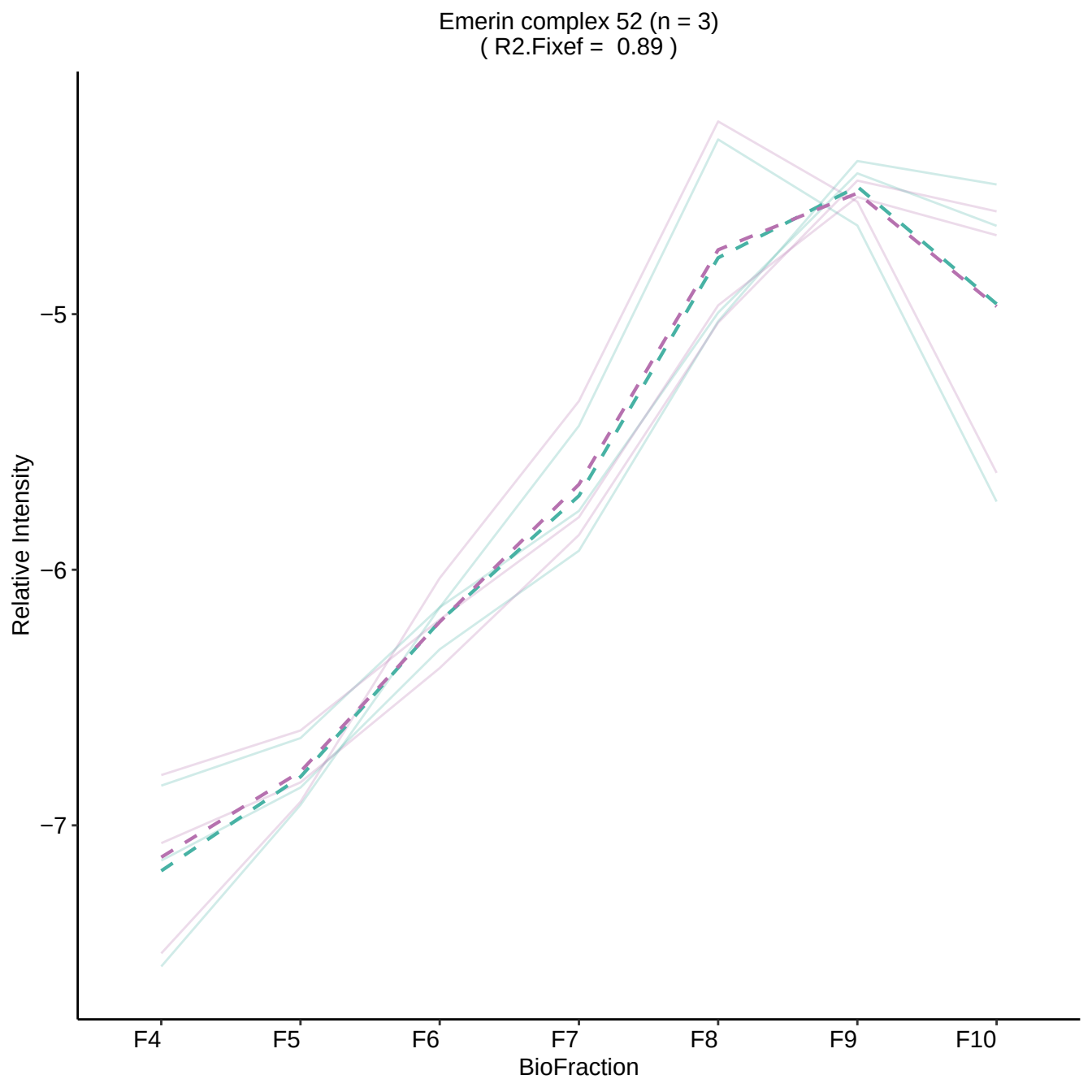

CSA−POLIIa complex (n = 4)
(R2.Fixef = 0.938)
−5.0
−5.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

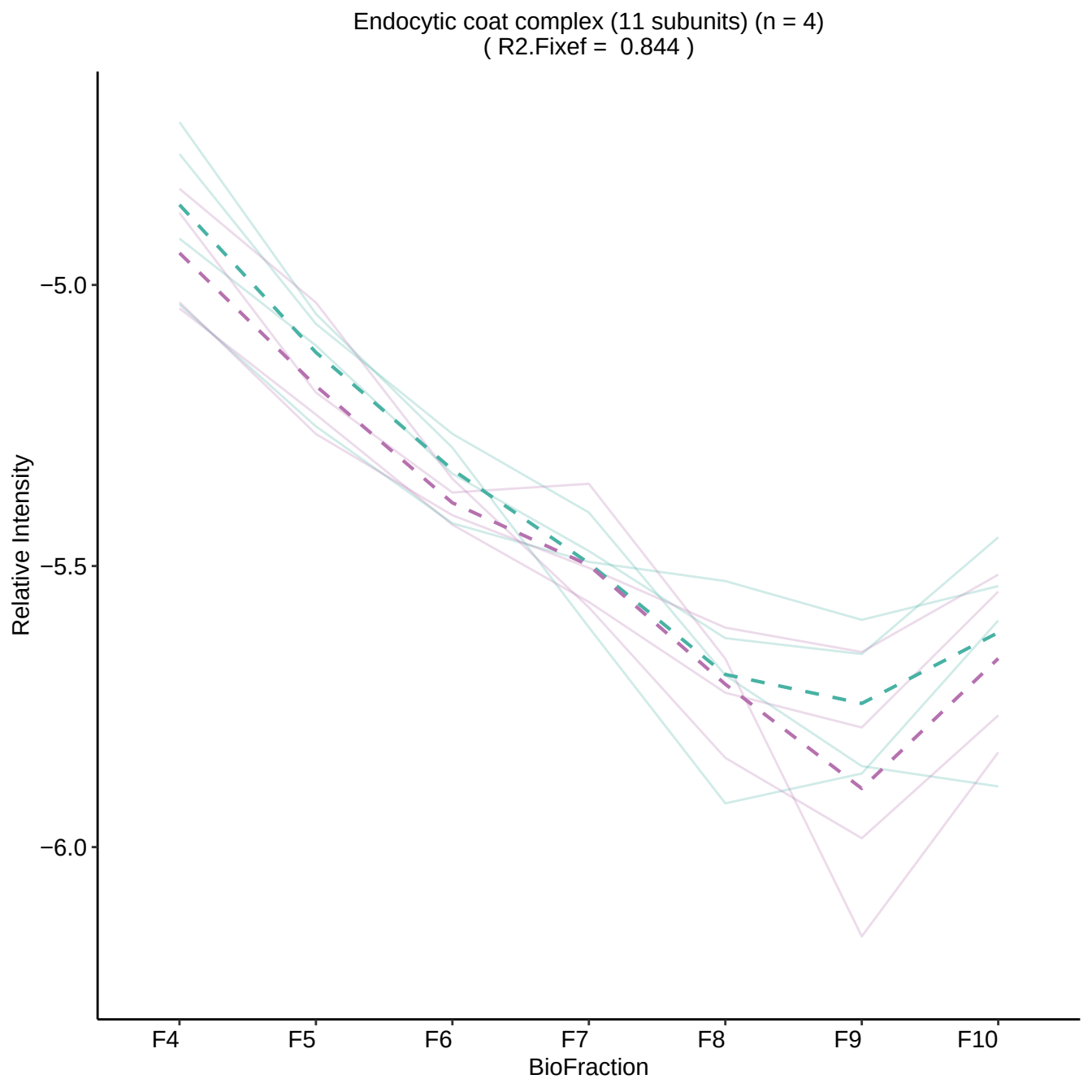

CTLH complex (n = 5)
(R2.Fixef = 0.983)
−4.0
−4.5
−5.0
−5.5
−6.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

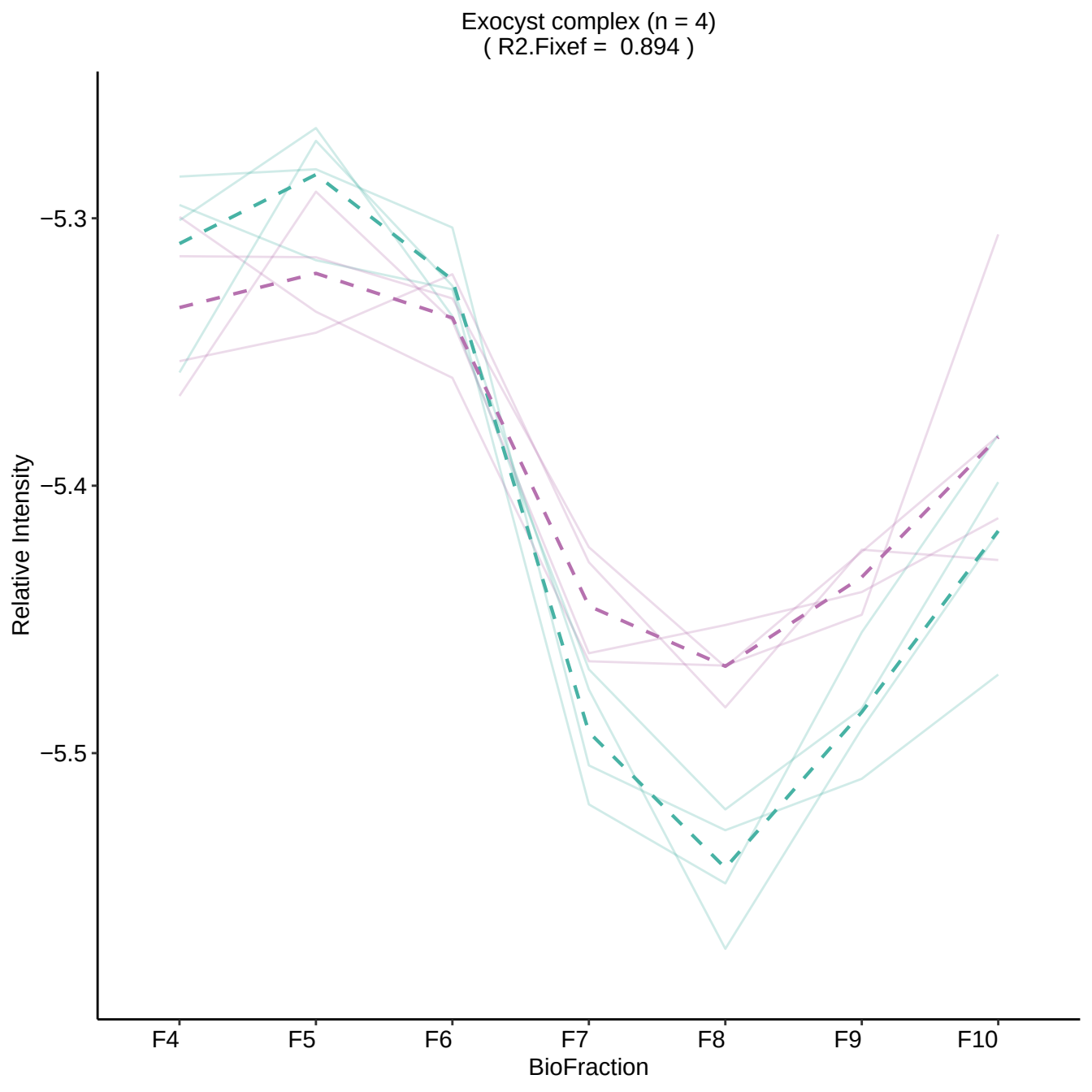

Cytochrome c oxidase, mitochondrial (n = 4)
(R2.Fixef = 0.872)
Relative Intensity
−3
−4
−5
−6
−7
−8
F4
F5
F6
F7
F8
F9
F10
BioFraction

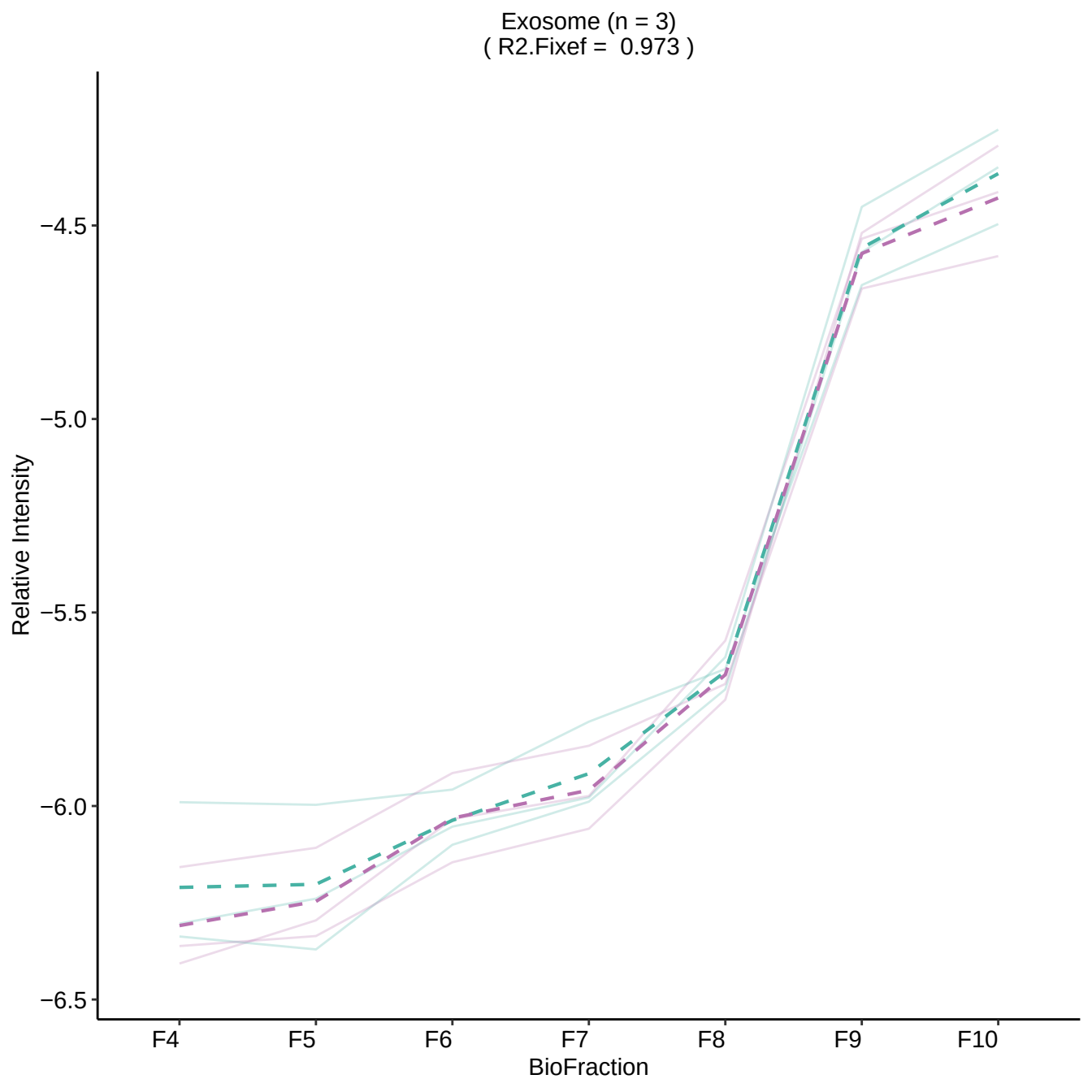

Dystrobrevin−syntrophin complex, brain−derived (n = 3)
(R2.Fixef = 0.877)
−5.0
−5.5
−6.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

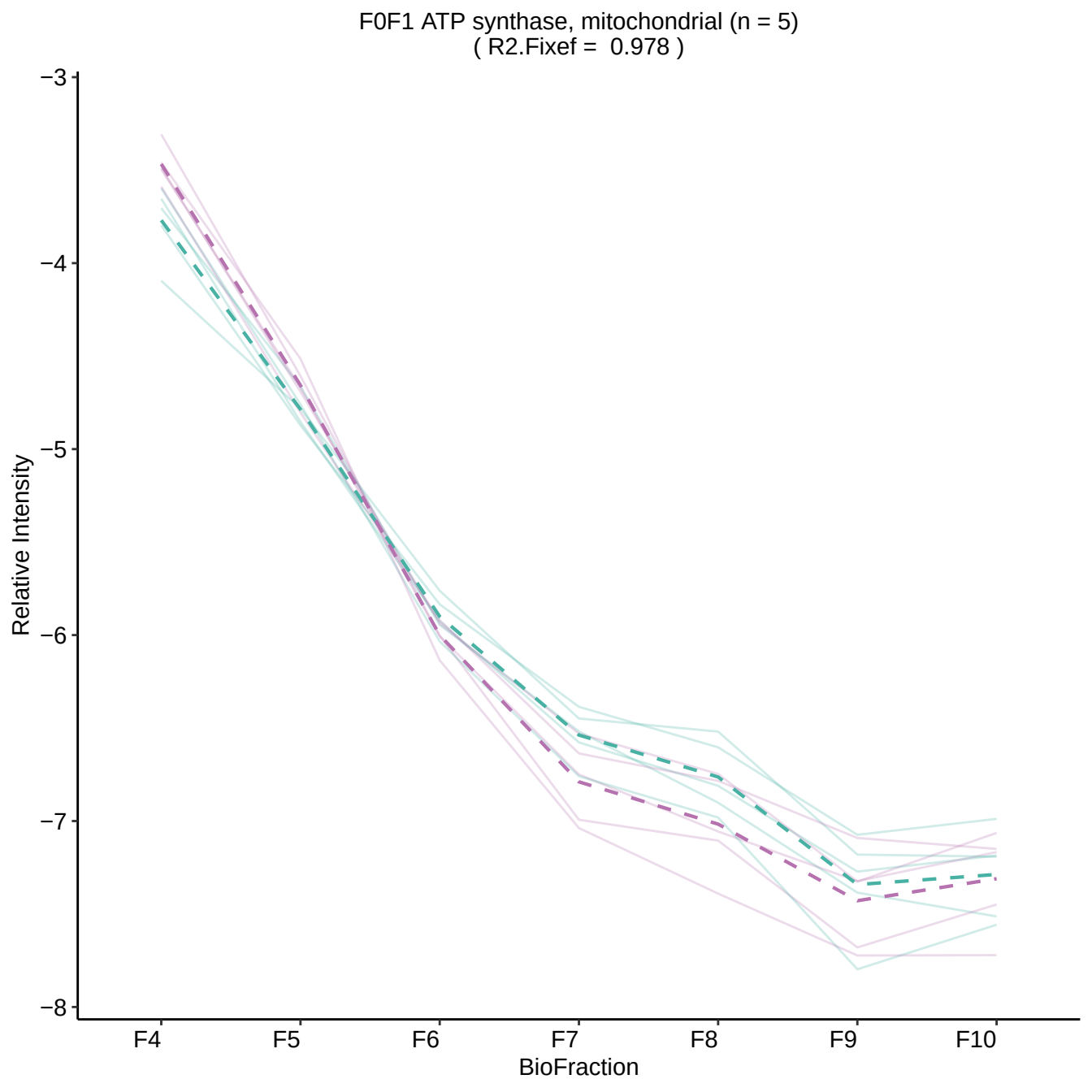

EIF2B1−EIF2B2−EIF2B3−EIF2B4−EIF2B5 complex (n = 3)
(R2.Fixef = 0.989)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

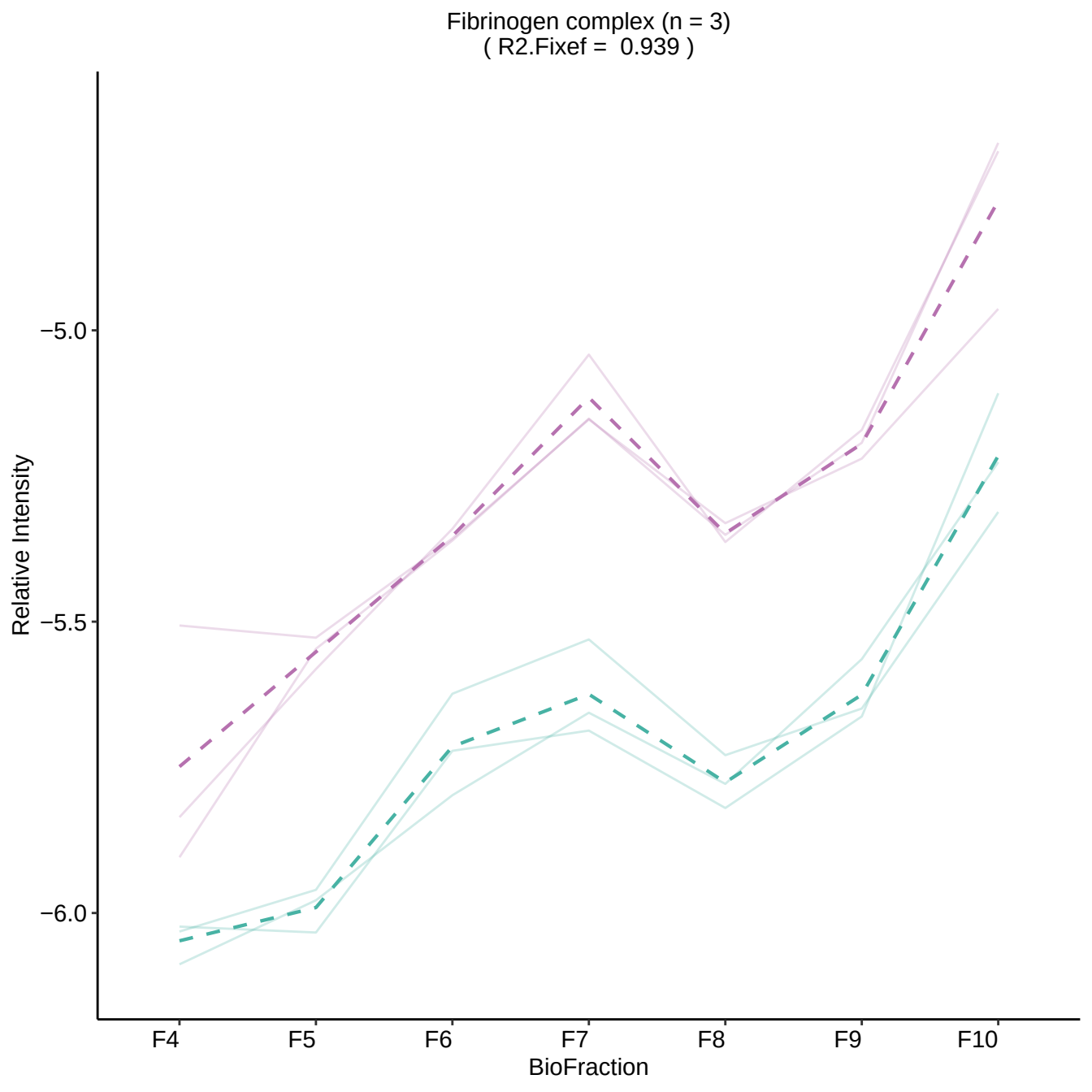

Emerin complex 52 (n = 3)
(R2.Fixef = 0.89)
Relative Intensity
−5
−6
−7
F4
F5
F6
F7
F8
F9
F10
BioFraction

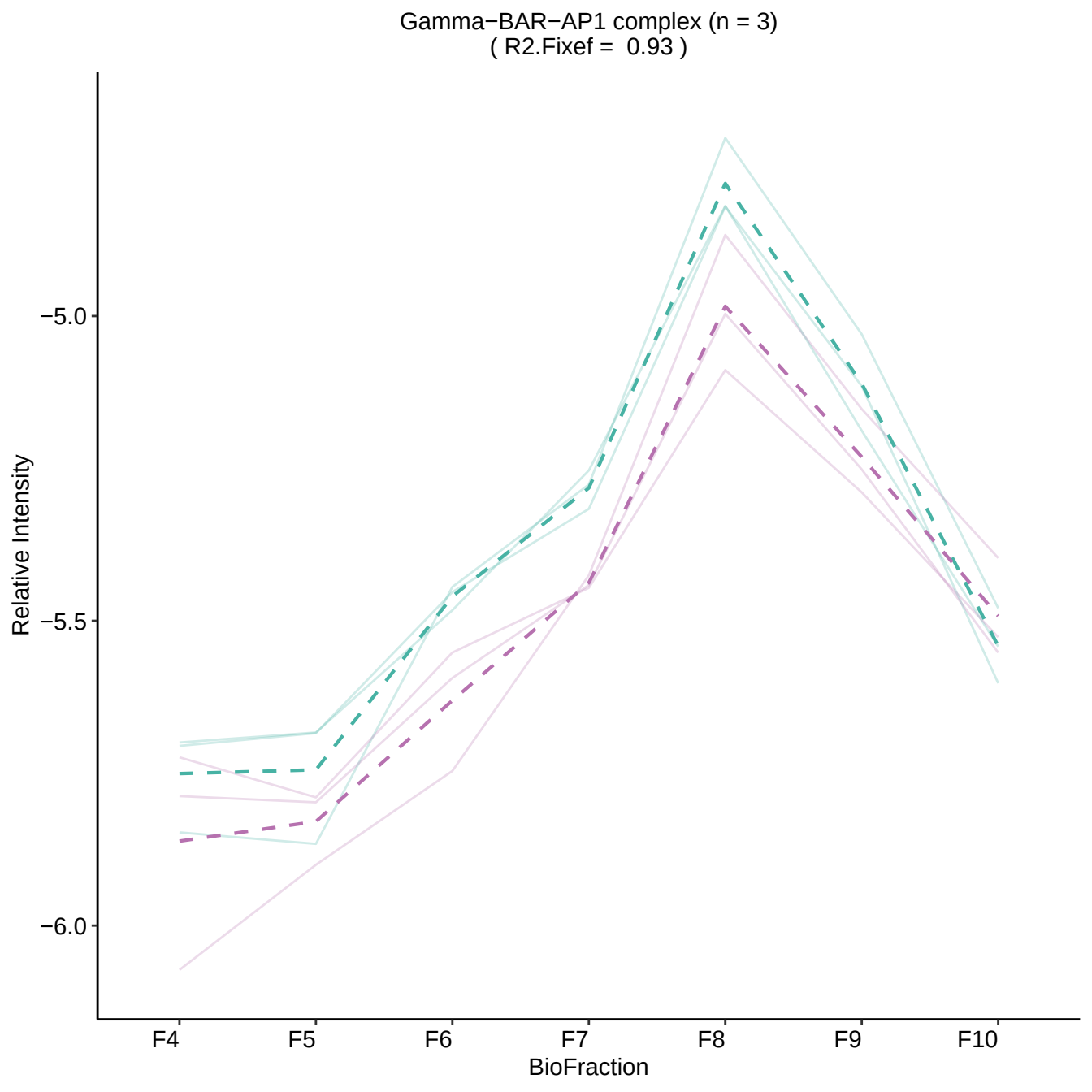

Endocytic coat complex (11 subunits) (n = 4)
(R2.Fixef = 0.844)
−5.0
−5.5
−6.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

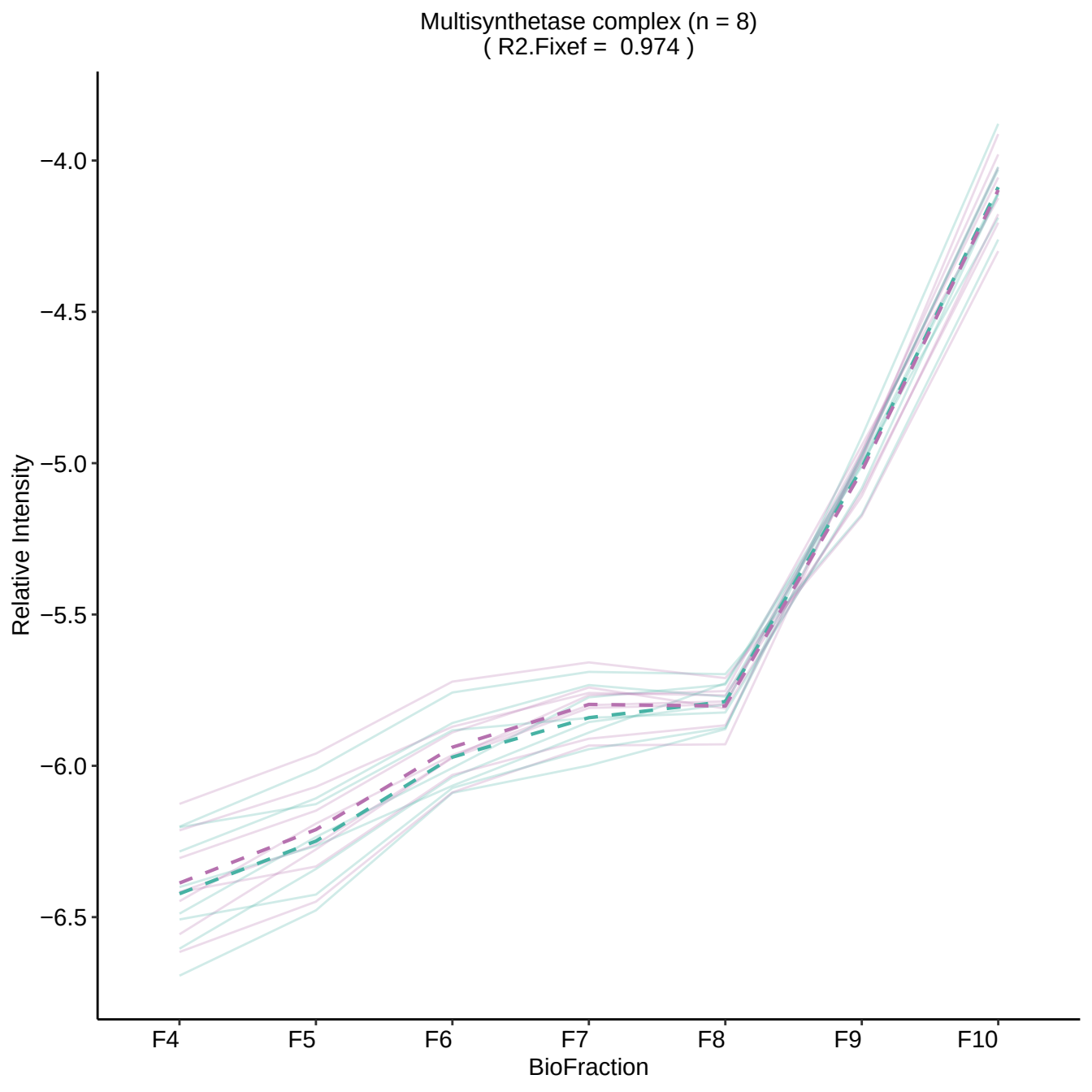

Exocyst complex (n = 4)
(R2.Fixef = 0.894)
−5.3
−5.4
−5.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

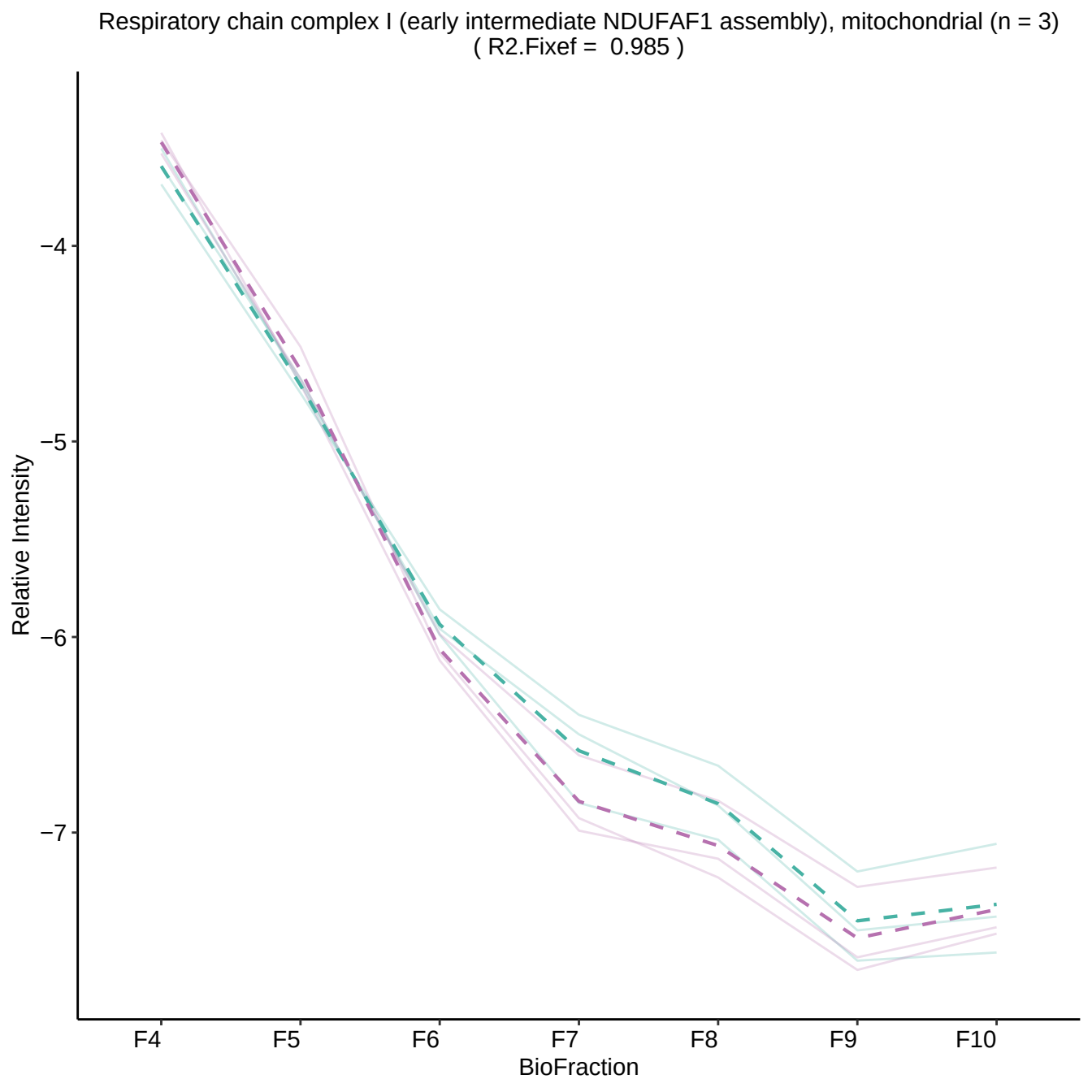

Exosome (n = 3)
(R2.Fixef = 0.973)
−4.5
−5.0
−5.5
−6.0
−6.5
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

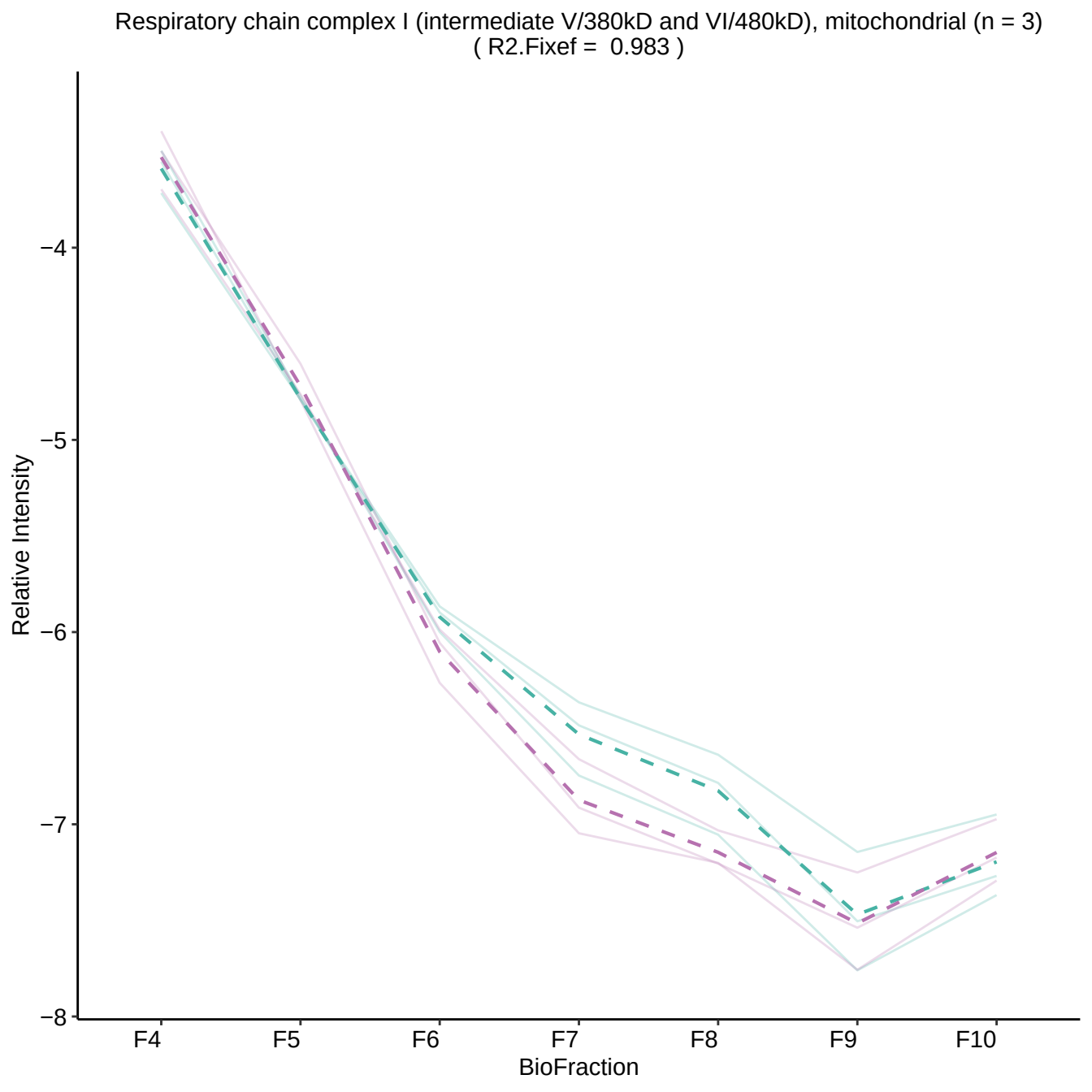

F0F1 ATP synthase, mitochondrial (n = 5)
(R2.Fixef = 0.978)
−3
−4
−5
−6
−7
−8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

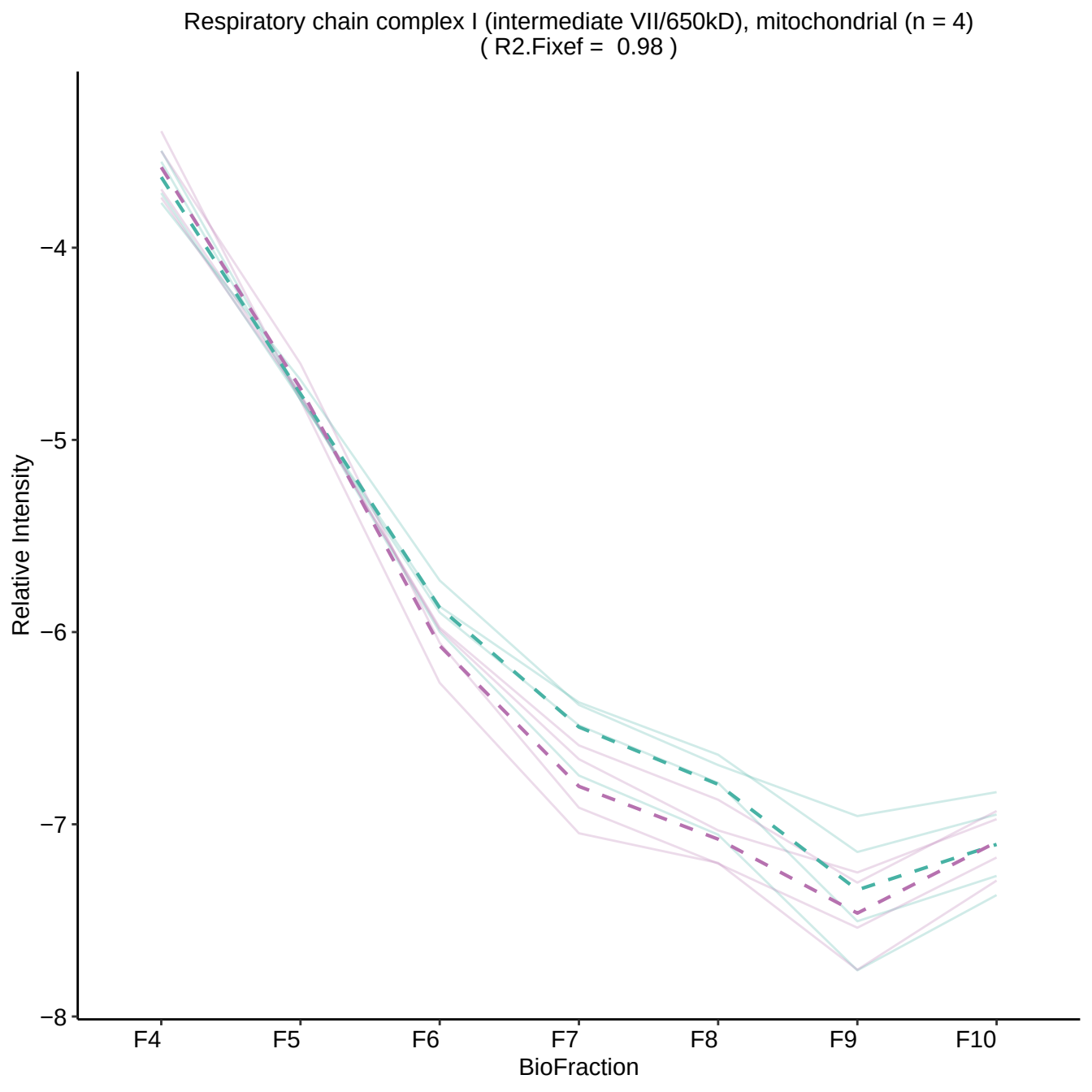

Fibrinogen complex (n = 3)
(R2.Fixef = 0.939)
Relative Intensity
−5.0
−5.5
−6.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

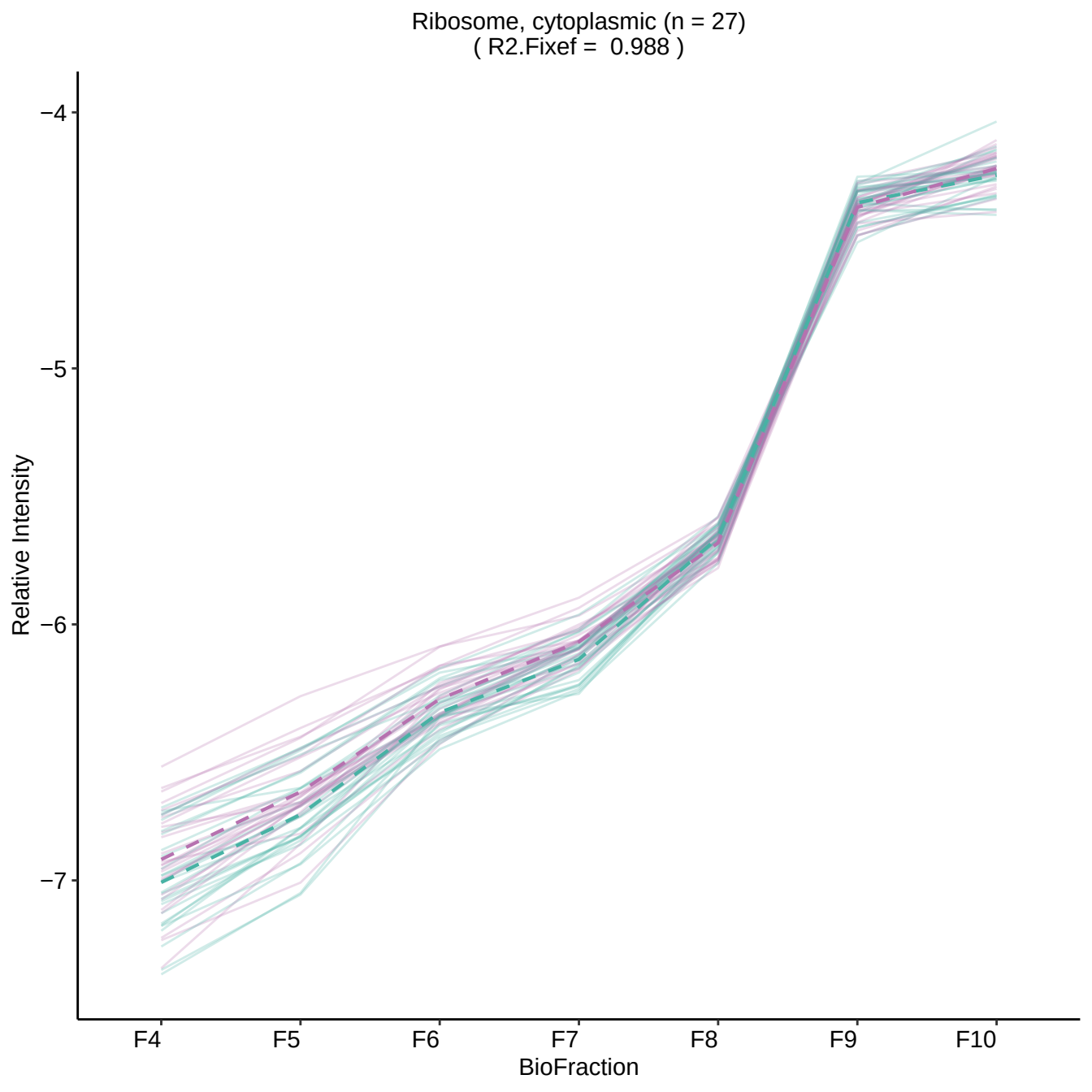

Gamma−BAR−AP1 complex (n = 3)
(R2.Fixef = 0.93)
Relative Intensity
−5.0
−5.5
−6.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

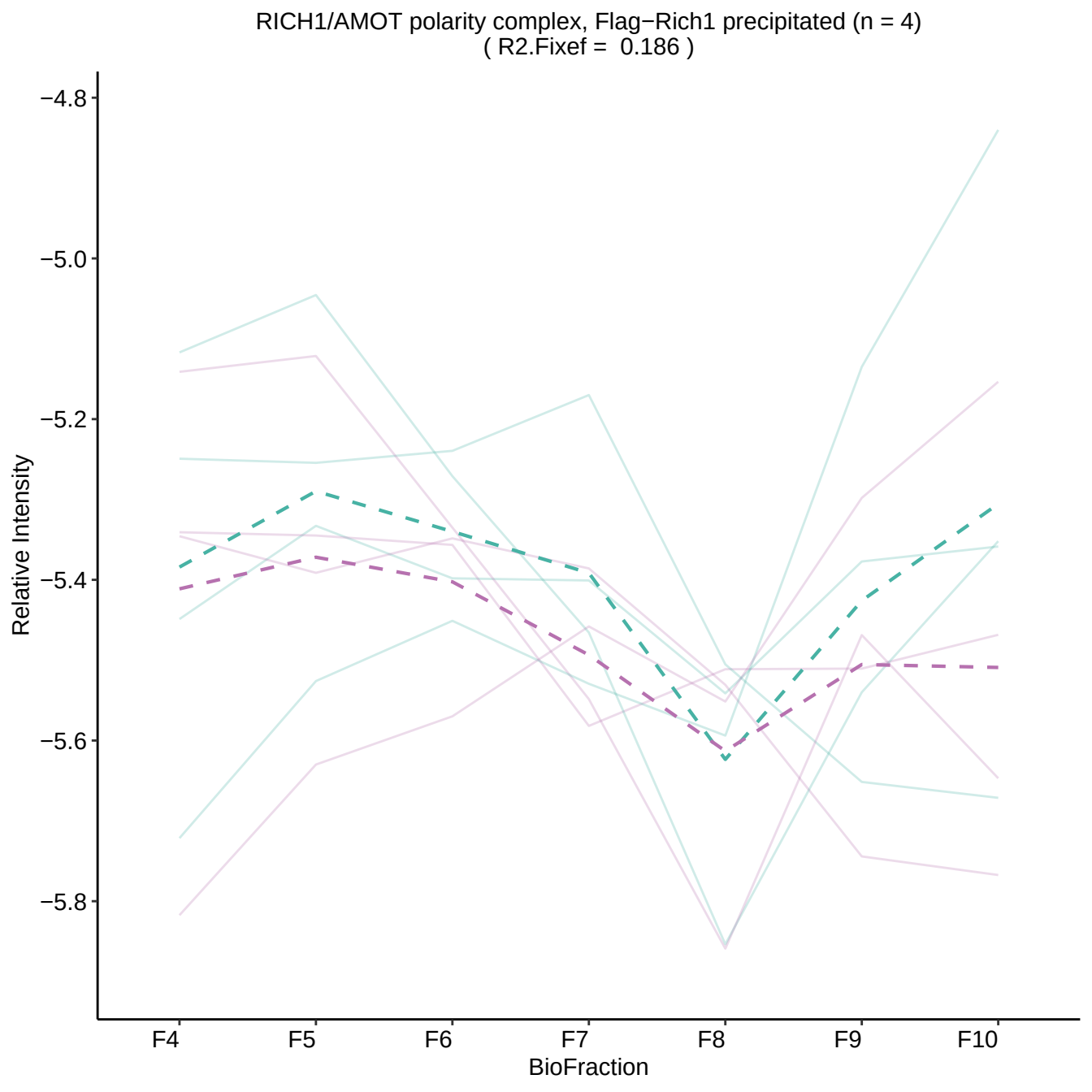

Multisynthetase complex (n = 8)
(R2.Fixef = 0.974)
Relative Intensity
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
F4
F5
F6
F7
F8
F9
F10
BioFraction

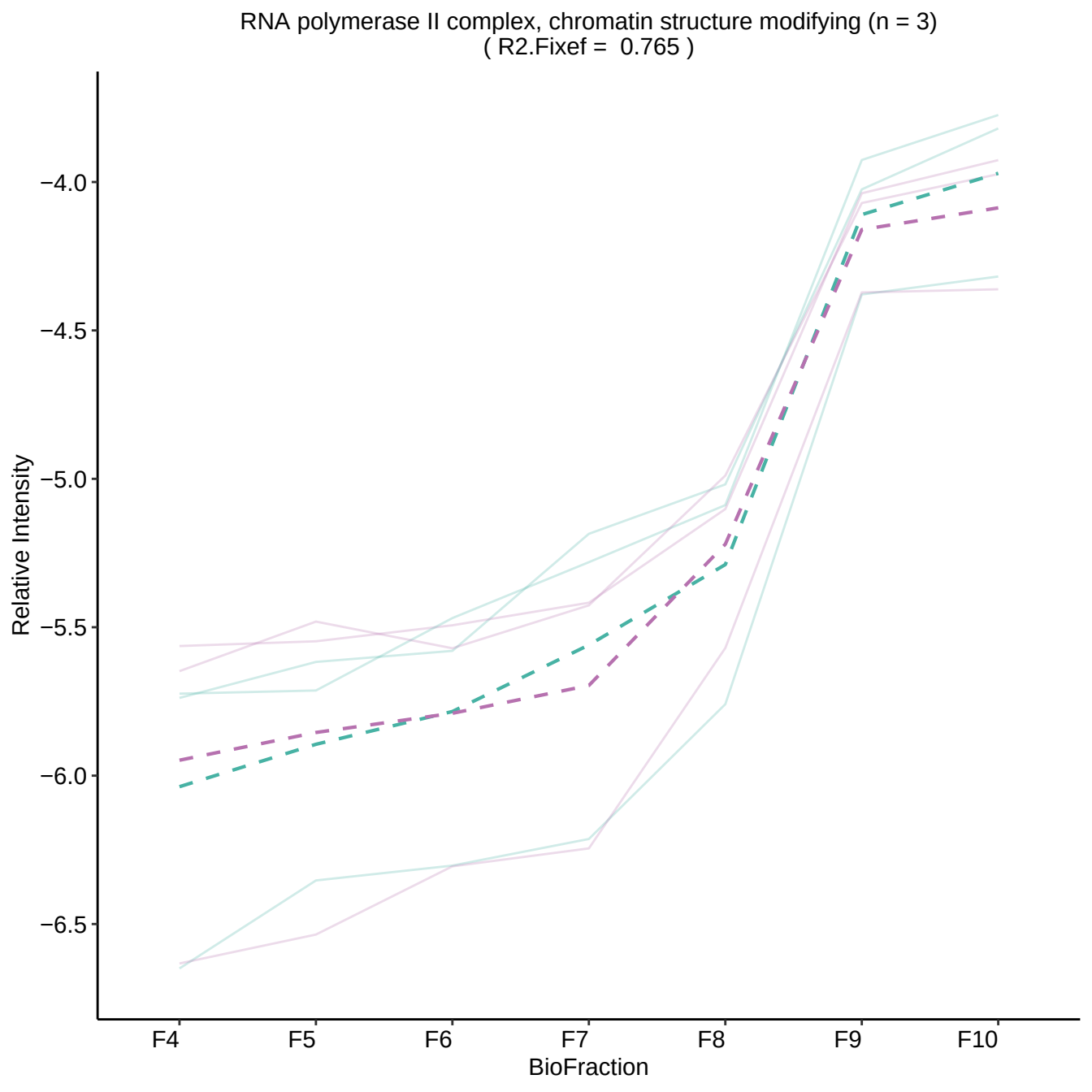

Respiratory chain complex I (early intermediate NDUFAF1 assembly), mitochondrial (n = 3)
(R2.Fixef = 0.985)
Relative Intensity
−4
−5
−6
−7
F4
F5
F6
F7
F8
F9
F10
BioFraction

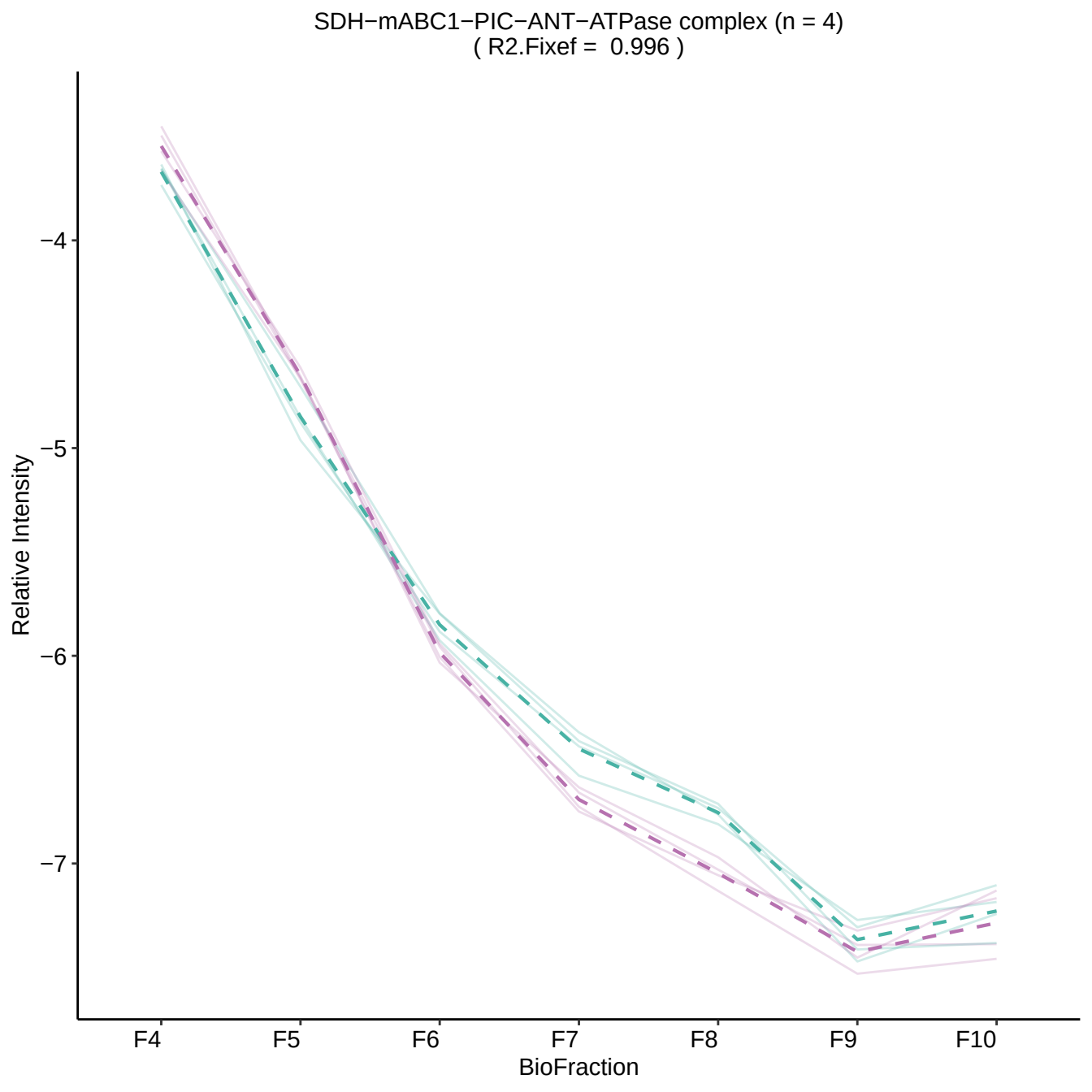

Respiratory chain complex I (intermediate V/380kD and VI/480kD), mitochondrial (n = 3)
(R2.Fixef = 0.983)
Relative Intensity
−4
−5
−6
−7
−8
F4
F5
F6
F7
F8
F9
F10
BioFraction

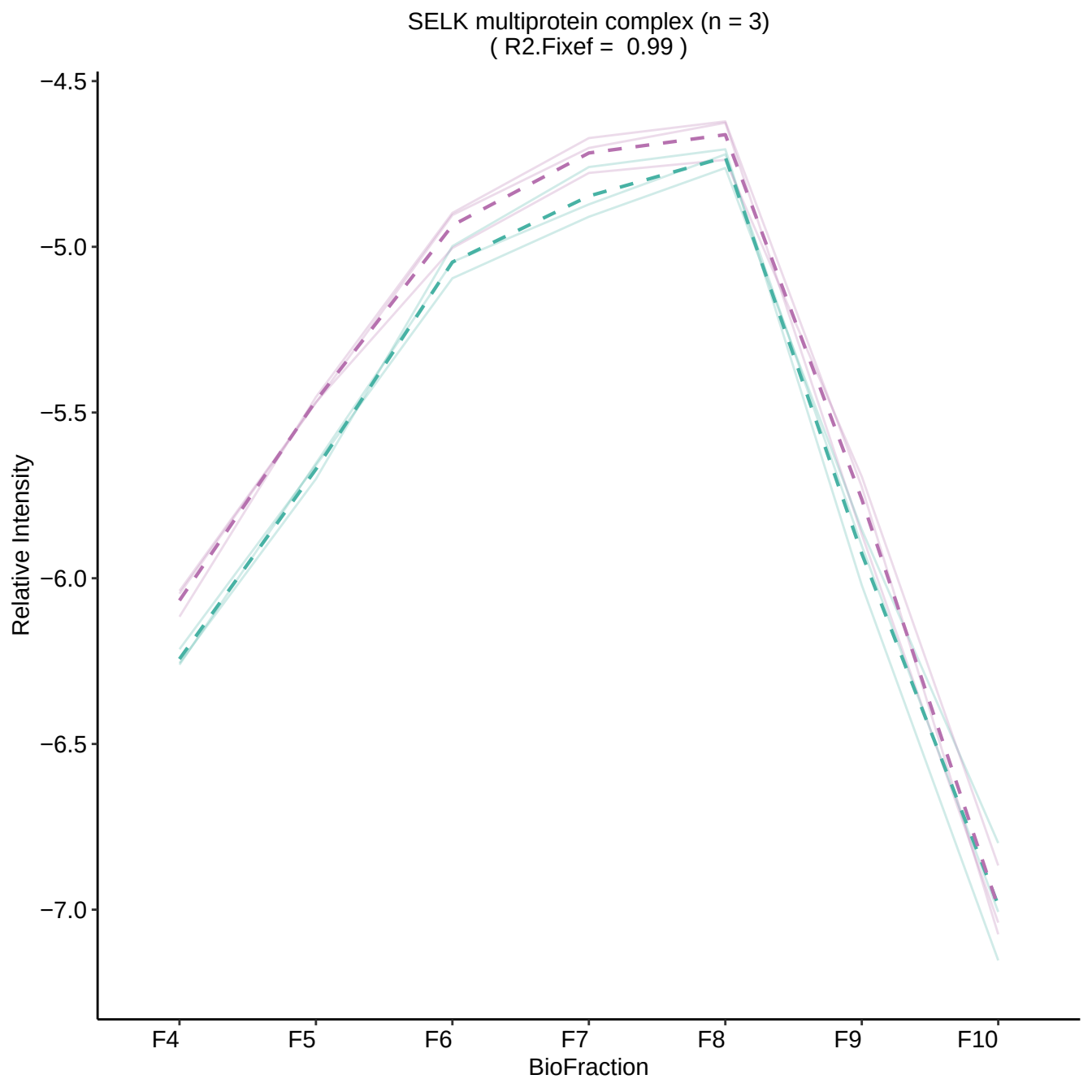

Respiratory chain complex I (intermediate VII/650kD), mitochondrial (n = 4)
(R2.Fixef = 0.98)
Relative Intensity
−4
−5
−6
−7
−8
F4
F5
F6
F7
F8
F9
F10
BioFraction

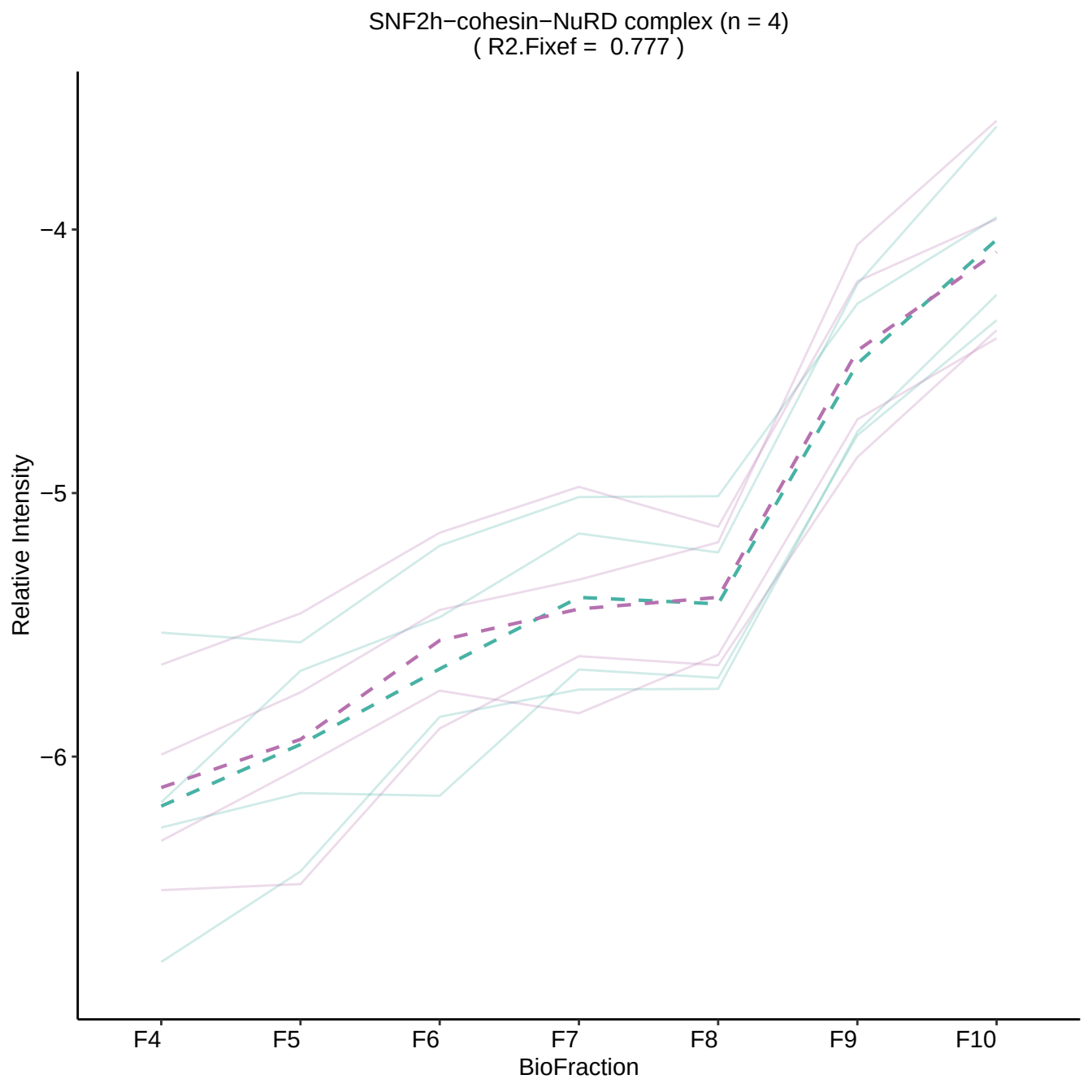

Ribosome, cytoplasmic (n = 27)
(R2.Fixef = 0.988)
−4
−5
−6
−7
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

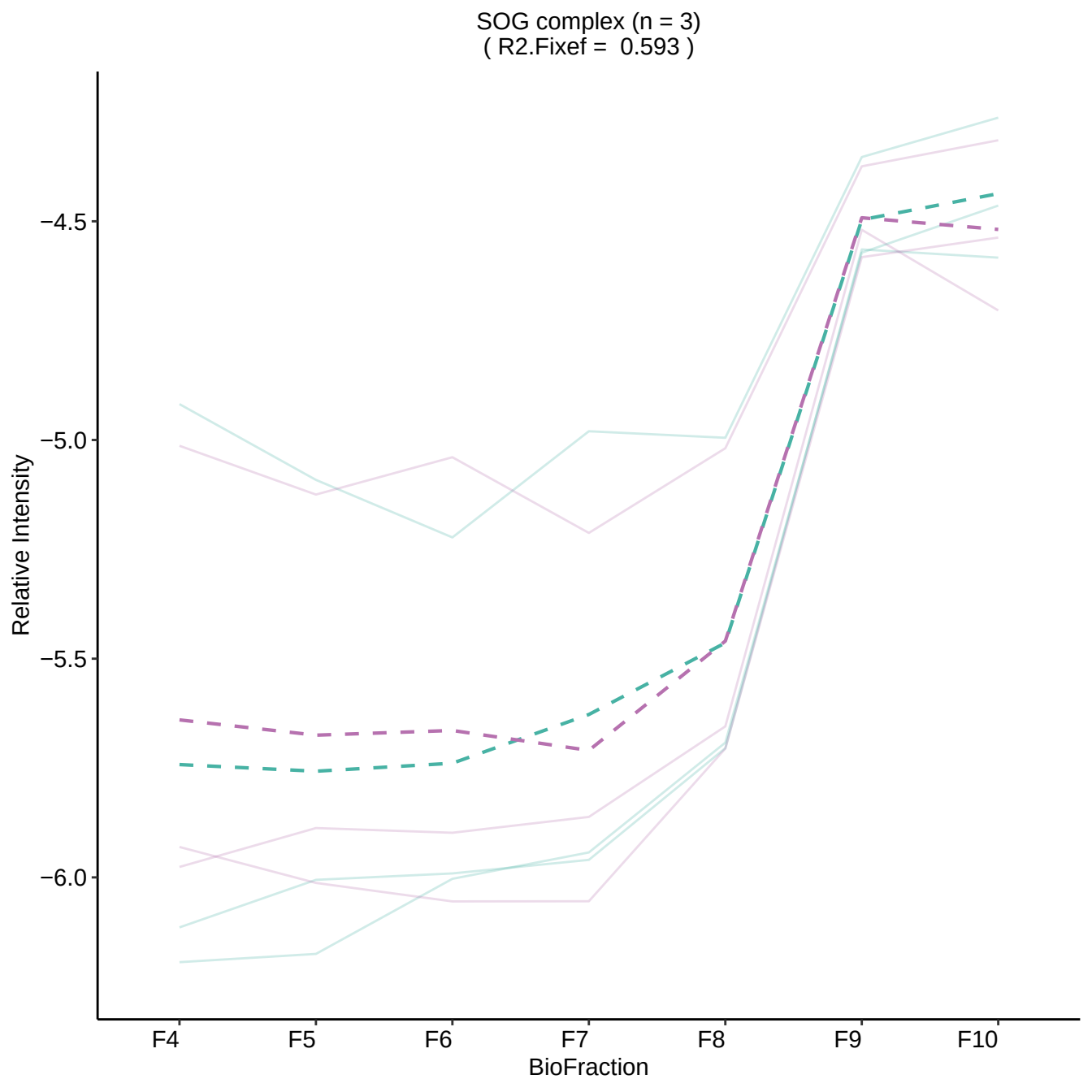

RICH1/AMOT polarity complex, Flag−Rich1 precipitated (n = 4)
(R2.Fixef = 0.186)
−4.8
−5.0
−5.2
−5.4
−5.6
−5.8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

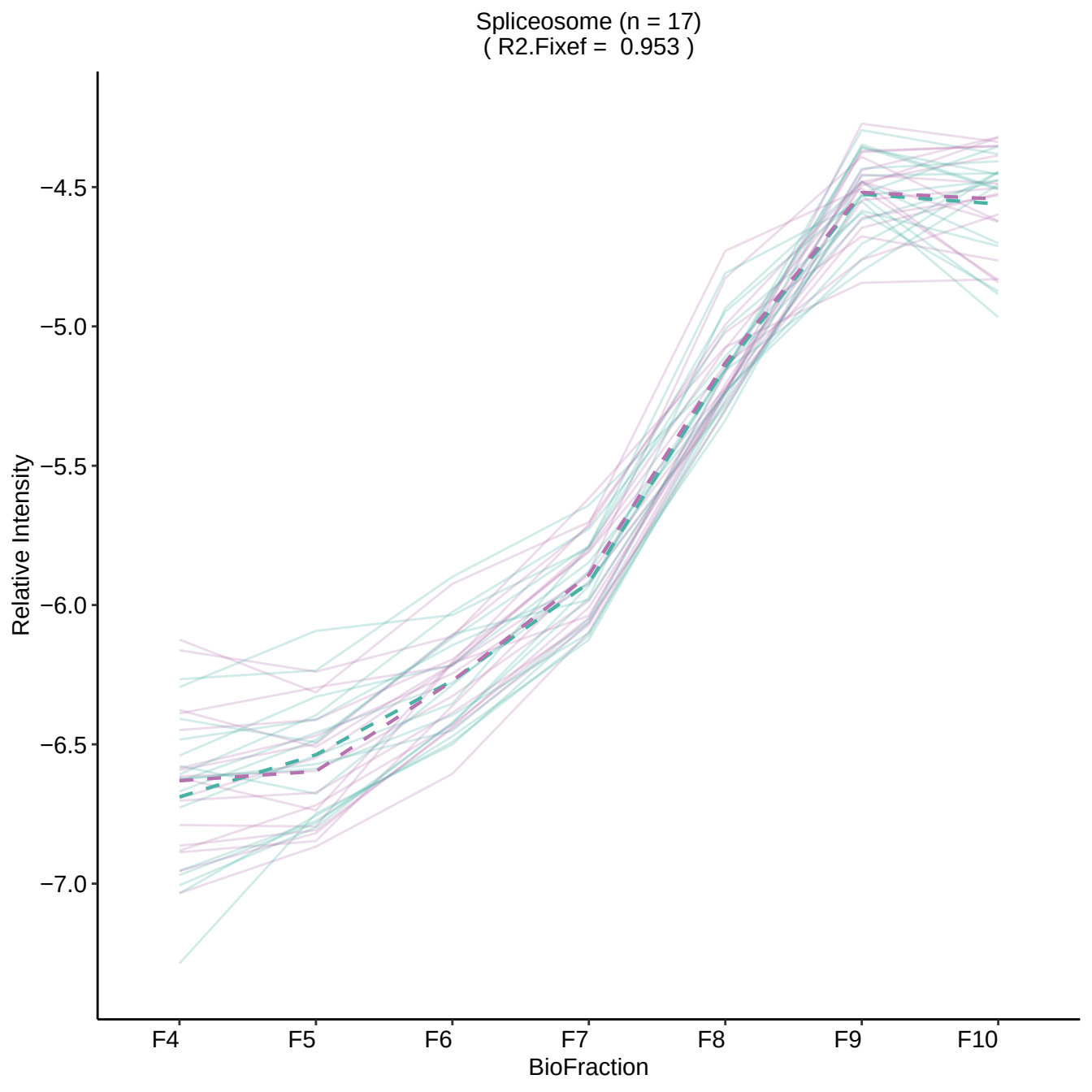

RNA polymerase II complex, chromatin structure modifying (n = 3)
(R2.Fixef = 0.765)
Relative Intensity
−4.0
−4.5
−5.0
−5.5
−6.0
−6.5
F4
F5
F6
F7
F8
F9
F10
BioFraction

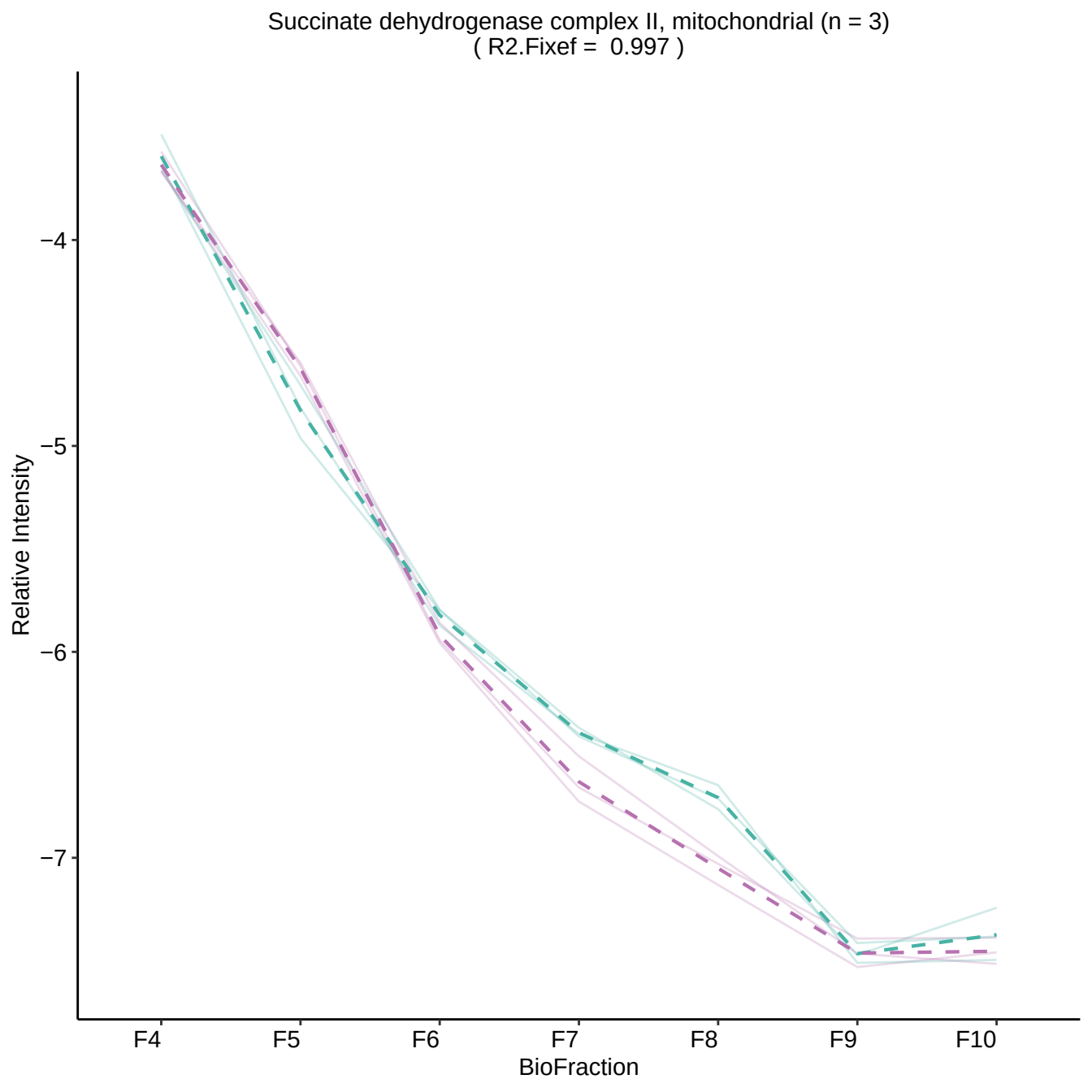

SDH−mABC1−PIC−ANT−ATPase complex (n = 4)
(R2.Fixef = 0.996)
Relative Intensity
−4
−5
−6
−7
F4
F5
F6
F7
F8
F9
F10
BioFraction

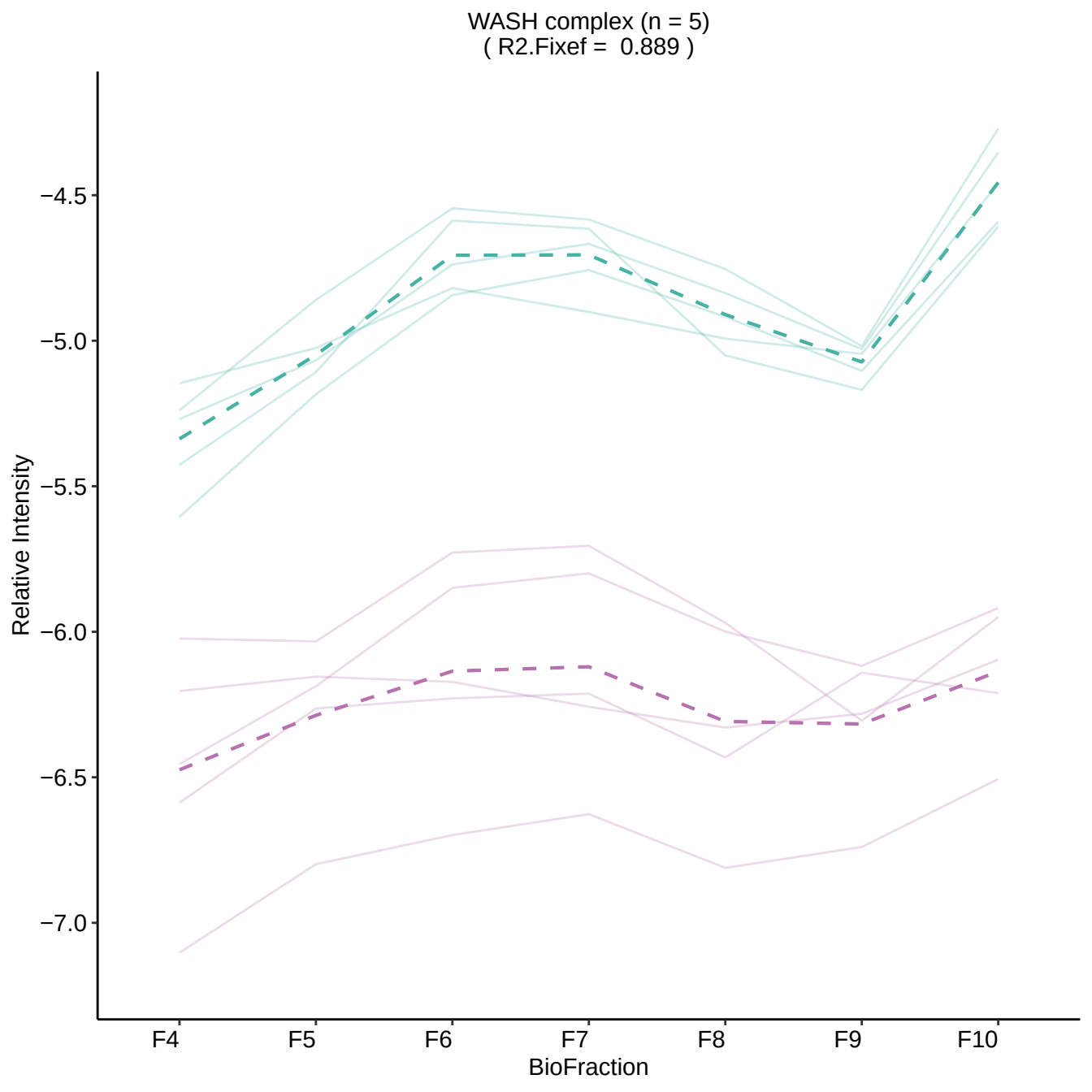

SELK multiprotein complex (n = 3)
(R2.Fixef = 0.99)
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction

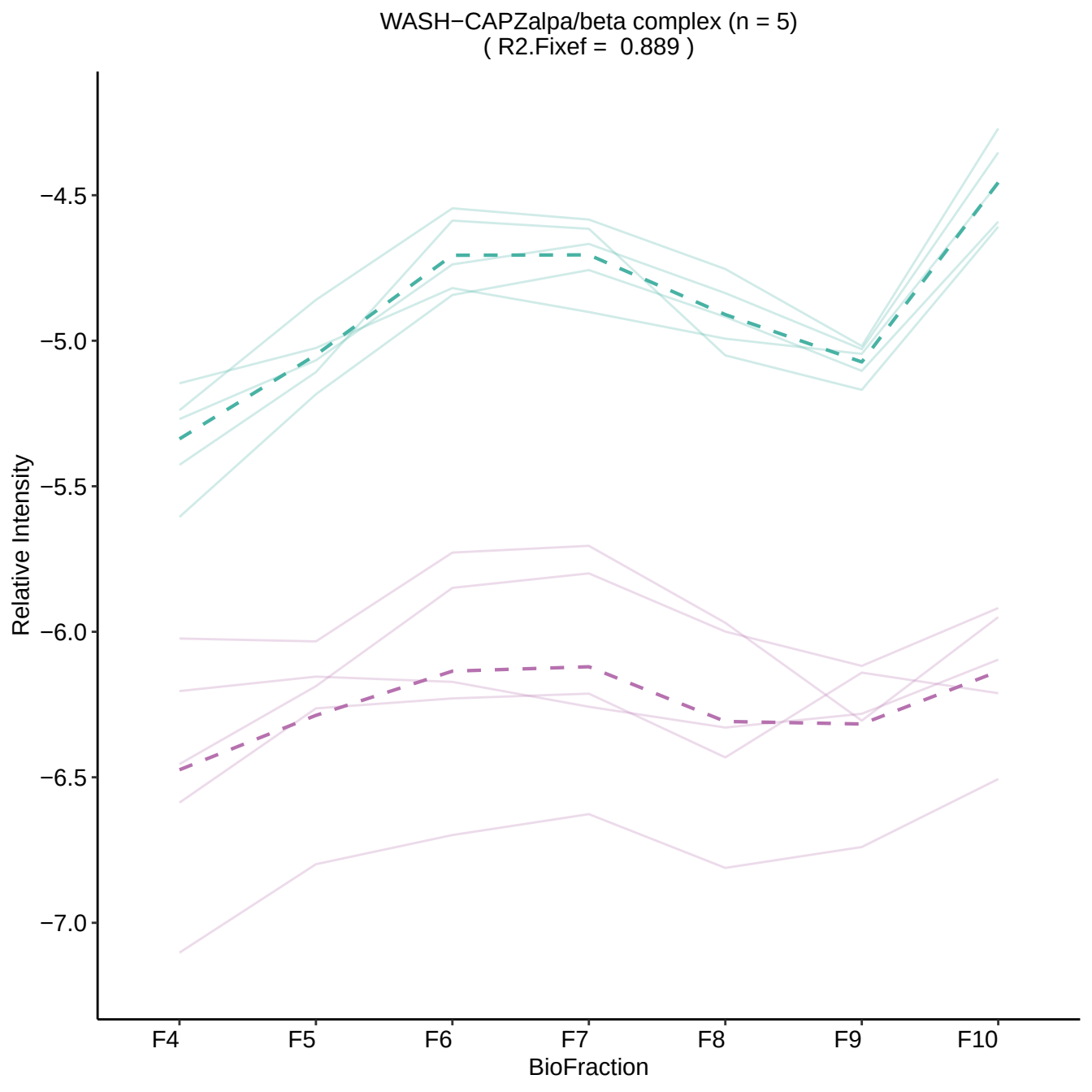

SNF2h−cohesin−NuRD complex (n = 4)
(R2.Fixef = 0.777)
Relative Intensity
−4
−5
−6
F4
F5
F6
F7
F8
F9
F10
BioFraction

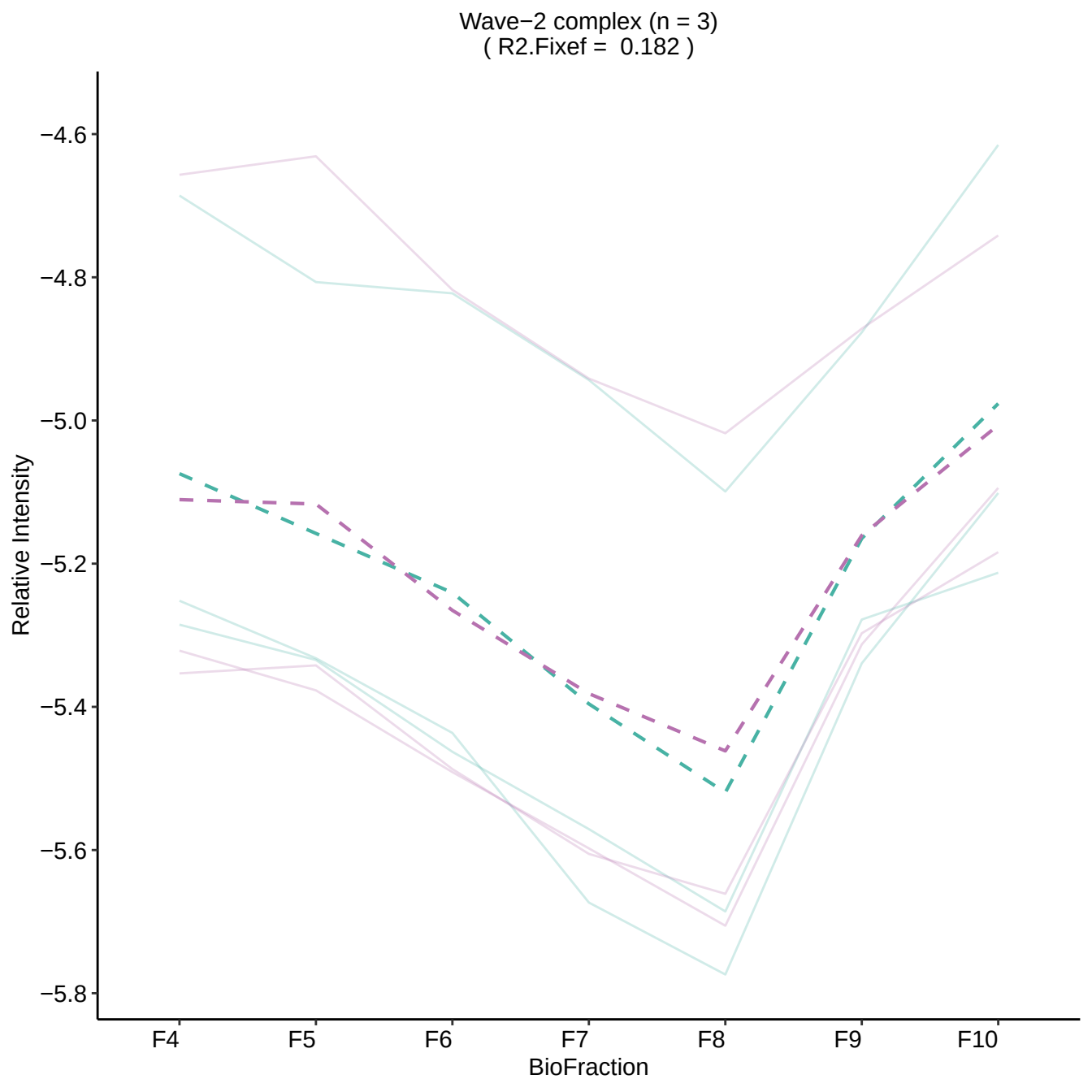

SOG complex (n = 3)
(R2.Fixef = 0.593)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

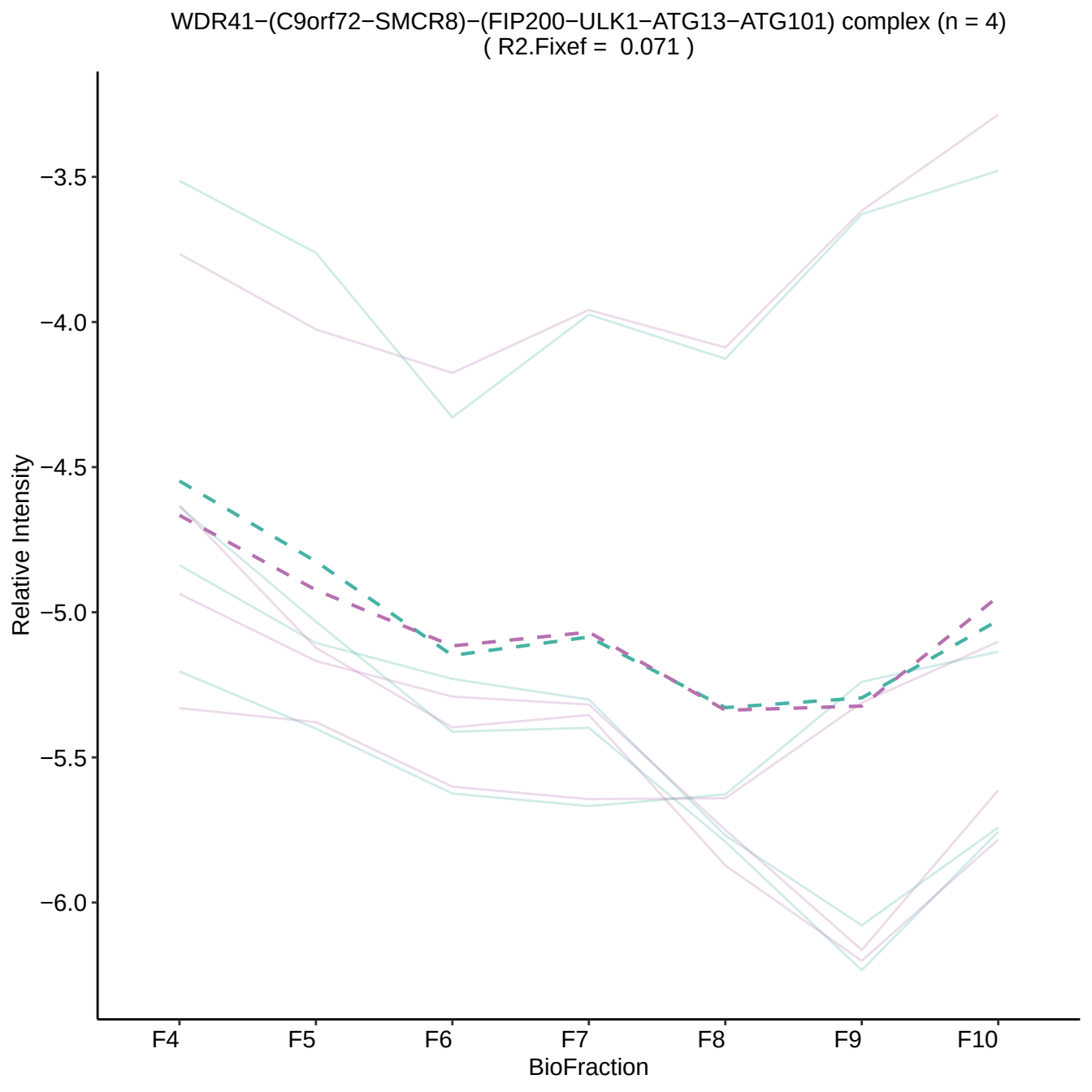

Spliceosome (n = 17)
(R2.Fixef = 0.953)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
F4
F5
F6
F7
F8
F9
F10
BioFraction

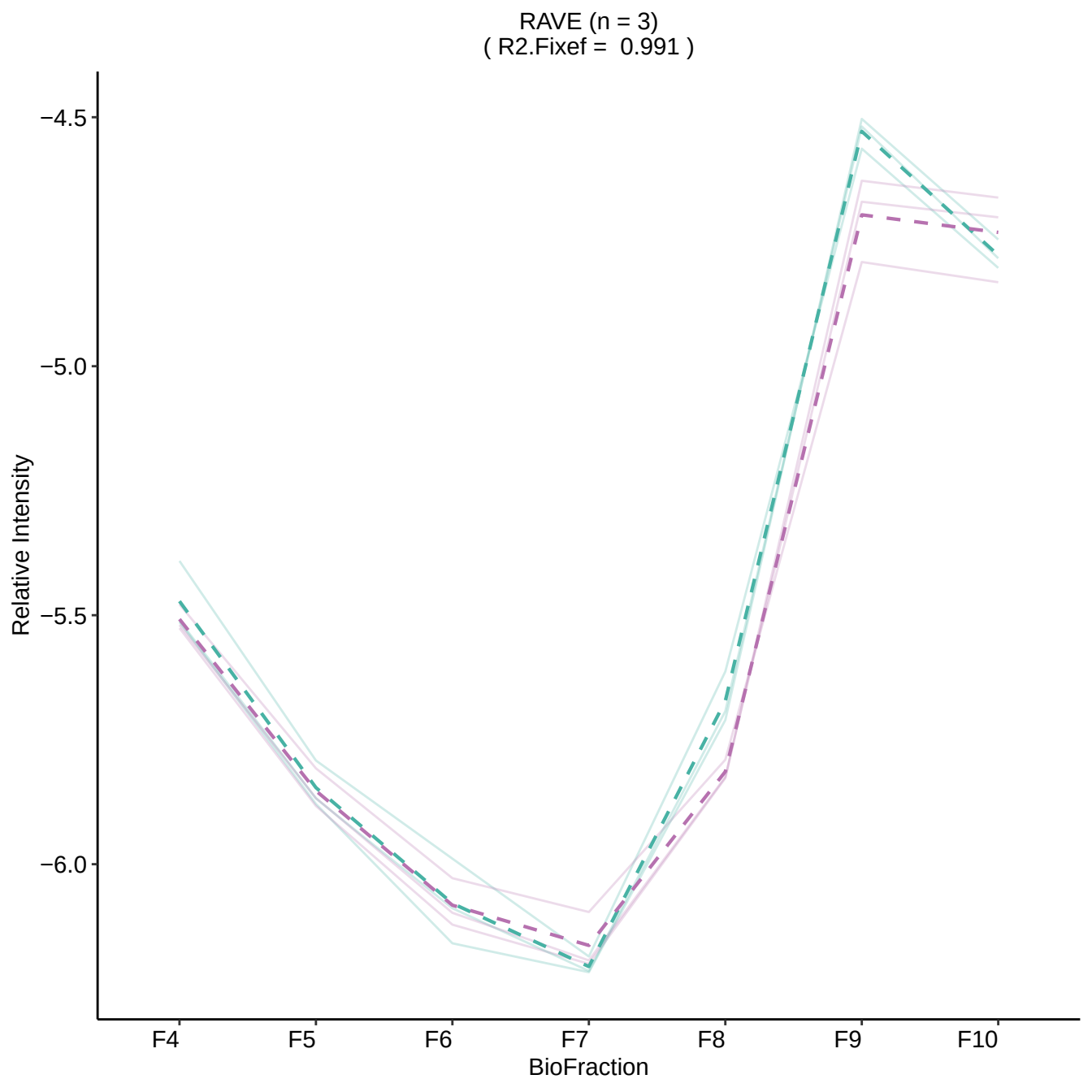

Succinate dehydrogenase complex II, mitochondrial (n = 3)
(R2.Fixef = 0.997)
Relative Intensity
−4
−5
−6
−7
F4
F5
F6
F7
F8
F9
F10
BioFraction


WASH complex (n = 5)
(R2.Fixef = 0.889)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
F4
F5
F6
F7
F8
F9
F10
BioFraction


WASH−CAPZalpa/beta complex (n = 5)
(R2.Fixef = 0.889)
Relative Intensity
−4.5
−5.0
−5.5
−6.0
−6.5
−7.0
F4
F5
F6
F7
F8
F9
F10
BioFraction


Wave−2 complex (n = 3)
(R2.Fixef = 0.182)
−4.6
−4.8
−5.0
−5.2
−5.4
−5.6
−5.8
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction


WDR41−(C9orf72−SMCR8)−(FIP200−ULK1−ATG13−ATG101) complex (n = 4)
(R2.Fixef = 0.071)
Relative Intensity
−3.5
−4.0
−4.5
−5.0
−5.5
−6.0
F4
F5
F6
F7
F8
F9
F10
BioFraction


RAVE (n = 3)
(R2.Fixef = 0.991)
−4.5
−5.0
−5.5
−6.0
Relative Intensity
F4
F5
F6
F7
F8
F9
F10
BioFraction