

Supplemental Table 1. Young IL 10tm mice metabolite measurements (full table, p-values unadjusted)

Metabolite	Control n=5	IL 10 tm n=6		P value
Cit	94.45	73.5167	↓	0.0036684
Orn	73.55	56.8167	↓	0.0100904
PC ae C38:2	4.95467	3.52205	↓	0.0103952
PC ae C38:3	1.97635	1.52756	↓	0.0129673
lysoPC a C14:0	8.68146	6.66934	↓	0.0206697
SM (OH) C14:1	1.72292	1.26118	↓	0.021117
PC ae C36:3	2.00604	1.7431	↓	0.0218773
lysoPC a C16:1	5.7321	4.32297	↓	0.025716
PC ae C36:2	9.64386	8.07114	↓	0.0276785
SM (OH) C22:2	1.85736	1.47848	↓	0.032493
PC aa C36:3	95.9061	81.4081	↓	0.0334162

Supplemental Table 2. Middle aged IL 10tm metabolite measurements (full table, p-values unadjusted)

Metabolite	Control n=12	IL 10 tm n=19		P value
Kynurenine / Trp *	0.0099273	0.0158913	↑	4.44E-05
PC ae C42:1 *	0.688724	0.927056	↑	0.0001366
PC aa C40:6	36.2821	53.1196	↑	0.0004424
PC aa C40:4	1.71318	2.33144	↑	0.0008275
PC aa C40:1	0.288054	0.346857	↑	0.0012507
lysoPC a C26:1	0.315676	0.434985	↑	0.0012549
PC ae C42:4	0.283372	0.385398	↑	0.0013905
PC ae C40:5	0.957944	1.27838	↑	0.0016083
PC ae C36:5	1.60583	2.41199	↑	0.001617
PC aa C42:5	0.226678	0.293754	↑	0.0019018
PC aa C32:3	0.177071	0.230479	↑	0.0023928
SM (OH) C14:1	1.80553	1.33645	↓	0.0027591
PC aa C34:3	10.4825	14.5773	↑	0.0028308
PC ae C36:4	2.50391	3.42155	↑	0.0028347
PC aa C38:4	116.879	156.955	↑	0.0032746
Kynurenine	1.21658	1.63516	↑	0.0035013
lysoPC a C28:1	0.636165	0.81616	↑	0.0035348

PC aa C38:6	76.1241	106.147	↑	0.0039363
PC aa C36:5	6.59467	9.30779	↑	0.0042725
PC ae C34:3	0.827801	1.14532	↑	0.0042915
PC ae C38:5	2.79781	3.75849	↑	0.0043954
PC ae C32:1	0.521774	0.664599	↑	0.0062371
PC ae C42:0	1.01761	1.17576	↑	0.0063958
PC ae C42:5	0.687499	0.817973	↑	0.0068315
PC aa C34:4	0.50192	0.662721	↑	0.0080307
PC ae C30:2	0.100309	0.119044	↑	0.0085778
PC ae C38:0	2.06255	2.77453	↑	0.0098392
Serotonin	9.05709	3.34308	↓	0.0125052
PC ae C30:1	0.208129	0.278508	↑	0.0126652
c4-OH-Pro	4.46417	5.58789	↑	0.0129227
PC aa C38:0	1.22125	1.56823	↑	0.0159307
PC aa C42:6	0.468822	0.552634	↑	0.0162391
PC aa C40:5	4.05247	5.21375	↑	0.0170496
PC ae C40:4	1.64796	2.09069	↑	0.0180223
PC ae C38:6	1.43781	1.84396	↑	0.0198495
PC ae C34:1	2.34154	2.89283	↑	0.019872
Gly	196.983	243.263	↑	0.0202694
PC aa C38:5	35.0897	44.7049	↑	0.0204442
PC ae C32:2	0.218689	0.26545	↑	0.0214631
PC aa C42:4	0.606987	0.665631	↑	0.022113
PC aa C36:4	127.586	162.928	↑	0.0225723
PC aa C36:6	0.349633	0.451909	↑	0.0270209
PC aa C42:1	0.206177	0.246872	↑	0.0301884
PC ae C38:4	3.91691	4.87044	↑	0.033847
PC aa C36:2	249.656	297.116	↑	0.0373519
SM C20:2	0.285843	0.348296	↑	0.0425698
lysoPC a C18:0	167.607	199.504	↑	0.0443904
PC aa C32:1	3.15613	4.09743	↑	0.0476246
PC aa C40:3	0.381395	0.45225	↑	0.047744
PC ae C36:0	0.39648	0.476158	↑	0.0478637
PC aa C38:3	24.9592	31.1117	↑	0.0492607

Supplemental Table 3. Old age IL 10tm mice metabolite measurements (full table, p-values unadjusted)

ID		Control n=20	IL 10 tm n=16		p-value
Kynurenine / Trp	*	0.0122069	0.0244054	↑	7.07E-07
alpha-AAA	*	6.325	2.98062	↓	2.98E-06
Trp	*	108.88	69.95	↓	5.46E-05
SM C18:0	*	4.54467	1.92527	↓	0.0001196
lysoPC a C20:4	*	81.891	44.5328	↓	0.0001547
lysoPC a C16:1	*	4.88079	2.87555	↓	0.0002673
lysoPC a C18:1		52.288	34.843	↓	0.0007301
PC ae C34:2		1.89739	2.73332	↑	0.0012113
lysoPC a C16:0		233.357	163.846	↓	0.0014066
PC aa C36:0		2.41212	1.65315	↓	0.0017975
SM C18:1		1.19859	0.760069	↓	0.0025144
lysoPC a C24:0		0.949746	0.710284	↓	0.0032113
Fisher ratio		1.95602	2.34761	↑	0.0034275
PC aa C40:3		0.272867	0.351153	↑	0.0037729
PC ae C42:1		0.563257	0.879288	↑	0.0038765
SM (OH) C14:1		1.35906	0.945649	↓	0.0043295
lysoPC a C18:2		164.512	125.249	↓	0.0044795
PC ae C42:3		0.895231	0.624062	↓	0.005439
PC ae C38:3		0.905353	1.24644	↑	0.0056297
Kynurenine		1.2398	1.62625	↑	0.0075272
C16:2-OH		0.0707002	0.079037	↑	0.0076157
PC aa C34:1		49.5688	36.2751	↓	0.010824
PC ae C42:0		0.866178	1.03258	↑	0.0110601
PC ae C36:2		4.78344	6.3887	↑	0.0114751
PC ae C36:3		1.0802	1.35751	↑	0.0117932
SM (OH) C16:1		0.397161	0.29129	↓	0.0125727
SM (OH) C24:1		0.473738	0.381408	↓	0.0170087
Thr		262.065	173.288	↓	0.0186021
PC aa C40:6		57.7794	40.2682	↓	0.0195769
PC aa C38:6		92.2034	64.7815	↓	0.0216498
PC aa C40:2		0.204337	0.254761	↑	0.0225663
PC ae C38:2		2.27925	3.18833	↑	0.0241852
lysoPC a C18:0		191.04	144.447	↓	0.0246072
Phe		74.175	90.5813	↑	0.0265114
Serotonin		4.71245	1.76533	↓	0.0271002

PC ae C38:0	1.90518	1.46673	↓	0.0316329
PC aa C36:3	44.266	55.8637	↑	0.0326819
PC aa C32:0	9.2091	7.48208	↓	0.0410201
C16:1-OH	0.156217	0.168695	↑	0.0429336
PC aa C42:5	0.205434	0.242672	↑	0.0449734
PC ae C30:2	0.0993353	0.0837001	↓	0.0455536
PC aa C24:0	0.33076	0.257438	↓	0.0475861
PC ae C44:5	0.289386	0.250126	↓	0.0477549
Tyr / Phe	0.823069	0.728002	↓	0.0479037

Supplemental Table 4. Demographics of young versus old including age, sex, BMI, systolic BP, Diastolic BP, IL6, TNF α , TNF α R1 (mean & SD).

	Young (n=50)	Old (n=116)	p-value
Age , mean(std)	25.6 (6.2)	77.6 (5.9)	<0.0001
Gender, male, n(%)	34 (68.0)	66 (56.9)	0.18
BMI	26.1 (6.6)	28.0 (5.9)	0.074
Systolic BP	113.9 (14.6)	132.8 (20.1)	<0.0001
Diastolic BP	73.3 (10.9)	73.4 (11.3)	0.96
IL6	0.69 (0.63)	2.04 (4.45)	0.046
TNF α	2.01 (1.16)	3.07 (1.28)	<0.0001
TNF α receptor1	793.65 (298.20)	1454.01 (659.77)	<0.0001

Supplemental Table 5. Average CV's for metabolites in 14 non-frail individuals CV= stdev (of all 4 visits)/ average (of all 4 visits)

Individual variability in blood metabolites can provide information which can help identify candidate metabolites for biomarkers of aging or disease. In order to understand the variability of the metabolites we measured, we performed metabolomics analysis on serum samples collected from fourteen pre-frail individuals four times over a six-month timespan at two-month intervals. This longitudinal collection allowed for the quantification of the variance of each metabolite over the six-month interval. We calculated the average (intra-assay) CV for each metabolite using the CV over the six-month period for each metabolite for each individual patient, and then averaging the CV's from all fourteen subjects. Most of the metabolites had relatively low CV with only 16 metabolites having CV's > 0.3

Metabolite	Average CV	Standard Deviation
PC aa C34:2	0.078	0.04
PC aa C36:2	0.112	0.051
Fisher ratio	0.114	0.044
Non essential AA	0.129	0.061
PC aa C36:4	0.134	0.074
Tyr / Phe	0.134	0.081
PC aa C34:1	0.137	0.064
PC aa C38:4	0.138	0.075
PC ae C42:5	0.14	0.098
Total AA	0.143	0.063
PC ae C44:3	0.148	0.082
PC aa C38:6	0.15	0.096
PC ae C42:1	0.151	0.083
PC ae C44:6	0.151	0.105
Glucogenic AA	0.153	0.118
PC ae C40:4	0.154	0.123
PC ae C34:1	0.155	0.09
PC aa C40:6	0.156	0.091
PC aa C42:5	0.156	0.075
PC ae C40:2	0.156	0.114
PC ae C44:5	0.157	0.09
PC aa C42:0	0.158	0.09
PC aa C36:3	0.158	0.091
PC ae C40:6	0.159	0.11

Phe	0.162	0.093
SM C16:1	0.162	0.11
PC ae C42:4	0.163	0.127
PC aa C40:5	0.163	0.08
PC ae C42:2	0.164	0.089
SM C24:1	0.165	0.105
PC ae C38:4	0.165	0.12
SM C16:0	0.167	0.093
PC aa C28:1	0.167	0.1
PC aa C38:5	0.168	0.097
PC aa C42:6	0.168	0.091
Orn / Arg	0.168	0.079
Gly	0.168	0.116
Val	0.171	0.074
PC ae C40:3	0.171	0.13
PC ae C36:4	0.171	0.104
PC aa C32:0	0.171	0.082
PC aa C42:1	0.172	0.107
PC ae C36:1	0.172	0.133
PC aa C38:3	0.173	0.088
PC aa C42:2	0.173	0.093
PC aa C40:3	0.174	0.08
PC ae C40:1	0.174	0.11
PC ae C40:5	0.175	0.114
SM C26:1	0.175	0.1
Gln	0.175	0.121
SM (OH) C16:1	0.175	0.105
SM (OH) C14:1	0.177	0.107
PC aa C38:0	0.178	0.107
Ala	0.179	0.126
PC ae C38:6	0.18	0.127
PC ae C38:5	0.181	0.114
PC ae C42:3	0.181	0.121
PC ae C38:0	0.183	0.12
PC aa C40:4	0.183	0.093
PC ae C34:2	0.184	0.125
PC ae C34:0	0.186	0.11
PC aa C42:4	0.187	0.064
PC ae C32:2	0.188	0.129
AAA	0.188	0.11
SM C24:0	0.188	0.101
PC ae C36:5	0.189	0.108
Total DMA / Arg	0.189	0.11

SM C18:0	0.189	0.086
SM C18:1	0.189	0.1
C2 / C0	0.189	0.133
C18:2	0.19	0.091
C0	0.19	0.112
PC ae C36:3	0.191	0.126
PC aa C32:3	0.192	0.113
SM (OH) C22:2	0.194	0.112
SM (OH) C24:1	0.194	0.106
PC aa C36:0	0.196	0.105
PC ae C38:3	0.196	0.126
lysoPC a C18:0	0.197	0.101
PC ae C32:1	0.198	0.109
lysoPC a C16:1	0.199	0.126
SM (OH) C22:1	0.199	0.109
PC ae C36:0	0.199	0.113
PC aa C34:3	0.199	0.089
C4	0.2	0.09
PC ae C36:2	0.2	0.174
Kynurenine / Trp	0.201	0.081
PC aa C36:1	0.202	0.107
lysoPC a C16:0	0.202	0.096
PC aa C40:2	0.205	0.112
Met-SO / Met	0.208	0.082
SM C26:0	0.208	0.113
lysoPC a C20:4	0.21	0.087
lysoPC a C17:0	0.21	0.148
Lys	0.211	0.122
PC aa C30:0	0.212	0.083
PC aa C36:6	0.213	0.111
ADMA / Arg	0.213	0.082
PC ae C38:2	0.214	0.115
Pro	0.214	0.071
Essential AA	0.216	0.115
Arg	0.218	0.117
PC ae C34:3	0.219	0.143
Cit / Arg	0.22	0.105
lysoPC a C18:1	0.221	0.117
SM C20:2	0.221	0.108
PC ae C30:2	0.223	0.101
PC aa C34:4	0.223	0.105
Trp	0.224	0.158

total DMA	0.226	0.142
Thr	0.226	0.09
PC ae C44:4	0.226	0.155
Ser	0.227	0.089
C18:1	0.227	0.126
Tyr	0.231	0.12
H1	0.231	0.139
Cit	0.234	0.092
PC aa C36:5	0.235	0.116
Orn	0.236	0.137
C2	0.238	0.131
PC aa C32:1	0.239	0.096
Cit / Orn	0.241	0.099
lysoPC a C20:3	0.241	0.081
PC aa C32:2	0.242	0.086
Kynurenine	0.247	0.152
SDMA / Arg	0.249	0.178
BCAA	0.253	0.127
C3	0.254	0.124
Glu	0.262	0.081
Asn	0.262	0.126
ADMA	0.267	0.176
PC aa C38:1	0.272	0.164
SDMA	0.28	0.233
Taurine	0.28	0.17
Serotonin / Trp	0.297	0.183
C5	0.301	0.215
Putrescine / Orn	0.302	0.194
lysoPC a C18:2	0.303	0.134
Serotonin	0.309	0.108
Met	0.31	0.176
PC ae C38:1	0.321	0.159
SM C22:3	0.326	0.308
Ile	0.33	0.168
Leu	0.331	0.169
Met-SO	0.349	0.211
Asp	0.385	0.15

Supplemental Table 6. Demographics of all older patients stratified by risk status into non-frail and frail groups

	Non-frail (n=83)	Frail (n=33)	p-value
Age , mean(std)	76.7 (5.7)	79.8 (5.9)	0.01
Gender, male, n(%)	39 (47.0)	27 (81.2)	0.0008
BMI	26.65 (3.45)	31.4 (8.5)	<0.0001
Systolic BP	133.03 (17.17)	132.4 (25.0)	0.066
Diastolic BP	74.19 (11.27)	71.5 (11.5)	0.53
IL6	1.96 (3.69)	2.22 (1.76)	0.22
TNFa	3.07 (1.35)	2.73 (0.93)	0.058
TNFa receptor1	1288.01 (580.40)	1862.70 (677.35)	0.0002
Walking time for 4m (sec)	3.75 (0.93)	7.77 (3.58)	<0.0001
Grip strength	26.65 (9.93)	19.04 (6.57)	<0.0001

Supplemental Table 7. Downstream kynurenine pathway metabolites & ratios ranked correlations with IL-6, TNF α , TNF α R1, and IFN γ levels, and frailty status, walking speed, and grip strength

IL-6				TNF α				TNF α R1				IFN γ			
Metabolite	Estimate	P-value	R ²	Metabolite	Estimate	P-value	R ²	Metabolite	Estimate	P-value	R ²	Metabolite	Estimate	P-value	R ²
KYN/PA	0.276	0.00995	0.109	TRP/5HT	3.324	0.00229	0.10390	KYN/PA	0.360	0.000776	0.1574	KA/KYN	-0.286	0.0101	0.108
				KYN/5HT	2.633	0.00281	0.07430	KYN/QA	0.338	0.00110	0.1546	KYN/AA	0.289	0.0210	0.119
				AA/5HT	0.923	0.0192	0.0759	KA/XA	0.350	0.00112	0.158				
				3HK/5HT	0.532	0.0315	0.0433	TRP/3HK	-0.330	0.00113	0.1303				
				XA/KA	-0.253	0.0323	0.0719	XA/TRP	0.298	0.00370	0.1185				
				5HT/KA	-0.232	0.0364	0.0769	KA/QA	0.295	0.00442	0.1155				
				TRP/KA	-0.220	0.0374	0.0607	KA	0.295	0.00580	0.1119				
				KYN/QA	0.236	0.0387	0.0649	KA/PA	0.281	0.00730	0.1089				
								TRP/AA	-0.306	0.0147	0.126				
								TRP/KA	-0.236	0.0252	0.0904				
								3HK/PA	0.230	0.0293	0.0621				
								XA/KYN	-0.218	0.0383	0.0603				
Frailty Status			Grip Strength				Walking Speed								
Metabolite	OR	P-value	Metabolite	Estimate	P-value	R ²	Metabolite	Estimate	P-value	R ²					
3HK/TRP	2.683	0.00251	TRP/QA	0.197	0.0211	0.1406	TRP/XA	0.308	0.00148	0.14020					
TRP/AA	0.281	0.0179	TRP/3HK	0.187	0.0350	0.1473	3HK/TRP	-0.308	0.00158	0.14580					
KA/PA	2.043	0.0314	TRP/XA	0.189	0.0405	0.1189	3HK/PA	-0.243	0.0154	0.1096					
PA/3HK	0.377	0.0321				TRP/KA	0.231	0.0179	0.0930						
KYN/3HK	0.418	0.0343				5HT/TRP	-0.199	0.0355	0.0424						
3HK/XA	1.797	0.0371				3HK	-0.205	0.0401	0.0720						
XA/AA	0.342	0.0374				PA/KA	0.201	0.0420	0.0938						
KA/KYN	1.845	0.0386													
3HK	1.786	0.0393													
KA/XA	1.928	0.0415													

Supplemental Table 8. Metabolite abbreviation list

Acylcarnitines			
C0	Carnitine	C10:1	Decenoylcarnitine
C2	Acetylcarnitine	C10:2	Decadienylcarnitine
C3	Propionylcarnitine	C12	Dodecanoylcarnitine
C3:1	Propenoylcarnitine	C12:1	Dodecenoylcarnitine
C3-OH	Hydroxypropionylcarnitine	C12-DC	Dodecanedioylcarnitine
C4	Butyrylcarnitine	C14	Tetradecanoylcarnitine
C4:1	Butenylcarnitine	C14:1	Tetradecenoylcarnitine
C4-OH(C3-DC)	Hydroxybutyrylcarnitine	C14:1-OH	Hydroxytetradecenoylcarnitine
C5	Valerylcarnitine	C14:2	Tetradecadienylcarnitine
C5:1	Tiglylcarnitine	C14:2-OH	Hydroxytetradecadienylcarnitine
C5:1-DC	Glutaconylcarnitine	C16	Hexadecanoylcarnitine
C5-DC(C6-OH)	Glutaryl carnitine (Hydroxyhexanoylcarnitine)	C16:1	Hexadecenoylcarnitine

C5-M-DC	Methylglutaryl carnitine	C16:1-OH	Hydroxyhexadecenoyl carnitine
C5-OH (C3-DC-M)	Hydroxyvaleryl carnitine (Methylmalonyl carnitine)	C16:2	Hexadecadienyl carnitine
C6 (C4:1-DC)	Hexanoyl carnitine (Fumaryl carnitine)	C16:2-OH	Hydroxyhexadecadienyl carnitine
C6:1	Hexenoyl carnitine	C16-OH	Hydroxyhexadecanoyl carnitine
C7-DC	Pimelyl carnitine	C18	Octadecanoyl carnitine
C8	Octanoyl carnitine	C18:1	Octadecenoyl carnitine
C9	Nonanoyl carnitine	C18:1-OH	Hydroxyoctadecenoyl carnitine
C10	Decanoyl carnitine	C18:2	Octadecadienyl carnitine

Amino Acids & Biogenic

Amines

Ala	Alanine	Ac-Orn	Acetylornithine
Arg	Arginine	ADMA	Asymmetric dimethylarginine
Asn	Asparagine	SDMA	Symmetric dimethylarginine
Asp	Aspartate	total DMA	Total dimethylarginine
Cit	Citrulline	alpha-AAA	alpha-Amino adipic acid
Gln	Glutamine	Carnosine	Carnosine
Glu	Glutamate	Creatinine	Creatinine
Gly	Glycine	Histamine	Histamine
His	Histidine	Kynurenine	Kynurenine
Ile	Isoleucine	Met-SO	Methionine sulfoxide
Leu	Leucine	Nitro-Tyr	Nitrotyrosine
Lys	Lysine	OH-Pro	Hydroxyproline
Met	Methionine	PEA	Phenylethylamine
Orn	Ornithine	Putrescine	Putrescine
Phe	Phenylalanine	Sarcosine	Sarcosine
Pro	Proline	Serotonin	Serotonin
Ser	Serine	Spermidine	Spermidine
Thr	Threonine	Spermine	Spermine
Trp	Tryptophan	Taurine	Taurine
Tyr	Tyrosine		
Val	Valine		

Monosaccharides

H1	Sum of Hexoses (including Glucose)
	SM sphingomyelin
	PC Phosphatidylcholine
	Aa acyl-acyl
	Ae acyl-alkyl

Supplemental Tables 9 & 10. The standards (**Supplemental Table 9**) and internal standards (**Supplemental Table 10**) were characterized using the MRM ion transitions and their voltages: declustering potentials (DP), collision energies (CE), and collision cell exit potentials (CXP) are listed below. Quinolinic Acid was characterized using spectra for picolinic acid and is reported as only the area under the curve ratio with the internal standard.

Supplemental Table 9.

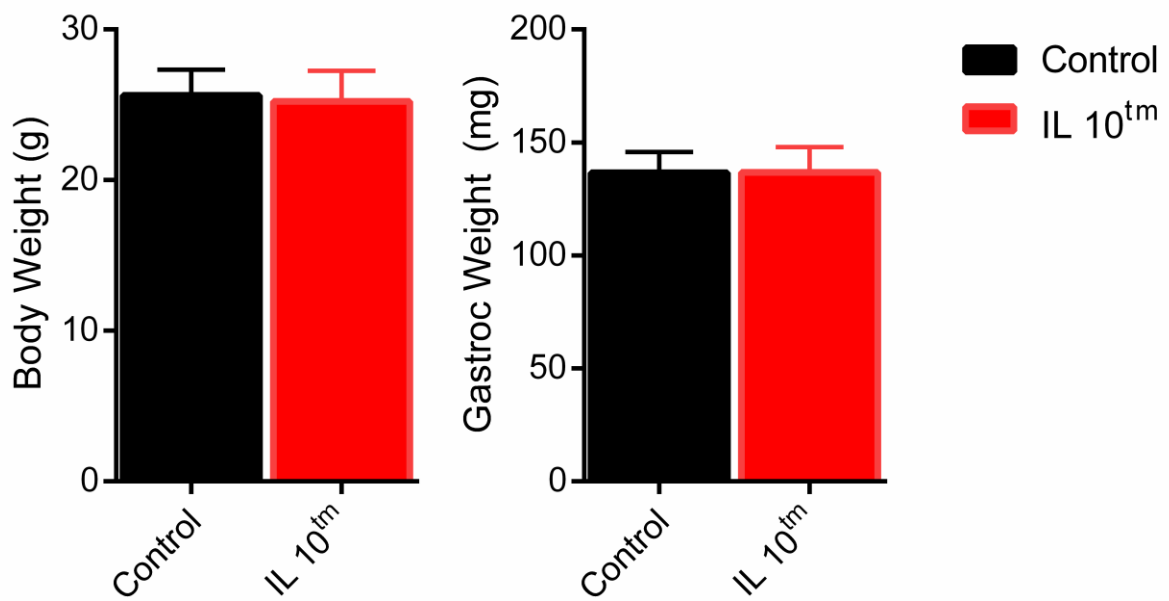
Analyte	Q1/Q3	DP	CE	CXP
3-Hydroxy Kynurenine	226.0/209.1	20	11	10
Kynurenine	209.0/94.0	20	19	15
Xanthurenic Acid	206.2/160.0	56	27	10
Kynurenic Acid	189.9/144.0	55	25	8
Picolinic Acid	124.0/78.0	10	25	10
Tryptophan	205.1/118.0	39	35	11
Anthranilic Acid	138.2/120.0	36	15	8
Serotonin	177.0/160.1	26	13	8

*Quinolinic Acid was characterized using spectra for picolinic acid and is reported as only the area under the curve ratio with the internal standard.

Supplemental Table 10.

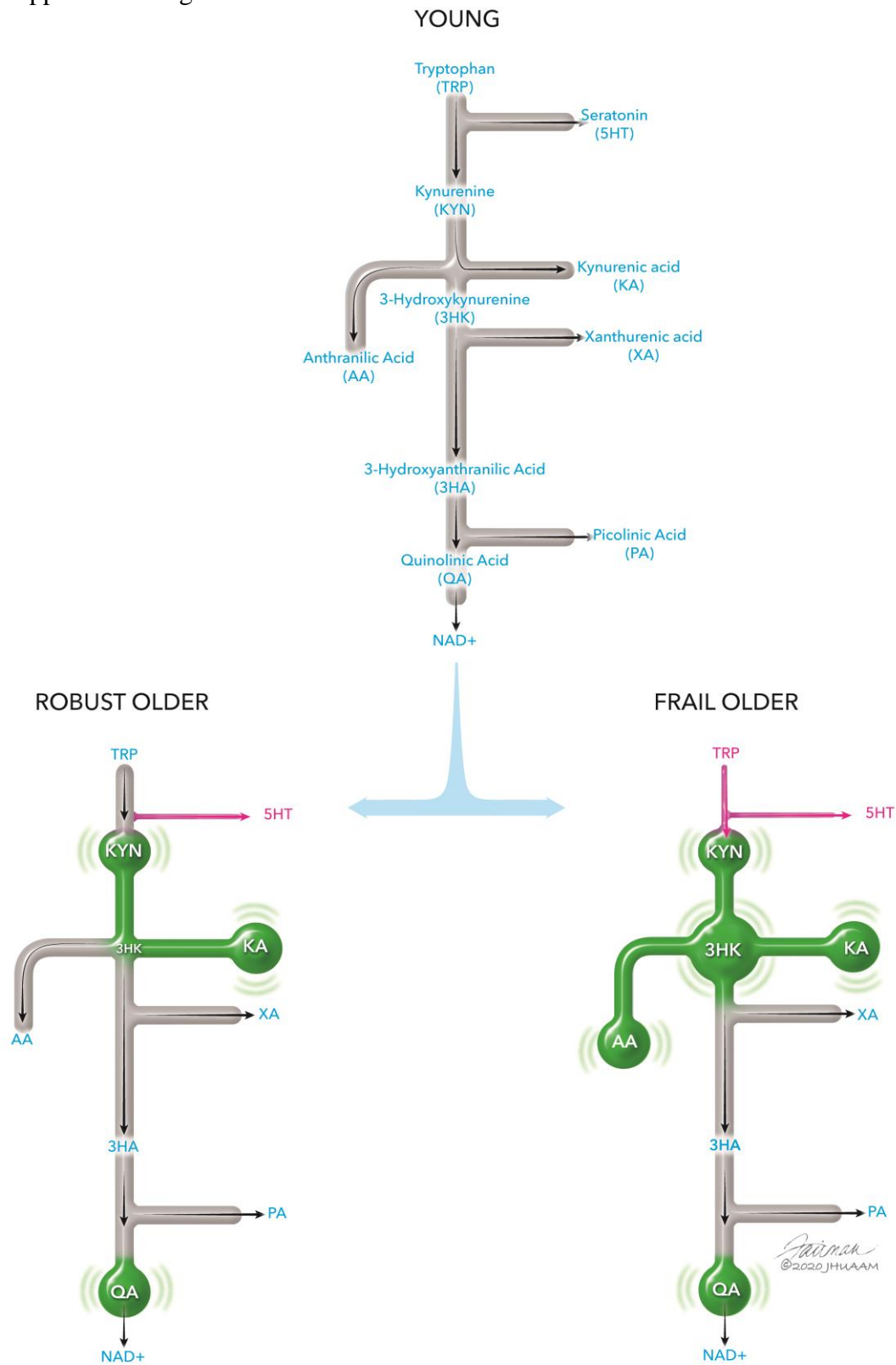
Internal Standard	Q1/Q3	DP	CE	CXP
D ₄ -KYN	231.0/98.0	20	19	15
D ₄ -XA	210.0/164.0	56	27	10
D ₅ -KA	195.0/149.2	55	27	8
D ₄ -PA	128.1/82.1	10	25	10
D ₅ -TRP	210.1/122.0	30	35	10
D ₃ -5HT	181.1/164.1	26	13	10

Supplemental Figure 1



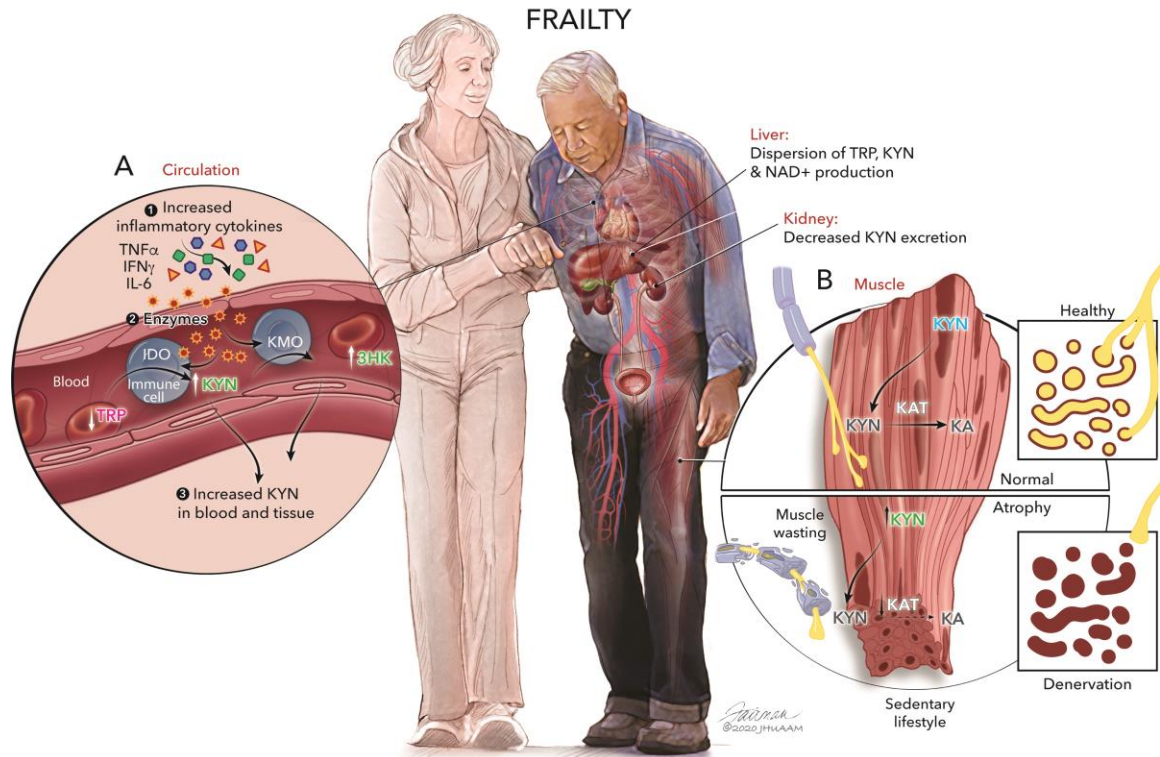
Supplemental Figure 1. Body weight and gastrocnemius weight from 3 month old IL 10tm and control mice (n=10 per group)

Supplemental Figure 2



Supplemental Figure 2. Summary of kynurenine pathway changes with age & frailty

Supplemental Figure 3



Supplemental Figure 3. Schematic summarizing the physiological connections between inflammation, kynurenines and neuromuscular integrity. (A) Increased levels of inflammation trigger IDO and KMO in immune cells to produce more kynurenines. (B) These kynurenines can damage neuromuscular junctions leading to decreased neuromuscular integrity.

